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**LA RELACIÓN ENTRE SUPERFETACIÓN, PLACENTACIÓN, RETENCIÓN DE
ESPERMA Y TAMAÑO DEL EMBRIÓN EN PECES VIVÍPAROS DE LA FAMILIA**

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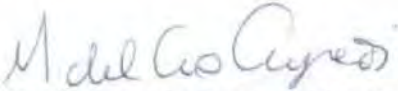
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RESUMEN

La superfetación es la capacidad de la hembra para llevar dentro de su ovario simultáneamente múltiples camadas de embriones en diferentes estadios de desarrollo. Este modo reproductor predomina en algunos géneros de peces vivíparos de la familia Poeciliidae (*Poeciliopsis*, *Heterandria* y *Neoheterandria*). La matrotrofia es un modo de aprovisionamiento materno en la cual la energía destinada al desarrollo del embrión es provista por la madre después de la fertilización y durante todo el desarrollo embrionario. En peces poecílidos, entre mayor es el nivel matrotrófico la superfetación aumenta. Tanto para la superfetación como para la matrotrofia existen hipótesis al respecto de su origen evolutivo. Sin embargo, son escasos los estudios que se enfocan en demostrar las consecuencias evolutivas (cambios o efectos en sus atributos de historias de vida) de la superfetación y la matrotrofia. Por lo que en la presente tesis probé diferentes hipótesis de las consecuencias evolutivas de la superfetación y la matrotrofia sobre características de historia de vida (tamaño de la cría) y reproductoras (nivel de placentación y retención de esperma) en algunas especies de la familia Poeciliidae.

En el primer capítulo, analicé la relación entre la matrotrofia y la superfetación con diferentes ultraestructuras placentarias. Esperaba que debido a que las hembras con superfetación requieren una regulación diferencial de nutrientes de embriones en distintos estadios de desarrollo, estas especies que superfetan tendrían una placenta ultraestructuralmente más compleja comparada con la placenta de especies que no superfetan. Esta hipótesis se cumplió parcialmente, ya que observé sólo una tendencia de algunas características morfológicas a ser más complejas (mayor número de microvellosidades en la célula folicular y mayor área del folículo materno) en las especies que superfetan en comparación con las especies que no superfetan. Con respecto a la matrotrofia, esperaba que debido a que ésta implica la transferencia activa de nutrientes por parte de la madre hacia sus embriones a lo largo de todo el desarrollo, las hembras de especies con altos niveles de matrotrofia tendrían una placenta más compleja (mayor valor en las diferentes ultraestructuras de la placenta) en comparación con las hembras de especies con bajos niveles de matrotrofia. Esta hipótesis se cumplió, porque observé que especies con altos niveles de matrotrofia tienen placentas más complejas (altos valores de las características ultraestructurales). Por lo que el área del folículo materno, el número de microvellosidades, el número de vesículas en la célula folicular y el tamaño de estas vesículas tienden a aumentar conforme aumenta el índice matrotrófico. Estas características ultraestructurales de la placenta posiblemente favorecen la transferencia activa de nutrientes entre la madre y el embrión.

En el segundo capítulo examiné la relación que existe entre las especies que superfetan y las que no superfetan con la capacidad de almacenar esperma. La hipótesis se debe a que las hembras de las especies que superfetan tienen que fertilizar constantemente a los ovocitos, se esperaban más y más grandes espermatecas en especies que superfetan en comparación con especies que no superfetan. Encontré espermatecas en ovarios de nueve de las doce especies que examiné (*Gambusia panuco*, *Xiphophorus hellerii*, *Poecilia mexicana*, *Poeciliopsis prolifica*, *P. infans*, *P.*

viriosa, *P. presidionis*, *P. gracilis* y *Heterandria formosa*). Las espermatecas están compuestas de epitelio ovárico y en general de forma ovalada o circular que rodean completamente a los espermatozoides para protegerlos. Las cabezas de los espermatozoides se encontraron en contacto con el epitelio ovárico y las colas hacia la luz de la espermateca. Además, encontré que existe una relación entre la presencia de superfetación con el número y tamaño de las espermatecas. Las especies que presentan superfetación tienen mayor número y más grandes espermatecas en comparación con las especies que no superfetan. Por lo tanto, las especies superfetadoras destinan una mayor proporción del volumen total del ovario a la producción de espermatecas en comparación con las especies sin superfetación.

En el tercer capítulo investigué la relación que existe entre la combinación de matrotrofia-superfetación con el tamaño de la cría. La hipótesis es que la evolución conjunta de la superfetación y la matrotrofia en la familia Poeciliidae ha facilitado la evolución de crías más grandes. Las hembras superfetadoras producen varias camadas pequeñas y más frecuentemente en diferentes estadios de desarrollo, en consecuencia con distinto tamaño, por lo que deberían liberar un poco de espacio dentro de su ovario y producir crías más grandes. Por otro lado, la matrotrofia implica que los embriones son pequeños en los primeros estadios de desarrollo y más grandes en los últimos estadios del desarrollo. Por lo que, una combinación de la presencia de superfetación y alta matrotrofia podría promover el incremento del tamaño de las crías al nacer. Sin embargo, los resultados de métodos comparados filogenéticos mostraron que altos niveles superfetación y matrotrofia no tienen una relación con el tamaño de las crías al nacer. Adicionalmente, los resultados de métodos comparados filogenéticos permitieron evaluar el nivel de heredabilidad filogenética del tamaño de la cría al nacer, mostrando que el tamaño de las crías en todas las especies que se analizaron tiene baja heredabilidad filogenética. Esto indica que la evolución del tamaño al nacer en poecílicos se debe a procesos de adaptación local.

ABSTRACT

Superfetation is the ability of females to simultaneously carry multiple broods of embryos at different developmental stages. This mode of reproduction predominates in some genera of viviparous fishes of the family Poeciliidae (*Poeciliopsis*, *Heterandria* and *Neoheterandria*). Matrotrophy is a mode of maternal provisioning in which females provide the nutrients to developing embryos after fertilization and during the embryo development. In poeciliid fishes, superfetation increases when the matrotrophy index is high. There are hypothesis with respect to the evolutionary origin of superfetation and matrotrophy. However, there are a few studies that considerer to prove evolutionary consequences (changes or effects on life history traits) of superfetation and matrotrophy. Hence, in this thesis I proved different hypothesis of evolutionary consequences of superfetation and matrotrophy on life history traits (offspring size) and reproductive characteristics (level of placentation and sperm storage) in some species of the family Poeciliidae.

In chapter one, I analyzed the relationship between matrotrophy and superfetation with different placental ultrastructures. I expected that because females with superfetation require differential regulation of embryo nutrients in distinct developmental stages, these superfetating species would have an ultrastructurally more complex placenta compared with placentas of non-superfetating species. This hypothesis was partially fulfilled, since I only observed a trend of some morphological characteristics to be more complex (greater microvilli number of the follicular cell and greater area of maternal follicle) in superfetating species compared with non-superfetating species. With respect to matrotrophy, I expected that because matrotrophy involves active transfer of nutrients from the mother to their embryos during the whole development, species with high matrotrophy would require a more specialized placenta (greater value in the different ultrastructures of the follicular placenta) compared with low matrotrophy. This hypothesis was fulfilled, because I observed that species with high degrees of matrotrophy have more complex placentas (high values of the ultrastructural characteristics). Therefore, area of maternal follicle, number of microvilli, number of vesicles, and size of these vesicles tend to increase with the matrotrophy index. These ultrastructural characteristics of the placenta possibly favor the active transfer of nutrients from the mother to the embryo.

In chapter two, I examined the relationship between superfetating species and non-superfetating species with the ability of sperm storage. The hypothesis is due to the fact females with superfetation having to fertilize their oocytes constantly, more and larger spermathecae were expected in species with superfetation compared to non-superfetating species. I found spermathecae in the ovaries of nine of the twelve species that I examined (*Gambusia panuco*, *Xiphophorus hellerii*, *Poecilia mexicana*, *Poeciliopsis prolifica*, *P. infans*, *P. viriosa*, *P. presidionis*, *P. gracilis* and *Heterandria formosa*). The spermathecae are composed of ovarian tissue and are in generally oval or circular structures that completely surround spermatozoa to protect them. The spermatozoa heads were found in contact with ovarian tissue and the tails towards the lumen of the spermatheca. Also, I found that there is a relationship between presence of superfetation with

number and size of the spermathecae. Superfetating species have a greater number and size spermathecae compared with non-superfetating species. Thus, species with superfetation allocate a greater proportion of the total volume of the ovary to spermathecae production compared with non-superfetating species.

In chapter three, I explored the relationship that exists between the combination of matrotrophy and superfetation with offspring size. The hypothesis is that the combined evolution of superfetation and matrotrophy in the family Poeciliidae has facilitated the evolution of larger offspring. Females with superfetation produce smaller broods and do so more often in different development stages, consequently with different size, so females with superfetation should release some space inside their ovary and produce larger offspring. On the other hand, matrotrophy involves that embryos are small in early development stages and large in late development stage. Therefore, a combination of superfetation and high degree of matrotrophy could promote the increase in offspring size. However, the results of phylogenetic comparative methods showed that high degree of superfetation-matrotrophy is not related with offspring size. Additionally, the results of phylogenetic comparative methods allowed to evaluate the level of phylogenetic heritability of the offspring size, showing that offspring sizes in all analyzed species have low phylogenetic heritability. That indicates that offspring size evolution in poeciliids is due to local adaption processes.

INTRODUCCIÓN GENERAL

Superfetación

La superfetación es la capacidad de una hembra para llevar dentro de su ovario y de forma simultánea múltiples camadas de embriones en diferentes estadios de desarrollo (Turner, 1937, 1940; Scrimshaw, 1994). En mamíferos, la superfetación ocurre en pocos grupos como mustélidos y lepóridos (e.g. *Lepus europaeus*, *Meles meles* y *Neovison vison*; Yamaguchi *et al.*, 2004, 2006; Roellig *et al.*, 2011; Corner *et al.*, 2015). Sin embargo, existen casos esporádicos en humanos, aunque esto sólo sucede como consecuencia de tratamientos de fertilidad (Pape *et al.*, 2008). Estos dos fetos simultáneos mantuvieron una diferencia de 4 semanas en su desarrollo (Tuppen *et al.*, 1999). Por otro lado, los moluscos como los bivalvos de la familia Sphaeriidae exhiben incubación secuencial (*secuencial brooding*) que es equivalente a la superfetación. Lo anterior se debe a que estos organismos presentan subconjuntos discretos de desarrollo en la camada (*brood masses*), con cada subconjunto encapsulado en su propio saco (Cooley y Foighil, 2000).

La superfetación es el modo reproducción que predomina en peces vivíparos (Turner, 1937, 1940; Scrimshaw, 1944; Thibault y Schultz, 1978; Zúñiga-Vega *et al.*, 2010) y ha evolucionado de manera independiente en tres familias (i.e. Clinidae, Poeciliidae y Zenarchopteridae; Reznick y Miles, 1989; Gunn y Thresher, 1991; Reznick *et al.*, 2007). Varios géneros de peces vivíparos de la familia Poeciliidae exhiben superfetación como *Poeciliopsis*, *Heterandria* y *Neoheterandria*, mientras que otros géneros no la presentan (e.g. *Alfaro*, *Brachyrhaphis* y *Xiphophorus*; Reznick y Miles, 1989; Zúñiga-Vega *et al.*, 2010).

De la misma manera, en la familia Zenarchopteridae algunos géneros como *Hemirhamphodon*, *Nomorhamphus* y *Dermogenys* poseen superfetación, mientras que otros géneros no (e.g. *Tondanichthys* y *Zenarchopterus*; Reznick *et al.*, 2007). Por lo que la distribución discontinua de la superfetación en las historias filogenéticas de las familias superfetadoras, sugiere que esta estrategia reproductiva ha evolucionado de forma independiente en diferentes linajes (Zúñiga-Vega *et al.*, 2010). Siendo posible medir el grado de superfetación en una hembra, definido como el número de camadas simultáneas dentro del ovario de la hembra, y el cual puede variar dependiendo de la especie. Por ejemplo, especies como *Heterandria formosa*, *Poeciliopsis prolifica* y *Poeciliopsis turneri* pueden tener un alto grado de superfetación, siendo el extremo de la misma las hembras de *Heterandria formosa*, las cuales pueden tener hasta ocho camadas de diferente estadio de desarrollo dentro de su ovario (Travis *et al.*, 1987). En contraste, *Poeciliopsis baenschii*, *P. viriosa* y *P. latidens* presentan un menor grado de superfetación, donde, hembras de *P. viriosa* pueden llevar solamente 2 camadas simultáneas de diferente estadio de desarrollo dentro de sus ovarios (Olivera-Tlahuel *et al.*, 2015). Esta variación en el grado de superfetación se puede presentar entre poblaciones de una misma especie (*Heterandria formosa*; Travis *et al.*, 1987; *Poeciliopsis prolifica*, Pires *et al.*, 2007; *P. turrubarensis*, Zúñiga-Vega *et al.*, 2007). Por ejemplo, en *Poeciliopsis turrubarensis*, las hembras de diferentes poblaciones pueden variar en su grado de superfetación, desde 2 a 4 camadas de diferente estadio de desarrollo dependiendo de las condiciones del ambiente (i.e. ambientes costeros o dulceacuícolas; Zúñiga-Vega *et al.*, 2007).

Superfetación y placentación

Al igual que la superfetación, la viviparidad ha evolucionado en múltiples ocasiones en los vertebrados (Blackburn, 1992). Durante la evolución de la oviparidad a la viviparidad, el aparato reproductor de la hembra desarrolló estructuras más complejas para la transferencia de nutrientes hacia el embrión de manera eficiente. Por lo que distintos tipos de placentas han evolucionado en distintas ocasiones a través del tiempo (Pollux *et al.*, 2009; Pollux *et al.*, 2014). En mamíferos, la placenta y sus características han sido ampliamente estudiadas (Mossman, 1937; Perry, 1981; Moll, 1985; Blackburn, 1999; Swain y Jones, 2000; Vogel, 2005; Carter y Mess, 2007). Sin embargo, en peces son relativamente pocos los estudios sobre la evolución de la placenta y sus características (e.g. los géneros *Heterandria* y *Poeciliopsis*; Rosen y Bailey 1963; Hrbek *et al.*, 2007; Pollux, *et al.*, 2009; Pollux *et al.*, 2014). Por ejemplo, en *Poeciliopsis gracilis* se ha mostrado que el incremento en la actividad del epitelio folicular que rodea al embrión y su alta vascularización facilita el desarrollo embrionario (Wourms, 1981; Wourms *et al.*, 1988; Constanz, 1989; Uribe *et al.*, 2004). Similarmente, en hembras de *Heterandria formosa* y de algunas especies de *Poeciliopsis* se observó la presencia de estructuras con función de absorción e intercambio de nutrientes en el epitelio folicular (i.e. microvellosidades, sistemas complejos de vesículas, complejos aparatos de Golgi con 4 o 5 cisternas y un extenso retículo endoplásmico; Turner 1940; Grove y Wourms, 1991; Pollux *et al.*, 2009).

En esta investigación, propongo la hipótesis de que las hembras de especies con superfetación desarrollaron características placentarias más complejas (valores elevados de las características ultraestructurales) a fin de mantener camadas simultáneas de diferentes

estadios de desarrollo dentro de su ovario. Hasta ahora, no se ha estudiado la posible relación entre las características placentarias y la presencia o ausencia de superfetación en peces vivíparos. La primera parte de esta tesis se enfocará en examinar morfológicamente las placentas de distintas especies de la familia Poeciliidae y la posible relación entre la complejidad placentaria y la presencia de matrotrofia y superfetación.

Superfetación y matrotrofia

Por otro lado, se ha demostrado que existe una relación positiva entre la superfetación y la matrotrofia. En peces poecílidos, entre mayor es el nivel matrotrófico la superfetación aumenta (Meredith *et al.*, 2011; Pollux *et al.*, 2014). La matrotrofia es un modo reproductor en la cual la energía destinada al desarrollo del embrión es provista por la madre después de la fertilización y durante todo el desarrollo embrionario (Wourms, 1981; Marsh-Matthews, 2011). Presentando, los embriones de especies matrotróficas escaso vitelo al momento de ser fertilizados (i.e. reserva de sustancias nutritivas). Por lo tanto, las especies matrotróficas se caracterizan por un aumento elevado del peso del embrión durante el desarrollo debido a que las madres les están transfiriendo nutrientes de manera activa (e.g. hembras de *Poecilia branneri* y *P. bifurca*; Pires *et al.*, 2010). En contraste a la matrotrofia, la lecitrotrofia es un modo reproductor en el cual la energía destinada al desarrollo del embrión es provista antes de la fecundación por medio del vitelo (Wourms, 1981; Marsh-Matthews, 2011). Por lo tanto, los embriones pierden peso durante su desarrollo embrionario debido a costos metabólicos (de 30 a 35 % de su peso; Wourms,

1981; Pires *et al.*, 2010). En la familia Poeciliidae, existe mucha variación entre estas dos estrategias reproductoras, desde especies completamente lecitotróficas hasta especies con matrotrofia extensa (Pollux *et al.*, 2014).

Al igual que lo que se espera con respecto a la superfetación, un organismo altamente matrotrofico debería tener una placenta mucho más desarrollada en comparación con especies con menor intensidad matrotrofica (Kwan *et al.*, 2015). Esta hipótesis se basa en el hecho de que en las especies altamente matrotroficas la madre proporciona todos los nutrientes necesarios para el crecimiento del embrión de manera activa a lo largo de todo el desarrollo (Mossman, 1937, 1987). Entonces, una placenta con ultraestructuras más complejas podría facilitar el intercambio de nutrientes entre la madre y el embrión. Recientemente, Kwan y colaboradores (2015) demostraron que especies de poecílicos con diferentes niveles de matrotrofia variaban en el grosor de la placenta folicular, el plegamiento del epitelio folicular y su cercana asociación con los capilares sanguíneos sugiere que estas características son requeridas para soportar el incremento del intercambio de sustancias de la madre hacia el embrión. En especies que son altamente matrotroficas el grosor de sus placentas foliculares es mayor en comparación con especies con menor nivel de matrotrofia. Sin embargo, no existen más estudios sobre la relación entre la matrotrofia y las características morfológicas y fisiológicas de las placentas. La primera parte de esta tesis se enfocará en examinar morfológicamente las placentas de distintas especies de la familia Poeciliidae y la posible relación entre la complejidad ultraestructural placentaria con la presencia de matrotrofia y superfetación.

Superfetación y retención de esperma

La retención de esperma es la capacidad de las hembras para almacenar esperma viable dentro de su aparato reproductor por largos periodos (Orr y Zuk, 2012). Este interesante fenómeno se presenta en algunos grupos de insectos (Simmons, 2001), reptiles (Birkhead y Møller, 1993), anfibios (Seber, 2002), peces óseos (Potter y Kramer, 2000), peces cartilagosos (Pratt, 1993) y mamíferos (Birkhead y Møller, 1993). El tiempo de retención del esperma puede variar entre estos grupos, desde sólo algunas horas hasta años después de que la hembra se aparee. Por ejemplo, los insectos (abejas y termitas reinas) pueden guardar el esperma durante varios años (Orr y Zuk, 2012). En contraste, las hembras de mamíferos (e.g. cerdos u ovejas) retienen el esperma sólo un par de horas (Orr y Zuk, 2012). Particularmente en peces óseos, la retención de esperma se ha documentado en las familias Cottidae, Embiotocidae, Sebastidae y Poeciliidae (Fraser y Renton, 1940; Darling *et al.*, 1980; Koya *et al.*, 2002; Vila *et al.*, 2007). En estos grupos de peces, el almacenamiento de los espermatozoides puede ocurrir en el ovario o en el gonoducto (Darling *et al.*, 1980; Koya *et al.*, 2002; Vila *et al.*, 2007). En *Helicolenus dactylopterus* (Sebastidae) la retención de esperma ocurre en cámaras de almacenamiento que se encuentran en el ovario (Vila *et al.*, 2007), para dichas familias, los espermatozoides se concentran en el interior de estas cámaras. Por otro lado, en *Xiphophorus maculatus* (Poeciliidae) se han observado espermatozoides asociados al epitelio del ovario y del gonoducto (Potter y Kramer, 2000).

En peces poecílicos la capacidad para almacenar esperma se ha estudiado relativamente poco y hasta la fecha sólo se ha documentado en *Heterandria formosa*,

Poecilia reticulata, *Xiphophorus maculatus* y *Poeciliopsis gracilis* (Turner, 1947; Contantz, 1989, Potter y Kramer, 2000; Kobayashi y Iwamatsu, 2002; Uribe *et al.*, 2004; Uribe y Grier, 2011). Únicamente en *Heterandria formosa* y *Poecilia reticulata* se han observado estructuras que almacenan el esperma temporalmente (denominadas 'solid plugs' y 'sperm storage micropockets', respectivamente) y que se forman debido al cierre de los pliegues ováricos que contienen a los espermatozoides (Fraser y Renton, 1940; Kobayashi y Iwamatsu, 2002).

Debido a que la superfetación implica una producción constante de embriones en diferentes estadios de desarrollo (Turner, 1937, 1940; Scrimshaw, 1994), se espera que las especies con superfetación requieran fertilizar constantemente a sus ovocitos. Por lo que, en esta investigación se propuso como hipótesis que en las hembras de especies con superfetación, existen estructuras que almacenen espermatozoides y que les permiten estar disponibles para fertilizar constantemente a diferentes ovocitos. Hasta la fecha, no existen trabajos que demuestren una relación entre la retención de esperma y la superfetación en peces de la familia Poeciliidae. Por lo tanto, el segundo tema de investigación de la presente tesis se enfocará en demostrar la presencia de estructuras de almacenamiento de esperma y su posible relación con la presencia de superfetación.

Superfetación y tamaño del embrión

Finalmente, la asociación entre la matrotrofia y la superfetación ha llevado a la hipótesis de que la evolución de uno de estos rasgos ha facilitado la evolución del otro

(Pollux *et al.*, 2009, 2014; Trexler y DeAngelis, 2010; Meredith *et al.*, 2011). La aparente coevolución de estas estrategias reproductoras puede llevar a importantes consecuencias en términos del esfuerzo reproductor. Por lo tanto, el significado adaptativo de la coevolución entre superfetación y matrotrofia podría encontrarse en algunas características de historias de vida como el tamaño de la cría al nacer.

Con base a lo anterior se espera que una hembra sin superfetación que produce grandes camadas de embriones debe sufrir de limitaciones tanto energéticas como de espacio dentro de su ovario, por lo que el tamaño individual de cada embrión debe ser relativamente pequeño (compromiso entre número y tamaño de las crías; Roff, 2002). En contraste, una hembra con superfetación reparte su esfuerzo reproductor produciendo pequeñas camadas que nacen frecuentemente (Reznick y Miles, 1989; Pollux *et al.*, 2009; Zuñiga-Vega *et al.*, 2010). Es decir, las hembras superfetadoras dentro de su ovario producen varias camadas en diferente estadio de desarrollo, en consecuencia con distinto tamaño, entonces deberían liberar un poco de espacio dentro de su ovario (en comparación con especies que carecen de superfetación). Por lo que la hipótesis de esta sección es que el espacio adicional que tienen las hembras superfetadoras en su ovario podrían usarlo para producir crías más grandes. Ya que la producción de crías más grandes favorece su adecuación en ciertos ambientes (Reznick y Endler, 1982; Reznick *et al.*, 1990; Bashey, 2008; Riesch *et al.*, 2010

Por otro lado, la matrotrofia implica que la hembra provee los nutrientes necesarios después de la fertilización y durante todo el desarrollo embrionario. Por lo que los embriones son pequeños en los primeros estadios de desarrollo debido a su escaso vitelo y

más grandes en los últimos estadios del desarrollo. Una combinación de la presencia de superfetación y alta matrotrofia podría promover el incremento del tamaño de las crías al nacer. Similar a los dos temas antes planteados, no existen estudios previos enfocados al estudio de los patrones de coevolución en las estrategias reproductoras de superfetación-matrotrofia, y sus consecuencias sobre las características de historias de vida como en el tamaño de las crías al nacer. El tercer tema de investigación de la presente tesis es examinar esta posible relación.

En resumen, existen varias interrogantes sobre las consecuencias morfológicas y reproductoras de la convergencia evolutiva de la superfetación. ¿Qué relación hay entre la superfetación y las características placentarias de peces poecílicos? ¿Existe una relación entre la intensidad matrotrofica y la complejidad de las placentas? ¿Existen estructuras morfológicas especializadas para la retención de esperma en estos peces vivíparos? ¿Ha habido un efecto de la coevolución de las estrategias reproductoras superfetación-matrotrofia sobre el tamaño del embrión al nacer? La búsqueda de respuestas a estas interrogantes es la base de la presente investigación. Por lo tanto, mi objetivo principal es conocer la relación que existe entre superfetación, placentación, matrotrofia, retención de esperma y tamaño del embrión en diferentes especies de la familia Poeciliidae. Los resultados de la presente tesis son una contribución importante al entendimiento de los mecanismos reproductores de los peces vivíparos de la familia Poeciliidae. Las posibles consecuencias o cambios en los atributos de la historia de vida de los peces vivíparos provocados por la superfetación no se habían estudiado hasta esta investigación. Por lo que abordé varios temas interesantes relacionados con la superfetación que van desde cambios

en el tamaño de la cría al nacer hasta cambios morfológicos ultraestructurales de la placenta folicular. Adicionalmente, utilicé métodos innovadores y actuales (i.e. métodos comparativos filogenéticos y estereología) para poder unir los enfoques ecológico, evolutivo y morfológico de los mecanismos de la reproducción en las especies estudiadas.

La familia Poeciliidae y sitios de estudio

Los poecílidos son peces de agua dulce y salobre que se distribuyen desde el este de Estados Unidos hasta el noroeste de Argentina (Miller et al., 2009). Esta familia está conformada por 22 géneros y 180 especies, en general son pequeños y llegan a medir únicamente de 31 a 70 mm (Miller et al., 2009). La familia Poeciliidae tiene fecundación interna y son vivíparos (excepto el género *Tomeurus*; Miller et al., 2009). Los machos tienen una aleta anal modificada (de los rayos 3 al 5) llamada gonopodio que permite fertilizar a la hembra (Miller et al., 2009). Por su parte, la hembra debido a la carencia del conducto de Müller tiene un ovario en donde ocurre tanto la fertilización como la gestación (Amoroso, 1968; Turner 1947; Wourms, 1981). El tipo de gestación de estos poecílidos es intrafolicular, es decir que los ovocitos se fecundan dentro de un folículo materno en el ovario y se mantienen ahí hasta justo antes de su nacimiento (Fig. 1A-B; Wourms et al., 1988). Además, los poecílidos tienen una placenta llamada tipo folicular que se encuentra formada por los tejidos maternos y embrionarios (Turner, 1940). Los tejidos embrionarios son la piel absorbente del embrión, saco pericárdico y el saco vitelino (Turner, 1940). Mientras que los tejidos maternos son el epitelio folicular y el tejido conjuntivo laxo que lo rodea (Turner, 1940;

Grove y Wourms, 1994; Fig. 1C). En la presente tesis me enfoqué en 12 especies de la familia Poeciliidae. Las especies superfetadoras son: *Heterandria formosa* (obtenida de un acuario privado de Estados Unidos), *Poeciliopsis infans*, *P. gracilis*, *P. presidionis*, *P. prolifica* and *P. viriosa*. Mientras que las especies no superfetadoras son: *Pseudoxiphophorus bimaculatus*, *Xiphophorus hellerii*, *Poecilia mexicana*, *Priapella intermedia* y *Gambusia panuco*. Estas especies fueron seleccionadas debido a que traté de muestrear la mayor cantidad de especies de poecílicos en un área grande de la República Mexicana (Fig. 2; tabla 1).

Preguntas de investigación y presentación de los capítulos que componen la tesis

Para lograr el objetivo principal, la presente tesis se enfoca en contestar las preguntas de investigación que mencioné anteriormente. Estas preguntas de investigación definen los objetivos particulares de cada capítulo.

Capítulo 1

¿En qué difieren las placentas de especies de peces que superfetan con respecto a las placentas de especies que no superfetan? ¿Altos niveles de matrotrofia están asociados con placentas más complejas?

Hipótesis:

a) Debido a que la superfetación requiere una regulación diferencial por parte de las hembras de las tasas de desarrollo de embriones en distintos estadios (Turner, 1937, 1940; Scrimshaw, 1944), las hembras de especies que superfetan tendrán una placenta más compleja (en su ultraestructura) comparada con la placenta de especies que no superfetan.

b) Debido a que la matrotrofía implica la transferencia activa de nutrientes por parte de la madre hacia sus embriones a lo largo de todo el desarrollo (Wourms, 1981; Marsh-Matthews, 2011), las hembras de especies con altos niveles de matrotrofía tendrán una placenta más compleja para cumplir esta función en comparación con las hembras de especies con lecitotrofía o matrotrofía incipiente.

Predicción:

Las placentas de hembras de especies con superfetación y altos niveles de matrotrofía, mostrarán características ultraestructurales que reflejen una mayor transferencia activa de nutrientes a través del epitelio folicular, en comparación con especies sin superfetación y bajos niveles de matrotrofía. Particularmente se espera observar mayor área y mayor grosor del folículo materno, presencia de microvellosidades y mayor cantidad de vesículas en las células foliculares. Estas características le confieren a la placenta mayor capacidad de eficiencia en el transporte de nutrientes desde la madre hacia sus embriones (Grove y Wourms, 1994).

Descripción general de los métodos:

Colecté tres hembras gestantes de especies superfetadoras y no superfetadoras de la familia Poeciliidae. Inmediatamente, se fijaron las muestras utilizando el protocolo específico para microscopía electrónica (Dykstra, 1993). Los métodos de microscopia electrónica permitieron observar y describir detalladamente la ultraestructura de los componentes maternos de la placenta folicular de todas las especies. Para cada especie, cuantifiqué algunas ultraestructuras placentarias que indican el grado de complejidad de la placenta folicular (mayor complejidad es debido a mayor valor de las ultraestructuras placentarias; Grove y Wourms, 1994). Mediante métodos filogenéticos comparativos realicé una comparación morfológica de las placentas foliculares entre las especies que superfetan y las que no superfetan. Además, también examiné mediante métodos filogenéticos comparativos la posible relación entre el nivel de matrotrofia de cada especie y el grado de complejidad de sus placentas.

Capítulo 2

¿Existen diferencias entre especies que superfetan y no superfetan en la capacidad de almacenar esperma?

Hipótesis:

Las hembras superfetadoras requieren fertilizar frecuentemente a distintos grupos de embriones, entonces las hembras de las especies que superfetan deberían tener

mayor capacidad de almacenar esperma en comparación con hembras de especies que no superfetan.

Predicción:

Las hembras de especies que superfetan deben tener más y más grandes (mayor volumen) estructuras para almacenar esperma dentro del ovario en comparación con hembras de especies que no superfetan.

Descripción general de los métodos:

Colecté tres hembras gestantes de especies superfetadoras y no superfetadoras de la familia Poeciliidae. Inmediatamente, se fijaron las muestras utilizando un protocolo histológico (Presnell *et al.*, 1997). A través de cortes seriados y de la técnica de tinción histológica con hematoxilina-eosina examiné la morfología general del ovario tanto de especies superfetadoras como de especies no superfetadoras. En los ovarios de algunas especies observé estructuras que tienen la función de almacenar espermatozoides (llamadas espermatecas). Por medio de métodos estereológicos cuantifiqué el número, volumen total y volumen promedio de las espermatecas para cada especie. Finalmente, utilicé métodos filogenéticos comparativos para analizar si existe una relación entre las características de las espermatecas (número, volumen total y volumen promedio) y la presencia o ausencia de superfetación.

Capítulo 3

¿La presencia conjunta de superfetación y matrotrofia promueven la producción de crías más grandes en especies de la familia Poeciliidae?

Hipótesis:

La evolución conjunta de la superfetación y la matrotrofia en la familia Poeciliidae ha facilitado la evolución de crías más grandes.

Predicciones:

a) Las hembras de especies que presentan superfetación y altos niveles de matrotrofia tienen crías al nacer más grandes en comparación con hembras de especies que carecen de superfetación y que exhiben lecitotrofia o matrotrofia incipiente.

b) El tamaño de las crías al nacer ha evolucionado de manera correlacionada con el grado de superfetación y matrotrofia.

Descripción general de los métodos:

Se realizó una búsqueda bibliográfica del tamaño (masa) en el último estadio de desarrollo de las crías de distintas especies de la familia Poeciliidae. Además, se disecaron otros especímenes provenientes de la Colección Nacional de Peces (Instituto de Biología, UNAM) para medir directamente el tamaño promedio del último estadio de desarrollo para otras especies adicionales. A través de métodos filogenéticos comparativos, analicé si existe una relación entre el tamaño al último estadio de desarrollo y el nivel de

superfetación y matrotrofia. También estimé la heredabilidad filogenética de estas tres características para conocer si evolucionaron aproximadamente a la misma tasa, lo que indicaría cambios similares y probablemente paralelos en la superfetación, la matrotrofia y el tamaño al nacer, a lo largo de la historia evolutiva de esta familia de peces vivíparos.

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LEYENDAS DE FIGURA

Fig. 1. Gestación intrafolicular de la familia Poeciliidae (*Poeciliopsis gracilis*). **A.** Se muestra el ovario tipo sacular con dos embriones en desarrollo. Las flechas indican el epitelio folicular que rodea a los embriones. (L) lumen del ovario, (V) vitelo embrionario, (E) embrión. **B.** Amplificación de un embrión en desarrollo. **C.** Amplificación del folículo materno que forma parte de la placenta folicular. Se muestra el folículo materno dividido en dos secciones, en la capa o epitelio folicular (CF) y el tejido conjuntivo (TC). Las líneas indican el grosor de cada sección.

Fig. 2. Mapa de los sitios de estudio de las especies en estudio de la familia Poeciliidae.

Tabla 1. Resumen de los datos y sitios de estudio de las 12 especies de la familia Poeciliidae.

ESPECIES	Coordenadas geográficas	Estado	Superfetación
<i>Gambusia panuco</i>	N 20° 41' W 103° 57'	San Luis Potosí	NO
<i>Poecilia butleri</i>	N 20° 9' W 103° 2'	Jalisco	NO
<i>Poecilia mexicana</i>	N 17° 26' W 95° 26'	Oaxaca	NO
<i>Priapella intermedia</i>	N 17° 9' W 95° 10'	Oaxaca	NO
<i>Pseudoxiphophorus bimaculatus</i>	N 17° 8' W 95° 7'	Oaxaca	NO
<i>Xiphophorus hellerii</i>	N 20° 34' W 104° 9'	Jalisco	NO
<i>Heterandria formosa</i>	Acuario privado	Estados Unidos	SI
<i>Poeciliopsis infans</i>	N 20° 34' W 104° 9'	Jalisco	SI
<i>Poeciliopsis gracilis</i>	N 21° 59' W 99° 15'	San Luis Potosí	SI
<i>Poeciliopsis presidionis</i>	N 22° 29' W 105° 21'	Nayarit	SI
<i>Poeciliopsis prolifica</i>	N 23° 03' W 105° 50'	Sinaloa	SI
<i>Poeciliopsis viriosa</i>	N 21° 02' W 104° 22'	Sinaloa	SI

Fig. 1.

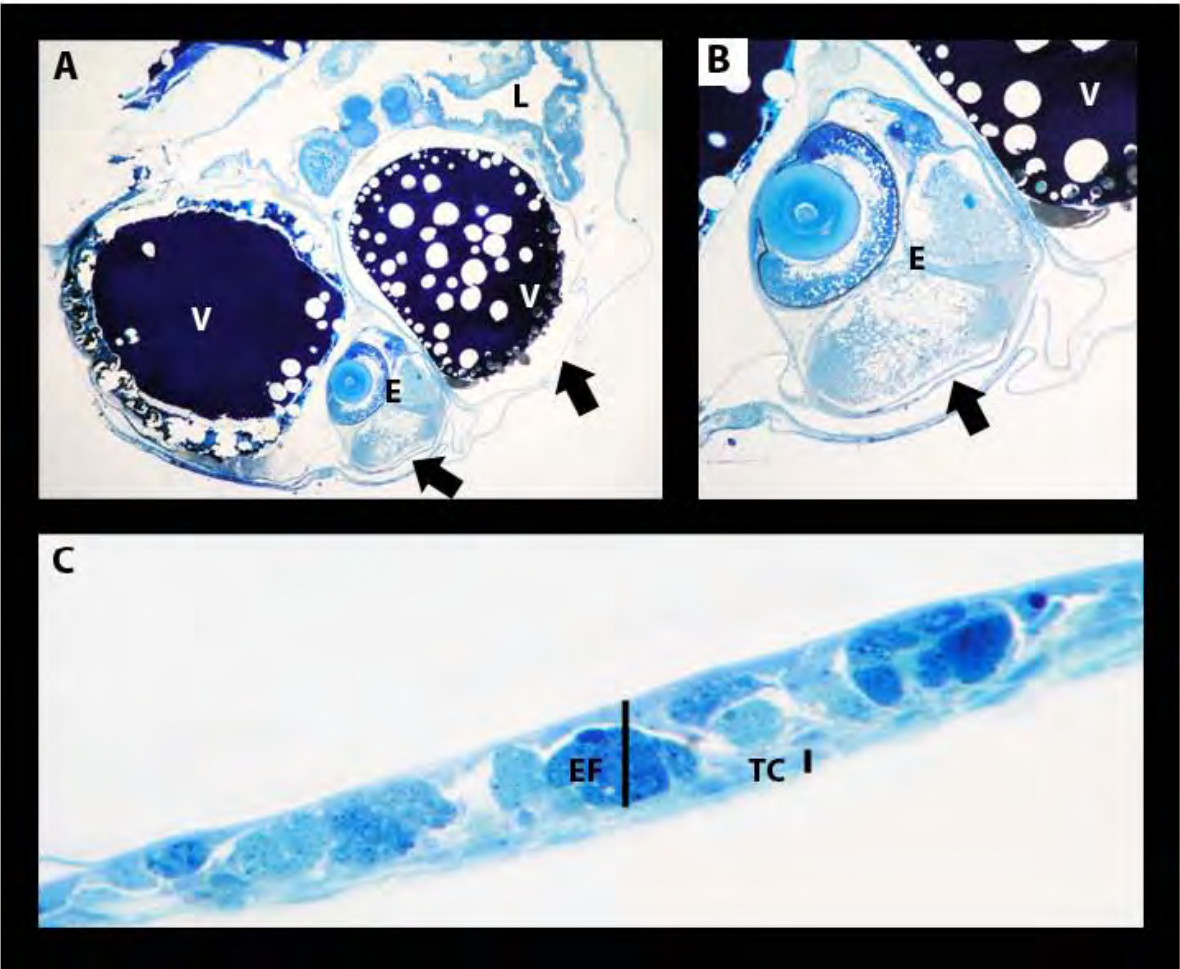
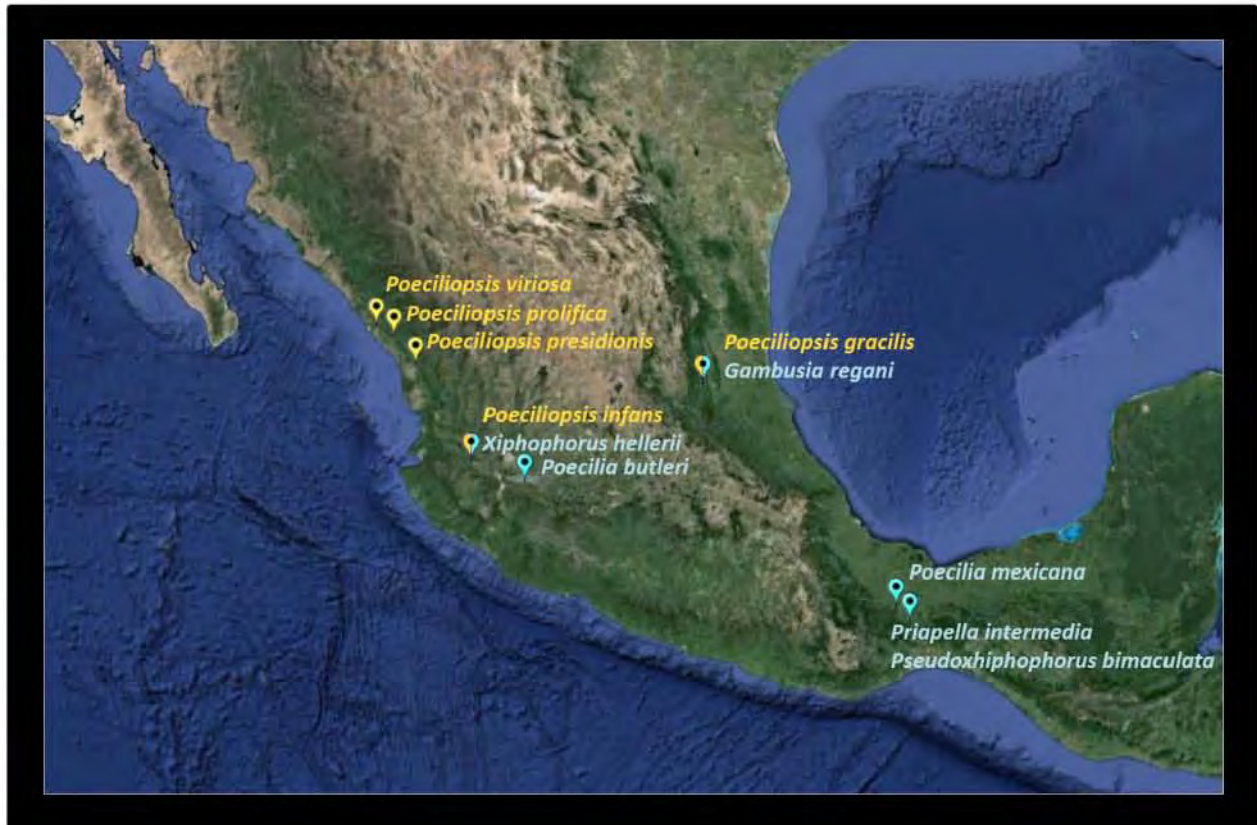


Fig. 2.



CAPÍTULO I

**Placental structures and their association with
matrotrophy and superfetation in poeciliid fishes**

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Maricela Villagrán-Santa Cruz y J. Jaime Zúñiga-Vega**

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**Placental structures and their association with matrotrophy and superfetation in
poeciliid fishes**

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Header: Degrees of placentation in poeciliids

Keywords: Matrotrophy; Poeciliidae; placentation; follicular placenta; superfetation

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20 **ABSTRACT**

21 Fish of the family Poeciliidae have a follicular placenta, the ultrastructure of which is little
22 known. In this article, we describe ultrastructure of the maternal follicular placenta of 11
23 species of poeciliids using electron microscopy. In addition, using stereological methods
24 and photomicrographs of placentas under electron microscopy, we examined and
25 quantified six ultrastructure characteristics that reflect the degree of complexity in
26 follicular placenta of each species. These characteristics included the number of vesicles,
27 vesicles area, number of microvilli, microvilli length, thickness of maternal follicle, and
28 follicular area. Additionally, using phylogenetic comparative methods, we evaluated the
29 relationship between the degree of matrotrophy and placental characteristics. We also
30 analyzed the possible relationship between the existence or lack of superfetation and
31 different placental characteristics. We found that all species had a similar morphological
32 structure; an inner follicular layer of epithelial nature as well as an outer follicular layer of
33 connective nature. In particular, *Poeciliopsis presidionis* presents *hyper-placentation*, or
34 more developed placental characteristics (thickness of maternal follicle and follicular area)
35 for the exchange of nutrients between the mother and embryos. We found a positive
36 relationship between the degree of matrotrophy and some placental characteristics, such
37 as follicular area, number of microvilli, and number and size of vesicles. Similarly, follicular
38 area and number and length of microvilli tended to be larger in species with superfetation
39 than those without superfetation. As such, we conclude that high degrees of matrotrophy
40 and superfetation are associated with placental characteristics that increase the efficiency
41 of nutrient transfer between mother and embryos.

42 INTRODUCTION

43 During the evolutionary transition from oviparity to viviparity, the female reproductive
44 system had to develop complex structures for the transfer of nutrients to the embryo
45 (Blackburn, 1992). As such, distinct placental types have evolved over time (Pollux et al.
46 2009; Pollux et al. 2014). The placenta is the close apposition or fusion of maternal and
47 fetal tissues for physiological exchange (Mossman, 1937). These tissues maximize the
48 transfer of nutrients to the embryo. For examples, in lizards and snakes, some species
49 have simple placentas, while others species (such as in the genera *Mabuya* and *Virginia*)
50 have complex placentas which allow the rapid transfer of nutrients (Blackburn and
51 Steward, 2011). It is therefore important to understand how different degrees of
52 placentation are related to other morphological, physiological, and reproductive
53 characteristics.

54 In fishes, viviparity implied that eggs had internal fertilization and were retained within
55 female reproductive system (Amoroso, 1968). The fishes lack Müllerian duct, in
56 consequence in the ovary occurs both fertilization and gestation (physiologically
57 equivalent to the mammalian uterus; Amoroso, 1968; Turner 1947; Wourms 1981). Then,
58 different types of placentas for the transfer of substances (nutrients, gases and waste)
59 were developed for the maintenance of the embryo within the ovary (Amoroso, 1968;
60 Blackburn, 2005, 2015). In particular, in the viviparous fish of the family Poeciliidae, the
61 placenta is follicular and was first described by Turner (1940). The main components of
62 the follicular placenta are the pericardial sac, the yolk sac, and the maternal follicle. The
63 yolk sac and pericardial sac are embryonic tissues which are intimately related to the

64 maternal follicular cells. This close relationship between maternal and embryonic tissues
65 allows the optimal exchange of nutrients and oxygen for the development of the embryo.
66 Additionally, Turner (1940) observed that the follicular cells had microvilli whose shape
67 varied among the species he analyzed (i.e. *Poeciliopsis sp.* and *Aulophallus elongatus*;
68 Turner, 1940). Microvilli are prolongations of the membrane of the follicular cells which
69 increase the surface area in contact with the embryo. Therefore, the increase in microvilli
70 surface apparently increases the efficiency of the exchange of substances between
71 mother and embryo. Likewise, Turner (1940) thus first inferred that the microvilli of the
72 follicular placenta could differ among species in accordance with the amount of nutrients
73 that the embryos receive directly from the mother during development.

74 Later, Grove and Wourms (1994) described the ultrastructure (characteristics
75 observed under an electron microscope) of the components of the follicular placenta of
76 *Heterandria formosa*. They demonstrated that the follicular cells had microvilli on their
77 apical surface (the zone closest to the embryonic tissue), a complex system of vesicles,
78 extensive endoplasmic reticulum and Golgi apparatus with 4 or 5 cisternae. As such, they
79 concluded that all of these structures are necessary for the exchange of nutrients of the
80 mother to the embryo. Then, in *Lebistes reticulatus (Poecilia reticulata)* it was observed
81 that the maternal follicular placenta consists of two layers: an inner layer and an outer
82 layer (Jollie and Jollie, 1964). The inner layer is conformed of epithelial follicular cells with
83 numerous vesicles, which confers a great activity of nutrients transport. In other words,
84 the vesicles permit the exchange and efficient movement of nutrients of the mother to
85 the embryo through the follicular placenta. In contrast, the outer follicular layer is

86 conformed of lax connective tissue with high vascularization, fibroblasts, collagen fibers
87 (Jollie and Jollie, 1964). Therefore, outer follicular layer has a protective and nutritive
88 function, because the capillaries are very close with follicular cells, suggesting an intense
89 transport of oxygen and nutrients to the embryo.

90 In addition to the variation in degree of placentation that can be found between
91 species in this family of fish, the amount of nutrient transfer through the placenta that
92 embryos experience over the course of their development varies widely among poeciliid
93 species (Pollux et al. 2009, 2014). There are lecithotrophic species, in which nutrient
94 transfer from mother to embryo occurs before the embryo is fertilized (Wourms, 1981;
95 Marsh-Matthews, 2011). In these lecithotrophic species, the energy reserve devoted to
96 embryonic development is stored in a nutritious substance known as yolk that is
97 deposited into the mature oocyte. In contrast, some species are matrotrophic, such that
98 most of the energy devoted to embryonic development is provided after fertilization and
99 is transferred directly from mother to embryo through the placenta during embryonic
100 development (Wourms, 1981; Marsh-Matthews, 2011). Therefore, embryos that are
101 highly matrotrophic (extensive matrotrophy) have little yolk and are characterized by a
102 large increase in weight during their development, since the mother is actively
103 transferring nutrients to them (Pires et al. 2010). Species with moderate matrotrophy are
104 those in which the mother deposits some yolk in the mature oocyte as well as transferring
105 some nutrients directly during embryonic development. In the Poeciliidae family, there is
106 therefore a continuum among species from completely lecithotrophic to extensively
107 matrotrophic (Pollux et al. 2014).

108 Recently, Kwan et al. (2015) demonstrated that among eight species of poeciliids,
109 the thickness of maternal follicle and vascularization and folding of the follicular
110 epithelium increased with increasing degrees of matrotrophy. To date, this is the only
111 study to examine the relationship between degree of matrotrophy and placental
112 complexity, even though many studies have assumed that higher degrees of matrotrophy
113 require more complex placentas (Reznick et al. 2002; Pollux et al. 2014; Ostrovsky et al.
114 2015). More comparative studies with larger numbers of species and which account for
115 phylogenetic relationships are still needed. As such, one of the main objectives of this
116 study was to analyze whether there is a relationship between degree of matrotrophy and
117 several ultrastructural characteristics of the maternal follicular placenta. The hypothesis is
118 that females that transfer more nutrients to their embryos during development (i.e. more
119 matrotrophic females) require a more complex placenta (Turner, 1940; Jollie and Jollie,
120 1964; Grove and Wourms, 1994; Kwan et al. 2015). Some of the characteristics that we
121 suggest could indicate a more highly developed placenta are the presence of numerous
122 and long microvilli of follicular cells, higher number and area of vesicles of follicular cells,
123 higher maternal follicle thickness, and higher follicular area.

124 On the other hand, superfetation is the capacity for females to
125 simultaneously carry within the reproductive tract litters of embryos at different
126 developmental stages (Turner, 1937, 1940; Scrimshaw, 1994). In poeciliids, the species of
127 some genera (e.g., *Poeciliopsis*, *Heterandria* and *Neoheterandria*) exhibit superfetation,
128 while those of other genera do not (e.g., *Alfaro*, *Brachyrhaphis* and *Xiphophorus*; Reznick
129 and Miles, 1989; Zúñiga-Vega et al. 2010). The phylogenetic distribution of this

130 reproductive strategy suggests that superfetation has developed independently on
131 multiple occasions (Olivera-Tlahuel et al. 2015). Because superfetation involves the
132 differential development of distinct groups of embryos, we propose the hypothesis that
133 the evolution of this reproductive strategy should be accompanied by the evolution of
134 more specialized placentas which facilitate the differential regulation of nutrients to the
135 distinct developmental stages. As such, the second objective of this study was to analyze
136 whether there is a relationship between degree of placentation and presence or absence
137 of superfetation.

138 We collected three visibly gestating females of superfetating and non-
139 superfetating poeciliid species in different rivers in Mexico. The superfetating species
140 were *Heterandria formosa*, *Poeciliopsis infans*, *P. gracilis*, *P. presidionis*, *P. prolifica* and *P.*
141 *viriosa* (table 1), and the non-superfetating species were *Pseudoxiphophorus bimaculatus*,
142 *Xiphophorus hellerii*, *Poecilia mexicana*, *Priapella intermedia* and *Gambusia panuco* (table
143 1). Each specimen was then anesthetized (using 3-aminobenzoic acid ethyl ester) and
144 immediately sacrificed by an overdose of the same anesthetic. The ovaries were dissected
145 from the collected females and immediately injected with Karnovsky fixative and were
146 placed by immersion on the same fixative for later processing in the laboratory.

147 **Electron microscopy**

148 In the laboratory, the three ovary samples of each species were processed as follows.
149 First, they were placed in 0.1 M sodium cacodylate buffer (0.1 M) for one night, then were
150 transferred to 1% osmium tetroxide for one hour. Second, the ovaries were dehydrated in

151 gradually increasing concentrations of alcohol (70 %, 80 %, 90 %, 96 % and 100 %). Third,
152 the processing was continued with acetonitrile for 20 minutes repeated at two different
153 proportions (1:1 acetonitrile:Epon and 1:2 acetonitrile:Epon). Fourth, the samples were
154 placed in increasing proportions of Epon until they were immersed in pure Epon. Finally,
155 the samples were placed in plastic molds and allowed to polymerize in a 60°C oven for 24
156 hours.

157 The block obtained and included in Epon was cut into 2-micron thick slices, a semi-
158 fine thickness. We used a Nova LKN BROMMA ultramicrotome to obtain serial sections of
159 each of the ovaries. They were then stained with toluidine blue and using light
160 microscopy, the follicular cells of ovarian follicles that contained an embryo in a relatively
161 advanced developmental stage (between stages 8 and 9; Haynes, 1995) were located.
162 Once follicular cells were located in each ovary, semi-fine sections were obtained. The
163 samples were then moved to the Instituto de Fisiología Celular at the Universidad
164 Nacional Autónoma de México (UNAM) and the Instituto de Pediatría to obtain fine (less
165 than 1 micron thick) sections.

166 The fine sections were observed using the electron microscopes at the Instituto de
167 Investigaciones Biomédicas and Instituto de Fisiología Celular at UNAM. Digital
168 photomicrographs were taken of the follicular cells of the different species included in the
169 study. Only 10 maternal follicles (one follicular cell and connective tissue) were randomly
170 selected from each ovary for posterior analysis, such that 30 photographs of maternal
171 follicles were obtained per species, since three ovaries were processed per species.

172 **Standardization of developmental stage**

173 During the course of embryonic development, follicular cells change in their
174 components as well as their morphology (Jollie and Jollie, 1964; Guerrero-Estévez, 2005).
175 At early stages, follicular cells are cuboidal and with low subjacent vascularization. On the
176 contrary, at intermediate and at the beginning of late developmental stages, follicular cells
177 are thinner and subjacent vascularization is high. At the most advanced stages of
178 development, the follicular cells are extremely thin and their association with
179 vascularization is limited. In order to avoid an erroneous comparative analysis, we
180 standardized the stage of embryonic development by selecting only embryos at the
181 beginning late developmental stages and close to the eye for the examination of maternal
182 follicle morphology (between stages 8 and 9; Haynes, 1995). In this way, we assured that
183 the differences among species reflected real morphological differences, and were not due
184 simply to chance differences in the stage of embryonic development.

185 **Stereology of the follicular epithelium**

186 The placenta is composed of both maternal and fetal tissue. However, in this article we
187 focus on maternal tissues (or maternal follicles). We describe the ultrastructure of the
188 maternal follicle and we consider the differences of these substructures (number and size)
189 as indicators of degree of complexity. For this, we used stereology, is a combination of
190 simple and efficient methods for quantifying morphological structures using histological
191 sections (Gundersen et al. 1988). However, stereology can also be applied to images
192 obtained from electron microscopy. Using stereological methods it is possible to count,

193 measure, and calculate the volume of structures of interest. In this study we used the
194 program ImageJ (Rasband, 1997-2016), which uses stereological tools to analyze
195 photomicrographs taken using electron microscopy. In particular, we quantified the
196 following characteristics of maternal follicle (one follicular cell and connective tissue): (1)
197 number of vesicles, NV; (2) area of vesicles, AV; (3) number of microvilli, NM; (4) length of
198 microvilli, LM; (5) thickness of maternal follicle, TF; (6) and area of maternal follicle, AF.
199 We considered these six variables indicators of the degree of complexity of the placenta.
200 In order to obtain a representative value for each of these variables for each species, we
201 averaged the values of the three ovaries analyzed per species.

202 **Matrotrophy index**

203 The variations between species of the family Poeciliidae in the amount of nutrients
204 transferred to embryos over the course of development (lecithotrophy to extensive
205 matrotrophy) can be measured using the matrotrophy index (MI; Worms et al. 1988). This
206 index represents the difference in dry mass between fertilization and the last stage of
207 embryonic development. If the embryo weighs 30-35% less (due to metabolic costs) at the
208 end of development than at oocyte fertilization, this indicates lecithotrophy (negative MI;
209 Worms et al. 1988). While, if the embryo at the last stage of development is of a similar
210 weight to that at oocyte fertilization, it has incipient matrotrophy (MI=0.6 to 0.7; Reznick
211 et al. 2002). In other words, the energy necessary for development was obtained mostly
212 from the yolk provided by the mother before oocyte fertilization. On the other hand, in a
213 moderately matrotrophic species, the embryo's mass increases with progressive
214 developmental stages ($0.8 < MI > 2$; Reznick et al. 2002), and the embryo obtains nearly all

215 its maternal provisioning during development through a placenta. The embryo therefore
216 weighs more at the last stage of development than at oocyte fertilization. Extensive
217 matrotrophy refers to species which completely lack yolk, such that all of the maternal
218 provisioning occurs during embryonic development and weight at birth may be up to five
219 times more than at oocyte fertilization (MI > 5 Reznick et al. 2002). There are extreme
220 cases, such as in *Poecilia branneri* and *Poecilia bifura*, where the embryo at the last stage
221 of development may weigh five times more than at fertilization (Pires et al. 2010). In each
222 of the study species we obtained data on the matrotrophy index from previous studies
223 (i.e. Pollux et al. 2014; Olivera-Tlahuel et al. 2015).

224 **Estimation of the relationship between placental complexity, matrotrophy, and** 225 **superfetation**

226 Prior to comparative analyses, all variables, including matrotrophy index, were log-
227 transformed to normalize the data. In addition, to explore whether maternal follicle size
228 affected the magnitude of the morphological characters we studied, area of maternal
229 follicle (AF) was included as an explanatory variable in linear regressions, in which each of
230 the other five ultrastructural characteristics (NV, AV, NM, LM and TF) was used as the
231 response variable. In the cases in which area of maternal follicle had a significant effect,
232 the residuals of these linear regressions were used in subsequent comparative analyses.

233 We performed two separate sets of analyses. In the first set, we examined the
234 relationship between matrotrophy index (MI) and our six morphological variables
235 indicating placental complexity (NV, AV, NM, LM, TF and AF). In the second set, we looked

236 for differences between species with and without superfetation in the same six variables.
237 In all cases we used two different evolutionary assumptions (i.e. Brownian motion, BM;
238 Phylogenetic Generalized Least Squares, PGLS).

239 First, we assumed that the traits evolved by accumulating gradual changes and
240 following a model of Brownian motion (BM), such that the phylogenetic relationships
241 between species can explain a large part of the variation observed in the characteristics
242 (Felsenstein, 1985). In the first set of analyses, we implemented this evolutionary model
243 using Phylogenetic Generalized Least Squares (PGLS; Martins and Hansen, 1997), treating
244 the matrotrophy index (MI) as a continuous independent variable and each of our six
245 characteristics of interest as a response variable. In the second analysis, we also
246 implemented a Brownian motion PGLS model to compare the six characteristics of interest
247 between species with and without superfetation, treating the presence/absence of
248 superfetation as a binary independent variable. PGLS is a regression technique which
249 incorporates phylogenetic information into the error term of the model (Martins and
250 Hansen, 1997).

251 In order to implement these comparative phylogenetic analyses, we used the
252 family Poeciliidae phylogeny published by Pollux et al. (2014), trimmed to our species of
253 interest. We created an ultrametric tree by transforming the original branch lengths to a
254 scale based on relative time (Fig. 1). To do this transformation we used a semiparametric
255 method based on penalized likelihood (Sanderson, 2002) implemented in the ‘ape’
256 package of the R program (Paradis et al. 2004; R Core Team, 2016).

257 Second, we assumed that traits evolved rapidly in the time and immediately
258 adapted to the local environment with no trace of their phylogenetic history. Under this
259 evolutionary assumption, phylogeny is unimportant, so we used Ordinary Least Squares
260 regression (OLS). For the first set of data, we used this approach to look for a possible
261 relationship between matrotrophy index (a continuous explanatory variable) and the six
262 morphological characteristics of interest (dependent variables). For the second set of
263 data, we used OLS to compare the different morphological characteristics between
264 species with and without superfetation (binary explanatory variable; Garland et al. 2005;
265 Lavin et al. 2008). We fit these PGLS and OLS regression models using the ‘ape’ and ‘nlme’
266 packages in R (Paradis et al. 2004; Pinheiro et al. 2016).

267 We used the Akaike Information Criterion (AIC) adjusted for small sample size (AICc;
268 Burnham y Anderson, 2002) to compare the fit of the PGLS-BM and OLS models to the
269 data for each of the morphological characteristics. The smallest AICc value indicates the
270 model with the best fit, and a difference in AICc values between models greater than two
271 units indicates a real difference in their capacity to explain the variation in the data.

272

273 **RESULTS**

274 **Morphological description of maternal follicle**

275 The follicular placenta in nearly all of the species was divided into two main layers,
276 the inner and outer follicular layers. The inner follicular layer is the closest to the

277 embryonic placental tissues. This layer is structured by follicular epithelium with their
278 ovoid nucleus and organelles. Ultrastructurally is possible observe the density and size of
279 microvilli toward the embryonic part, and inside of follicular cells their nuclei, vesicles,
280 mitochondria and lipid droplets. We observed vesicles distributed randomly in the
281 follicular cells. Externally and in close contact with this inner follicular layer is found the
282 external follicular layer. In contrast, the outer follicular layer is composed mainly of blood
283 capillaries immersed in an extracellular matrix with collagen fibers. Presumably, the
284 function of this layer is to protect and nourish follicular cells and the embryo. In the
285 placentas of nearly all of the species we identified specific structures for nutrient
286 transport and cellular communication. In particular, in the follicular cells we found
287 prolongations of the membrane, or microvilli, oriented toward the embryo (Fig. 3A and
288 3C). In this same cells we also found a higher number of vesicles, mitochondria, and the
289 presence of cell junctions (i.e. desmosomes; 3A-C). All of these characteristics apparently
290 confer great nutrient exchange (Fig. 3B and 2C). In contrast, we found a large number of
291 blood capillaries in the external layer in close relation with follicular cells, which suggests a
292 high potential for gases and nutrients exchange (Fig. 3D).

293 We also observed the interaction between maternal and embryonic (pericardial
294 sac, yolk sac and tissue embryo) placental tissues (*Poeciliopsis prolifica*, MI=5.4; *Priapella*
295 *intermedia*, MI=0.6; Fig. 3E). However, both the follicular cells of the mother and the
296 embryonic tissue (pericardial sac) present microvilli, suggesting close physiological
297 communication. Specifically in *Poeciliopsis prolifica* and *Priapella intermedia*, we could
298 observe a large and uniform number of microvilli in the embryonic tissue (pericardial sac).

299 In fact, in these two species the microvilli are of different shape (Fig. 3E and 3F). In
300 addition, in these species we observed a large number of blood capillaries that
301 presumably transport oxygen and nutrients to embryo. This network of blood capillaries
302 forms part of the embryonic pericardial sac.

303 In non-superfetating species (*Gambusia panuco*, MI=0.79; *Pseudoxiphophorus*
304 *bimaculatus*, MI=0.65; *Poecilia mexicana*, MI=0.63; *Priapella intermedia*, MI=0.6;
305 *Xiphophorus hellerii*, MI=0.61), like in superfetating species (*Heterandria formosa*, MI=35;
306 *Poeciliopsis gracilis*, MI=0.84; *P. infans*, MI= 1.05; *P. prolifica*, MI=5.4; *P. viriosa*, MI=0.93),
307 the follicular placenta consists of an inner and an outer follicular layer. In almost all
308 species (except in two superfetating species: *Poeciliopsis presidionis* MI= 21.5 and *P.*
309 *prolifera* MI=5.4), on the embryo side there is a layer without an apparent cellular
310 structure called the follicular “ribbon” (described previously by Kwan et al. 2015). The
311 follicular placentas of non-superfetating species were similar (Fig. 4A-E). In the maternal
312 side, the follicular cells of inner follicular layer presented various types of cell junctions,
313 nucleus, vesicles, mitochondria, and other organelles (Fig. 4A-E). In the outer follicular
314 layer, we observed collagen fibers, fibroblasts, and blood capillaries. Only some of the
315 species without superfetation presented microvilli of follicular cells (*Poecilia mexicana*,
316 MI=0.63; *Priapella intermedia*, MI=0.6; *Xiphophorus hellerii*, MI=0.61). On average the
317 non-superfetating species had 1.3 to 2 microvilli per cell, which is lower than in
318 superfetating species (on average 7 microvilli per follicular cell). In contrast, non-
319 superfetating species as *Gambusia panuco* (MI=0.79) and *Pseudoxiphophorus bimaculatus*
320 (MI=0.65) did not present microvilli, which could suggest less nutrient transfer from

321 mother to embryos. In general non-superfetating species had a lower follicular cell
322 thickness (6.46 μm) than superfetating species (9.87 μm ; not statistically significant).

323 All of the superfetating species *Heterandria formosa*, MI=35; *Poeciliopsis gracilis*,
324 MI=0.84; *P. infans*, MI= 1.05; *P. prolifica*, MI=5.4; *P. viriosa*, MI=0.93; *P. presidionis*,
325 MI=21.5) had follicular placentas with similar morphological characteristics (Fig. 5-6). Like
326 non-superfetating species, the follicular placenta of all superfetating species was
327 composed of two layers (inner and outer follicular layers). In five of the six superfetating
328 species, the inner follicular layer is formed by flat follicular epithelium with the nucleus,
329 organelles, numerous vesicles, and microvilli (in the zone closest to the embryo; Fig. 5D-E).
330 In the outer layer we found various arrangements of fibers and blood capillaries (Fig. 5A-
331 E). In superfetating species the thickness of maternal follicle was greater than non-
332 superfetating species, but only was a trend (average of 9.87 μm and 6.46 μm ,
333 respectively).

334 Unlike in all of the other (superfetating and non-superfetating) species, in
335 *Poeciliopsis presidionis* (a superfetating specie; MI=21.5) the follicular placenta is arranged
336 differently (Fig. 6). In this species, the epithelium of the inner follicular layer folds, forming
337 peaks and valleys (Fig. 6A). We found that the epithelium has columnar cells and blood
338 capillaries of outer follicular layer protrude towards the epithelium especially on the peaks
339 (Fig. 6A). In contrast, the outer follicular layer maintains its lax connective tissue
340 configuration and is composed of with high vascularization, fibroblasts, collagen fibers
341 organized in several layers (Fig. 6A). In the inner follicular layer, the follicular cells have
342 many vesicles, mitochondria and other organelles. In addition, the inner epithelium

343 presents fat droplets which stained strongly with uranyl (Fig. 6B-C). The thickness of
344 maternal follicle is clearly larger (average of 18.5 μm) than the other superfetating species
345 (average of 9.87 μm). Due to the large differences in the characteristics of the follicular
346 placenta in *P. presidionis* (MI=21.5) compared to the other species examined, we did not
347 include this species in the statistical comparisons between superfetating and non-
348 superfetating species.

349 **Relationship between degree of placentation, superfetation, and matrotrophy**

350 The electron microscopy and stereological methods we used allowed the clear
351 quantification of six follicular placenta characteristics (number of vesicles, NV; area of
352 vesicles, AV; number of microvilli, NM; length of microvilli, LM; thickness of maternal
353 follicle, TF; area of maternal follicle, AF; table 1). In the regressions in which we used AF as
354 an explanatory variable to examine the effect of maternal follicle on the other
355 morphological characteristics, NM and TF yielded significant results ($P < 0.05$ in both
356 cases), so the residuals from these regressions were used in subsequent analyses of these
357 two variables.

358 According to the models with the best fit, the number of vesicles (NV) and the area
359 of vesicles (AV) had a positive trend with the matrotrophy index (PGLS-BM model for NV:
360 slope = 0.05, $P = 0.098$; OLS model for AV: slope = 0.20, $P = 0.098$; table 2, Fig. 7). In the
361 particular case of the number of microvilli (NM), the best-fitting model (OLS) indicated
362 that the relationship between this variable and the matrotrophy index was positive but
363 not significant (table 2). However, assuming Brownian motion, this positive relationship

364 was significant (slope = 0.17, $P = 0.052$; Fig. 7). Under neither of the evolutionary
365 assumptions was there a statistical relationship between LM or TF and matrotrophy index
366 (table 2, Fig. 7). According to the OLS model (which had the best fit for this variable) the
367 area of maternal follicle (AF) had a significant positive relationship with the matrotrophy
368 index (slope = 0.16, $P = 0.006$, table 2; Fig. 7). This indicates that as a specie's matrotrophy
369 index increases, so does its area of maternal follicle.

370 On the other hand, comparing the morphological ultrastructures of the placenta
371 between superfetating (*Heterandria formosa*, *Poeciliopsis gracilis*, *P. infans*, *P. prolifica*, *P.*
372 *viriosa*) and non-superfetating species (*Gambusia panuco*, *Pseudoxiphophorus*
373 *bimaculatus*, *Poecilia mexicana*, *Priapella intermedia*, *Xiphophorus hellerii*), the number of
374 microvilli (NM) and area of maternal follicle (AF) were greater in species with
375 superfetation (species with superfetation, NM= 7, AF= 159,464. 84 μm^2 ; species without
376 superfetation NM= 1.3, AF= 66,833. 64 μm^2 ; Fig. 8) and in both cases the difference
377 between superfetating and non-superfetating species was marginally significant in the
378 best-supported models (NM: $P = 0.074$, AF: $P = 0.059$; table 3). In none of the other
379 morphological variables and under neither of the evolutionary assumptions was there a
380 significant difference between superfetating and non-superfetating species (table 3).
381 However, in figure 8 we show that, in general, superfetating species had a trend toward
382 positive values of most of the placental morphological characteristics we examined.

383

384

385 **DISCUSSION**

386 The follicular placenta of all species we analyzed, presented the same structures
387 described previously by Jollie and Jollie (1964) in *Poecilia reticulata*. Specifically, we refer
388 to the presence of an inner and an outer follicular layer. In the inner layer we identified
389 numerous vesicles, microvilli on the apical surface (zone closest to embryonic tissue), cell
390 junctions, and extensive endoplasmic reticulum. Grove and Wourms (1994) observed
391 these same structures in the follicular placenta of *Heterandria formosa*. As such, we can
392 conclude that the inner follicular layer in all species has a high degree of metabolic activity
393 and is responsible for the transfer of nutrients from the mother to the embryo. For its
394 part, the outer follicular layer is composed of lax connective tissue with high
395 vascularization (blood capillaries), fibroblasts, and collagen fibers (Jollie and Jollie, 1964).
396 As such, this outer layer appears to serve a protective function for the embryo and could
397 facilitate gas exchange. An additional characteristic we noticed was the presence of a
398 follicular ribbon or egg envelope (porous fertilization membrane), which was observed in
399 some species of the genus *Heterandria* and *Poeciliopsis* (*Heterandria formosa*, *Poeciliopsis*
400 *gracilis*, *P. latidens*, *P. occidentalis*, *P. turrubarensis* and *P. viriosa*; Grove and Wourms,
401 1994; Kwan et al. 2015). This structure is found between the inner follicular layer and the
402 absorbent tissue of the embryo. In our study the follicular ribbon was observed in almost
403 all species, except two species with superfetation and high matrotrophy index
404 (*Poeciliopsis presidionis* MI= 21.5 and *P. prolifica* MI=5.4). This agrees with Kwan et al.
405 (2015) that observed the follicular ribbon only in species with lecithotrophy and moderate

406 matrotroph (except in species with high matrotrophy index; *P. prolifica*, *P. turneri* y *P.*
407 *retropinna*).

408 On the other hand, in *Poeciliopsis presidionis*, an extensively matrotrophic species
409 (MI=21.5, Reznick et al. 2002), we observed hyper-placentation. This implies that the
410 placenta of this specie has a totally different arrangement than that observed in the other
411 ten species. This type of hyper-placentation consists of folds of the inner follicular layer
412 and is composed of columnar-type cells. We found that the outer follicular layer follows
413 the inner follicular layer fold (papillae-shaped) taking the capillaries close to the
414 epithelium. In addition, the outer follicular layer presents connective tissue with several
415 layers of collagen fibers. This type of placental morphology has been found in other
416 species with extensive matrotrophy, such as *P. turneri* and *P. retropinna*. (Kwan et al.
417 2015). A similar arrangement to the type of hyper-placentation observed in these poecillid
418 species has also been observed in the uterine epithelium of the omphalallantoic placenta
419 in snakes (*Virginia striatula*; Blackburn and Steward, 2011). The association between a
420 high matrotrophy index and follicular hyper-placentation indicates that the folding of the
421 inner follicular layer (formed by columnar cells) maximizes the transfer of nutrients from
422 the mother to the embryo due to a high secretory activity (Igarashi, 1961; Villagrán-Santa
423 et al., 2005; Blackburn and Steward, 2011).

424 Several studies have used the matrotrophy index as an indicator of the degree of
425 placentation (e.g. Reznick et al. 2002; Pollux et al. 2014; Ostrovsky et al. 2015). These
426 studies have assumed that the species with a high matrotrophy index (which is simply the

427 quotient of the embryo's weight at birth divided by the egg's initial weight at fertilization)
428 necessarily possess more specialized placentas. However, none of these studies used
429 morphological evidence to validate this assumption. Recently, Kwan et al. (2015) were the
430 first to demonstrate that, indeed, species with higher degrees of matrotrophy (i.e. higher
431 values of the matrotrophy index) presented thicker and more structurally complex
432 follicular placentas. In addition, they identified certain structures of the follicular placenta
433 in these matrotrophic species, such as microvilli, which facilitate the exchange of nutrients
434 between mother and embryo.

435 In this study, we present additional evidence that species with high degrees of
436 matrotrophy have more complex placentas (higher values of the ultrastructural
437 characteristics). In particular, we observed that area of maternal follicle, number of
438 microvilli, number of vesicles, and size of these vesicles tend to increase with the
439 matrotrophy index. These positive relationships indicate that species with elevated
440 degrees of matrotrophy possess more specialized placentas *Heterandria formosa*, MI = 35;
441 *Poeciliopsis prolifica*, MI = 5.4, which favor the active and continuous transfer of nutrients
442 from the mother to the embryos during their development. In other words, more and
443 larger vesicles permit the efficient exchange and movement of nutrients, such that the
444 follicular cells present are more metabolically active (high micropinocytotic activity; Jollie
445 and Jollie, 1964). In addition, the presence of microvilli indicates an increase in the surface
446 area of contact between the follicular placenta and the embryo. So, the microvilli
447 apparently increase the efficiency of the exchange of substances between the mother and
448 embryo (Turner, 1940).

449 It is important to note that the species with the highest values of the matrotrophy
450 index (i.e. *Heterandria formosa*, MI = 35; *Poeciliopsis prolifica*, MI = 5.4) also had the
451 largest number and size of vesicles, highest number of microvilli, and highest follicular
452 area. This suggests that the specialization or complexity of the placentas in the species
453 studied only occurs at a matrotrophy index larger than five (species with extensive
454 matrotrophy, MI > 5; Reznick et al. 2002).

455 Finally, we should emphasize that the relationship between superfetation and
456 placental complexity had not been previously examined in the family Poeciliidae, and our
457 study is the first to demonstrate that superfetating species have placental ultrastructures
458 which reflect a higher degree of placental complexity (high exchange of substances) than
459 non-superfetating species. On average, superfetating species (*Heterandria formosa*,
460 *Poeciliopsis gracilis*, *P. infans*, *P. prolifica* and *P. viriosa*) had higher area of maternal
461 follicle and more microvilli than non-superfetating species (*Heterandria formosa*,
462 *Poeciliopsis gracilis*, *Poeciliopsis infans*, *Poeciliopsis prolifica*, *Poeciliopsis viriosa*). In
463 addition, superfetating species also showed a trend (though non-significant) toward
464 higher values of number and size of vesicles and length of microvilli compared to non-
465 superfetating species. These results support (despite the statistical power of the sample
466 size, 10 species) the hypothesis that species with superfetation should have a more
467 complex placenta given the need to differentially regulate the development of embryos at
468 different developmental stages (Zúñiga-Vega et al. 2010). This exploration of placental
469 complexity and its relationship with different levels of matrotrophy and superfetation still
470 needs to be carried out in a greater number of species.

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480

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577

578 **FIGURE LEGENDS**

579 **Fig. 1.** Phylogeny of the ten species used in the study. Black rectangles indicate presence
580 of superfetation. Modified from Pollux et al. (2014).

581 **Fig. 2.** Follicular placenta of *Poeciliopsis infans* (superfetating). Shown are the inner (IFL)
582 and outer (OFL) follicular layers. The arrow indicates a blood capillaries. Nucleus of

583 follicular cell (NF). The white lines represent the thickness of each layer. The scale bar
584 shows 2 μm .

585 **Fig. 3. A.** Sections of follicular placenta of *Poeciliopsis infans* (superfetating). Shown are
586 vesicles (V) and microvilli (MV) dispersed within the cells of the inner follicular layer. Inner
587 follicular layer (IFL), outer follicular layer (OFL). **B.** Section of the inner follicular layer of
588 *Poeciliopsis viriosa* (superfetating). Shown are vesicles (V), mitochondria (M) and cell
589 junctions (CJ) of the follicular cell. **C.** Section of the inner follicular layer of *Heterandria*
590 *formosa* (superfetating). Shown are vesicles (V) and microvilli (MV) in the apical side of
591 follicular cell. **D.** Section of the follicular placenta of *Poeciliopsis infans* (superfetating).
592 Blood capillary is shown (indicated by arrow) in outer follicular layer by close contact with
593 inner follicular layer. There are two types of nuclei, nucleus of follicular cell (NF) and
594 nucleus of connective tissue (NC). **E.** Interaction between maternal (IFL) and embryonic
595 tissues (E) in *Poeciliopsis prolifica* (superfetating). A large number of microvilli (MV) are
596 evident, nucleus of follicular cell and abundant endoplasmic reticulum (ER) are shown. The
597 embryo epithelium of pericardial sac shows several and thin microvilli. **F.** Vascularized
598 embryonic tissue (pericardial sac) of *Priapella intermedia* (non-superfetating). A large
599 number of microvilli are shown (MV). Arrows indicate blood capillaries. The scale bar
600 shows 2 μm in panels A, D, E, F and 200 nm in panel B.

601 **Fig. 4.** Different follicular placentas in species without superfetation **A.** *Gambusia panuco*.
602 **B.** *Pseudoxiphophorus bimaculatus*. **C.** *Poecilia mexicana*. **D.** *Priapella intermedia*. **E.**
603 *Xiphophorus hellerii*. Inner follicular layer (IFL), outer follicular layer (OFL), nucleus of

604 follicular cells (NF),), nucleus of follicular cells (NC), nucleus of endothelial cells (NE),
605 vesicles (V), cell junctions (CJ) and follicular ribbon (R). The scale bar shows 2 μm .

606 **Fig. 5.** Different follicular placentas in species with superfetation **A.** *Heterandria formosa*.
607 **B.** *Poeciliopsis gracilis*. **C.** *Poeciliopsis infans*. **D.** *Poeciliopsis prolifica*. **E.** *Poeciliopsis viriosa*.
608 Inner follicular layer (IFL), outer follicular layer (OFL), nucleus of follicular cells (NF),
609 nucleus of endothelial cells (NE), vesicles (V), cell junctions (CJ) and follicular ribbon (R).
610 The scale bar shows 2 μm .

611 **Fig. 6.** Follicular hyper-placenta of *Poeciliopsis presidionis* (superfetating). The inner
612 follicular layer (IFL) is folded and its cells are type columnar. The outer follicular layer (OFL)
613 is composed of with high vascularization, fibroblasts, collagen fibers organized in several
614 layers. Large quantities of lipid droplets (LD) are seen, strongly stained with uranyl. The
615 follicular nuclei are columnar (FN) and the arrow indicates blood capillaries. Nucleus of
616 follicular cells (NF). The scale bar indicates 2 μm .

617 **Fig. 7.** Relationships between matrotrophy and different morphological characteristics of
618 the placentas of 10 species of poeciliid fish. The broken line indicates the relationship
619 assuming Brownian motion (i.e. accounting for phylogeny using phylogenetic generalized
620 least squares, PGLS-BM). The solid line shows the predicted relationship using ordinary
621 least squares (OLS), which does not account for phylogeny.

622 **Fig. 8.** Average of the values of different morphological characters of the placentas for
623 species without superfetation (gray bars) and with superfetation (black bars). Error bars
624 show one standard error.

TABLE 1. Summary of data and results of the morphological characteristics of placentas of 10 species from the Poeciliidae family. MI indicates the matrotrophic index. SF indicates the presence or absence of superfetation. NV indicates the average number of vesicles per follicular cell. AV represents the average area of these vesicles. NM indicates the mean number of microvilli per follicular cell. LM indicates the average length of these microvilli. TF indicates the mean thickness of maternal follicle. AF represents the average area of maternal follicle.

SPECIES	Geographic coordinates	MI	SF	NV	AV (μm^2)	NM	LM (μm)	TF (μm)	AF (μm^2)
<i>Gambusia panuco</i>	N 20° 41' W 103° 57'	0.79	NO	14.28	168.428	0	0	10.45	114228.75
<i>Pseudoxiphophorus bimaculatus</i>	N 17° 8' W 95° 7'	0.65	NO	21.07	38.237	0	0	5.64	62860.51
<i>Heterandria formosa</i>	Private aquarium	35	YES	48.83	535.205	13.60	0.85	16.92	352053.58
<i>Poecilia mexicana</i>	N 17° 26' W 95° 26'	0.63	NO	19.77	336.019	1.33	0.49	5.00	48473.19
<i>Poeciliopsis gracilis</i>	N 21° 59' W 99° 15'	0.84	YES	18.57	99.865	3.68	0.78	8.84	89901.33

<i>Poeciliopsis infans</i>	N 20° 34' W 104° 9'	1.05	YES	21.23	332.560	3.11	0.62	9.23	153007.38
<i>Poeciliopsis prolifica</i>	N 23° 03' W 105° 50'	5.4	YES	22.07	527.738	8.10	0.86	6.87	99651.54
<i>Poeciliopsis viriosa</i>	N 21° 02' W 104° 22'	0.93	YES	20.57	33.717	5.00	0.75	7.50	102710.40
<i>Priapella intermedia</i>	N 17° 9' W 95° 10'	0.6	NO	36.43	53.694	1.50	1.22	4.86	38902.24
<i>Xiphophorus hellerii</i>	N 20° 34' W 104° 9'	0.61	NO	20.67	312.286	1.33	1.71	6.39	69703.54

TABLE 2. Results of two regression models with different evolutionary assumptions examining the relationship between the matrotrophic index and different morphological characteristics of the placentas of 10 species from the Poeciliidae family. These two models used Phylogenetic Generalized Least Squares, assuming a Brownian motion evolutionary model (PGLS-BM) and an Ordinary Least Squares model, which assumes that characteristics adapt immediately to their environment. In all cases, the independent variable was the matrotrophic index and the dependent variables the average number of vesicles per follicular cell (NV), area of those vesicles (AV), average number of microvilli per follicular cell (NM) and the average length of these microvilli (LM), the average thickness of maternal follicle (TF) and the average area of maternal follicle (AF). The fit of each model was evaluated using the Akaike Information Criterion adjusted for small sample size (AICc). Also shown is the difference in AICc ($\Delta AICc$) between each model and the model with the best fit. For each characteristic, the models are listed from best to worst fit.

Response variable	Regression model	AICc	$\Delta AICc$	Slope	<i>P</i>
NV	PGLS-BM	3.62	0.00	0.052	0.098
	OLS	4.29	0.67	0.075	0.043
AV	OLS	23.52	0.00	0.195	0.097
	PGLS-BM	24.95	1.42	0.198	0.097
NM	OLS	16.91	0.00	0.082	0.266

	PGLS-BM	19.49	2.57	0.171	0.052
LM	OLS	4.13	0.00	0.227	0.503
	PGLS-BM	39.94	35.81	0.131	0.639
TF	PGLS-BM	-9.95	0.00	-0.009	0.452
	OLS	-9.86	0.09	-0.006	0.641
AF	OLS	9.74	0.00	0.164	0.006
	PGLS-BM	15.33	5.59	0.061	0.323

TABLE 3. Results of two regression models with different evolutionary assumptions examining the relationship between the presence or absence of superfetation and different morphological characteristics of the placentas of 10 species from the Poeciliidae family. These two models used Phylogenetic Generalized Least Squares, assuming a Brownian motion evolutionary model (PGLS-BM) and an Ordinary Least Squares model, which assumes that characteristics adapt immediately to their environment. In all cases, the independent variable was the matrotrophic index and the dependent variables the average number of vesicles per follicular cell (NV), area of those vesicles (AV), average number of microvilli per follicular cell (NM) and the average length of these microvilli (LM), the average thickness of maternal follicle (TF) and the average area of maternal follicle (AF). The fit of each model was evaluated using the Akaike Information Criterion adjusted for small sample size (AICc). Also shown is the difference in AICc ($\Delta AICc$) between each model and the model with the best fit. For each characteristic, the models are listed from best to worst fit. The regression slope shown denotes the difference between species with and without superfetation.

Response variable	Regression model	AICc	$\Delta AICc$	Slope	<i>P</i>
NV	PGLS-BM	2.80	0.00	0.127	0.363
	OLS	6.48	3.68	0.058	0.579
AV	OLS	24.23	0.00	0.188	0.554

	PGLS-BM	24.85	0.62	0.224	0.679
NM	OLS	13.01	0.00	0.310	0.074
	PGLS-BM	20.12	7.11	0.286	0.483
LM	PGLS-BM	36.44	0.00	1.037	0.363
	OLS	38.46	2.02	1.082	0.183
TF	OLS	-11.96	0.00	-0.023	0.492
	PGLS-BM	-12.22	0.26	-0.009	0.870
AF	PGLS-BM	9.73	0.00	0.446	0.059
	OLS	11.20	1.47	0.344	0.034

Fig. 1

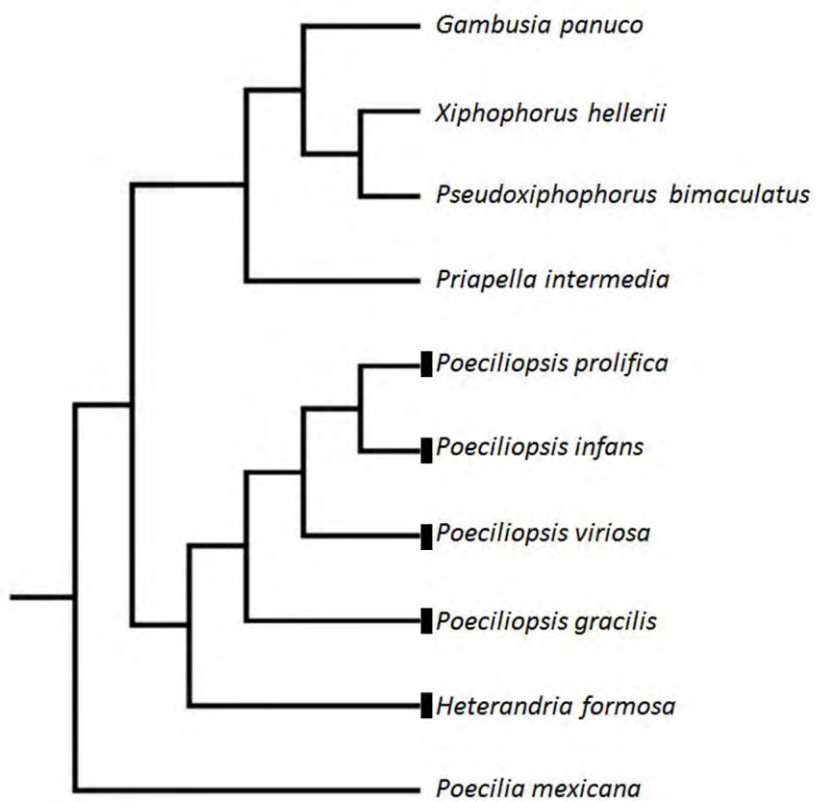


Fig. 2

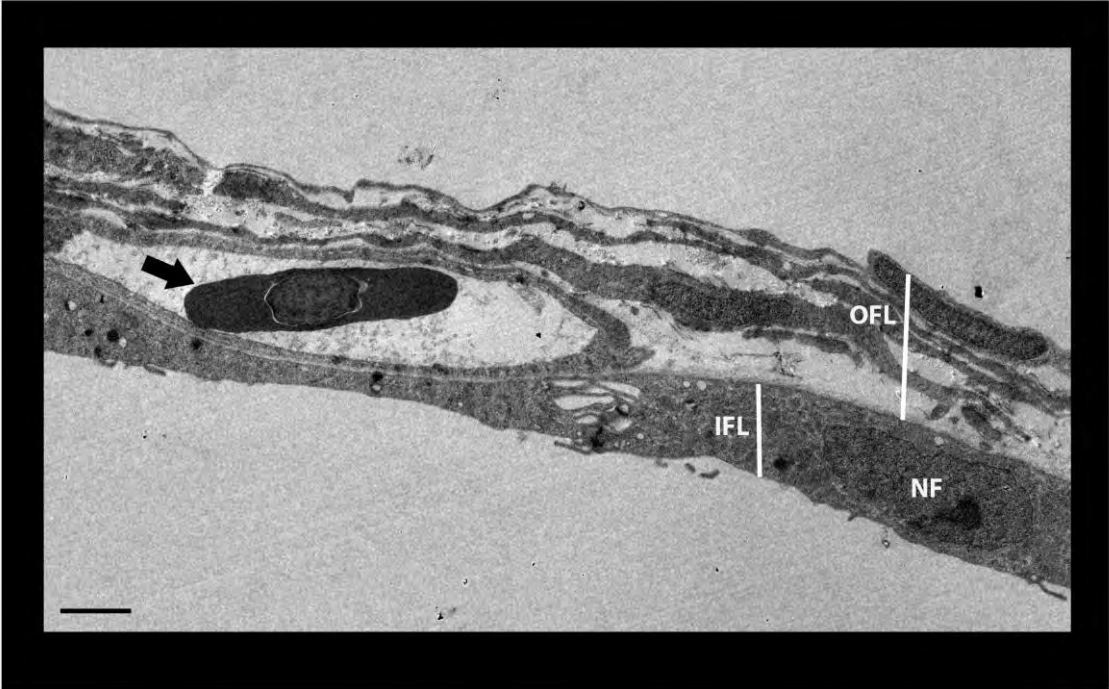


Fig. 3

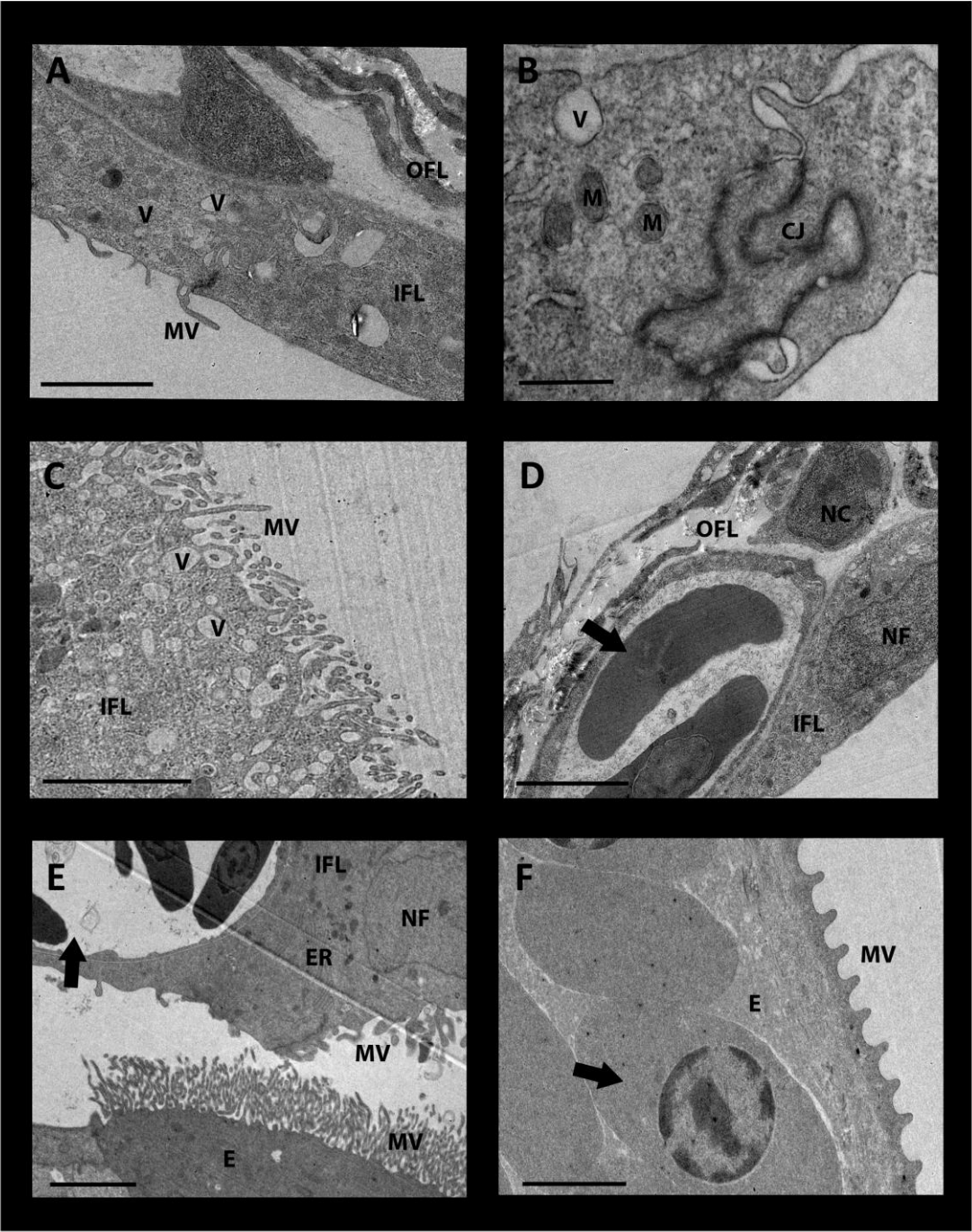


Fig. 4

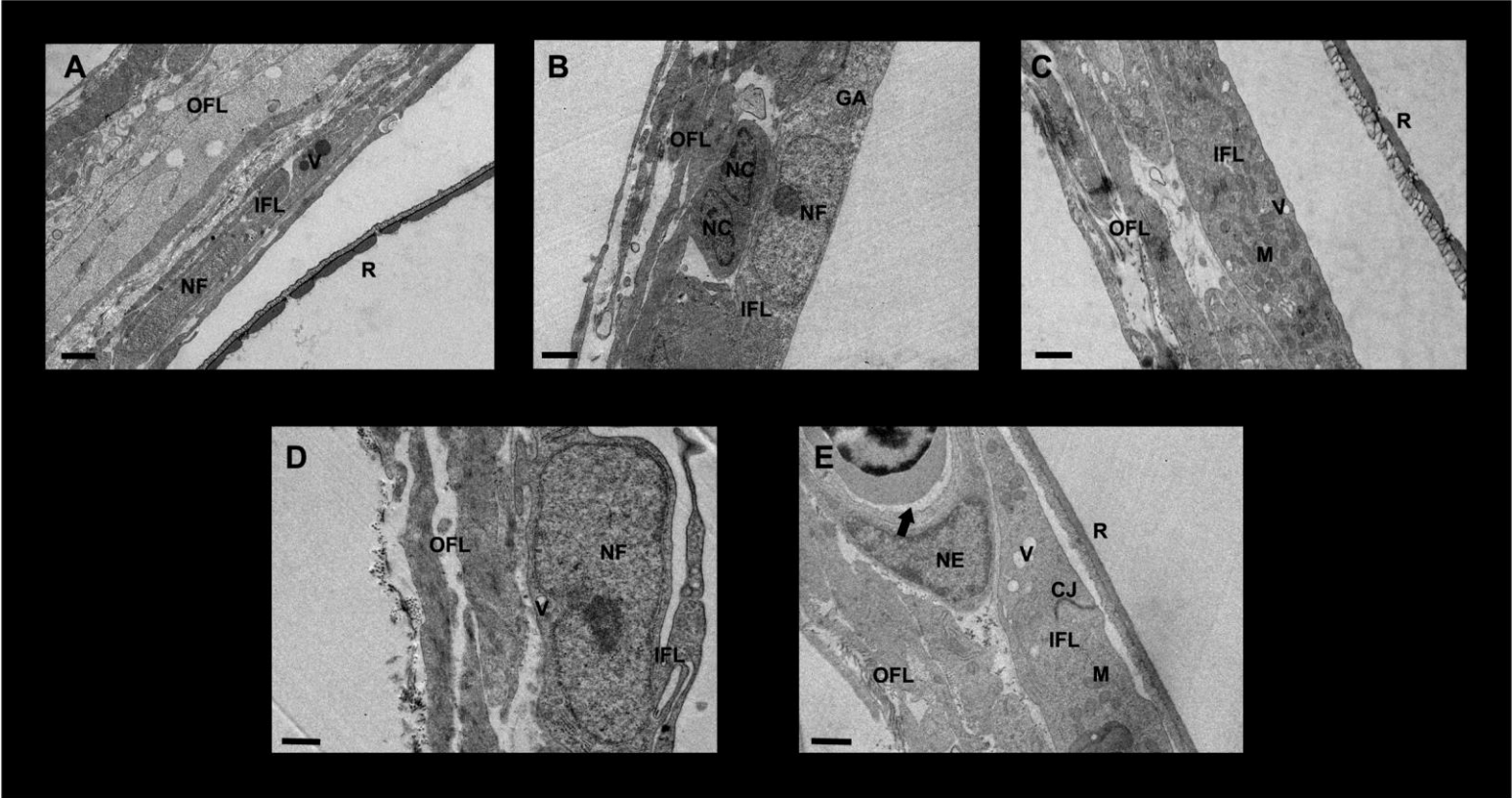


Fig. 5

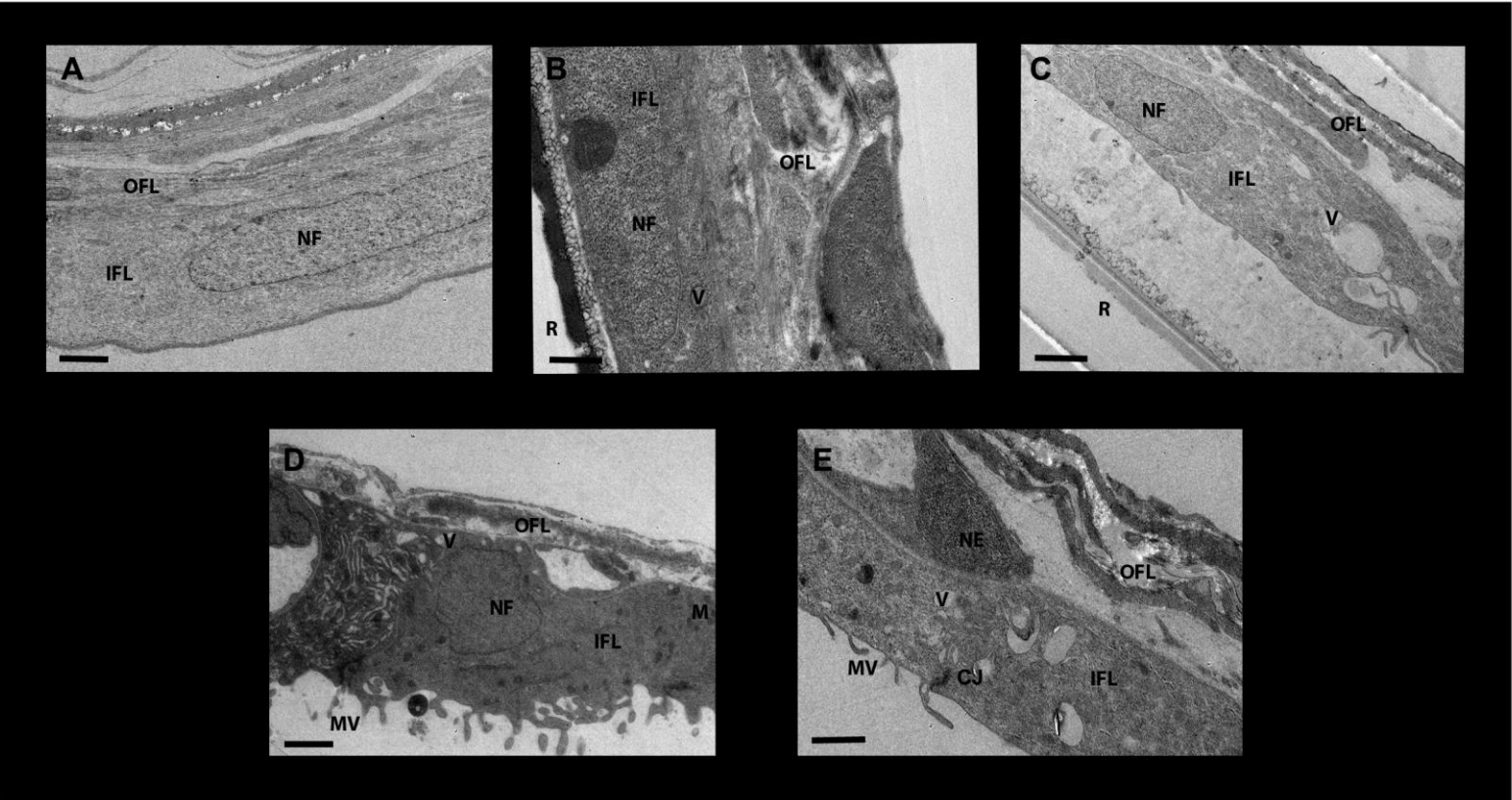


Fig. 6

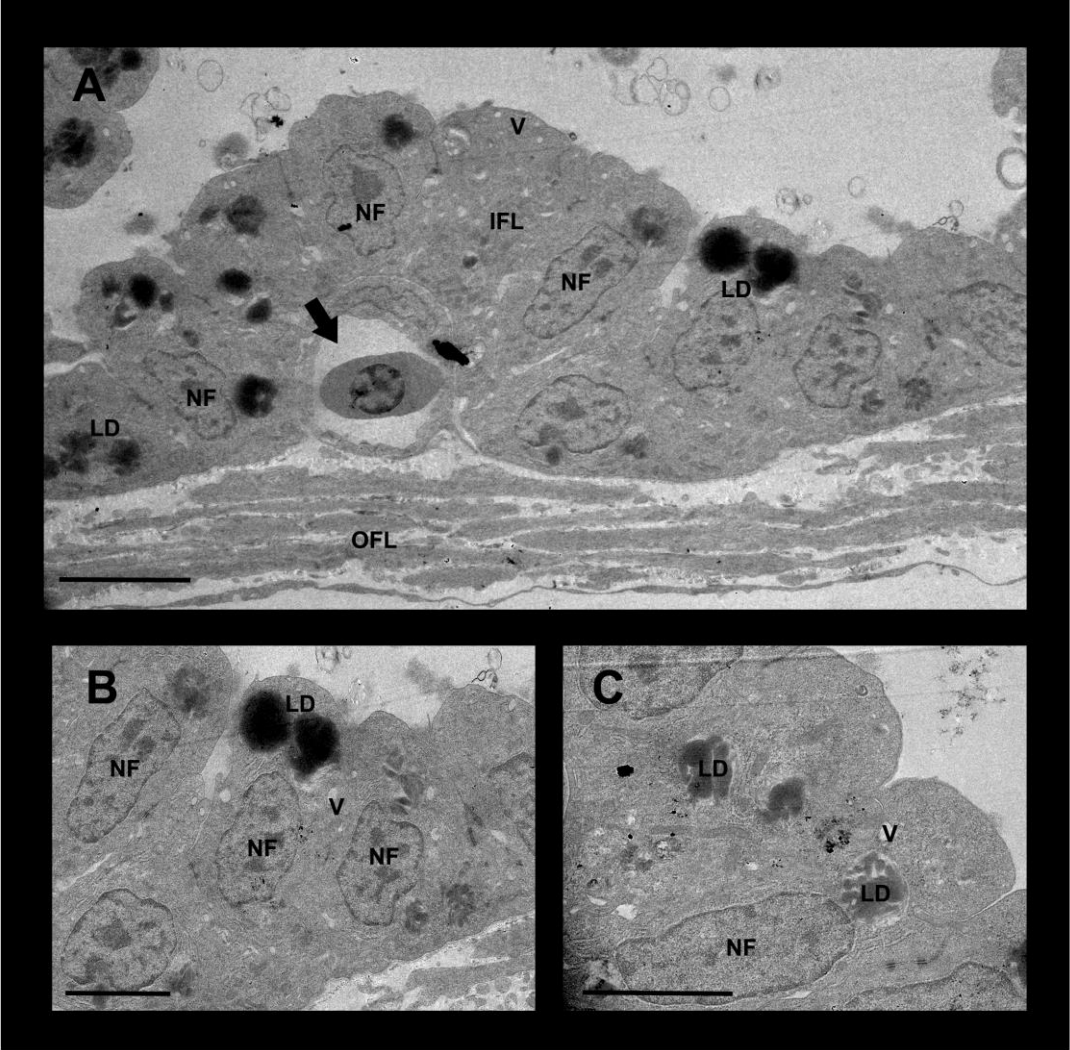


Fig. 7

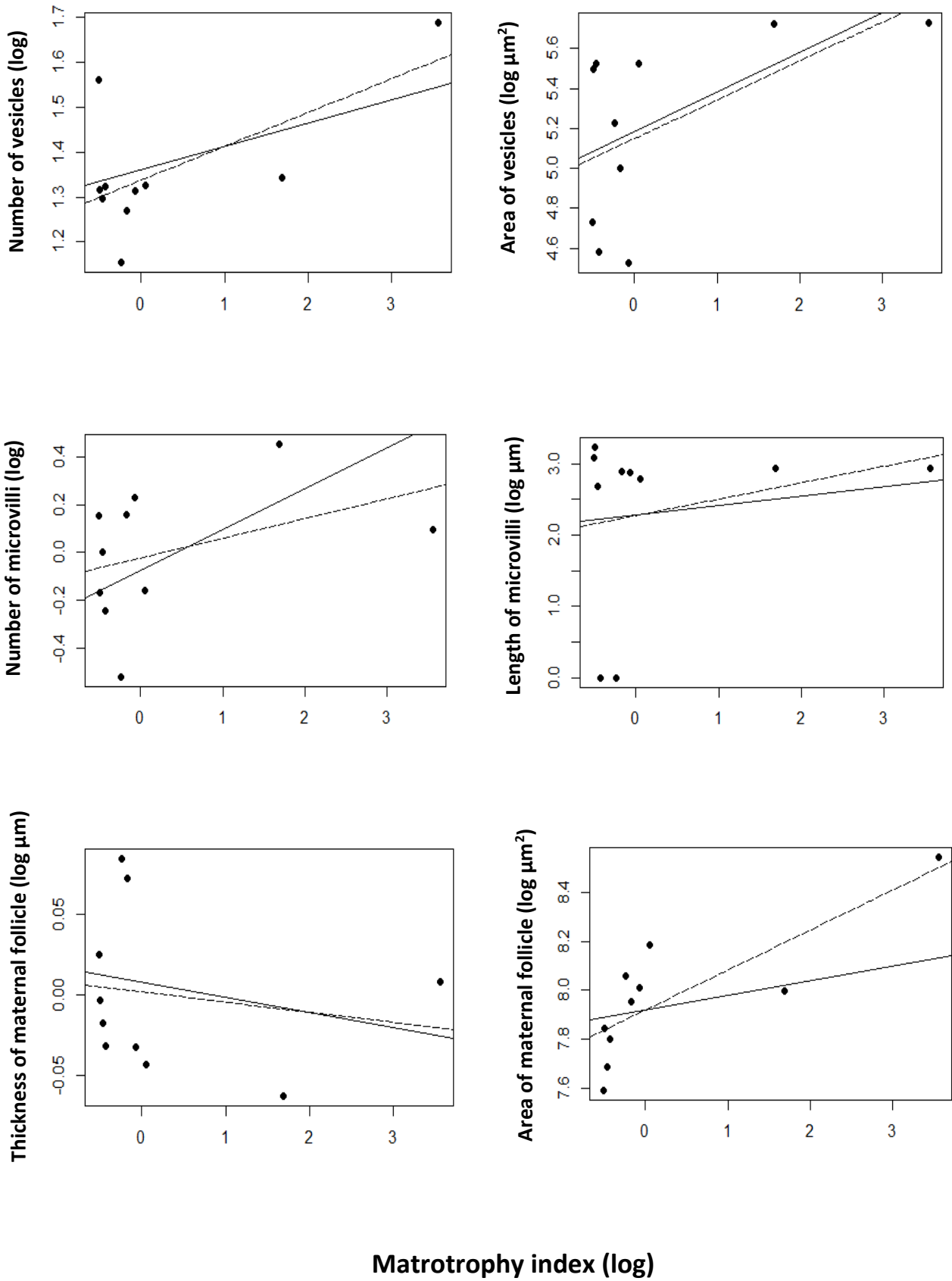
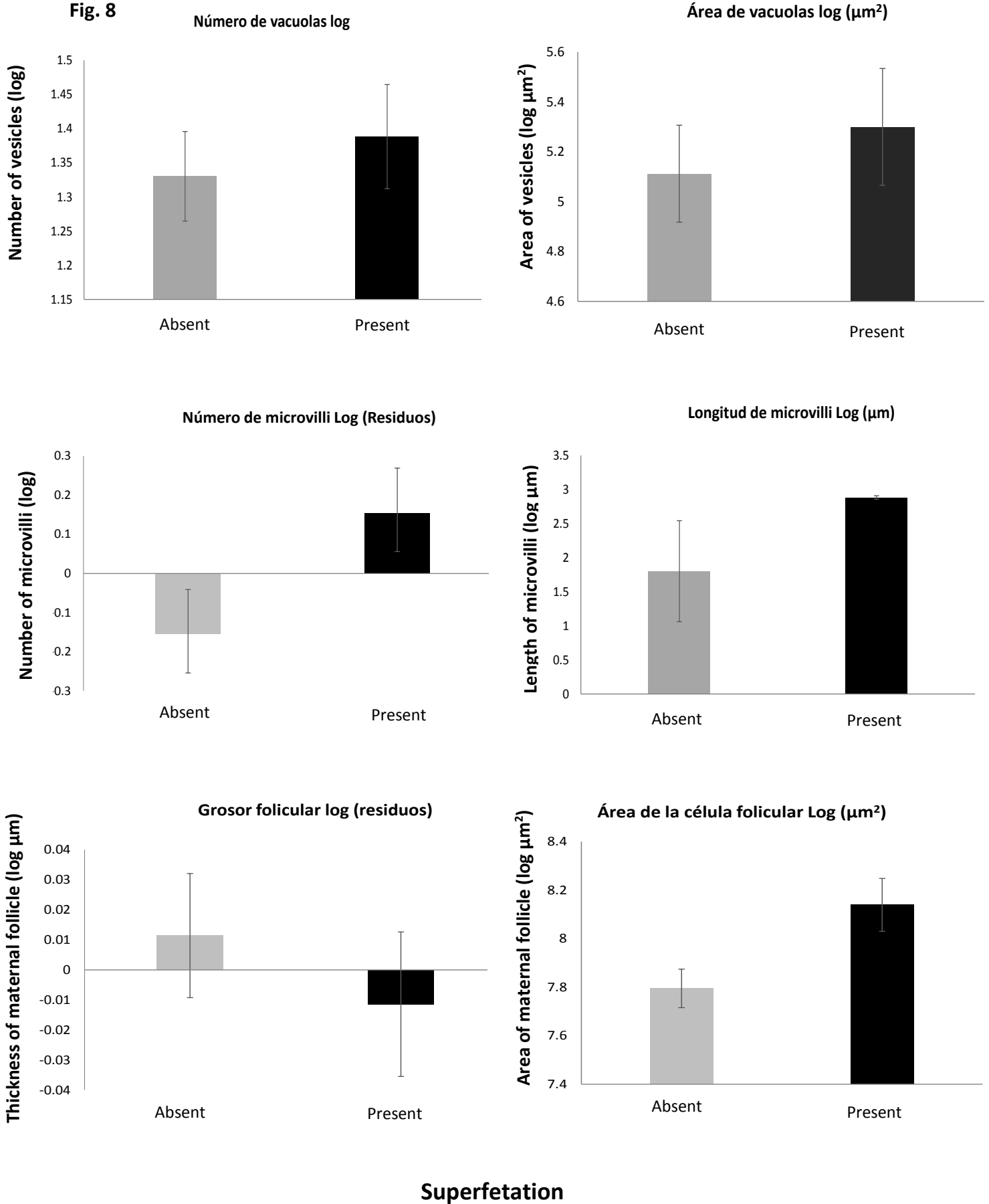


Fig. 8



CAPÍTULO II

**Morphological structures for potential sperm storage in
poeciliid fishes. Does superfetation matter?**

**Claudia Olivera-Tlahuel, Maricela Villagrán-Santa Cruz,
Norma A. Moreno-Mendoza, J. Jaime Zúñiga-Vega**

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RESEARCH ARTICLE

Morphological structures for potential sperm storage in poeciliid fishes. Does superfetation matter?

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Abstract

Sperm storage within the female reproductive tract has been reported as a reproductive strategy in several species of vertebrates and invertebrates. However, the morphological structures that allow for sperm to be stored and kept viable for long periods are relatively unknown in osteichthyes. We use histological and stereological tools to identify and quantify sperm storage structures (spermathecae) in 12 species of viviparous Poeciliidae. We found spermathecae in nine species, six of which exhibit superfetation (the ability of females to simultaneously carry within the ovary two or more broods of embryos at different stages of development). These spermathecae are folds of ovarian tissue that close around spermatozoa. We compared the number and size (volume) of spermathecae between species with and without superfetation. Species that exhibit superfetation had a significantly higher number of spermathecae than species that do not exhibit this reproductive strategy. In addition, we found that the mean volume of spermathecae and total volume of spermathecae present in the ovary are marginally higher in species with superfetation. Our results contribute to the understanding of the morphological structures that allow for sperm storage in viviparous osteichthyes and suggest a positive relationship between superfetation and the capacity of females to store sperm.

KEYWORDS

Poeciliidae, sperm storage, spermathecae, superfetation, viviparity

1 | INTRODUCTION

In many animal species, females are able to store viable sperm within their reproductive tracts for long periods of time (Orr & Zuk, 2012). This phenomenon is present in some groups of insects (Simmons, 2001), reptilian sauropsids (Birkhead & Møller, 1993; Sever & Hamlett, 2002), amphibians (Sever, 2002), osteichthyes (Potter & Kramer, 2000), cartilaginous fishes (Pratt, 1993), and mammals (Birkhead & Møller, 1993). Among these diverse taxa, the retention time of sperm can vary from only a few hours to years after mating. For example, insects such as bee and termite queens can store sperm for nearly their entire lives (several years) (Orr & Zuk, 2012). Conversely, the majority of female mammals retain sperm for only a few hours (Orr & Zuk, 2012). There is also variation among taxa in the specific location within the female

reproductive tract where spermatozoa are stored. For example, in reptilian sauropsids, sperm storage occurs in the oviduct (Villagrán-Santa Cruz, Mendoza-Cruz, Granados-González, Rheubert, & Hernández-Gallegos, 2017), whereas in bats spermatozoa are stored in the uterus (Birkhead & Møller, 1993).

In the osteichthyes, where internal fertilization has evolved independently several times (Gross & Shine, 1981; Mank, Promislow, & Avise, 2005), sperm retention has only been documented for Anablepidae, Cottidae, Embiotocidae, Sebastidae, and Poeciliidae (Darling, Noble, & Shaw, 1980; Fraser & Renton, 1940; Koya, Munehara, & Takano, 2002; Muñoz, Koya, & Casadevall, 2002; Turner, 1938; Vila, Sabat, Hernandez, & Munoz, 2007). In these groups of fish, sperm storage may occur in the ovary or in the gonoduct (Darling et al., 1980; Koya et al., 2002; Vila et al., 2007). For example, in *Helicolenus*

dactylopterus (Sebastidae) sperm retention occurs in storage chambers within the ovary that are lined by a cuboidal epithelium supported by muscle cells (Vila et al., 2007). The spermatozoa are concentrated inside these chambers, which are connected via a duct to the lumen of the ovary. In *Xiphophorus maculatus* (Poeciliidae), spermatozoa have been observed in association with the ovary and gonoduct epithelium (Potter & Kramer, 2000).

In poeciliid fishes, the ability to store sperm has been suggested because of the long time (several months) during which isolated females are able to produce young (Clark & Aronson, 1951; Greven, 2011; Hildemann & Wagner, 1954; Hubbs, 1964, 1997). However, the morphological structures where spermatozoa may be stored have only been examined in *Heterandria formosa*, *Poecilia formosa*, *Poecilia reticulata*, *X. maculatus*, and *Poeciliopsis gracilis* (Constantz, 1989; Kobayashi & Iwamatsu, 2002; Potter & Kramer, 2000; Turner, 1947; Uribe, De la Rosa-Cruz, Guerrero-Estévez, García Alarcón, & Aguilar-Morales, 2004; Uribe & Grier, 2011; Uribe, Grier, De la Rosa-Cruz, & Schartl, 2016). Temporary sperm storage structures (called “solid plugs” and “sperm storage micropockets”) have been observed in *H. formosa* and *P. reticulata*; these structures are formed by the closure of ovarian folds that contain spermatozoa (Fraser & Renton, 1940; Kobayashi & Iwamatsu, 2002).

In this study, our first objective was to locate sperm storage structures, which we will refer to as “spermathecae”, in viviparous species of Poeciliidae. The term “spermathecae” has been previously used to describe specialized sperm storage structures in other groups such as amphibians and insects (Dallai, 1975; Gupta & Smith, 1969; Pool & Hoage, 1973; Tombes & Roppel, 1972). For example, in salamanders, spermatozoa are stored in a pigmented structure (spermatheca) that opens into the oviducts (Pool & Hoage, 1973). Similarly, in beetles, spermathecae are abdominal invaginations near the reproductive tract which store viable sperm (Tombes & Roppel, 1972).

The study of morphological structures for potential sperm storage in poeciliid fishes may provide insight into some of the mechanisms and processes associated with superfetation, which is a particular mode of reproduction that occurs in some poeciliid species. Superfetation is the ability of females to simultaneously carry in the ovary multiple broods of embryos at different developmental stages (Turner, 1937, 1940; Zúñiga-Vega, Macías-García, & Johnson, 2010). Apparently, superfetation has evolved independently several times within the Poeciliidae (Meredith, Pires, Reznick, & Springer, 2011; Pires, Banet, Pollux, & Reznick, 2011; Pollux, Pires, Banet, & Reznick, 2009). Some species have a high degree of superfetation; for example, *H. formosa* can carry up to eight simultaneous broods, each at a different developmental stage (Scrimshaw, 1944). Other species, such as *Poeciliopsis turrubarensis*, have a lower degree of superfetation, and can only carry two or three simultaneous broods (Zúñiga-Vega, Reznick, & Johnson, 2007). Many other species, such as *Gambusia vittata*, do not exhibit superfetation, and females only produce one group of embryos at a time (Pires et al., 2011).

Many studies have attempted to find adaptive explanations for superfetation (Frías-Alvarez & Zúñiga-Vega, 2016; Macías-García & González-Zuñiga, 2005; Olivera-Tlahuel, Ossip-Klein, Espinosa-Pérez, & Zúñiga-Vega, 2015; Pollux et al., 2009; Travis, Farr, Henrich, & Cheong,

1987; Zúñiga-Vega et al., 2007, 2010). However, no study to date has focused on describing the morphological and physiological mechanisms that may allow superfetation to occur. For this reason, the second objective of this study is to explore the possibility of a relationship between the capacity for sperm storage and the presence of superfetation in members of the family Poeciliidae.

This second objective is derived from the hypothesis that superfetating species need to frequently fertilize mature oocytes, which are constantly developing in order to form new broods (Macías-García & González-Zuñiga, 2005; Pollux, Meredith, Springer, & Reznick, 2014). In this sense, superfetation would depend, to some extent, on morphological and physiological mechanisms that would allow females constant access to spermatozoa. As such, we expected to observe more and larger spermathecae in species with superfetation compared to non-superfetating species, where we expected to observe fewer, smaller, or even no spermathecae, because females without superfetation only develop one brood of embryos at a time and would therefore not need such frequent access to spermatozoa.

In addition, the potential association between sperm storage within spermathecae and superfetation might as well be adaptive. Most superfetating species do not exhibit apparent signs of precopulatory female choice (Macías-García & González-Zuñiga, 2005; Pollux, Meredith, Springer, & Reznick, 2014). Males are smaller than females, lack ornaments, and do not exhibit courtship behaviors (Magurran, 2011). Thus, the concurrent presence of both superfetation and sperm storage may facilitate cryptic (post-copulatory) female choice because sperm competition may take place within spermathecae and spermatozoa of the highest quality could then be used to fertilize different broods (Evans & Pilastro, 2011). Previous studies in poeciliids have demonstrated that ovarian fluid mediates sperm competition by increasing the longevity of some particular spermatozoa (Gasparini & Evans, 2013; Gasparini & Pilastro, 2011). These physiological processes may take place within spermathecae.

2 | MATERIALS AND METHODS

2.1 | Study species and histological protocol

We collected gestating females of 12 viviparous species of Poeciliidae (Table 1). The majority of these females were collected from different rivers in Mexico (collection of individuals was conducted under ethical approval from the National Commission of Aquaculture and Fisheries from Mexico; permits DGOPA.07010.210612.1749 and PPF/DGOPA-233/2013). The only exceptions were females of *H. formosa*, which were acquired from a private aquarium in the United States. Six of our study species exhibit superfetation, whereas the other six do not (Table 1). Given that in Mexico most of the species with superfetation belong to the genus *Poeciliopsis* (Miller, Minckley, & Norris, 2005), our sample of superfetating species includes five from this genus.

We collected three visibly pregnant females of each species, which were anesthetized (using 3-aminobenzoic acid ethyl ester) and sacrificed (by means of an overdose of the same anesthetic) upon capture to extract ovaries. Immediately following their extraction, the ovaries

TABLE 1 Summary of data and results for spermathecae of 12 species of the family Poeciliidae

Species	Geographic coordinates	N	SF	Ovary volume (mm ³)	Number of spermathecae	Total volume of all spermathecae (mm ³)	Volume of each spermatheca (mm ³)	Relative volume of all spermathecae	Relative volume of each spermatheca
<i>Gambusia panuco</i>	N 20° 41' W 103° 57'	3	NO	274.8	0.7	0.61	0.31	0.0022	0.0011
<i>Poecilia butleri</i>	N 20° 9' W 103° 2'	3	NO	955.6	0.0	-	-	-	-
<i>Poecilia mexicana</i>	N 17° 26' W 95° 26'	3	NO	637.0	4.7	2.67	0.63	0.0050	0.0010
<i>Priapella intermedia</i>	N 17° 9' W 95° 10'	3	NO	218.4	0.0	-	-	-	-
<i>Pseudoxiphophorus bimaculatus</i>	N 17° 8' W 95° 7'	3	NO	457.2	0.0	-	-	-	-
<i>Xiphophorus hellerii</i>	N 20° 34' W 104° 9'	3	NO	441.6	2.0	4.04	0.67	0.0058	0.0010
<i>Heterandria formosa</i>	Private aquarium	3	YES	47.8	7.0	0.89	0.13	0.0196	0.0027
<i>Poeciliopsis infans</i>	N 20° 34' W 104° 9'	3	YES	222.0	17.7	24.74	1.59	0.1140	0.0074
<i>Poeciliopsis gracilis</i>	N 21° 59' W 99° 15'	3	YES	297.1	15.3	43.25	2.63	0.1761	0.0129
<i>Poeciliopsis presidionis</i>	N 22° 29' W 105° 21'	3	YES	240.4	4.7	1.61	0.35	0.0119	0.0034
<i>Poeciliopsis prolifica</i>	N 23° 03' W 105° 50'	3	YES	195.3	5.0	5.47	1.46	0.0408	0.0076
<i>Poeciliopsis viriosa</i>	N 21° 02' W 104° 22'	3	YES	631.8	8.0	4.69	0.55	0.0118	0.0012

SF indicates presence or absence of superfetation. All values represent averages per species (across all individual ovaries). Relative volumes represent proportions with respect to total ovary volume. N = sample size.

were fixed in Bouin's solution for 30 min and then stored in 70% ethanol until processing. In the laboratory, each ovary was dehydrated prior to embedding in paraffin. Then, the entire ovaries were serially sectioned longitudinally in 6- μ m section thickness using a rotary microtome (HistoStat 820). Finally, sections were stained using hematoxylin-eosin (Presnell, Schreiber, & Humason, 1997). We carried out this protocol for a total of 36 individual ovaries (three per each species).

2.2 | Identification and stereology of spermathecae

Each of the histological preparations (all serial sections of each ovary) was examined using a light microscope in order to detect as many spermathecae as possible. We defined a spermatheca as a closed structure (complete closure of the ovarian folds) that contains spermatozoa originating from the ovarian lumen after a mating event. The criterion for identifying a spermatheca was the presence of spermatozoa accumulated in a specific region completely surrounded by ovarian tissue (i.e., surrounded by a layer of epithelial cells, which was in turn surrounded

by a layer of conjunctive tissue). We did not consider structures with small openings into the ovarian lumen to be spermathecae, but rather deep ovarian folds.

Each spermatheca was serially photographed (at 6- μ m intervals) under 40x magnification. Photographs were then analyzed stereologically using the "Volumest" plug-in for the program ImageJ (Schneider, Rasband, & Eliceiri, 2012) to quantify the volume of each spermatheca. For each of the three ovaries that we examined per species, we quantified the total number of spermathecae, the total volume of all spermathecae, and the average volume per spermatheca. Then, for each of these variables we calculated the average across the three ovaries to obtain a representative value for each species.

2.3 | Estimating differences between species with and without superfetation

We searched for differences in the number, total volume, and average volume of spermathecae between species with and without

superfotation using two different evolutionary assumptions. First, we assumed that these traits evolved under Brownian motion (BM) and, hence, phylogenetic relatedness would account for most of the observed interspecific variation (Felsenstein, 1985). We implemented this evolutionary model in our comparison between species with and without superfotation by means of phylogenetic generalized least squares (PGLS) (Martins & Hansen, 1997), treating the presence/absence of superfotation as a binary (dummy) independent variable. PGLS is a regression technique that incorporates phylogenetic information into the error structure of the model. We used the phylogeny of the Poeciliidae published by Pollux et al. (2014), trimmed to our focal species. We created an ultrametric tree by transforming the original branch lengths to a scale based on relative time (Figure 1). To do so, we used a semiparametric smoothing method based on penalized likelihood (Sanderson, 2002) implemented in the R package “ape” (Paradis, Claude, & Strimmer, 2004; R Core Team, 2016). We used a smoothing parameter equal to 0.1, which is a conservative value (Revell & Reynolds, 2012). We used our resulting ultrametric tree to estimate how many times superfotation has evolved throughout the phylogenetic history of our study species by means of a likelihood-based ancestral reconstruction method implemented in the program MESQUITE 3.0 (Maddison & Maddison, 2009).

Second, we assumed that traits evolve fast and adapt immediately to their local environments, leaving behind no trace of their phylogenetic history. Under this evolutionary assumption, phylogeny is unimportant and, thus, we used ordinary least-squares regression (OLS) to compare number, total volume, and average volume of spermathecae between species with and without superfotation (Garland, Bennett, & Rezende, 2005; Lavin, Karasov, Ives, Middleton, & Garland, 2008). We fitted PGLS and OLS regression models using the R packages “ape” and “nlme” (Paradis et al., 2004; Pinheiro, Bates, DebRoy, Sarkar, & R Core Team, 2016).

We compared the fit of these two evolutionary models using the Akaike information criterion (AIC) adjusted for small sample sizes (AICc) (Burnham & Anderson, 2002). The smallest AICc value indicates the best-fitting model. A difference between models in AICc values larger than 2 units indicates a real difference in their fit to the data. Originally, we considered using a third evolutionary model that would estimate the amount of phylogenetic signal in the relationship between superfotation and our response variables. This alternative model would indicate how important the phylogeny is in explaining the effect of superfotation, rather than assuming that phylogeny either explains most of the data (PGLS assuming BM) or is unimportant (OLS). However, such alternative model requires the estimation of one additional parameter (i.e., a measure of phylogenetic signal such as λ) (Freckleton, Harvey, & Pagel, 2002; Pagel, 1999) and our sample size (12 species) was too small to allow a precise estimation of this additional parameter.

Before analyses, we log-transformed the number of spermathecae. This variable was not statistically affected by the volume of the ovary (slope = -0.007, $p = 0.33$). In contrast, total and average volume of spermathecae were expressed as proportions of the total volume of

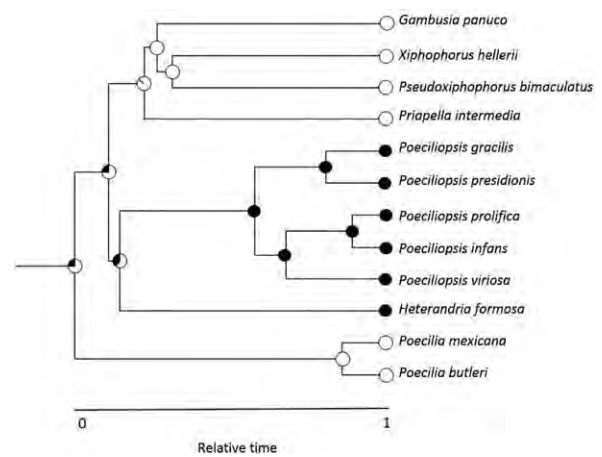


FIGURE 1 Ultrametric phylogenetic tree of the study species from the family Poeciliidae. Branch lengths are expressed in relative time. In the ancestral nodes, circles depict proportional likelihoods for the presence (black) and absence (white) of superfotation

the ovary to make these two variables fully comparable among species. These proportions were arcsine-transformed before analyses to meet assumptions of normality and homogeneity of variances (Zar, 2010). To test the effect of superfotation we used our full data set (12 species) for the number of spermathecae and a reduced data set (nine species) for total and average volume of spermathecae, because we did not find spermathecae in three species (Table 1).

3 | RESULTS

3.1 | Morphology of spermathecae

We found spermathecae in the ovaries of nine of the twelve species we examined (*Gambusia panuco*, *Xiphophorus hellerii*, *Poecilia mexicana*, *Poeciliopsis prolifica*, *Poeciliopsis infans*, *Poeciliopsis viriosa*, *Poeciliopsis presidionis*, *P. gracilis*, and *H. formosa*; Figures 2–4). In general, we observed that the spermathecae were not found in a specific area within the ovary, but rather were distributed more or less randomly, though always near the ovarian lumen. The spermathecae were frequently located next to blood vessels. For example, in *G. panuco*, *P. prolifica*, *P. infans*, and *P. viriosa* (Figures 2a–c and 3a–i), a large number of erythrocytes are visible near the spermathecae.

We confirmed that the spermathecae are composed of ovarian tissue and are generally oval or circular structures that completely surround spermatozoa to protect and store them within the ovary (Figures 2–4). In Figures 2–4, we present representative images of the proximal (a, d, g), middle (b, e, h), and distal sections of these structures (c, f, i). In all cases, the spermathecae were formed by thickened ovarian epithelium (brackets in Figures 2–4) that opens and closes to contain and protect spermatozoa. In Supporting Information, we provide additional serial sections of one spermatheca from each species to show their morphologies in greater detail (Supporting Information Figure S1).

The spermatozoa were oriented in a particular way within the spermathecae, with the heads in direct contact with the thickened

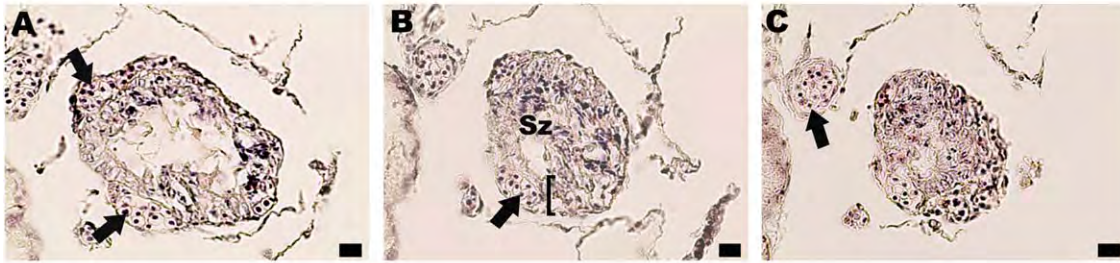
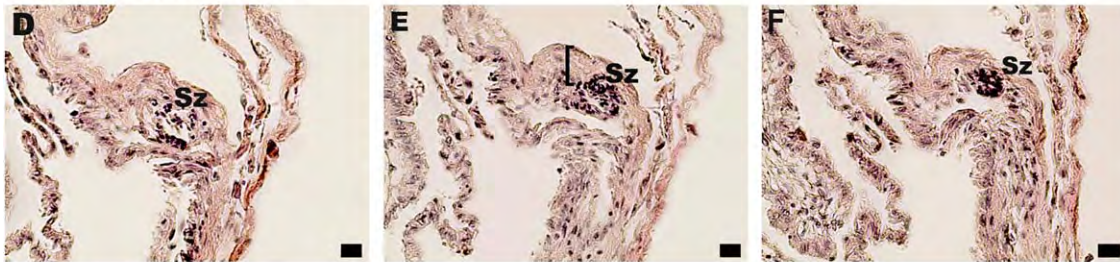
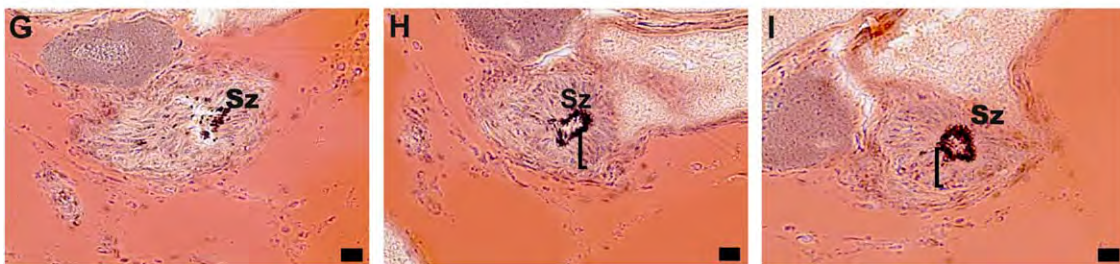
Gambusia panuco***Xiphophorus hellerii******Poecilia mexicana***

FIGURE 2 Sections of ovarian tissue stained using the hematoxylin-eosin technique that show the spermathecae observed in *G. panuco*, *X. hellerii*, and *P. mexicana* (species without superfetation). Heads of spermatozoa within the spermathecae stain blue (Sz). The different panels show representative images of the proximal (a, d, g), middle (b, e, h), and distal (c, f, i) sections of the spermathecae. Arrows point out erythrocytes. Brackets indicate thickened ovarian wall. Scale bar = 10 μ m

ovarian tissue and the tails towards the lumen of the spermatheca (Figures 2h,i and 4g–i). However, we did observe some spermathecae in which spermatozoa were dispersed around the interior (e.g., *P. prolifica*, see Supporting Information, Figure S1).

3.2 | Quantification of the spermathecae and comparison between species with and without superfetation

All of the ovaries we examined from superfetating species had spermathecae. In contrast, the ovaries of three of the species that do not superfetate (*Priapella intermedia*, *Pseudoxiphophorus bimaculatus*, and *Poecilia butleri*) lacked spermathecae (Table 1). In addition, in two of the species that do not superfetate (*X. hellerii* and *G. panuco*), we found spermathecae in only one of the three ovaries we examined (6 and 2 spermathecae, respectively). We did not find apparent differences in the morphology of spermathecae between superfetating and non-superfetating species (Figures 2–4 and Supporting Information, Figure S1).

The maximum number of spermathecae that we observed per individual ovary was 22 in one female of *Poeciliopsis infans* (a species with superfetation). The absolute volume of each spermatheca varied widely, from 0.11 mm³ in one female of *H. formosa* to 3.42 mm³ in one female of *P. gracilis*. However, much of this variation in the size of spermathecae can be explained by variation in ovary size (*H. formosa* is a small species and *P. gracilis* is relatively large). After correcting for ovary size (expressing volume of spermathecae as proportions of ovary volume), the average relative volume of each spermatheca varied from 0.001 in *X. hellerii* and *P. mexicana* (both species without superfetation) to 0.0129 in *P. gracilis* (a species with superfetation). Table 1 shows average values per species for number, total volume, and average volume of spermathecae.

We identified three groups of species that clearly differ in the relative volume of all spermathecae (Table 1). First, species with the smallest proportion of total ovary volume (<.01) devoted to the production of spermathecae. This group is composed of the three species without superfetation in which we observed spermathecae (*G. panuco*, *P. mexicana*, and *X. hellerii*). The second group includes four superfetating

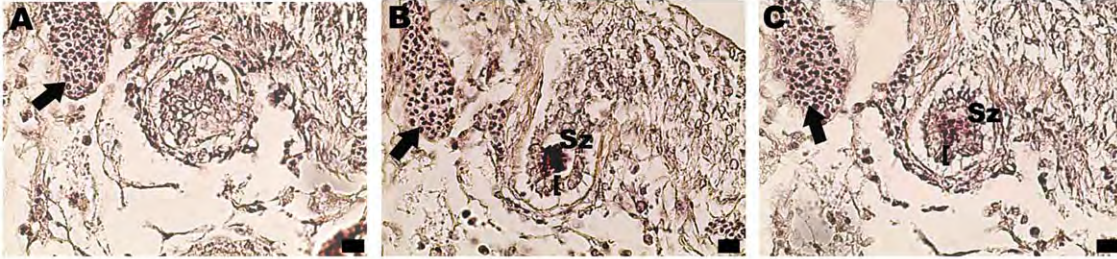
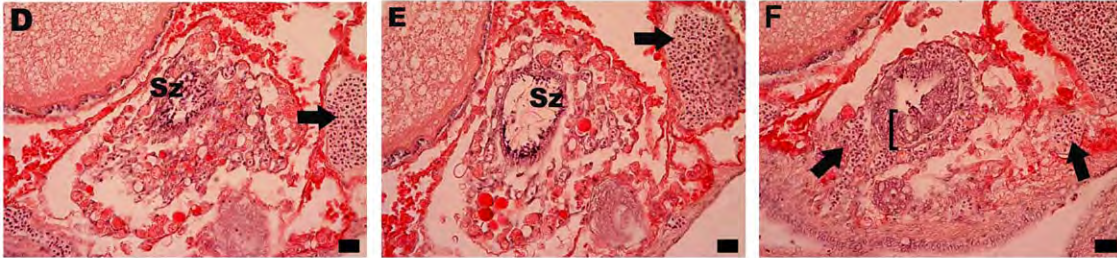
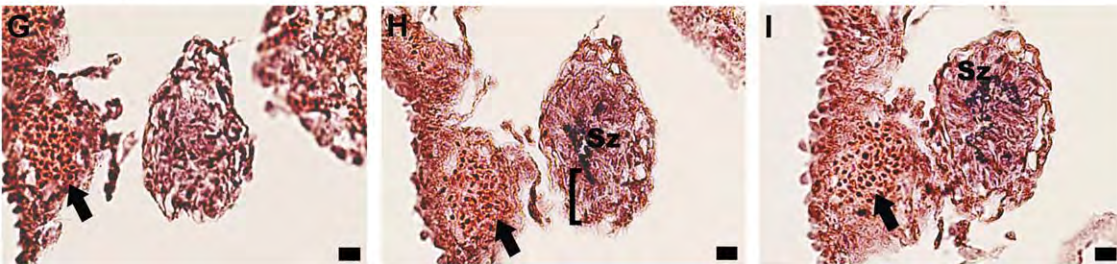
Poeciliopsis prolifica***Poeciliopsis infans******Poeciliopsis viriosa***

FIGURE 3 Sections of ovarian tissue stained using the hematoxylin-eosin technique that show the spermathecae observed in *P. prolifica*, *P. infans*, and *P. viriosa* (species with superfetation). Heads of spermatozoa within the spermathecae stain blue (Sz). The different panels show representative images of the proximal (a, d, g), middle (b, e, h), and distal (c, f, i) sections of the spermathecae. Arrows point out erythrocytes. Brackets indicate thickened ovarian wall. Scale bar = 10 μ m

species with a total relative volume of spermathecae between 0.01 and 0.04 (*H. formosa*, *P. presidionis*, *P. prolifica*, and *P. viriosa*). The third group includes the two superfetating species with the largest proportions of ovary volume devoted to spermathecae (0.11 and 0.18 in *P. infans* and *P. gracilis*, respectively; Table 1).

We found some uncertainty with respect to how many times superfetation has evolved throughout the phylogenetic history of our study species. Both presence and absence of superfetation were almost equally likely in the common ancestor of *Poeciliopsis* and *Heterandria* (the two genera where superfetation is present; Figure 1). If this common ancestor exhibited superfetation, then this reproductive strategy only evolved once. In contrast, if this common ancestor lacked superfetation, then this mode of reproduction evolved twice independently: once on the branch leading to the common ancestor of all *Poeciliopsis* species and once on the branch leading to *H. formosa* (Figure 1).

In all three response variables, the model that assumed phylogenetic independence (OLS) outperformed the model that assumed a Brownian motion model of evolution (Δ AICc > 2 in all three cases; Table 2). Therefore, phylogeny does not explain the effect of superfetation

on number, total volume, and average volume of spermathecae. According to OLS regression, the total number of spermathecae was significantly higher in species with superfetation than in species without superfetation ($p = 0.001$; Table 2, Figure 5a). Also according to OLS regressions, total and average volume of spermathecae (after correcting for ovary size) were higher in species with superfetation than in species without superfetation, although these differences were only marginally significant ($p = 0.09$ and 0.06, respectively; Table 2 and Figure 5b,c).

4 | DISCUSSION

4.1 | Spermathecae in poeciliid fishes

The first work in poeciliid fishes to demonstrate temporary sperm storage in a structure formed by the closing of ovarian folds that contain spermatozoa (a structure they called a “solid plug”) was carried out in *H. formosa* (Fraser & Renton, 1940). Later, Kobayashi and Iwamatsu (2002) described special sperm retention structures in *P. reticulata*,

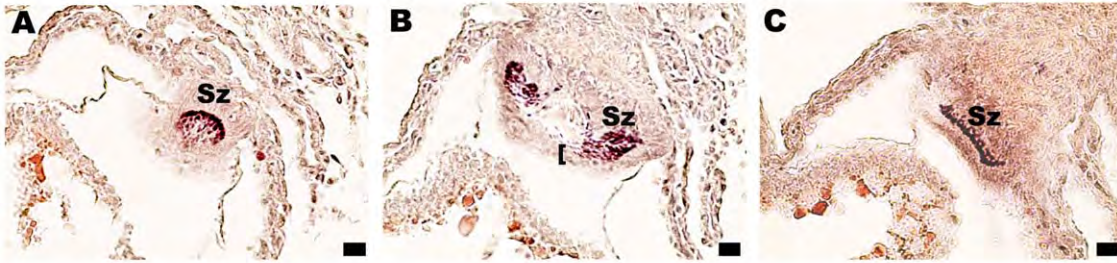
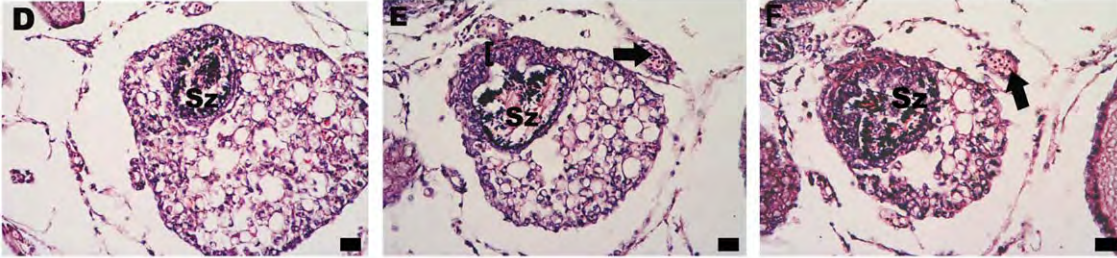
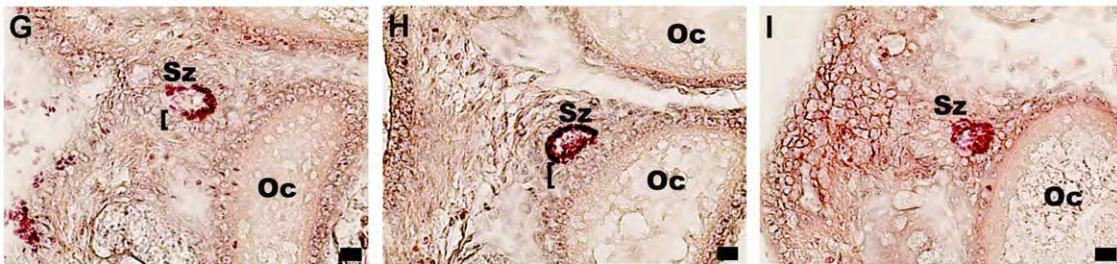
Poeciliopsis presidionis***Poeciliopsis gracilis******Heterandria formosa***

FIGURE 4 Sections of ovarian tissue stained using the hematoxylin-eosin technique that show the spermathecae observed in *P. presidionis*, *P. gracilis*, and *H. formosa* (species with superfetation). Heads of spermatozoa within the spermathecae stain blue (Sz). The different panels show representative images of the proximal (a, d, g), middle (b, e, h), and distal (c, f, i) sections of the spermathecae. Arrows point out erythrocytes. Brackets indicate thickened ovarian wall. Scale bar = 10 μm . Oocytes (Oc)

which they termed “sperm storage micropockets.” In the present study, we describe similar structures in nine species of the Poeciliidae (*G. panuco*, *X. hellerii*, *P. mexicana*, *P. prolifica*, *P. infans*, *P. viriosa*, *P. presidionis*, *P. gracilis*, and *H. formosa*). However, we refer to these structures as “spermathecae” in order to be consistent with the term used for analogous structures in other taxonomic groups (Dallai, 1975; Pool & Hoage, 1973; Tombes & Roppel, 1972). Similar to “sperm storage micropockets” and “solid plugs”, the spermathecae we observed are composed of ovarian folds, which close, completely surrounding the spermatozoa. However, spermatozoa may be temporarily stored as well in folds of the ovarian epithelium that do not close, but that instead connect directly to the ovarian lumen or to developing oocytes. These funnel-like folds have been called “dellen” (Greven, 2011; Mendoza, 1943). In fact, we observed spermatozoa within such deep ovarian folds (which we did not consider to be spermathecae) in most of the ovaries that we examined (Figure 6).

The spermathecae of all of the species we studied are composed of thickened ovarian epithelium. In general, the heads of the spermatozoa are oriented toward the ovarian epithelium and the tails toward

the lumen of the spermatheca. This particular orientation of the spermatozoa could indicate that the internal ovarian tissue nourishes the spermatozoa to keep them viable. For example, in *X. maculatus*, the cells of the ovarian tissue that interact with spermatozoa exhibit abundant ribosomes, microvilli, appendices, and desmosomes, which are characteristics of cells that transport nutrients (Potter & Kramer, 2000). The blood vessels that we observed surrounding the spermathecae may also contribute nutrients to the spermatozoa stored within. Blood vessels transport oxygen and nutrients necessary for cells, so their proximity to spermathecae is additional evidence that the ovarian tissue of the female provides the necessary media to keep spermatozoa viable for a relatively prolonged period.

We still do not know how long spermathecae last within the ovary. However, there are studies that provide information about how long spermatozoa remain viable within the ovary, which is indirect evidence of the time between the formation and disappearance of spermathecae. For example, sperm may be stored for up to nine months in *H. formosa* and up to 10 months in *H. dactylopterus* (Fraser & Renton, 1940; Hubbs, 1964, 1997; Scrimshaw, 1944; Vila et al., 2007).

TABLE 2 Results from phylogenetic generalized least squares assuming a Brownian motion model of evolution (PGLS-BM) and ordinary least squares regression (OLS) fitted to the number, total volume, and average volume of spermathecae in ovaries of poeciliid fishes

Response variable	Regression model	AICc	Δ AICc	Regression slope	<i>p</i>
Number of spermathecae	OLS	31.48	0.00	1.70	0.001
	PGLS-BM	40.56	9.08	1.69	0.15
Total volume of spermathecae	OLS	3.27	0.00	0.16	0.09
	PGLS-BM	8.47	5.20	0.13	0.54
Average volume of spermathecae	OLS	-18.37	0.00	0.04	0.06
	PGLS-BM	-15.29	3.08	0.03	0.41

Total and average volume of spermathecae were treated as proportions of total ovary volume. In all cases, the explanatory variable was the presence or absence of superfetation. The fit of each model to the data was evaluated using the Akaike Information Criterion adjusted for small sample sizes (AICc), with the smallest value indicating the best-fitting model. Differences in AICc values between models (Δ AICc) are also shown. The regression slope represents the estimated difference between species with and without superfetation.

Other vertebrates exhibit structures analogous to the spermathecae of poeciliid fishes (Dallai, 1975; Holt & Lloyd, 2010; Palmer, Rostal, Grumbles, & Mulvey, 1998; Suarez, 2008). For example, in lizards, specialized tubules for sperm storage are found in the oviduct of females (Villagrán-Santa Cruz et al., 2017). In some amphibians, such as caecilians, particular cloacal structures function as sites of sperm storage (Kuehnelt & Kupfer, 2012). In fish from other families, such as *H. dactylopterus* (Sebastidae), sperm retention occurs in storage chambers in the ovary (Vila et al., 2007).

4.2 | Does superfetation matter?

We have presented evidence that superfetation is positively related to the number and size of spermathecae. Independent of phylogenetic history, the superfetating species in our sample had more and larger spermathecae than non-superfetating species. As such, the former species devote a larger proportion of their ovarian volume to the formation of spermathecae than the latter. This difference between species with and without superfetation is substantial: the largest relative volumes of all spermathecae occurred in two species with superfetation and were on average over 30 times greater than the smallest relative volumes, which occurred in non-superfetating species. In addition, in three of our six non-superfetating species we found no spermathecae (Table 1). These results suggest a relationship between sperm storage within spermathecae and superfetation.

A potential caveat is our sample size. We examined three individual ovaries from each of the 12 species. Thus, the observed differences between species with and without superfetation should be interpreted with caution. In the case of number of spermathecae, the observed difference was significant at the conventional α level of 0.05. However, the differences in total and average volume of spermathecae could be considered significant only if we use a less conservative α level of 0.1, which may be appropriate for small sample sizes because the statistical power to detect differences is low. Regardless, in all three variables

variation among individual ovaries (within each species) was lower than the variation that we observed among species (Supporting Information, Figure S2) and, therefore, the observed differences between species with and without superfetation are likely to remain with the inclusion of additional individuals.

In general, spermathecae are considered a morphological mechanism that allows a continuous supply of spermatozoa, facilitating reproduction when there is a low abundance of males or when reproductive cycles are asynchronous (Birkhead & Møller, 1993; Gist & Congdon, 1998). However, in populations of poeciliid fishes, males are abundant and females are receptive year-round (Frias-Alvarez, Garcia, Vázquez-Vega, & Zúñiga-Vega, 2014; Houde, 1997). Therefore, it is unlikely that in poeciliids these are the main functions of sperm storage within spermathecae. Based on our evidence of higher prevalence of spermathecae in superfetating species, we propose that sperm storage is an important mechanism for the development of superfetation, because it allows a constant supply of viable spermatozoa with which to fertilize frequently developing broods. However, this hypothesis associating superfetation with sperm storage assumes that the frequency of copulations is the same in species with and without superfetation, since superfetating species could acquire a constant supply of spermatozoa simply by copulating more frequently.

Another hypothesis that could explain the higher prevalence of spermathecae in species with superfetation suggests the presence of post-copulatory sexual selection. Males of some species of the family Poeciliidae exhibit elaborated ornaments and courtship behaviors, and females actively choose reproductive partners before copulation (Basolo, 1990; Bisazza, 1993). In contrast, males of other species lack these ornaments or courtship behaviors and their reproductive success is based on forced copulations (Magurran, 2011). Superfetation is more common in species where sexual coercion prevails and, as such, could be a reproductive strategy to compensate for the apparent lack of female mate choice (Macías-García & González-Zuarth, 2005; Pollux et al., 2014). When females are unable to choose a high-quality mate

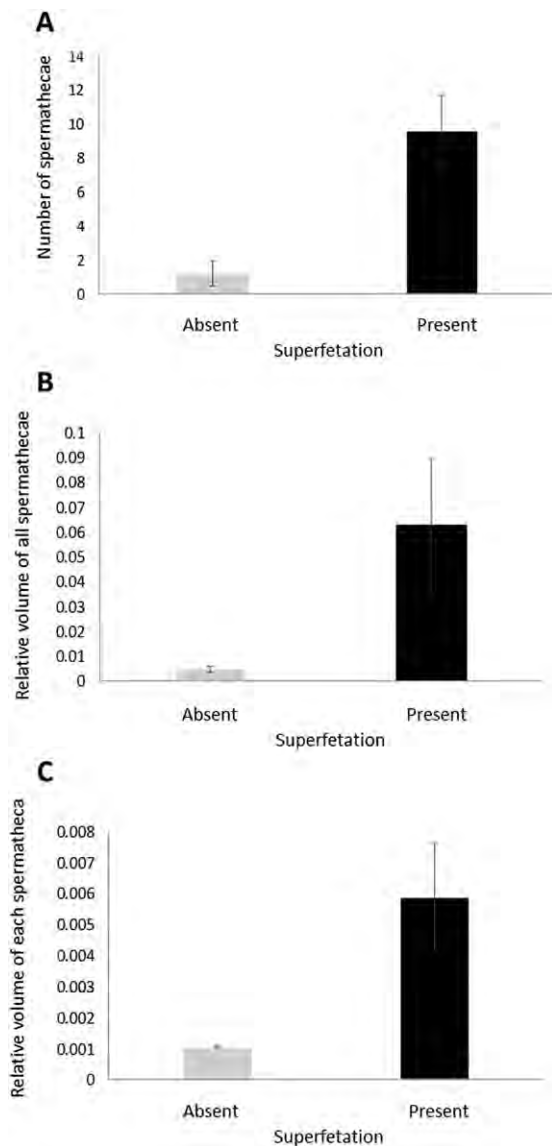


FIGURE 5 Total number (a), total volume (b), and average volume (c) of spermathecae in ovaries of poeciliid species with (black bars) and without (gray bars) superfetation. Volumes in b and c are expressed as proportions of total ovary volume. Error bars represent one standard error

before copulation, they could use mechanisms of post-copulatory sexual selection (i.e., cryptic female choice) to bias paternity in favor of some males and against others by means of physiological processes that occur within their ovaries (Birkhead & Pizzari, 2002; Evans & Pilastro, 2011). Sperm storage in spermathecae would have a fundamental role for this purpose in species with superfetation. Higher quality spermatozoa would have a higher likelihood of surviving within the spermathecae and would be stored for some period of time until being used by the females. In addition, sperm competition could occur within the spermathecae. Previous studies in poeciliids have documented sperm competition as well as an important influence of the female, through the ovarian fluid, on the motility and survival of spermatozoa (Gasparini & Evans, 2013; Gasparini & Pilastro, 2011; Gasparini, Simmons, Beveridge, & Evans, 2010).

Another possible advantage of sperm retention within spermathecae in species with superfetation is greater potential for storing sperm of different males, favoring the fertilization of oocytes by different genotypes and thus increasing their progeny's genetic variability (Macías-García & González-Zuarth, 2005; Pollux et al., 2014). If females that superfetate usually lack mechanisms for pre-copulatory sexual selection, and therefore cannot assure the genetic quality of their young, they could at least increase their genetic variability. In this way, the ability to store sperm in spermathecae could facilitate multiple paternity, particularly in species with superfetation.

Finally, it is important to emphasize that we also found spermathecae (although fewer and smaller) in some species that do not superfetate. This indicates that cryptic female choice, sperm competition, and multiple paternity could also occur in non-superfeting species. In fact, multiple paternity has been reported in species with superfetation (e.g., *H. formosa*) (Soucy & Travis, 2003) as well as in species without superfetation (*Gambusia holbrooki*, *Poecilia latipinna*, and *P. reticulata*) (Kelly, Godin, & Wright, 1999; Travis, Trexler, & Mulvey, 1990; Zane, Nelson, Jones, & Avise, 1999). However, our evidence of higher prevalence of spermathecae in superfetating species suggests that these processes are more important and more frequent in the reproduction of species without apparent signs of pre-copulatory sexual selection, as is the case of the majority of species that exhibit superfetation (Macías-García & González-Zuarth, 2005; Pollux et al., 2014).

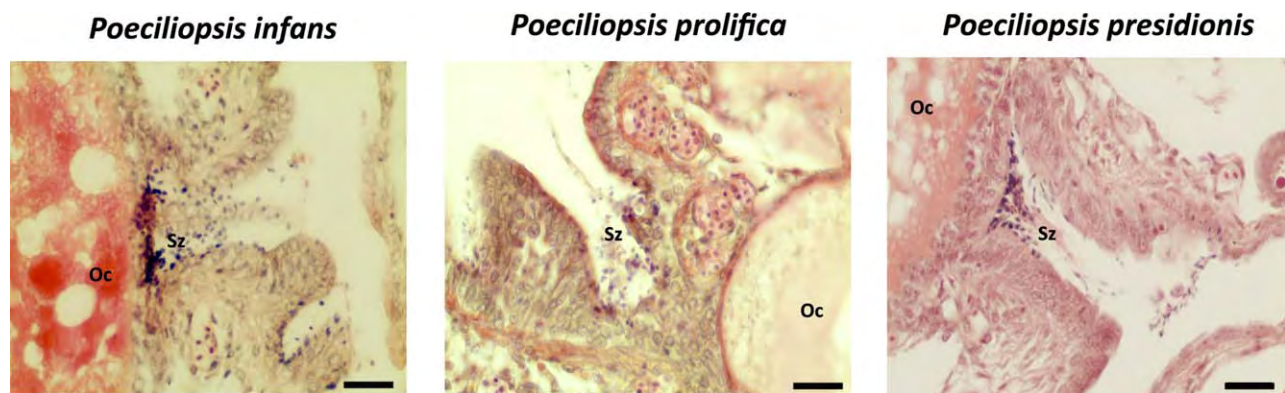


FIGURE 6 Sections of ovarian tissue stained using the hematoxylin-eosin technique that show spermatozoa (Sz) within folds of the ovarian epithelium observed in *P. infans*, *P. prolifica*, and *P. presidionis*. Heads of spermatozoa within the ovarian folds stain blue. Scale bar = 10 µm. Oocytes (Oc)

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AUTHOR CONTRIBUTIONS

C.O.T. and J.J.Z.V. designed the study and conducted fieldwork. C. O.T., M.V.S.C., and N.A.M.M. processed and analyzed all the samples. C.O.T. and J.J.Z.V. conducted all comparative analyses. All authors wrote the manuscript and gave final approval for publication.

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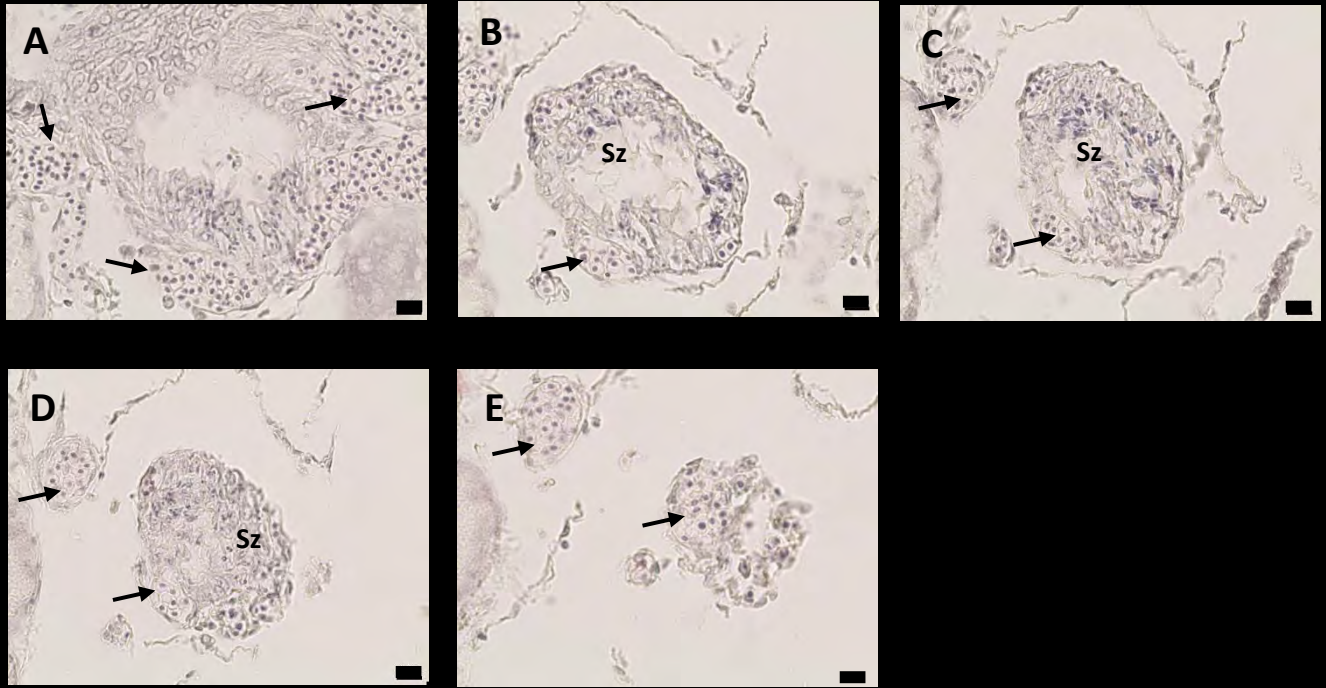
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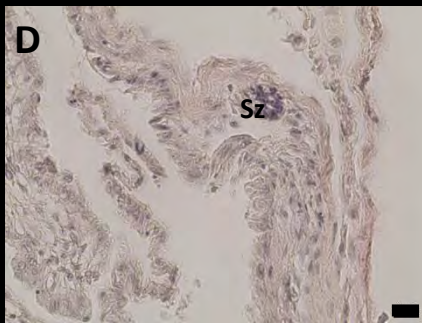
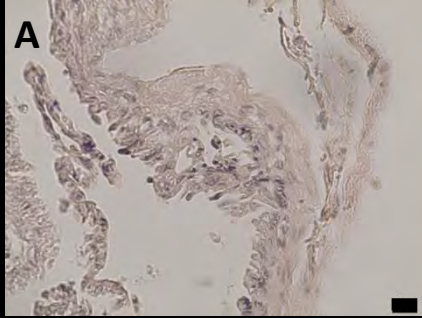
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Fig. S1. Hematoxylin-eosin stained serial sections of spermathecae for each study species. For *Poeciliopsis gracilis* we show spermathecae from two different females. Heads of spermatozoa (Sz) within spermathecae stain blue. Arrows point out erythrocytes. Scale bar = 10 μ m.

Gambusia panuco



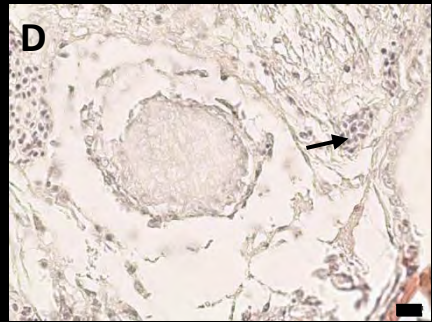
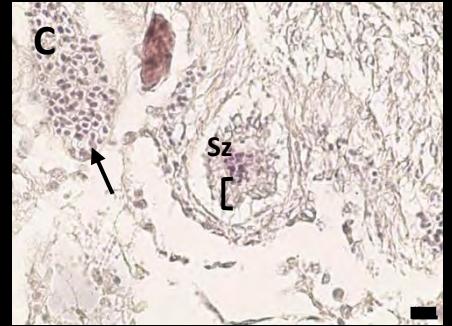
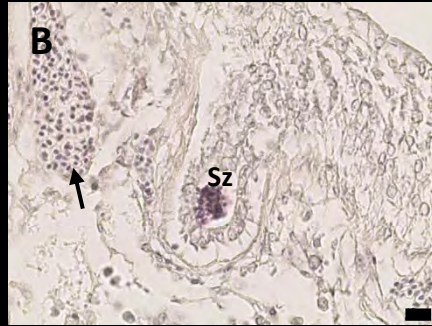
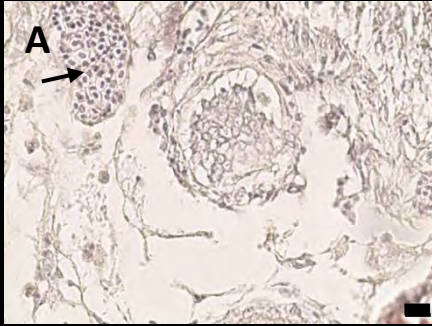
Xiphophorus hellerii



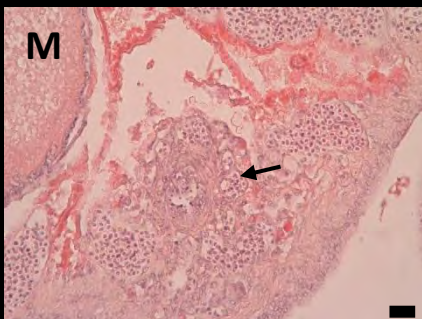
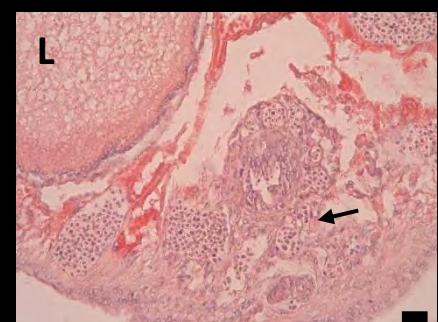
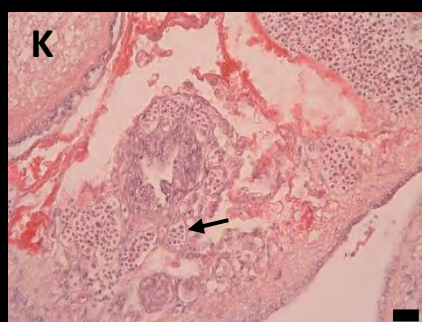
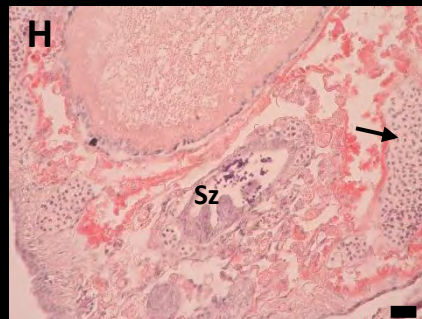
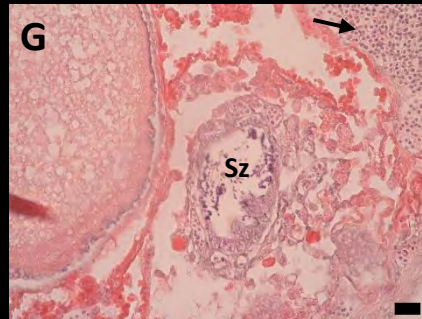
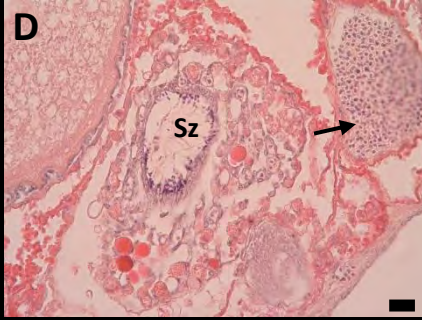
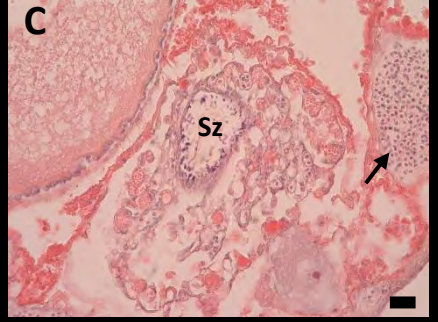
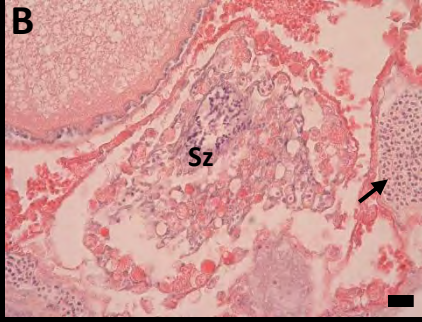
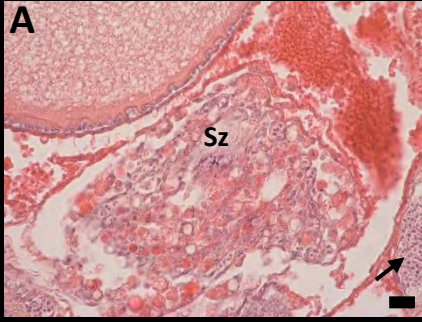
Poecilia mexicana



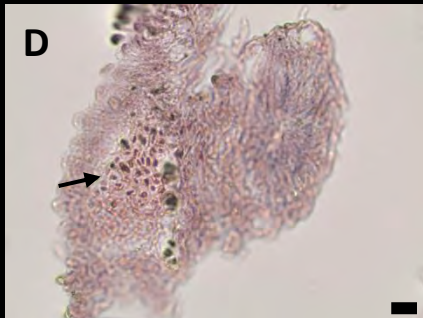
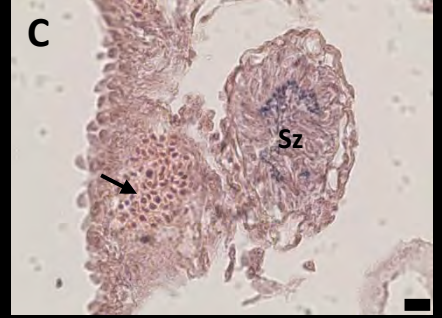
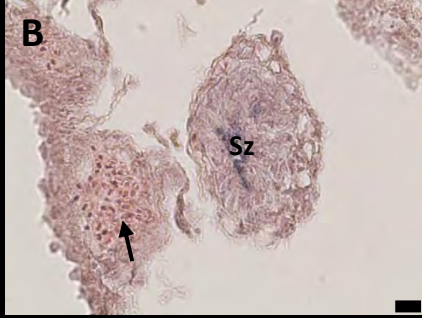
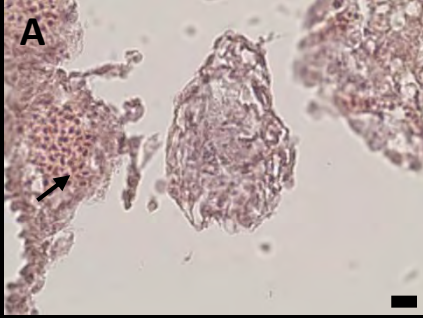
Poeciliopsis prolifica



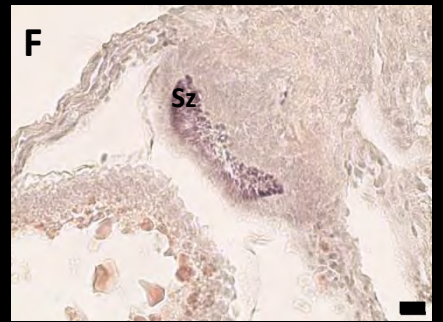
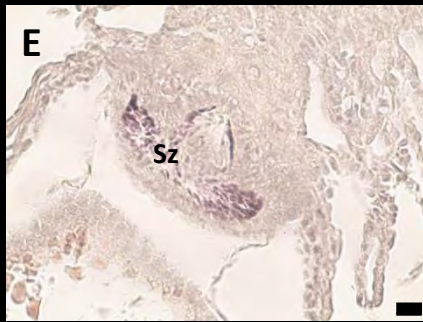
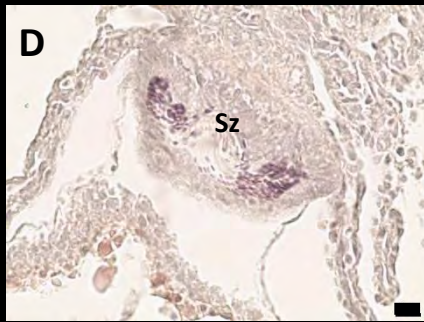
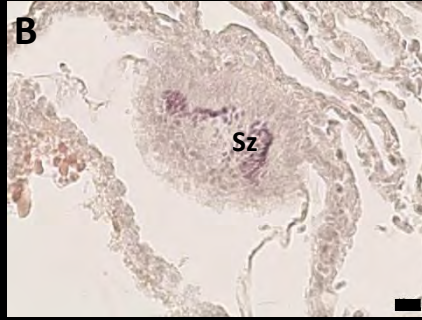
Poeciliopsis infans



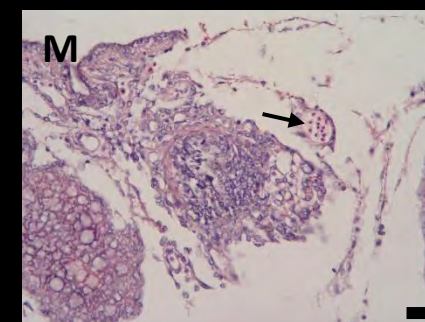
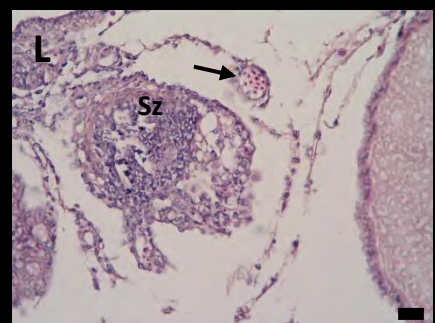
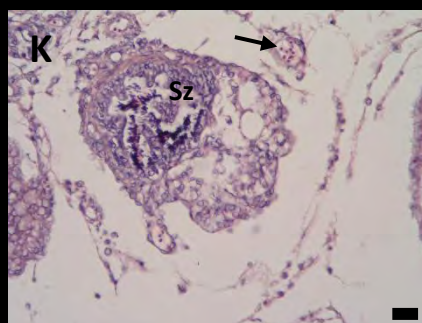
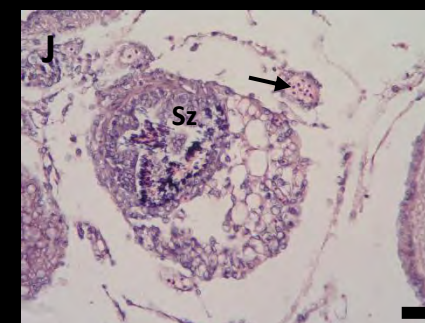
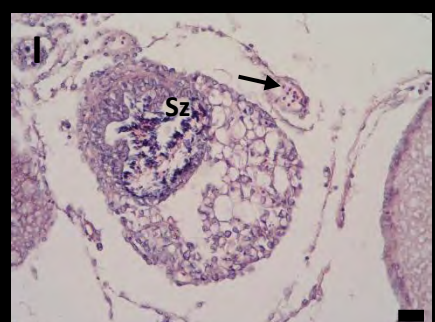
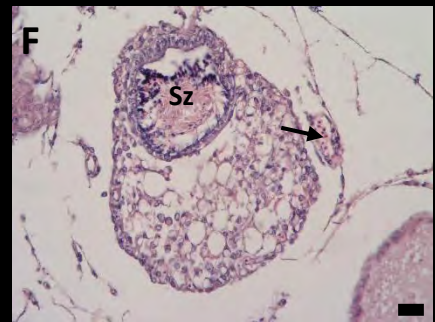
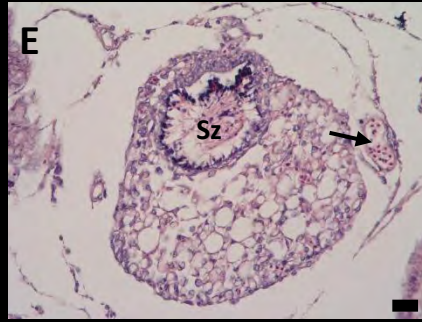
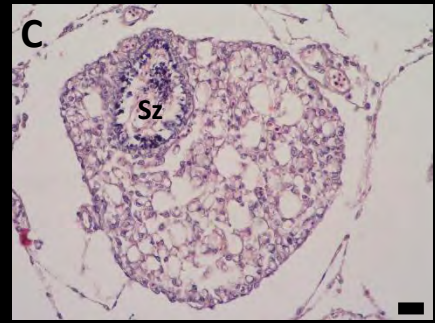
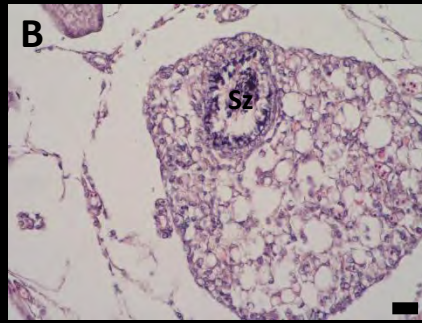
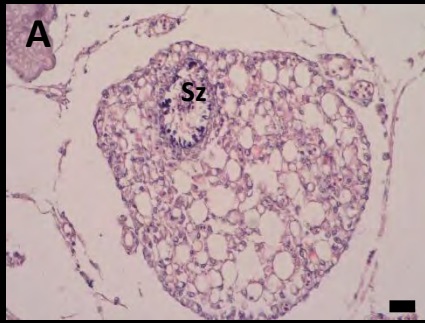
Pociliopsis viriosa



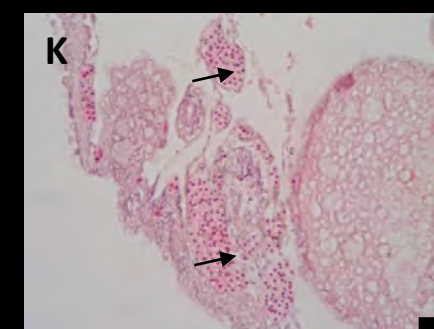
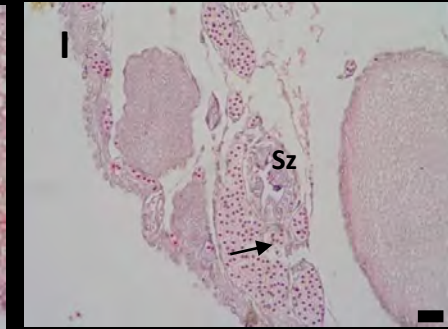
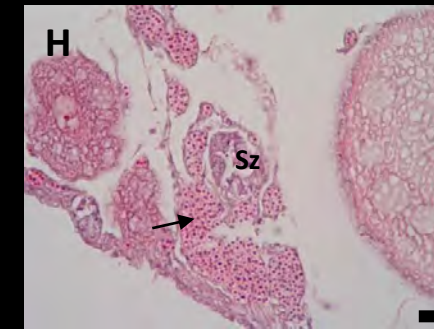
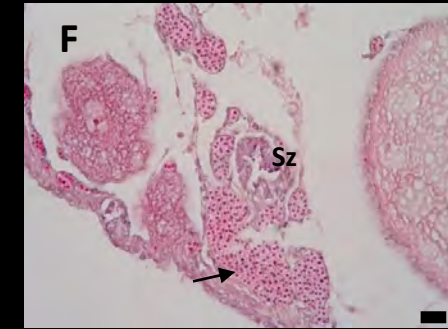
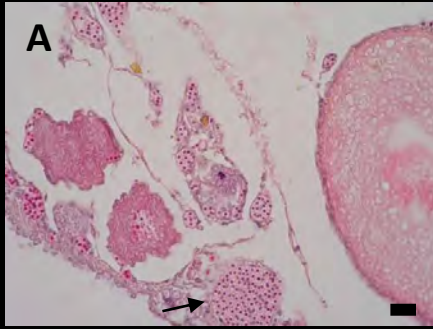
Poeciliopsis presidionis



Poeciliopsis gracilis - female 1



Poeciliopsis gracilis – female 2



Heterandria formosa

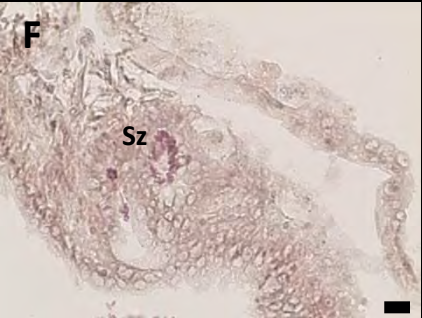
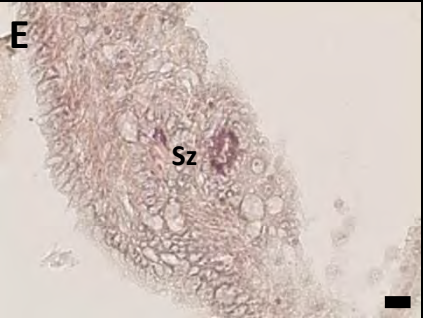
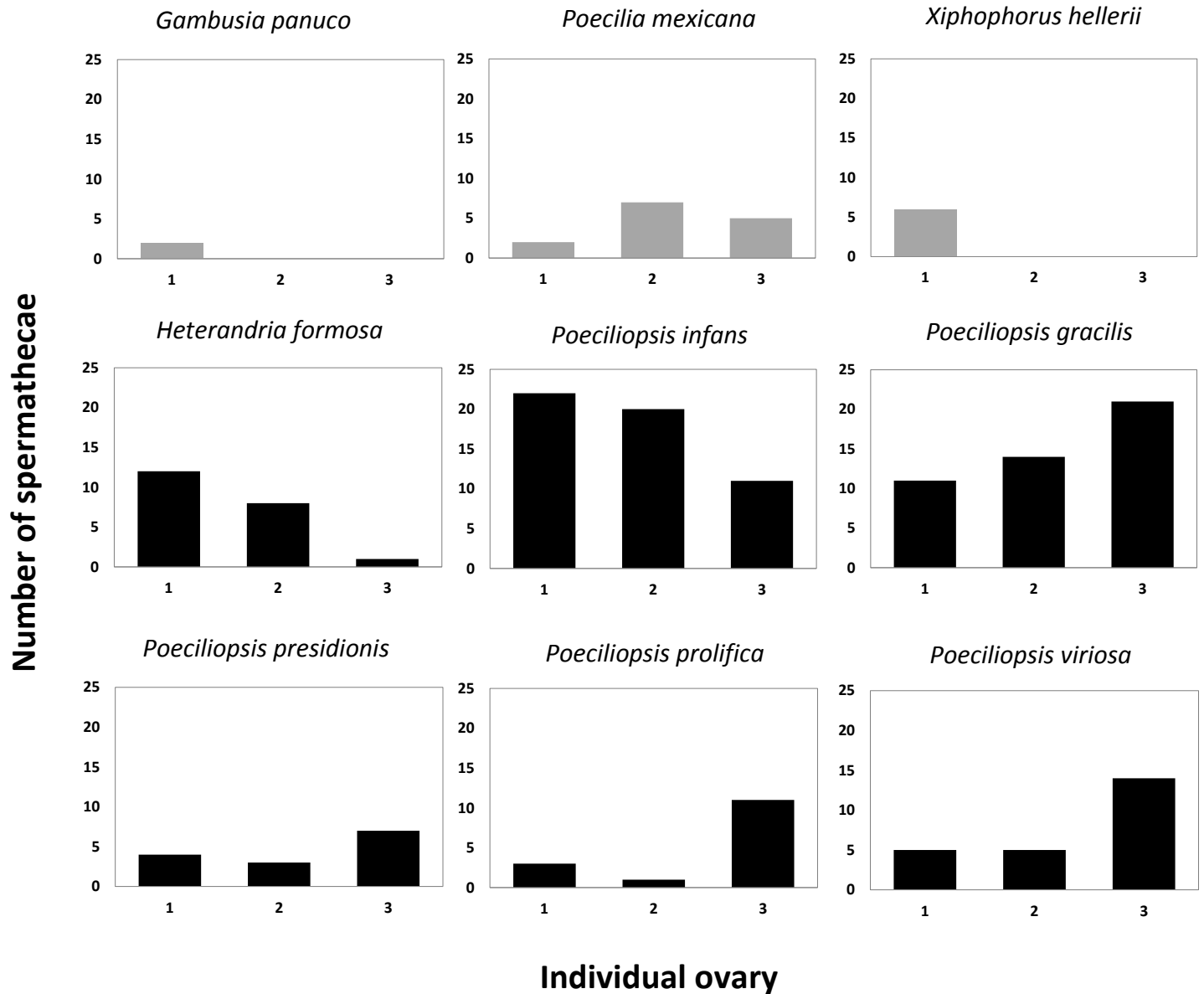
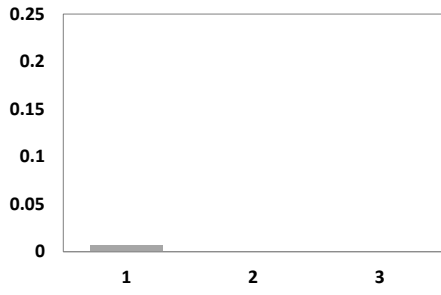


Fig. S2. Variation among individual ovaries in total number, total volume, and average volume of spermathecae for nine of the 12 species that we examined. Total and average volume of spermathecae are expressed as proportions of total ovary volume. Grey bars correspond to species without superfetation and black bars correspond to species with superfetation. We did not find spermathecae in the remaining three species (*Poecilia butleri*, *Priapella intermedia*, and *Pseudoxiphophorus bimaculatus*).

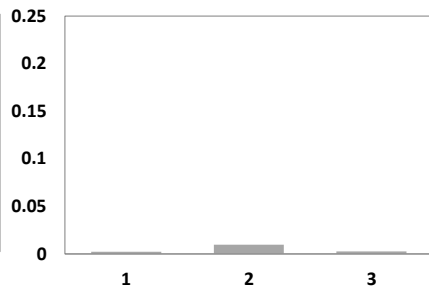


Relative volumen of all spermathecae

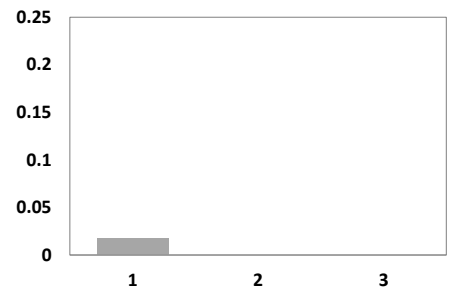
Gambusia panuco



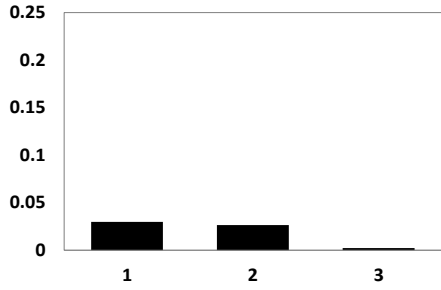
Poecilia mexicana



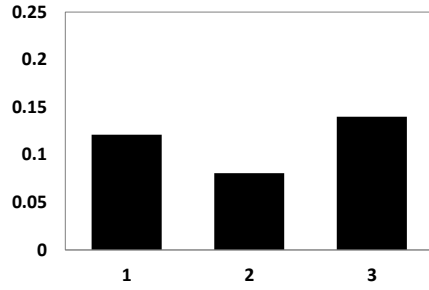
Xiphophorus hellerii



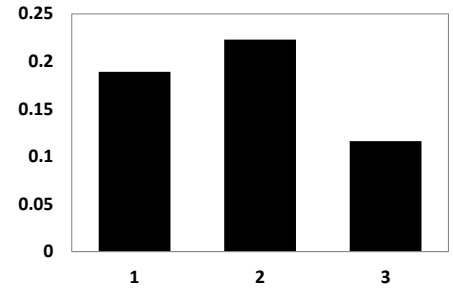
Heterandria formosa



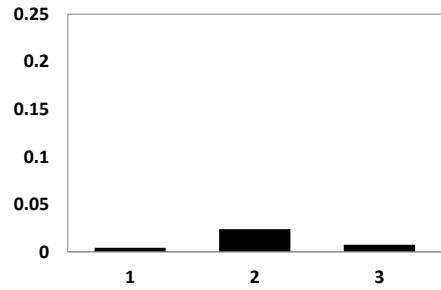
Poeciliopsis infans



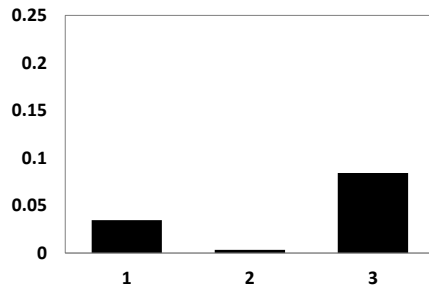
Poeciliopsis gracilis



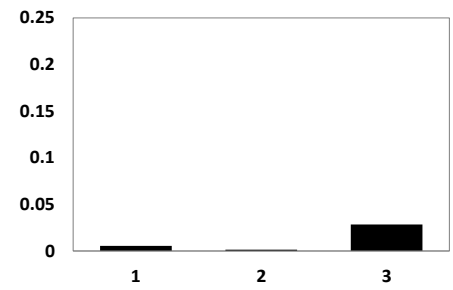
Poeciliopsis presidionis



Poeciliopsis prolifica

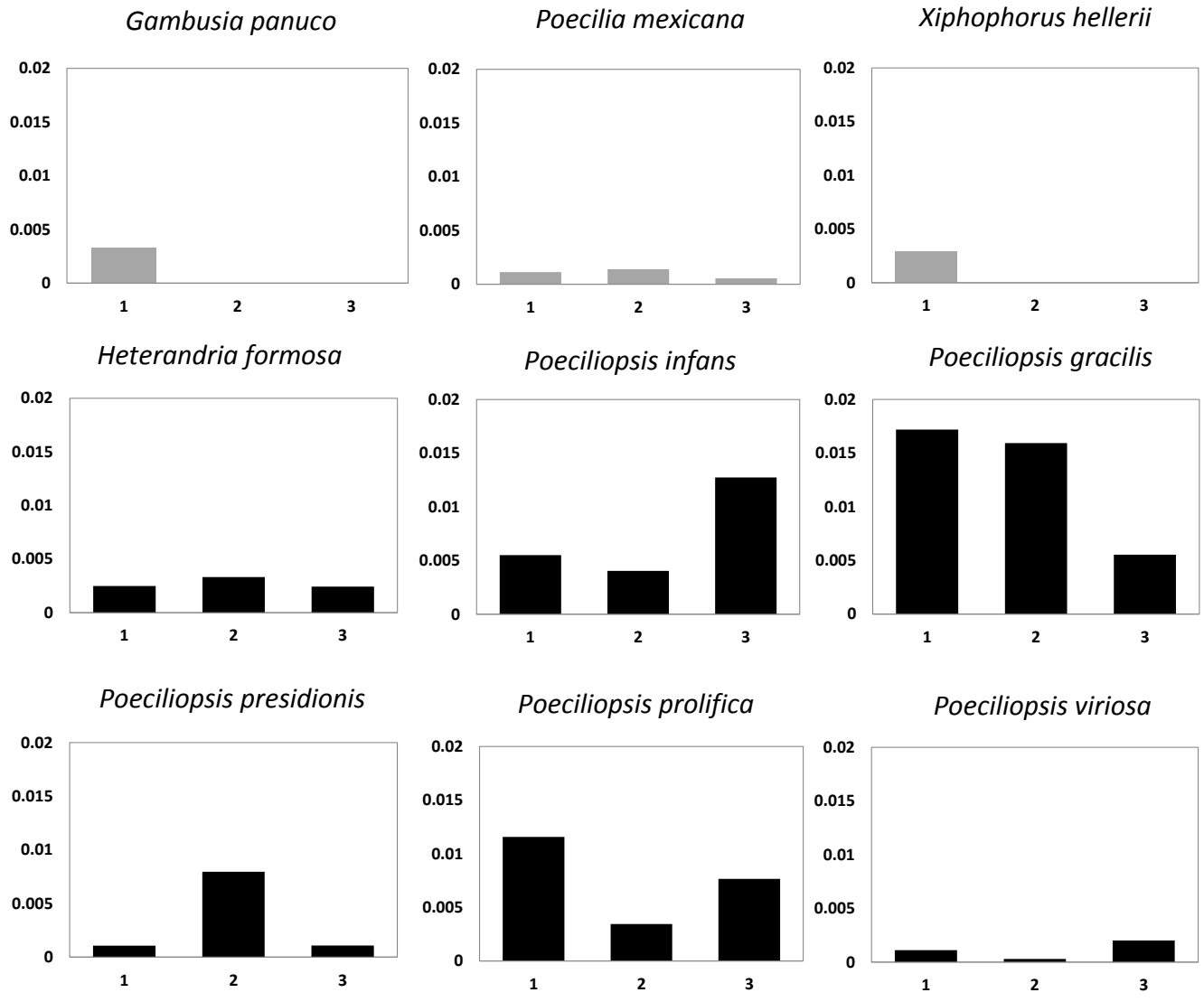


Poeciliopsis viriosa



Individual ovary

Relative volumen of each spermatheca



Individual ovary

CAPÍTULO III

**Have superfetation and matrotrophy facilitated the
evolution of larger offspring in poeciliid fishes?**

**Claudia Olivera-Tlahuel, Alison G. Ossip-Klein, Héctor S.
Espinosa-Pérez, and J. Jaime Zúñiga-Vega**

Biological Journal of the Linnean Society, 2015, 116: 787-804.



Have superfetation and matrotrophy facilitated the evolution of larger offspring in poeciliid fishes?

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Superfetation is the ability of females to simultaneously carry multiple broods of embryos, with each brood at a different developmental stage. Matrotrophy is the post-fertilization maternal provisioning of nutrients to developing embryos throughout gestation. Several studies have demonstrated that, in viviparous fishes, superfetation and matrotrophy have evolved in a correlated way, such that species capable of bearing several simultaneous broods also exhibit advanced degrees of post-fertilization provisioning. The adaptive value of the concurrent presence of both reproductive modes may be associated with the production of larger newborns, which in turn may result in enhanced offspring fitness. In this study, we tested two hypotheses: (1) species with superfetation and moderate or extensive matrotrophy give birth to larger offspring compared with species without superfetation or matrotrophy; (2) species with higher degrees of superfetation and matrotrophy (i.e. more simultaneous broods and increased amounts of post-fertilization provisioning) give birth to larger offspring compared with species with relatively low degrees of superfetation and matrotrophy (i.e. fewer simultaneous broods and lesser amounts of post-fertilization provisioning). Using different phylogenetic comparative methods and data on 44 species of viviparous fishes of the family Poeciliidae, we found a lack of association between offspring size and the combination of superfetation and matrotrophy. Therefore, the concurrent presence of superfetation and moderate or extensive matrotrophy has not facilitated the evolution of larger offspring. In fact, these traits have evolved differently. Superfetation and matrotrophy have accumulated gradual changes that largely can be explained by Brownian motion, whereas offspring size has evolved fluidly, experiencing changes that probably resulted from selective responses to the local conditions. © 2015 The Linnean Society of London, *Biological Journal of the Linnean Society*, 2015, **116**, 787–804.

ADDITIONAL KEYWORDS: lecithotrophy – offspring size – phylogenetic comparative methods – Poeciliidae – simultaneous broods – viviparous fishes.

INTRODUCTION

Superfetation is the ability of females to simultaneously carry multiple broods of embryos at different developmental stages (Turner, 1937, 1940; Scrimshaw, 1944). This mode of reproduction involves the occurrence of fertilization and development of a new brood before a preceding brood is born and can be

clearly identified by the presence of groups of embryos within a single female that are in discretely different and well-separated stages of development (Pires *et al.*, 2011a; Roellig *et al.*, 2011). Superfetation is a relatively rare phenomenon that occurs in some viviparous species of three phylogenetically distant families of fishes: Clinidae (Gunn & Thresher, 1991), Zenarchopteridae (Reznick, Meredith & Collette, 2007a) and Poeciliidae (Turner, 1937, 1940; Scrimshaw, 1944; Schultz, 1961; Thibault & Schultz,

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1978; Pollux *et al.*, 2009, 2014). Some species of the bivalve family Sphaeriidae exhibit brood masses that contain developmentally discrete subsets of embryos, with each subset encapsulated in a different brood sac. This phenomenon has been termed 'sequential brooding' but is equivalent to superfetation (Cooley & Foighil, 2000).

Within the fish family Poeciliidae, superfetation occurs in some genera such as *Poeciliopsis*, *Poecilia*, *Heterandria* and *Neoheterandria*, whereas it is absent in other genera (e.g. *Brachyrhaphis*, *Gambusia*, *Xhiphophorus* and *Alfaro*; Pires, Arendt & Reznick, 2010; Zúñiga-Vega, Macías-García & Johnson, 2010). The phylogenetic distribution of superfetation within the family Poeciliidae suggests that it has evolved independently multiple times (Reznick & Miles, 1989; Pollux *et al.*, 2009; Pires *et al.*, 2010; Meredith *et al.*, 2011). In addition, the number of simultaneous broods present within females (i.e. degree of superfetation) varies widely among superfetating species. For example, *Heterandria formosa* exhibit a high degree of superfetation (females can carry up to eight different broods), whereas *Poecilia parae* exhibit a relatively low degree of superfetation (the maximum number of simultaneous broods is two; Travis *et al.*, 1987; Pires *et al.*, 2010). The multiple independent origins, along with the observed interspecific variation, suggest that superfetation might convey certain adaptive advantages that are not yet completely understood (Zúñiga-Vega *et al.*, 2010).

In addition to superfetation, the amount of nutrients that are transferred from mothers to embryos during development varies widely among viviparous species of the family Poeciliidae (Reznick, Mateos & Springer, 2002; Pollux *et al.*, 2009, 2014). Some species provide all the necessary nutrients for development before fertilization in the form of yolk without additional post-fertilization nutrient transfer. Hence, embryos lose mass during development due to metabolic costs (embryo mass decreases ~ 35% from fertilization to birth). This mode of maternal provisioning is called lecithotrophy (Wourms, 1981; Marsh-Matthews, 2011). In contrast, other species provide nutrients to developing embryos after fertilization through specialized morphological structures (i.e. placentas; Turner, 1940; Lombardi & Wourms, 1985; Mossman, 1991). This mode of maternal provisioning is called matrotrophy (Wourms, 1981; Marsh-Matthews, 2011). The relative amount of pre- and post-fertilization maternal provisioning varies widely among species (Pollux *et al.*, 2009, 2014). Incipient matrotrophy refers to species in which females provide only small amounts of nutrients to developing embryos after fertilization in addition to the yolk that was previously allocated to eggs. In such cases

embryo mass decreases only slightly (10–15%) or remains constant throughout development. Moderate matrotrophy refers to species in which embryo mass increases moderately throughout development, reflecting a relatively larger provisioning from the mother after fertilization, although still relying to some extent on pre-fertilization yolk allocation. Extensive matrotrophy refers to species in which yolk is absent from mature ova, the mother throughout gestation actively provides all the nutrients that are necessary for development, and the mass at birth is at least five times larger than the egg mass at fertilization (Reznick *et al.*, 2002; Marsh-Matthews, 2011; Pires *et al.*, 2011a). In some species, the increase in embryo size throughout development reaches 60 000% (*Poecilia branneri* and *P. bifurca*; Pires *et al.*, 2010). There exists a continuum in the modes of maternal provisioning among poeciliid species, from strict lecithotrophy to extensive matrotrophy (Pollux *et al.*, 2009, 2014).

Some studies have demonstrated an evolutionary link between superfetation and matrotrophy (Reznick & Miles, 1989; Pollux *et al.*, 2009, 2014; Trexler & DeAngelis, 2010; Meredith *et al.*, 2011). Most species with superfetation also exhibit some degree of post-fertilization maternal provisioning, although a few exceptions exist (e.g. *Priapichthys festae*, *Neoheterandria tridentiger*, *N. elegans* and *Poeciliopsis monacha* exhibit superfetation and lecithotrophy; Reznick & Miles, 1989; Pollux *et al.*, 2009). This association has led to the hypothesis that the evolution of one of these two traits has facilitated the evolution of the other (Pollux *et al.*, 2009, 2014; Trexler & DeAngelis, 2010; Meredith *et al.*, 2011). The apparent coevolution of these reproductive strategies may have important consequences in terms of reproductive output. Hence, the adaptive significance of matrotrophy and superfetation could be associated with their patterns of covariation with life-history traits, such as number and size of offspring. For instance, species with superfetation tend to produce smaller broods and do so more often than non-superfetating species (Reznick & Miles, 1989; Pollux *et al.*, 2009). This mode of reproduction allows females to give birth continuously, with the interval between broods being as short as a few days. In contrast, species without superfetation, which for the most part exhibit lecithotrophy or only incipient matrotrophy, tend to produce larger broods and give birth less frequently, with many days between broods (Fraser & Renton, 1940; Thibault & Schultz, 1978; Cheong *et al.*, 1984; Leips *et al.*, 2009). The production of larger broods in non-superfetating species could result in smaller offspring because viviparous females have limited space within their reproductive tracts and, hence, they should

experience the widely recognized trade-off between number and size of offspring (Smith & Fretwell, 1974; Brockelman, 1975; Stearns, 1976).

In contrast, superfetating species that exhibit certain amounts of post-fertilization nutrient transfer might be able to increase the size of individual offspring. When matrotrophic females transfer an amount of nutrients during development that results in at least a slight increase in embryo mass from fertilization to birth (i.e. moderate matrotrophy), early-stage embryos are smaller than late-stage embryos. In combination with superfetation, where smaller broods imply fewer full-term embryos at any given time, this amount of post-fertilization nutrient transfer should promote less space requirements within the reproductive tract because these females bear a mix of early (small) and late (large) embryos compared with non-superfetating females that, at some point, would bear a high number of large full-term embryos. Therefore, the additional body volume that may be available to females exhibiting superfetation and at least moderate matrotrophy could be used to produce larger newborns. The combined effects of superfetation and matrotrophy on the potential reduction of space requirements within the reproductive tract could be even stronger in those species that exhibit extensive matrotrophy because at fertilization the ova is remarkably smaller than late-stage embryos. Thus, superfetation and extensive matrotrophy promote the presence of very small early-stage embryos, few large-late stage embryos and, potentially, low ovarian volume. This in turn could more easily allow for increases in the size of individual neonates.

Producing large newborns may increase their probability of surviving to maturity, at least under particular ecological conditions (Reznick, 1982; Reznick, Bryga & Endler, 1990; Bashey, 2008; Gordon *et al.*, 2009; Riesch *et al.*, 2010a). Therefore, if the coevolution of superfetation and matrotrophy has facilitated the evolution of larger offspring in poeciliid fishes, then the adaptive significance of the combined presence of these two reproductive modes could be associated with enhanced offspring fitness.

In this study we used modern phylogenetic comparative methods to analyse the evolutionary relationship between offspring size and the combined presence of superfetation and matrotrophy in 44 species of viviparous fishes of the family Poeciliidae. We focused on testing two main hypotheses. (1) Species with superfetation and at least moderate matrotrophy give birth to larger offspring compared with species without superfetation, species with superfetation and lecithotrophy, and species with superfetation and incipient matrotrophy (after accounting for differences among species in the size of adult females).

(2) Species with higher degrees of superfetation and matrotrophy (i.e. more simultaneous broods and increased post-fertilization maternal provisioning) give birth to larger offspring compared with species with relatively low degrees of superfetation and matrotrophy (i.e. fewer simultaneous broods and less post-fertilization provisioning). In other words, offspring size should increase as the number of simultaneous broods and the amount of post-fertilization nutrient transfer increase. This hypothesis is based on the observation that superfetating species with more simultaneous broods and relatively high levels of matrotrophy tend to produce fewer newborns per brood compared with superfetating species with fewer simultaneous broods and lower levels of matrotrophy. For example, *Xenodexia ctenolepis* (dry weight at birth is over three times larger than unfertilized ova) and *Poecilia branneri* (dry weight at birth is over 86 times larger than unfertilized ova) produce on average 4.13 and 4.77 simultaneous broods, with 4.40 and 4.13 newborns per brood, respectively (Reznick *et al.*, 2007b; Pires *et al.*, 2010). In contrast, *Poeciliopsis monacha* (embryo dry weight decreases during development) and *P. tur-rubarensis* (embryo dry weight barely increases during development) produce on average 2 and 2.12 simultaneous broods, with 11.80 and 7.64 newborns per brood, respectively (Thibault & Schultz, 1978; Zúñiga-Vega, Reznick & Johnson, 2007). For those species with advanced degrees of superfetation and matrotrophy, more smaller broods probably imply very few large full-term embryos, more small embryos of earlier stages, less total space requirements within the reproductive tract and, therefore, the possibility of producing considerably large offspring.

MATERIAL AND METHODS

DATA COLLECTION

We conducted an extensive literature search to extract data on average standard length (mm) of reproductive females, embryo dry mass (mg) at the last stage of development [i.e. stage 11 according to Haynes (1995) or the equivalent stage in other classification methods of embryonic development], degree of superfetation (average number of simultaneous broods at different developmental stages) and matrotrophy index (MI) for species of the family Poeciliidae (Table 1). We considered dry mass at the last stage of development as an appropriate proxy for size (mass) at birth (Haynes, 1995). MI is a standard measure of the amount of post-fertilization maternal provisioning and is calculated as the dry mass of the offspring at birth divided by the dry mass of the egg

Table 1. Female size (standard length), offspring size (mass), degree of superfetation (average number of simultaneous broods per female) and matrotrophy index (MI) for 44 species of the fish family Poeciliidae

Species	Female size (mm)	Offspring size (mg)	Offspring size (residuals)	Superfetation (no. of broods)	MI	PC1	Reference(s)
<i>Brachyrhaphis rhabdophora</i>	37.11	1.64	-0.458	1.00	0.77	-0.623	Johnson & Belk (2001), Pollux <i>et al.</i> (2014)
<i>Brachyrhaphis episcopi</i>	30.08	2.43	0.823	1.00	0.83	-0.621	Jennions & Telford (2002)
<i>Brachyrhaphis holdridgei</i>	22.61	0.6	-0.485	1.00	0.66	-0.628	Pollux <i>et al.</i> (2014), This study
<i>Heterandria formosa</i>	18.88	0.58	-0.244	3.09	35.00	2.439	Schrader & Travis (2012), Leips & Travis (1999), Pollux <i>et al.</i> (2014)
<i>Poeciliopsis lucida</i>	32.5	0.71	-1.066	3.00	2.00	0.968	Thibault & Schultz (1978), Pollux <i>et al.</i> (2014)
<i>Poeciliopsis pleurospilus</i>	45.01	1.04	-1.612	1.35	0.50	-0.367	This study
<i>Poeciliopsis occidentalis</i>	34.20	1.66	-0.235	1.64	1.12	-0.114	Pollux <i>et al.</i> (2014), This study
<i>Poeciliopsis prolifica</i>	24.63	0.59	-0.636	3.30	5.40	1.343	Pires, McBride & Reznick (2007), Pollux <i>et al.</i> (2014)
<i>Poeciliopsis infans</i>	28.75	0.82	-0.694	1.79	1.05	-0.003	Frías-Alvarez <i>et al.</i> (2014), Molina-Moctezuma (2015)
<i>Poeciliopsis monacha</i>	32.5	1.26	-0.516	2.00	0.61	0.140	Thibault & Schultz (1978), Pollux <i>et al.</i> (2014)
<i>Poeciliopsis viriosa</i>	37.93	0.43	-1.726	2.23	0.93	0.329	Pollux <i>et al.</i> (2014), This study
<i>Poeciliopsis turneri</i>	48.5	3.39	0.496	3.00	41.40	2.642	Thibault & Schultz (1978), Pollux <i>et al.</i> (2014)
<i>Poeciliopsis gracilis</i>	30.34	0.92	-0.705	1.88	0.84	0.057	Frías-Alvarez <i>et al.</i> (2014), Molina-Moctezuma (2015)
<i>Poeciliopsis turrubarensis</i>	39.38	1.58	-0.677	2.12	1.05	0.251	Zúñiga-Vega <i>et al.</i> (2007)
<i>Poeciliopsis latidens</i>	39.71	1.04	-1.240	1.12	0.86	-0.526	Pollux <i>et al.</i> (2014), This study
<i>Poeciliopsis fasciata</i>	30.78	1.92	0.264	1.34	0.91	-0.355	Pollux <i>et al.</i> (2014), This study
<i>Poeciliopsis baenschi</i>	25.62	1.28	-0.015	1.64	0.98	-0.122	Molina-Moctezuma (2011), This study
<i>Heterophallus milleri</i>	23.76	1.22	0.055	1.00	0.74	-0.624	Riesch <i>et al.</i> (2011)
<i>Gambusia panuco</i>	26.86	1.16	-0.222	1.00	0.79	-0.622	A. Ader, J. J. Zúñiga-Vega & J. Johnson, unpubl. data
<i>Gambusia vittata</i>	24.53	1.16	-0.059	1.00	1.29	-0.601	Weldele, Zúñiga-Vega & Johnson (2014)
<i>Gambusia sexradiata</i>	24.19	1.81	0.615	1.00	1.74	-0.582	Riesch <i>et al.</i> (2010a), R. Riesch, unpubl. data
<i>Gambusia eurystoma</i>	23.63	5.74	4.584	1.00	0.76	-0.624	Riesch <i>et al.</i> (2010a), R. Riesch, unpubl. data
<i>Gambusia hubbsi</i>	28.39	3.22	1.731	1.00	0.86	-0.619	Riesch <i>et al.</i> (2013)
<i>Gambusia yucatanana</i>	23.45	0.8	-0.344	1.00	0.53	-0.633	This study
<i>Gambusia holbrooki</i>	31.7	0.97	-0.750	1.00	0.64	-0.629	Meffe (1990), Pollux <i>et al.</i> (2014)
<i>Gambusia affinis</i>	35	1.9	-0.051	1.00	0.62	-0.629	Stearns (1983), Pollux <i>et al.</i> (2014)

Table 1. Continued

Species	Female size (mm)	Offspring size (mg)	Offspring size (residuals)	Superfetation (no. of broods)	MI	PC1	Reference(s)
<i>Gambusia speciosa</i>	28.58	0.94	-0.562	1.00	0.45	-0.637	This study
<i>Gambusia aurata</i>	23.82	0.66	-0.510	1.00	0.82	-0.621	This study
<i>Belonesox belizanus</i>	99.4	6.9	0.450	1.00	0.70	-0.626	Turner & Snelson (1984), Pollux <i>et al.</i> (2014)
<i>Pseudoxiphophorus jonesii</i>	48.10	4.36	1.494	1.00	0.65	-0.628	This study
<i>Priapella chamulae</i>	30.3	2.31	0.688	1.00	0.71	-0.626	Riesch <i>et al.</i> (2012)
<i>Priapella olmecae</i>	44.38	2.96	0.356	1.00	0.76	-0.623	This study
<i>Poecilia mexicana</i>	41.63	3.26	0.846	1.00	0.63	-0.629	Riesch, Plath & Schlupp (2010b), Pollux <i>et al.</i> (2014)
<i>Poecilia sulphuraria</i>	26.4	3.73	2.380	1.00	0.69	-0.627	Riesch <i>et al.</i> (2010a)
<i>Poecilia butleri</i>	41.77	1.64	-0.784	1.00	2.30	-0.558	Zúñiga-Vega <i>et al.</i> (2011)
<i>Poecilia</i> (<i>Micropoecilia</i>) <i>bifurca</i>	14.45	0.48	-0.035	1.20	55.05	1.836	Pires <i>et al.</i> (2010), Pollux <i>et al.</i> (2014)
<i>Poecilia</i> (<i>Micropoecilia</i>) <i>branneri</i>	19.16	0.62	-0.224	4.77	86.41	5.915	Pires <i>et al.</i> (2010), Pollux <i>et al.</i> (2014)
<i>Poecilia</i> (<i>Micropoecilia</i>) <i>parae</i>	23.33	0.67	-0.465	1.03	6.75	-0.346	Pires <i>et al.</i> (2010), Pollux <i>et al.</i> (2014)
<i>Poecilia</i> (<i>Micropoecilia</i>) <i>picta</i>	21.3	0.75	-0.243	1.00	0.78	-0.623	Pires <i>et al.</i> (2010), Pollux <i>et al.</i> (2014)
<i>Poecilia</i> (<i>Acanthophaelus</i>) <i>reticulata</i>	17.8	0.86	0.111	1.00	0.66	-0.628	Pires <i>et al.</i> (2010), Pollux <i>et al.</i> (2014)
<i>Poecilia</i> (<i>Acanthophaelus</i>) <i>wingei</i>	27.1	1.86	0.461	1.00	0.84	-0.620	Pires <i>et al.</i> (2010), Pollux <i>et al.</i> (2014)
<i>Phalloceros caudimaculatus</i>	29.62	0.8	-0.775	1.00	2.14	-0.565	Arias & Reznick (2000), Pollux <i>et al.</i> (2014)
<i>Phalloceros anisophallos</i>	29.9	0.79	-0.804	1.00	2.80	-0.537	Almeida-Silva & Mazzoni (2014)
<i>Xenodexia ctenolepis</i>	47	3.57	0.781	4.13	3.38	1.896	Reznick <i>et al.</i> (2007b), Pollux <i>et al.</i> (2014)

Offspring size corresponds to the dry mass at the last stage of development. Given that larger species produce significantly larger offspring, we also show the residuals from a linear regression between female length and offspring mass. The column PC1 shows the species-specific scores in the first principal component that represents a combined measure of superfetation and matrotrophy.

at fertilization (Wourms, Grove & Lombardi, 1988; Trexler, 1997; Reznick *et al.*, 2002; Marsh-Matthews, 2011). Thus, MI represents the change in embryonic dry mass during development, with an MI value higher than 1 indicating an increase in embryo mass and lower than 1 indicating a decrease in embryo mass. An MI equal to 1 indicates that embryo mass remains constant from fertilization to parturition. When information was available for more than one

population of a single species we used average values as input in our comparative analyses.

For some species (i.e. *Brachyrhaphis rhabdophora*, *B. episcopi*, *Gambusia vittata*, *G. panuco*, *Heterandria formosa*, *Poecilia butleri*, *Poeciliopsis baenschii*, *P. gracilis*, *P. infans* and *P. turrubarensis*), the authors who have published life-history descriptions provided us with their raw data, and we conducted a regression between stage of development and

log-transformed embryo dry mass. From this regression, we estimated the dry mass at the last developmental stage as a proxy for size at birth. In addition, for three of these species (*B. episcopi*, *G. panuco* and *P. baenschi*), we also used this regression to estimate the dry mass at fertilization [stage 4 according to Haynes (1995) or the equivalent stage in other classification methods] and then calculated an MI value as explained above (for most species MI values were available in Pollux *et al.*, 2014).

In addition, we quantified the same variables (standard length of reproductive females, dry mass of embryos at the last stage of development and degree of superfetation) from preserved females of the following species: *Brachyrhaphis holdridgei*, *Poeciliopsis fasciata*, *P. latidens*, *P. occidentalis*, *P. pleurospilus*, *P. viriosa*, *Gambusia yucatanana*, *G. speciosa*, *G. aurata*, *Pseudoxiphophorus jonesii* and *Priapella olmecae* (Table 1). These specimens were preserved in ethanol at the National Collection of Fishes (Instituto de Biología, Universidad Nacional Autónoma de México). We dissected 20 reproductive females per species. Before dissection, we measured the standard length (mm) of all females with a digital caliper. Upon dissection, we removed the ovary and separated embryos into distinct broods based on stage of development (according to Haynes, 1995). We quantified the degree of superfetation as the number of broods at different developmental stages within each female. We dried each brood of embryos at 55 °C overnight, and then weighed embryos to the nearest 0.01 mg. To obtain an estimate of the dry mass at the last stage of development, we ran a regression between developmental stage and log-transformed embryo dry mass. In addition, for those species whose MI was not available in the literature (*G. yucatanana*, *G. speciosa*, *G. aurata*, *Poeciliopsis pleurospilus*, *Pseudoxiphophorus jonesii* and *Priapella olmecae*) we also used this regression to estimate dry mass at fertilization and then calculated an MI as explained above. In total, we obtained data for 44 poeciliid species (Table 1).

ACCOUNTING FOR POTENTIAL CONFOUNDING FACTORS

Given that larger species could produce larger offspring and/or have higher degrees of superfetation or matrotrophy simply as a consequence of a larger body size (Reznick & Miles, 1989), we conducted linear regressions (using species-specific average values) between female length and offspring mass, between female length and superfetation, and between female length and MI. Offspring mass increased significantly with species-specific female size ($F = 31.7$, d.f. = 1, 42, $P < 0.0001$). Therefore, in all comparative analyses we used the residuals of

this regression as size-corrected estimates of offspring size. In contrast, neither superfetation nor matrotrophy were affected by species-specific female size (superfetation: $F = 0.003$, d.f. = 1, 42, $P = 0.96$; matrotrophy: $F = 1.7$, d.f. = 1, 42, $P = 0.19$) and therefore we used the raw (unadjusted) values of number of simultaneous broods and MI.

Another potential confounding factor is the way in which specimens were originally preserved. Some species were preserved in formalin, whereas others were preserved in ethanol (Supporting Information, Table S1). Ethanol can extract lipids from the tissues and therefore might bias the estimates of offspring mass and MI (Shields & Carlson, 1996). We tested this potential confounding effect by means of general linear models using offspring dry mass and MI as response variables, the preservation method (formalin or ethanol) as a main factor and female length as a covariate. We also included in these models the interaction between preservation method and female length. Neither the preservation method nor its interaction with female length significantly affected offspring dry mass (preservation method: $F = 0.45$, d.f. = 1, 40, $P = 0.51$; interaction: $F = 0.10$, d.f. = 1, 40, $P = 0.76$) or MI (preservation method: $F = 0.14$, d.f. = 1, 40, $P = 0.71$; interaction: $F = 0.31$, d.f. = 1, 40, $P = 0.58$). Again here, female length had a highly significant effect on offspring mass ($F = 12.07$, d.f. = 1, 40, $P = 0.001$), whereas it did not affect MI ($F = 1.29$, d.f. = 1, 40, $P = 0.26$). Therefore, most of the variation among species in offspring mass was explained by the size of adult females, with no detectable effect of the preservation method on either offspring mass or MI.

To further examine the potential confounding effect of the preservation method, we implemented all our comparative analyses using only the 33 species that were preserved in ethanol (Table S1). All results were qualitatively similar to those obtained using the complete data set (44 species). Therefore, we report here the results obtained with the 44 species and those from the subset of 33 species in the Supporting Information.

A COMBINED MEASURE OF SUPERFETATION AND MATROTROPHY

We hypothesized that the combined effect of superfetation and at least moderate matrotrophy may allow for increases in size at birth in poeciliid fishes. Therefore, we conducted a principal components analysis on our species-specific values for degree of superfetation and MI. This analysis resulted in a single principal component with an eigenvalue > 1 (1.59) in which both superfetation and MI had high loadings (0.89 for both variables). Hence, we consid-

ered this principal component as a combined measure of superfetation and matrotrophy. Species with high positive scores corresponded to those with the highest degrees of superfetation and the largest amounts of post-fertilization maternal provisioning (e.g. *Poecilia branneri*: 4.77 simultaneous broods and MI = 86.41), whereas those with relatively high negative scores corresponded to lecithotrophic species without superfetation (e.g. *Gambusia speciosa*: lacks superfetation and MI = 0.45; Table 1). This combined measure of superfetation and matrotrophy was used as input in our comparative analyses.

PHYLOGENY AND BRANCH LENGTHS

We used a well-resolved phylogeny of the family Poeciliidae that was reconstructed by Pollux *et al.* (2014) using a large molecular data set (20 nuclear and eight mitochondrial markers). We pruned their topology to our genera and species of interest. Although this phylogenetic reconstruction did not include all the 44 species that we considered in our study, we used it as the main representation of the relationships among genera. However, it did not include the genus *Heterophallus*. Therefore, the phylogenetic position of *H. milleri* with respect to other genera was based on Doadrio *et al.* (2009). Phylogenetic relationships within genera, among those species that were not included in the topology of Pollux *et al.* (2014), were obtained from the following studies: Mojica, Meyer & Barlow (1997) (*Brachyrhaphis*), Lydeard, Wooten & Meyer (1995), Langerhans *et al.* (2012) (*Gambusia*), Ptacek & Breden (1998) (*Poecilia*), Meredith *et al.* (2010) (*Poecilia*, subgenus *Micropoecilia*), Morales-Cazan & Albert (2012) (*Poeciliopsis*) and Meyer, Schories & Scharl (2011) (*Priapella*). The composite phylogeny that we used for comparative analyses is shown in Fig. 1.

Phylogenetic comparative methods require information about the expected amount of evolutionary change along each branch of the phylogenetic tree (i.e. branch lengths; Harvey & Pagel, 1991). Given that we based our comparative analyses on a composite phylogeny, we lacked branch lengths in units of DNA sequence divergence. Therefore, we used a conservative approach that consisted of generating 1000 possible sets of branch lengths in units of time by means of computer simulation, assuming a standard branching process of speciation (Martins, 1996). We conducted each analysis on each of these 1000 phylogenies and report average results. For analyses for which the calculation of average results across 1000 phylogenies was not automated, we generated five sets of branch lengths and report the average results across those five phylogenies.

PHYLOGENETIC COMPARATIVE METHODS

We used the program COMPARE 4.6b (Martins, 2004) to implement different phylogenetic comparative methods. To compare offspring size (adjusted for female length) between species with different degrees of superfetation and matrotrophy we used the adaptation-inertia model of Hansen (1997). This comparative method accounts for phylogenetic relatedness when analysing the relationship between a phenotypic trait (e.g. offspring size) and a particular selective force. Hansen (1997) described this as a model of evolutionary change based on strong stabilizing selection in which phenotypes evolve towards an adaptive optimum. This model accounts for competing selective forces that might result in different optimal phenotypic values depending on the particular evolutionary context. We implemented this method twice, with each implementation representing a different hypothesis. First, we tested the hypothesis that superfetation accompanied by at least moderate matrotrophy would be enough to facilitate the evolution of larger offspring. For this purpose, we classified our species into one of the following two groups, which we considered alternative reproductive modes: (1) species with superfetation and at least moderate matrotrophy (i.e. all superfetating species with an MI > 1), and (2) species without superfetation, species with superfetation and lecithotrophy (MI < 0.8) and species with superfetation and incipient matrotrophy ($0.8 \leq \text{MI} \leq 1$). We treated these two groups as alternative evolutionary forces and estimated how much of the variation across species in offspring size can be explained by the relative time each species has evolved with either of these two reproductive modes (r^2).

In the second implementation we tested the hypothesis that the combination of superfetation and extensive matrotrophy would provide even further opportunities for the evolution of larger offspring. Thus, we classified our species into one of the following two groups (alternative reproductive modes): (1) species with superfetation and extensive matrotrophy (i.e. all superfetating species with an MI > 5), and (2) all other species. Again, we estimated the proportion of interspecific variance in offspring size that can be explained by these two alternative evolutionary forces (r^2).

This method also allowed us to estimate optimal values of offspring size for species that exhibit these different combinations of the presence/absence of superfetation and relative amount of matrotrophy. Based on our hypotheses, we expected a larger optimal size at birth for superfetating species that exhibit moderate or extensive matrotrophy compared with that for non-superfetating species, superfetating

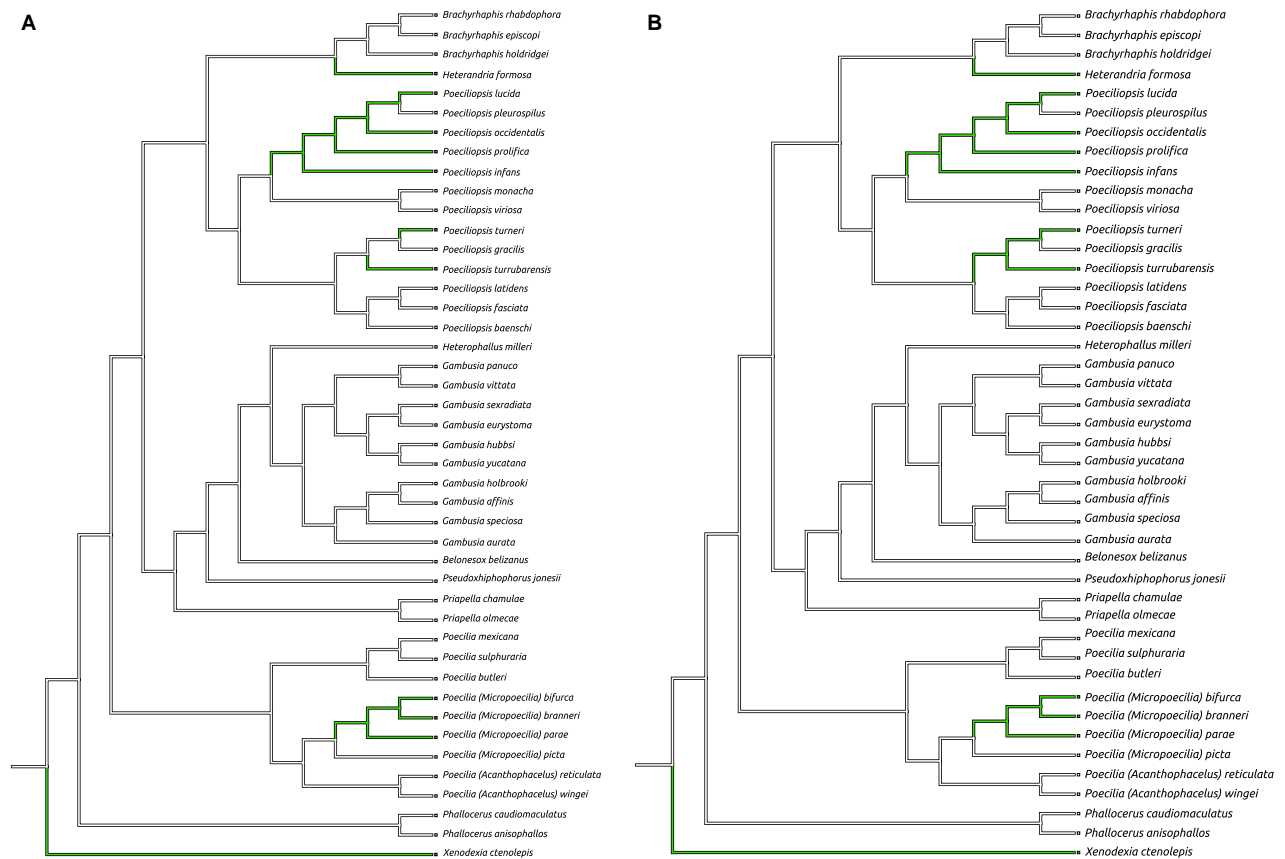


Figure 1. Composite phylogeny showing the relationships among 44 species of the family Poeciliidae as well as estimated ancestral states of the combined presence of superfetation and at least moderate matrotrophy (i.e. matrotrophy index > 1) based on parsimony. Green branches indicate presence of superfetation and at least moderate matrotrophy, whereas white branches indicate absence of superfetation, superfetation with lecithotrophy or superfetation with incipient matrotrophy. Both trees (A, reconstruction 1; B, reconstruction 2) were equally parsimonious.

species with lecithotrophy and superfetating species with incipient matrotrophy. Given that we adjusted our data to account for interspecific variation in female length, we actually expected a positive optimal value for superfetating species with moderate or extensive matrotrophy (a positive residual indicates a larger offspring size than that expected for a given female length) and a negative optimal value for non-superfetating species, superfetating species with lecithotrophy and superfetating species with incipient matrotrophy (a negative residual indicates a smaller offspring size than that expected for a given female length).

Hansen's (1997) method requires previous knowledge about the evolutionary history of the putative selective agent across the phylogeny. Thus, we reconstructed ancestral states for (1) the presence of superfetation with at least moderate matrotrophy and (2) the presence of superfetation with extensive matrotrophy using maximum likelihood and parsimony

approaches implemented in the program MES-QUITE 3.0 (Maddison & Maddison, 2009).

To estimate the magnitude of the relationship between offspring size and our combined measure of superfetation and matrotrophy we used different phylogenetic comparative methods. The first was Felsenstein's (1985) independent contrasts (FIC). This method solves the statistical problem of non-independence in the data due to shared ancestry and assumes that traits evolve along a phylogeny at a constant rate. Therefore, this mode of evolution can be described by Brownian motion, in which phenotypes diverge under random genetic drift. Hence, the expected phenotypic difference between sister species grows proportional to the time since they shared a common ancestor. From this method, we obtained a phylogenetically corrected correlation coefficient (r). We compared this with a Pearson correlation coefficient calculated using the uncorrected data (i.e. a non-phylogenetic approach referred to as TIPS;

Martins & Garland, 1991). The TIPS method assumes that traits adapt immediately to the local environment, leaving behind no trace of phylogenetic relationships.

We also used the phylogenetic generalized least squares method (PGLS) (Martins & Hansen, 1997), which uses an Ornstein–Uhlenbeck (OU) process as the underlying model of evolution. An OU process assumes that traits evolve under stabilizing selection. Similar to the FIC approach, PGLS estimates the relationship between two traits by means of a phylogenetically corrected correlation coefficient, r . In addition, PGLS estimates an additional parameter, α , which may vary between 0 and 15.5 (in COMPARE). When α is small, the rate of local adaptation is very slow and phenotypes appear to evolve by Brownian motion, which results in high phylogenetic signal. In contrast, when α is large the rate of adaptive change is fast and phenotypes evolve without a strong phylogenetic signal. Given that PGLS is a least squares regression model, we used our combined measure of superfetation and matrotrophy as predictor variable (X) and size-corrected offspring size as response variable (Y).

Finally, we used the phylogenetic mixed model (PMM) (Lynch, 1991; Housworth, Martins & Lynch, 2004), which also estimates the correlation between two traits after accounting for phylogenetic relatedness among species. The PMM assumes that phenotypes are the result of a linear combination of gradually accumulated evolutionary changes that have occurred along the phylogenetic history of the species and rapid evolutionary changes that probably resulted from adaptive responses to sudden changes in the environment. Thus, this model estimates the relative proportion of gradual Brownian-motion-like change (h^2 , defined as phylogenetic heritability) for both traits (offspring size and the combined measure of superfetation and matrotrophy). When h^2 approaches one, most of the interspecific variation in the phenotypes is explained by the phylogenetic history. When h^2 approaches zero, phenotypes show little trace of phylogenetic signal and evolutionary changes have occurred faster than expected by Brownian motion (Housworth *et al.*, 2004). Using PMM, we also calculated h^2 separately for degree of superfetation and MI.

RESULTS

ANCESTRAL RECONSTRUCTIONS OF SUPERFETATION AND MATROTROPHY

The parsimony analysis resulted in two equally parsimonious ancestral reconstructions of superfetation and at least moderate matrotrophy, referred to

herein as reconstructions 1 and 2 (Fig. 1). In reconstruction 1, the combination of superfetation and MI values > 1 originated in several occasions: once within the subgenus *Micropoecilia*, three times within the genus *Poeciliopsis*, with one loss (in *P. pleurospilus*), and once in *Hererandria formosa* (Fig. 1A). Reconstruction 2 was quite similar, with the only difference being two origins (instead of three) within *Poeciliopsis* and two losses (in *P. pleurospilus* and *P. gracilis*) (Fig. 1B). In both reconstructions we found uncertainty in whether the common ancestor of the family exhibited superfetation and at least moderate matrotrophy or not. If it did, then *Xenodexia ctenolepis* retained the ancestral state. If not, an additional independent origin occurred in *X. ctenolepis*.

The maximum likelihood reconstruction also indicated an origin of superfetation with at least moderate matrotrophy within the subgenus *Micropoecilia*, two origins within *Poeciliopsis* and two losses (in *P. pleurospilus* and *P. gracilis*), one additional origin in *H. formosa*, and high uncertainty with respect to the presence or absence of this reproductive mode in the common ancestor of the entire family (proportional likelihoods: 0.53 for superfetation and at least moderate matrotrophy and 0.47 for any other strategy) (Fig. 2). Again, *X. ctenolepis* represents either an additional independent origin or retention of an ancestral state. Note that the proportional likelihoods of the ancestral nodes in the group comprising *P. turneri*, *P. gracilis* and *P. turrubarensis* indicated high uncertainty with respect to the presence or absence of superfetation and MI values > 1 (Fig. 2). Given all this uncertainty, we decided to implement Hansen's (1997) method using both parsimony reconstructions 1 and 2 (the maximum likelihood reconstruction summarized these two) and assuming either presence or absence of superfetation and at least moderate matrotrophy in the most ancestral node. All results were qualitatively similar. Thus, for simplicity we report here only the results using reconstruction 1 and assuming that the ancestor of the family exhibited both superfetation and an MI > 1 .

Regarding the ancestral reconstructions of superfetation and extensive matrotrophy, the parsimony analysis resulted in a single most parsimonious tree and identified four evolutionary origins (Fig. 3A). The first occurred within the subgenus *Micropoecilia*, particularly in the common ancestor of *Poecilia* (*Micropoecilia*) *bifurca*, *P. (M.) branneri*, and *P. (M.) parae*, the second in *Poeciliopsis turneri*, the third in *P. prolifica* and the fourth in *Heterandria formosa*. The maximum likelihood reconstruction gave identical results with minimum uncertainty with respect to the presence of superfetation and extensive matrotrophy in ancestral nodes (Fig. 3B).



Figure 2. Composite phylogeny showing the relationships among 44 species of the family Poeciliidae as well as estimated ancestral states of the combined presence of superfetation and at least moderate matrotrophy (i.e. matrotrophy index > 1) based on maximum likelihood. Circles depict proportional likelihoods for the presence (green) and absence (white) of superfetation and at least moderate matrotrophy.

POTENTIAL EVOLUTIONARY RELATIONSHIPS BETWEEN OFFSPRING SIZE AND THE COMBINATION OF SUPERFETATION AND MATROTROPHY

Interspecific variation in offspring size could not be explained well by the evolution of superfetation and at least moderate matrotrophy (Hansen's $r^2 = 0.05$) (Table 2). Contrary to what we expected, the optimal offspring size (after adjusting for female length) for species with superfetation and MI values > 1 was -0.38 , whereas that for non-superfeting species, superfeting species with lecithotrophy and superfeting species with incipient matrotrophy was 0.16.

Similarly, the evolution of superfetation accompanied by extensive matrotrophy did not explain the interspecific variation in offspring size (Hansen's $r^2 = 0.01$) (Table 2). Again, the estimated optimal values were opposite to our prediction. The optimal offspring size for species with superfetation and MI

values > 5 was -0.26 , whereas that for all other species was 0.05.

The evolutionary relationship between degrees of superfetation and matrotrophy treated as a combined continuous variable (i.e. the species-specific scores in the first principal component) and offspring size was very weak, as indicated by the small, non-significant correlation coefficients obtained from FIC, PGLS and PMM (0.23, 0.07 and -0.21 , respectively; $P > 0.13$ in all cases) (Table 2). A non-phylogenetic approach also indicated a weak correlation (TIPS $r = -0.10$, $P = 0.50$). According to PGLS, the combination of superfetation and matrotrophy explained only 8% of the interspecific variation in offspring size ($r^2 = 0.08$). All these analyses conducted on the subset of 33 species that were preserved in ethanol yielded equivalent results (Supporting Information, Table S2).

PHYLOGENETIC SIGNAL IN SUPERFETATION, MATROTROPHY AND OFFSPRING SIZE

PGLS estimated a large value of the parameter α (15.4), which indicates that the species-specific offspring sizes have been pulled strongly towards optimal values. Therefore, the evolution of this trait cannot be explained very well by the phylogenetic history alone. Consistent with this result, the phylogenetic heritability of offspring size was low ($h^2 = 0.22$), retaining only weak evidence of the phylogeny. In contrast, the combination of superfetation and matrotrophy has tracked the phylogeny more closely, resulting in a remarkably high phylogenetic heritability ($h^2 = 0.86$) (Table 2). Separately, superfetation and MI values also had high phylogenetic heritabilities ($h^2 = 0.81$ and 0.87, respectively). We obtained similar estimates of phylogenetic signal for offspring size, superfetation and matrotrophy from the subset of 33 species that were preserved in ethanol (Table S2).

DISCUSSION

SUPERFETATION AND MATROTROPHY HAVE NOT FACILITATED THE EVOLUTION OF LARGER OFFSPRING

We have demonstrated here that the combination of superfetation and matrotrophy does not exhibit an evolutionary relationship with offspring size, regardless of whether superfetation and matrotrophy were treated as a dichotomous variable (presence or absence of superfetation and moderate or extensive matrotrophy) or as a continuous variable (degrees of superfetation and matrotrophy combined into a single measure). Thus, neither of our two hypotheses was supported by our results. In fact, the estimated

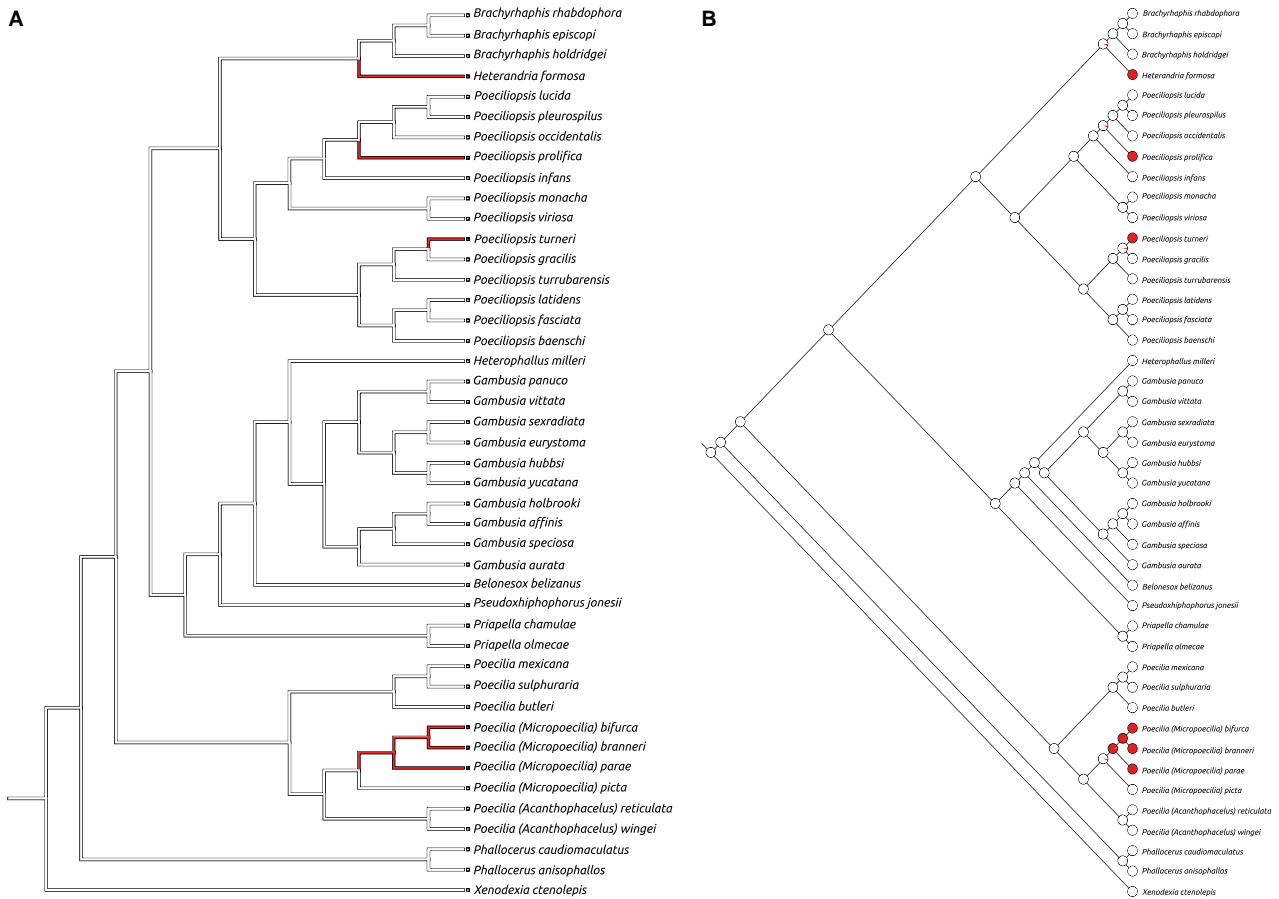


Figure 3. Composite phylogeny showing the relationships among 44 species of the family Poeciliidae as well as estimated ancestral states of the combined presence of superfetation and extensive matrotrophy (i.e. matrotrophy index > 5) based on (A) parsimony and (B) maximum likelihood. Red branches in A indicate presence of superfetation and extensive matrotrophy. Circles in B depict proportional likelihoods for the presence (red) and absence (white) of superfetation and extensive matrotrophy.

Table 2. Phylogenetic estimates of the magnitude of the evolutionary relationship between offspring size and the combination of superfetation and matrotrophy calculated using different phylogenetic comparative methods

Hansen's method					
SF and MI > 1	SF and MI > 5	FIC	TIPS	PGLS	PMM
$r^2 = 0.05$	$r^2 = 0.01$	$r = 0.23$	$r = -0.10$	$r = 0.07$ $r^2 = 0.08$ $\alpha = 15.4$	$r = -0.21$
					Offspring size $h^2 = 0.22$ SF and MI $h^2 = 0.86$ SF $h^2 = 0.81$ MI $h^2 = 0.87$

FIC, Felsenstein's independent contrasts; TIPS, non-phylogenetic approach; PGLS, phylogenetic generalized least squares method assuming an Ornstein-Uhlenbeck model of evolution; PMM, phylogenetic mixed model; SF, superfetation; MI, matrotrophy index; r , correlation coefficient; r^2 , proportion of the variation among species in offspring size explained by superfetation and matrotrophy; α , a measure of the strength of selection pulling offspring size towards optimal values; h^2 , phylogenetic heritability.

optimal sizes at birth for species with superfetation and moderate or extensive matrotrophy were smaller than those for species without superfetation, species with superfetation and lecithotrophy, and species with superfetation and incipient matrotrophy (although this effect was weak). This result was contrary to expectation. Pires *et al.* (2011b) found a similar result using only six species of the genus *Poeciliopsis*, all of which exhibit superfetation. Larger MI values were associated with smaller offspring. Reznick & Miles (1989), in a preliminary examination that did not account for phylogenetic relatedness, also found that superfetation (which they noted was strongly linked to matrotrophy) was not associated with larger offspring.

Our estimates of phylogenetic heritability also confirmed that offspring size has evolved at a different pace in comparison with superfetation and matrotrophy, rather than in a correlated way as we expected. Offspring size has evolved flexibly at a faster pace, leaving behind little trace of the phylogeny. In contrast, the combination of superfetation and matrotrophy has evolved at a rate that has been proportional to the time elapsed between speciation events. When we analysed the degree of superfetation and MIs separately, we confirmed that both traits evolved for the most part by Brownian motion. This is the main reason why most of the interspecific variation in the degrees of superfetation and matrotrophy can be explained by the phylogenetic history. However, we highlight that the particular combination of superfetation and extensive matrotrophy has arisen recently and independently in three distinct species (*Heterandria formosa*, *Poeciliopsis prolifica* and *P. turneri*) with little evidence of the presence of this particular reproductive mode in ancestral nodes. In contrast, within the subgenus *Micropoecilia*, superfetation and extensive matrotrophy arose in the common ancestor of *Poecilia* (*Micropoecilia*) *bifurca*, *P. (M.) branneri* and *P. (M.) parae* and these three species inherited this reproductive mode (Fig. 3). Apparently, these distinct and relatively recent evolutionary events that resulted in the co-occurrence of superfetation and extensive matrotrophy did not promote parallel changes in offspring size.

We note here an important caveat. Superfeting species appear to produce smaller broods (Reznick & Miles, 1989; Pollux *et al.*, 2009), thus potentially providing additional space within the reproductive tract if at least moderate matrotrophy is present. However, the potential for this additional space within the reproductive tract, which we hypothesized could be used to produce larger offspring, depends on total fecundity remaining similar to that of non-superfeting and lecithotrophic species. If the selective forces that favour the presence of superfetation and

matrotrophy also promote increased fecundity, then the additional space could be used to produce more embryos (by means of increasing the number of simultaneous broods) rather than larger offspring.

We found no evidence of an effect of superfetation and matrotrophy on offspring size at the interspecific level. However, several studies have demonstrated that these three traits may vary widely among populations within particular species (Trexler, 1985; Riesch, Martin & Langerhans, 2013; Frías-Alvarez *et al.*, 2014). Therefore, a large offspring size as a result of increased levels of superfetation and matrotrophy at the intraspecific level is still a possibility that deserves future field and experimental tests.

OFFSPRING SIZE AS A LOCAL ADAPTATION

If offspring size is not influenced by the combination of superfetation and moderate or extensive matrotrophy, then other factors must have caused the observed variation in size at birth among species of the family Poeciliidae. Our results indicated a low phylogenetic signal in offspring size because there was more interspecific variation than expected by Brownian motion. Therefore, the evolution of offspring size in poeciliid fishes appears to be driven primarily by local adaptation to particular environmental conditions.

Several studies in poeciliids demonstrated adaptive responses of offspring size to local conditions. For example, Leips & Travis (1999) found that offspring of *Heterandria formosa* from high-density populations were 45% larger than offspring from low-density populations. High population density is characterized by high levels of competition and, under these conditions, large offspring are more likely to survive to maturity (Schrader & Travis, 2012). This same positive effect of population density and intraspecific competition on offspring size has been observed in the guppy, *Poecilia reticulata* (Bashey, 2006, 2008).

Predation is also an important selective force for the evolution of offspring size (Gordon *et al.*, 2009; Schrader & Travis, 2012). The size at birth of *Poecilia reticulata* differs between populations inhabiting two types of predation environments. Female *P. reticulata* from populations sympatric with predators that feed on small juvenile fish produce larger newborns relative to females from populations that do not experience this predation pressure. Larger newborns reach adult sizes faster than smaller newborns and spend less time at body sizes susceptible to higher predation rates. In contrast, when coexisting with predators that prey on large adult fish, females of *P. reticulata* increase their reproductive

output by producing many smaller offspring (Reznick, 1982; Reznick & Endler, 1982; Reznick *et al.*, 1990). Two species in the genus *Brachyrhaphis* (*B. episcopi* and *B. rhabdophora*) show a similar pattern of predator-mediated divergence in offspring size (Johnson & Belk, 2001; Jennions & Telford, 2002). Additional studies in *Gambusia hubbsi* and *Poecilia vivipara* also have suggested local selective effects of predation on offspring size (Downhower, Brown & Matsui, 2000; Gomes & Monteiro, 2007). Apparently, high rates of cannibalism on juveniles may also select for larger offspring sizes as observed in *Poeciliopsis monacha* (Thibault, 1974; Weeks & Gaggiotti, 1993).

Additionally, toxic compounds in the water (e.g. hydrogen sulfide) could be selective agents for offspring size. Species such as *Poecilia sulphuraria* and *Gambusia eurystoma* that inhabit water bodies with high concentrations of hydrogen sulfide produce relatively large newborns. This is presumably because a large body size results in a relatively low surface/volume ratio, which in turn implies that less surface area per volume of body tissue is exposed to the toxin. In addition, larger offspring probably have lower metabolic rates and, hence, lower oxygen consumption compared with smaller offspring (Riesch *et al.*, 2010a). Some studies have documented effects of water salinity on offspring size with contradictory results. In *Gambusia affinis*, *G. holbrooki* and *Heterandria formosa*, water bodies with high salinities appear to promote smaller offspring (Stearns & Sage, 1980; Brown-Peterson & Peterson, 1990; Alcaraz & García-Berthou, 2007; Martin *et al.*, 2009). In contrast, in *Poecilia vivipara* high salinity is associated with larger offspring (Gomes & Monteiro, 2007). The mechanisms by which salinity favours large offspring in some species and small offspring in other species remain unknown. In addition to factors that promote adaptive changes, offspring size might also exhibit phenotypic plasticity as a result of food availability (Grether & Kolluru, 2011). In *Brachyrhaphis episcopi*, females produce larger offspring during the wet season when food is more abundant (Jennions *et al.*, 2006). All these studies have demonstrated that in poeciliid fishes, offspring size may exhibit adaptive and plastic responses to a wide array of environmental conditions, thereby explaining the fluid evolutionary changes that we have detected here.

ADAPTIVE VALUE OF SUPERFETATION AND MATROTROPHY

The combination of superfetation and moderate or extensive matrotrophy has not facilitated the evolution of larger offspring in poeciliid fishes. Several

authors have suggested an evolutionary link between superfetation and matrotrophy and have proposed other potential benefits of their concurrent presence (Reznick & Miles, 1989; Pollux *et al.*, 2009, 2014; Trexler & DeAngelis, 2010; Meredith *et al.*, 2011). First, matrotrophy may increase total fecundity because initial egg size is small and females may fertilize a relatively large number of eggs without a large initial energetic investment. In contrast, lecithotrophic species are limited in the initial number of eggs that females can fertilize because a large amount of resources are needed to produce fully yolked eggs (Trexler, 1997; Trexler & DeAngelis, 2003, 2010; Marsh-Matthews *et al.*, 2005; Marsh-Matthews, 2011). Thus, total fecundity may be limited in lecithotrophic species. However, matrotrophic females need constant food supply to actively transfer nutrients to developing embryos. Otherwise, matrotrophy could become a disadvantage if enough resources are not available to sustain developing offspring. Hence, matrotrophy could increase total fecundity in environments where food resources are constant and abundant. According to a theoretical model developed by Trexler & DeAngelis (2003, 2010), in these predictable and stable environments, the previous presence of superfetation would facilitate the evolution of matrotrophy. Recently, some studies have conducted empirical tests of this 'resource-availability' hypothesis (Trexler, 1997; Marsh-Matthews & Deaton, 2006; Banet & Reznick, 2008; Banet, Au & Reznick, 2010; Pollux & Reznick, 2011; Riesch *et al.*, 2013).

Second, matrotrophy and superfetation could reduce the locomotory cost imposed by viviparity. During pregnancy, females gain mass and volume, which negatively affects escape response and swimming performance due to increased drag forces on the female body (Plaut, 2002; Ghalambor, Reznick & Walker, 2004; Langerhans & Reznick, 2010). Matrotrophy could reduce this reproductive burden because mature ova are small and thus initial space requirements are low (Thibault & Schultz, 1978; Marsh-Matthews, 2011). Superfetation might provide additional advantages because it further reduces reproductive allocation by allowing females to divide the total number of developing embryos into smaller broods (Thibault & Schultz, 1978; Zúñiga-Vega *et al.*, 2010). Therefore, both matrotrophy and superfetation could result in thinner bodies without a reduction in the total number of offspring produced because females bear a combination of small early-stage embryos and fewer large late-stage embryos compared with lecithotrophic and non-superfeting species that must bear many large embryos (Miller, 1975; Thibault & Schultz, 1978; Zúñiga-Vega *et al.*, 2007, 2010; Pollux *et al.*, 2009; Pires *et al.*, 2011a).

Selective environments where a streamlined body shape is needed such as high-velocity water systems or habitats where fish must swim fast to escape from abundant predators might promote the evolution of both reproductive strategies. Zúñiga-Vega *et al.* (2007) provided evidence that supports this hypothesis using *Poeciliopsis turrubarensis* as a model system. However, they focused only on the adaptive value of superfetation without any emphasis on spatial variation in matrotrophy. Additional tests of this 'locomotor cost' hypothesis are needed, analysing the effects of the concurrent presence of matrotrophy and superfetation on body streamlining and fitness in these particular selective environments.

Third, moderate and extensive matrotrophy have coevolved with specialized morphological structures that facilitate the active transfer of nutrients from mother to embryos (i.e. placentas; Turner, 1940; Jollie & Jollie, 1964; Knight *et al.*, 1985; Grove & Wourms, 1994; Kwan *et al.*, 2015). This enhanced physiological communication through placental tissues could facilitate the rise of parent-offspring conflicts during pregnancy (i.e. individual embryos demand a greater investment and mothers attempt to optimize the allocation to each offspring; Trivers, 1974; Zeh & Zeh, 2000, 2008; Crespi & Semeniuk, 2004; Schrader & Travis, 2008). Parent-offspring conflicts during embryo development may drive a shift from pre-copulatory female mate choice to post-copulatory mechanisms of sexual selection (Pollux *et al.*, 2014). The capacity to control the amount of resources that are devoted to each offspring through matrotrophy and placentas would allow females to allocate more resources to those offspring that carry the best paternal genes. In addition, superfetation could further enhance post-copulatory female selection by facilitating multiple paternity because each brood may be fertilized by a different male (Travis, Trexler & Mulvey, 1990; Zane *et al.*, 1999; Soucy & Travis, 2003). This potential to produce temporally overlapping, mixed-paternity broods might also increase genetic variability of the offspring (Macías-García & González-Zuarth, 2005). Pollux *et al.* (2014) used phylogenetic comparative methods and a large data set (94 species) to suggest that the correlated evolution of superfetation and matrotrophy in the family Poeciliidae has allowed placental females to gain control over paternity, thereby relying to less extent on pre-copulatory mechanisms of sexual selection.

Additional theoretical models have attempted to explain the fitness benefits of superfetation and matrotrophy, although considering them separately (Thibault, 1974; Downhower & Brown, 1975; Travis *et al.*, 1987; Zúñiga-Vega *et al.*, 2010; Pires *et al.*, 2011b; Bassar, Auer & Reznick, 2014). However,

their correlated evolution strongly suggests that the adaptive benefits of superfetation depend, at least to some degree, on the concurrent presence of matrotrophy and/or vice versa (Reznick & Miles, 1989; Pollux *et al.*, 2009, 2014; Trexler & DeAngelis, 2010; Meredith *et al.*, 2011). Therefore, further theoretical and empirical studies must focus on the combination of both traits when examining their potential adaptive value.

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SUPPORTING INFORMATION

Additional Supporting Information may be found in the online version of this article at the publisher's website:

Table S1. List of the species that we used for comparative analyses showing the method of preservation (ethanol or formalin).

Table S2. Phylogenetic estimates of the magnitude of the evolutionary relationship between offspring size and the combination of superfetation and matrotrophy calculated using different phylogenetic comparative methods. These estimates were obtained from analyses conducted on data from the 33 species that were preserved in ethanol.

Table S1. List of the species that we used for comparative analyses showing the method of preservation (ethanol or formalin).

Species	Preservation method
<i>Brachyrhaphis rhabdophora</i>	Ethanol
<i>Brachyrhaphis episcopi</i>	Formalin
<i>Brachyrhaphis holdridgei</i>	Ethanol
<i>Heterandria formosa</i>	Formalin
<i>Poeciliopsis lucida</i>	Ethanol
<i>Poeciliopsis pleurospilus</i>	Ethanol
<i>Poeciliopsis occidentalis</i>	Ethanol
<i>Poeciliopsis prolifica</i>	Formalin
<i>Poeciliopsis infans</i>	Ethanol
<i>Poeciliopsis monacha</i>	Ethanol
<i>Poeciliopsis viriosa</i>	Ethanol
<i>Poeciliopsis turneri</i>	Ethanol
<i>Poeciliopsis gracilis</i>	Ethanol
<i>Poeciliopsis turrubarensis</i>	Ethanol
<i>Poeciliopsis latidens</i>	Ethanol
<i>Poeciliopsis fasciata</i>	Ethanol
<i>Poeciliopsis baenschi</i>	Ethanol
<i>Heterophallus milleri</i>	Formalin
<i>Gambusia panuco</i>	Ethanol
<i>Gambusia vittata</i>	Ethanol
<i>Gambusia sexradiata</i>	Formalin
<i>Gambusia eurystoma</i>	Formalin

<i>Gambusia hubbsi</i>	Formalin
<i>Gambusia yucatana</i>	Ethanol
<i>Gambusia holbrooki</i>	Ethanol
<i>Gambusia affinis</i>	Ethanol
<i>Gambusia speciosa</i>	Ethanol
<i>Gambusia aurata</i>	Ethanol
<i>Belonesox belizanus</i>	Ethanol
<i>Pseudoxhiphophorus jonesii</i>	Ethanol
<i>Priapella chamulae</i>	Formalin
<i>Priapella olmecae</i>	Ethanol
<i>Poecilia mexicana</i>	Formalin
<i>Poecilia sulphuraria</i>	Formalin
<i>Poecilia butleri</i>	Ethanol
<i>Poecilia (Micropoecilia) bifurca</i>	Ethanol
<i>Poecilia (Micropoecilia) branneri</i>	Ethanol
<i>Poecilia (Micropoecilia) parae</i>	Ethanol
<i>Poecilia (Micropoecilia) picta</i>	Formalin
<i>Poecilia (Acanthophaelus) reticulata</i>	Ethanol
<i>Poecilia (Acanthophaelus) wingei</i>	Ethanol
<i>Phalloceros caudiomaculatus</i>	Ethanol
<i>Phalloceros anisophallos</i>	Ethanol
<i>Xenodexia ctenolepis</i>	Ethanol

Table S2. Phylogenetic estimates of the magnitude of the evolutionary relationship between offspring size and the combination of superfetation and matrotrophy calculated using different phylogenetic comparative methods. These estimates were obtained from analyses conducted on data from the 33 species that were preserved in ethanol.

Hansen's method		FIC	TIPS	PGLS	PMM
SF and MI > 1	SF and MI > 5				
$r^2 = 0.002$	$r^2 = 0.01$	$r = 0.21$	$r = 0.13$	$r = 0.15$ $r^2 = 0.02$ $\alpha = 12.04$	$r = 0.07$
					Offspring size $h^2 = 0.52$ SF and MI $h^2 = 0.98$ SF $h^2 = 0.88$ MI $h^2 = 0.86$

FIC, Felsenstein's independent contrasts; TIPS, non-phylogenetic approach; PGLS, phylogenetic generalized least squares method assuming an Ornstein-Uhlenbeck model of evolution; PMM, phylogenetic mixed model; SF, superfetation; MI, matrotrophy index; r , correlation coefficient; r^2 , proportion of the variation among species in offspring size explained by superfetation and matrotrophy; α , a measure of the strength of selection pulling offspring size towards optimal values; h^2 , phylogenetic heritability.

DISCUSIÓN GENERAL

La superfetación es una estrategia reproductora que evolucionó independientemente en diferentes géneros de la familia Poeciliidae (Turner, 1937; Reznick y Miles, 1989; Gunn y Thresher, 1991; Scrimshaw, 1994; Reznick *et al.*, 2007). Es interesante estudiar esta estrategia debido a las ventajas adaptativas que le confiere a las especies que la presentan (e.g. forma más hidrodinámica, reparten su esfuerzo reproductor y mayor número de crías; Zúñiga-Vega *et al.*, 2007, 2010). Recientemente, se demostró que la superfetación y la matrotrofia han evolucionado en conjunto (Meredith *et al.*, 2011; Pollux *et al.*, 2014). La matrotrofia es un importante aprovisionamiento materno que ha evolucionado y ha sido estudiado ampliamente en poecílicos. Sin embargo, varios estudios únicamente se han enfocado en conocer el efecto por separado de la matrotrofia y la superfetación sobre diferentes aspectos reproductores en poecílicos (i.e. ornamentaciones, longitud de gonopodio e índice de selección sexual; Pollux *et al.*, 2014). Por lo que un aspecto interesante por conocer son las consecuencias sobre los mecanismos morfológicos y fisiológicos, que permiten que funcionen estas dos estrategias en conjunto. Por lo tanto, mi tesis se enfocó en conocer algunas de las consecuencias de la superfetación y la matrotrofia en diferentes aspectos de la vida de los poecílicos. En especial, examiné las consecuencias en aspectos reproductores (como el nivel de complejidad de la placenta folicular y retención de esperma) y cambios en las historias de vida (como el tamaño del embrión al nacer).

¿En qué difieren las placentas de especies de peces que superfetan con respecto a las placentas de especies que no superfetan? ¿Altos niveles de matrotrofia están asociados con placentas más complejas?

En el primer capítulo observé que la placenta folicular en todas las especies de poecílicos se dividió en dos capas principales que anteriormente habían sido observadas por Jollie y Jollie (1964): la capa folicular interna y la capa folicular externa. La capa folicular interna contiene las células foliculares aplanadas con elementos citoplasmáticos evidentes como vesículas, aparato de Golgi, extenso retículo endoplásmico y mitocondrias que indican alta actividad metabólica. La capa folicular externa se compone principalmente de tejido conjuntivo laxo con vasos sanguíneos y fibrillas de colágeno, estas estructuras ayudan a la protección de las células foliculares y el embrión. En las placentas de todas las especies identificamos estructuras que aparentemente facilitan el transporte de nutrientes y gases. Particularmente, en la capa folicular interna se encuentran prolongaciones de la membrana o microvellosidades orientados hacia el embrión que incrementan la superficie de contacto para el intercambio de sustancias (como nutrientes, gases y desechos). Estas estructuras de la placenta folicular fueron observadas anteriormente en *Poecilia reticulata* y *Heterandria formosa* (Jollie y Jollie, 1964; Grove y Wourms, 1994).

Poeciliopsis presidionis fue un caso particular, la estructura de su placenta es diferente a las otras especies observadas con anterioridad (*Poecilia reticulata*, Jollie y Jollie, 1964; *Heterandria formosa*, Grove y Worms, 1994). No en lo referente a que está constituida por las mismas dos capas (interna y externa), sino en cuanto al grosor y estructura de ambas capas. Pude examinar con detalle la organización general de este tipo

de híper-placenta (estructura con características placentarias más desarrolladas para el intercambio de nutrientes entre madre y sus embriones). La híper-placenta consiste primordialmente en el plegamiento del epitelio de la capa folicular interna, con un tipo celular que no es plano sino columnar. El plegamiento de la capa folicular interna es seguido por el plegamiento de la capa folicular externa, destacando la entrada de los capilares sanguíneos hasta el límite epitelial. Otra diferencia muy marcada entre esta especie y todas las demás es que la capa externa presenta mayor grosor del tejido conjuntivo y múltiples capas de fibrillas de colágeno. Este tipo de morfología placentaria se ha observado también en otras especies con matrotrofía extensa como *Poeciliopsis turneri* y *P. retropinna* por medio de microscopía óptica (Kwan *et al.*, 2015). Sin embargo, estas últimas especies no fueron analizadas detalladamente con microscopía electrónica como en el presente estudio, por lo que no me fue posible comparar sus componentes celulares con los que yo observé en *P. presidionis*. Con base a mis resultados y los de Kwan *et al.* (2015) es posible concluir que a partir de ciertos valores de matrotrofía (*P. presidionis* MI=21.5, *P. turneri* MI=41.4, *P. retropinna* MI=117; Reznick *et al.*, 2002; Kwan *et al.*, 2015) exista la necesidad de estructuras placentarias muy complejas (híper-placentas) para facilitar la transferencia de sustancias activas entre la madre y el embrión a lo largo del desarrollo.

Hasta la fecha no se había examinado la posible relación entre la superfetación y la complejidad de las placentas en especies de la familia Poeciliidae. Mi estudio es el primero en demostrar que las especies superfetadoras aparentemente tienen características de las placentas que reflejan un mayor nivel de complejidad en comparación con las especies que no superfetan. Las especies con superfetación mostraron mayor área del folículo materno,

más microvellosidades y de mayor longitud en comparación con las especies sin superfetación. Mi primera hipótesis en esta sección argumenta que la superfetación requiere una regulación diferencial por parte de las hembras de las tasas de desarrollo de embriones en distintos estadios (Turner, 1937, 1940; Scrimshaw, 1944). Por lo tanto, las hembras de especies que superfetan deberían tener una placenta más compleja comparada con especies que no superfetan. Esta hipótesis se cumplió parcialmente, ya que observé únicamente una tendencia de características morfológicas más desarrolladas y complejas en las especies que superfetan en comparación con las especies que no superfetan.

Por otro lado, con respecto a la asociación entre matrotrofia y el nivel de complejidad de las placentas encontré evidencia de que altos niveles de matrotrofia también se relacionan con placentas más complejas. Es decir, el área del folículo materno, el número de microvellosidades, el número de vesículas y el tamaño de estas vesículas tienden a aumentar conforme aumenta el índice matrotrófico. Estas ultraestructuras placentarias sugieren la transferencia activa y continua de nutrientes entre la madre y el embrión (Turner, 1940; Jollie y Jollie, 1964; Grove y Wourms, 1994). La segunda hipótesis planteada para esta sección fue que, debido a que la matrotrofia implica la transferencia activa de nutrientes por parte de la madre hacia sus embriones a lo largo de todo el desarrollo (Wourms, 1981; Marsh-Matthews, 2011), las hembras de especies con altos niveles de matrotrofia tendrían una placenta más compleja en comparación con las hembras de especies con lecitotrofia o matrotrofia incipiente. Mis resultados apoyaron esta hipótesis debido que las especies con altos niveles de matrotrofia presentaron características placentarias más complejas y presumiblemente ayudan a maximizar la

transferencia activa de nutrientes. Consistente también con esta hipótesis, Kwan y colaboradores (2015) mostraron que el grosor del folículo materno crece conforme aumenta el nivel matrotrófico en ocho especies de poecílidos. En resumen, con base en el presente trabajo es posible concluir que existe un mayor nivel de complejidad en las placentas de especies que exhiben alta matrotrofia.

Antes de esta investigación no había estudios sobre la superfetación y su relación con la complejidad de la placenta. Por lo que el primer capítulo de esta tesis tiene una relevancia importante porque pude identificar las posibles consecuencias de la superfetación y la matrotrofia sobre la estructura celular de la placenta folicular materna. Sin embargo, aún quedan varios aspectos por conocer sobre la placentación en poecílidos. Primero, en mi proyecto de investigación se trabajó con un tamaño de muestra pequeño, lo que implica que nuestras relaciones biológicas pueden tener poco poder estadístico. Se necesitan estudios adicionales con más especies y con un mayor número de muestras por especie para fortalecer la evidencia de las relaciones observadas entre la superfetación, la matrotrofia y el nivel de placentación. Segundo, en la presente investigación tomé en cuenta sólo algunas características ultraestructurales que facilitan la transferencia de nutrientes (microvellosidades y vesículas) y algunas medidas de la placenta folicular materna (área y grosor del folículo materno; de acuerdo con Jollie y Jollie, 1964 y Grove y Wourms, 1994). Sin embargo, hace falta examinar otras ultraestructuras placentarias como la cantidad de mitocondrias y aparatos de Golgi y su posible asociación con la presencia de altos niveles de superfetación y matrotrofia. Tercero, en esta tesis mostré cómo algunas ultraestructuras de la placenta folicular, específicamente en la parte materna de las

placentas, se encuentran relacionadas con la superfetación y matrotrofia. Sin embargo, hasta la fecha no se han investigado las características ultraestructurales placentarias del tejido absorbente del embrión. Al igual que la placenta folicular materna, el tejido absorbente del embrión debería tener características ultraestructurales que optimicen la transferencia de nutrientes, tales como numerosas microvellosidades y vesículas.

¿Existen diferencias entre especies que superfetan y no superfetan en la capacidad de almacenar esperma?

En el segundo capítulo propuse que debido a que la superfetación implica una producción constante de embriones en diferentes estadios de desarrollo (Turner, 1937, 1940; Scrimshaw, 1994), se espera que las especies con superfetación requieran fertilizar constantemente a sus ovocitos. Por lo que, en las hembras de especies con superfetación deberían existir estructuras que almacenen espermatozoides, y que les permitan estar disponibles para fertilizar constantemente a diferentes grupos de embriones.

Encontré espermatecas solamente en ovarios de nueve de las 12 especies que examiné (*Gambusia panuco*, *Xiphophorus hellerii*, *Poecilia mexicana*, *Poeciliopsis prolifica*, *P. infans*, *P. viriosa*, *P. presidionis*, *P. gracilis* y *Heterandria formosa*). Estructuras similares que almacenan esperma debido al cierre de los pliegues ováricos se habían reportado con anterioridad en *Heterandria formosa* y *Poecilia reticulata* (Fraser y Renton, 1940; Kobayashi y Iwamatsu, 2002). En general, observamos que las espermatecas no se encuentran en un área específica dentro del ovario, sino que están distribuidas aleatoriamente, aunque

siempre en los límites cercanos a la luz del ovario. Las espermatecas están compuestas de epitelio ovárico engrosado y frecuentemente se localizan junto a túbulos de vasos sanguíneos. En general, las cabezas de los espermatozoides se encontraron orientadas hacia el epitelio ovárico y las colas hacia la luz de la espermateca. Este particular arreglo de los espermatozoides, podría indicar que el tejido ovárico interno nutre a los espermatozoides para mantenerlos viables. Potter y Kramer (2000) reportaron que las células del tejido ovárico que interactúan directamente con los espermatozoides presentan abundantes ribosomas, microvellosidades y desmosomas, que son características de células que transportan nutrientes.

Por otra parte, en la presente tesis encontré que existe una relación entre la presencia de superfetación con el número y el tamaño de las espermatecas. Las especies que presentan superfetación tienen mayor número de espermatecas y más grandes espermatecas en comparación con las especies que no superfetan, sin importar su historia filogenética. Por lo tanto, las especies superfetadoras destinan una mayor proporción del volumen total del ovario a la producción de espermatecas en comparación con las especies sin superfetación. Adicionalmente, en tres de las seis especies que no superfetan no encontré la presencia de espermatecas. La hipótesis que puse a prueba con respecto a este tema de investigación es que debido a la necesidad de fertilizar frecuentemente a distintos grupos de embriones, las hembras de las especies que superfetan deberían tener mayor capacidad de almacenar esperma en comparación con hembras de especies que no superfetan. Mis resultados apoyaron esta hipótesis y sugieren que el almacenamiento de esperma dentro de las espermatecas favorece la superfetación, ya que presumiblemente

facilita la disponibilidad constante de espermatozoides para fertilizar a las camadas que se desarrollan frecuentemente.

La relación entre la superfetación con el mayor número de las espermatecas puede deberse a que en las especies superfetadoras puede haber selección sexual post-copulatoria dentro de las espermatecas (elección femenina críptica, Birkhead y Pizzari, 2002; Evans y Pilastro, 2011). Es decir, estudios reportados recientemente han demostrado que en especies con superfetación es muy común la coerción sexual, por lo que una estrategia de elección post-copulatoria por parte de la hembra compensaría la falta de las hembras para escoger a su pareja (Macías-García y González-Zuarth, 2005; Pollux *et al.*, 2014). Por lo que la retención de esperma en una espermateca puede ser fundamental, debido a que los espermatozoides más fuertes sobrevivirán en las espermatecas hasta ser usados por las hembras. Además, la competencia de esperma puede ocurrir en las espermatecas. Previos estudios han documentado la influencia de la hembra sobre la competencia espermática, a través del fluido ovárico que provoca cambios en la motilidad y supervivencia de los espermatozoides (Gasparini *et al.*, 2010; Gasparini y Pilastro, 2011; Gasparini y Evans, 2013).

Este estudio es el primero que examina la asociación entre la capacidad de almacenar esperma y la presencia de superfetación. A partir de estos resultados surgieron otros temas que no se han resuelto. Por ejemplo, cuál es la posible función tienen los vasos sanguíneos que se encontraron rodeando a las espermatecas, sugiero que podría ser que sean necesarios para aportar nutrientes a los espermatozoides que se almacenan dentro de las espermatecas, debido a que los vasos sanguíneos transportan oxígeno y

nutrientes necesarios para las células (Schaller et al., 2008). Su cercanía a las espermatecas representa evidencia adicional de que el tejido ovárico de las hembras provee los medios necesarios para mantener viables a los espermatozoides durante periodos de tiempo relativamente prolongados. También se necesitan más estudios para conocer exactamente, cuáles son las sustancias nutritivas que reciben los espermatozoides dentro de las espermatecas, y cuáles son los mecanismos fisiológicos que permiten el transporte de estos nutrientes hacia el interior de las espermatecas. Otro aspecto importante por explorar es cuál es la duración de las espermatecas dentro del ovario. Existen estudios que sugieren una duración aproximada de la retención de esperma con base en simples observaciones sobre el tiempo que una hembra aislada de machos puede seguir produciendo crías. Sin embargo, ninguno de estos trabajos hizo una examinación fisiológica o anatómica (Clark y Aronson, 1951; Hildemann y Wagner, 1954; Hubbs, 1964, 1997; Greven, 2011).

¿La presencia conjunta de superfetación y matrotrofia promueven la producción de crías más grandes en especies de la familia Poeciliidae?

En el tercer capítulo, examiné la hipótesis de que la evolución conjunta de la superfetación y la matrotrofia en la familia Poeciliidae ha facilitado la evolución de crías más grandes. Debido a que las hembras con superfetación, reparten su esfuerzo reproductor produciendo pequeñas camadas que nacen frecuentemente (Reznick y Miles, 1989; Pollux et al., 2009; Zuñiga-Vega et al., 2010). Por lo que las hembras superfetadoras pudieran tener

mayor espacio disponible dentro de su ovario en comparación con hembras que no superfetan. Además, la matrotrofia implica que los embriones son pequeños en los primeros estadios de desarrollo debido a su escaso vitelo y más grandes en los últimos estadios del desarrollo (Wourms, 1981; Marsh-Matthews, 2011). En consecuencia, una combinación de la presencia de superfetación y alta matrotrofia podría promover el incremento del tamaño de las crías al nacer. Sin embargo, encontré que altos niveles superfetación y matrotrofia no tienen una relación con el tamaño de las crías al nacer. Por lo tanto, la hipótesis no fue apoyada por mis resultados. De hecho, los resultados que obtuve fueron contrarios a los que esperaba. El tamaño óptimo al nacer estimado para especies con superfetación y matrotrofia moderada o extensa fue menor comparado con el tamaño al nacer estimado para especies sin superfetación, especies con superfetación y lecitrotrofia y especies con superfetación y matrotrofia incipiente. Un estudio que es consistente con estos resultados es el de Pires y colaboradores (2011), quienes encontraron que altos valores de matrotrofia están asociados a crías de menor tamaño al nacer en seis especies del género *Poeciliopsis*. Sus resultados soportan la hipótesis de la facilitación de historias de vida (i.e. la matrotrofia facilita la evolución de algunos rasgos de la historia de vida), por lo que la evolución de alta matrotrofia promueve la maduración temprana de los embriones provocando crías de tamaño pequeño (Pires *et al.*, 2011).

Otro resultado importante es que el tamaño de la cría en las especies que se analizaron tiene baja heredabilidad filogenética. Esto indica que la evolución del tamaño al nacer en poecílicos parece deberse principalmente a procesos de adaptación local a condiciones particulares del medio ambiente. Varios estudios apoyan este resultado, ya que

han demostrado que ciertos factores ambientales como alta competencia, alta depredación o presencia de sustancias tóxicas en el agua tienen como consecuencia cambios predecibles y diferencias entre sitios (intraespecíficas) en el tamaño óptimo de las crías (i.e. *Heterandria formosa*, *Poecilia reticulata*, *P. sulphuraria*, *Brachyrhaphis episcopi*, *B. rhabdophora*, *Gambusia eurystoma*; Reznick, 1982; Reznick y Endler, 1982; Reznick *et al.*, 1990; Leips y Travis, 1999; Johnson y Belk, 2001; Jennions y Telford, 2002, Bashey, 2006, 2008; Riesch *et al.*, 2010).

Aún hacen falta estudios enfocados en entender el significado adaptativo de la presencia conjunta de altos niveles de superfetación y matrotrofia y sus posibles efectos sobre otras características fisiológicas, conductuales y de historias de vida. Por ejemplo, una hipótesis que aún no se ha puesto a prueba, es que la combinación entre la matrotrofia y superfetación podrían reducir los costos locomotores asociados a la viviparidad (Pollux *et al.*, 2009). Durante el embarazo, las hembras ganan peso y volumen, lo cual tiene efectos negativos en su respuesta de escape y en su rendimiento al nadar (Plaut, 2002; Ghalambor *et al.*, 2004; Langerhans y Reznick, 2010). Entonces, las hembras con matrotrofia y superfetación podrían repartir su esfuerzo reproductor teniendo varias camadas de embriones en diferente estadio de desarrollo. Estas hembras tendrían pocos embriones de tamaño pequeño por estar en estadios tempranos, pocos embriones de tamaño mediano por estar en estadios intermedios y pocos embriones de tamaño grande por estar en estadios de desarrollo más avanzados. En consecuencia, esta repartición de camadas y los diferentes tamaños embrionarios provocaría la disminución del volumen total ocupado por

los embriones en comparación con especies que carecen de superfetación y matrotrofia (Pollux *et al.*, 2009; Reznick y Miles, 1989)

Además, sería interesante comparar los resultados que hemos observado con especies de la familia Poeciliidae con otras familias de peces que también presentan superfetación (Clinidae y Zenarchopteridae; Gunn y Thresher, 1991; Reznick *et al.*, 2007). Otras interrogantes que aún genera la superfetación y que necesitan ser estudiadas son: ¿Cuáles son los procesos fisiológicos que subyacen a la capacidad de procrear simultáneamente a varios grupos de embriones de diferentes estadios? particularmente, ¿Qué hormonas regulan este desarrollo embrionario diferencial? ¿Cómo se relaciona la superfetación con la selección sexual post-copulatoria? ¿Las especies con superfetación tienen mayores niveles de paternidad múltiple?

En la presente tesis logré examinar una variedad de consecuencias adaptativas de la superfetación y matrotrofia que van desde la evaluación de aspectos reproductores como la complejidad de algunas ultraestructuras placentarias (e.g. el área de folículo materno, el número y tamaño de microvellosidades, el número y tamaño de vesículas y el grosor del folículo materno) y la retención de esperma (presencia, tamaño y número de espermatecas), hasta rasgos en las historias de vida como el tamaño de la cría en algunas especies de la familia Poeciliidae. Estas consecuencias adaptativas podrían ser la base de nuevas investigaciones de los peces poecílicos y su peculiar modo reproductor que es la viviparidad.

CONCLUSIONES

- En todas las especies de poecílidos la placenta folicular se dividió en dos capas foliculares: la interna y la externa. La capa folicular interna contiene un epitelio de células foliculares. Mientras que la cara externa es de tejido conjuntivo laxo que predominan capilares sanguíneos y fibras de colágena en toda la capa. *Poeciliopsis presidionis* fue la especie que sobresalió debido a que esta presenta un plegamiento prominente en la capa folicular interna. Por su parte, la capa folicular externa tiene varias capas de fibra de colágena y sus capilares sanguíneos protruyen hacia la capa folicular interna.
- Demostré que las especies superfetadoras tienen características de las placentas que reflejan un mayor nivel de complejidad (mayor área folicular, más microvellosidades y de mayor longitud) en comparación con las especies que no superfetan.
- Encontré una asociación positiva entre la intensidad matrotrofica y el nivel de complejidad de las placentas. El área del folículo materno, el número de microvellosidades, el número de vesículas y el tamaño de estas vesículas aumentan conforme aumenta el índice matrotrofico. Estas características placentarias sugieren que la transferencia de nutrientes es activa entre la madre y el embrión.
- Observé espermatecas en los ovarios de nueve de las 12 especies que se examinaron (*Gambusia panuco*, *Xiphophorus hellerii*, *Poecilia mexicana*, *Poeciliopsis prolifica*, *P. infans*, *P. viriosa*, *P. presidionis*, *P. gracilis* y *Heterandria formosa*).

- Este estudio es el primero que examina la asociación entre la capacidad de almacenar esperma y la presencia de superfetación. Las especies que presentan superfetación tienen mayor número de espermatecas y espermatecas más grandes en comparación con las especies que no superfetan
- Encontré que altos niveles superfetación y matrotrofia no tienen una relación con el tamaño de las crías al nacer.
- El tamaño de la cría en las especies que se analizaron tiene baja heredabilidad filogenética. Esto indica que la evolución del tamaño al nacer en poecílicos parece deberse principalmente a procesos de adaptación local.
- Aún hacen falta estudios enfocados en entender el significado adaptativo de la presencia conjunta de altos niveles de superfetación y matrotrofia y sus posibles efectos sobre otras características fisiológicas, conductuales y de historias de vida.
- Logré examinar una variedad de consecuencias adaptativas de la superfetación y matrotrofia que van desde la evaluación de aspectos reproductores como la complejidad ultraestructural de la placenta y la retención de esperma, hasta rasgos en las historias de vida como el tamaño de la cría en algunas especies de la familia Poeciliidae.

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