



UNIVERSIDAD NACIONAL AUTÓNOMA DE MÉXICO
POSGRADO EN CIENCIAS BIOMÉDICAS
INSTITUTO DE ECOLOGÍA

**PREPARACIÓN INMUNOLÓGICA Y ASPECTOS DE LA
GENÉTICA CUANTITATIVA DE LA RESPUESTA INMUNE DE
*Aedes Aegypti***

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A mis padres

Miguel Moreno y María Esther García

A la Universidad Nacional Autónoma de México

AGRADECIMIENTOS

Al Dr. Alejandro Córdoba por la dirección, consejos y todo el apoyo durante el tiempo que duro el desarrollo de la tesis. Muchas gracias por la amistad.

Al Dr. Humberto Lanz por permitirme ser parte integral de su grupo de trabajo en el Instituto Nacional de Salud Pública, y dejarme desarrollar libremente las ideas en su modelo de estudio. Por todas las correcciones y consejos en los experimentos y las interpretaciones de resultados. Por el apoyo académico, económico y sobre todo personal. Muchas gracias por la confianza y amistad.

Al Dr. Juan Núñez y Dra. Robyn Hudson por haber aceptado ser parte del comité tutorial. Muchas gracias por todas las sugerencias que hicieron crecer enormemente este trabajo.

A los miembros del Jurado de Examen, Dr. Mario H. Rodríguez, Dr. Raúl Cueva, Dra. Ingeborg Becker, y Dr. Juan Fornoni muchas gracias por el tiempo invertido en la revisión de la tesis, y por las acertadas sugerencias que mejoraron el escrito de la tesis.

Al Dr. Salvador Hernández por los protocolos. Al grupo de Malaria-Dengue por las sugerencias y buenos ratos.

A los amigos y compañeros del INSP: Martha, Javier, Renaud, Lupita, Raúl, Toñita, Didier, Rebeca, Avigail, Priscila, Marisol, Raquel, Benito, Luis, Inci. A Dolores Méndez por todo el apoyo en trámites y por la amistad.

A los amigos y compañeros del Instituto de Ecología, UNAM: Dr. Carlos Cordero, Rafa, Nubia, Víctor, Lizeth, Ceci, Jorge, Isaac, Raúl, Memo, Daniel, Daniela, Adriana, Ángela, Jesús, Robert, Ana Leshner, Karla, Allari.

A los grandes amigos (y colegas) que hice en Cuernavaca: Alex “Veracruz”, Alex “El Jefe”, Fabi, Chalio, El Vera, Valeria, Jimena, Che y Brenda, Karlita, Bernardo, a Cassandra y toda su familia, Uli, Brenda y Richard, Yadira y Richard, More y su familia, muchas gracias por su amistad y abrirme las puertas de sus casas.

A Peter, Axa, Magali, Emmanuel, Carmen, Horacio, Ana, Adriana, Manuel, Adrian, gracias.

A CONACYT (Beca No. 172947), Instituto de Ecología, Posgrado en Ciencias Biomédicas, UNAM.

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CAPÍTULO I. INTRODUCCIÓN GENERAL

Teoría Inmunoecológica

El sistema inmune de los animales es un medio de defensa que ha evolucionado para proteger al organismo de los efectos ocasionados por agentes infecciosos. Estudios recientes (ver Schmid-Hempel 2005) han revelado las células y moléculas que participan en el reconocimiento, eliminación y/o tolerancia a los agentes patógenos y sus efectos. Actualmente se sabe que estas células y moléculas actúan de manera conjunta entre sí, y con otros sistemas (ej. nervioso, reproductivo; Tu, Flat & Tatat 2005). Por esta razón se ha propuesto que la actividad inmune está íntimamente relacionada con las estrategias de ciclo de vida de los organismos involucradas con la supervivencia y reproducción diferencial (i.e. adecuación) en un ambiente dado. A este conjunto de estrategias se le conoce como la historia de vida de un organismo, y en conjunto maximizan la probabilidad de supervivencia y reproducción (Stearns, 1992). Estas estrategias están asociadas a características como el tamaño, edad a la maduración, tasa de crecimiento, desarrollo y reproducción. Existen también características asociadas a las características de historia de vida, como la coloración corporal, comportamientos de apareamiento, producción hormonal, y que se clasifican comúnmente como características morfológicas, fisiológicas y conductuales que también están relacionadas con la supervivencia y reproducción diferencial del organismo.

Hamilton y Zuk (1982) y Folstad y Karter (1992), fueron los primeros en proponer que la capacidad inmune de los organismos podía estar relacionada con las características de historias de vida (e intermediarias) antes mencionadas. Estos autores propusieron que la resistencia a parásitos en vertebrados está relacionada con la expresión de características sexuales secundarias relacionadas con la elección de pareja. A partir de entonces surge formalmente una nueva rama

de la biología conocida como *inmunoeología* (“ecological immunity”). La inmunoeología trata de explicar cuáles son los procesos microevolutivos (presiones selectivas, flujo de genes entre poblaciones, efectos de deriva génica y mutaciones) y cómo, en combinación con factores ambientales (ej. cantidad y calidad de recursos, humedad, temperatura) y de interacción con otros organismos (ej. competencia, depredación, parasitismo), ha evolucionado la forma en que los organismos montan una respuesta inmune en contra de patógenos.

La evidencia encontrada en un amplio número de insectos, muestra que existe un costo evolutivo de la habilidad de los organismos de montar y mantener una respuesta inmune para minimizar los efectos de una infección, a esta habilidad se le conoce como inmunocompetencia (Owens & Wilson 1999). Este costo es debido a que las características dirigidas a lidiar con los patógenos puedan estar relacionadas negativamente con características de historias de vida. La respuesta inmune al covariar negativamente con otras características de historia de vida puede estar generando la aparición de disyuntivas (o trade-offs en inglés) (la disyuntiva más común es supervivencia vs. reproducción). Las disyuntivas ocurren cuando dos o más características no pueden ser favorecidas por el mecanismo de la selección natural (Núñez-Farfán 1993), esto a pesar de que las características en disyuntiva puedan estar genéticamente correlacionadas (Cheveraud et al, 1983) y contribuyendo a incrementar el éxito reproductivo del organismo. De forma tal que las disyuntivas podrán limitar la evolución de las características que se encuentran en disyuntiva. La base de las disyuntivas radica en que los recursos requeridos para la generación de la respuesta inmune pueden ser recursos que también son requeridos para la expresión de otras características involucradas en la supervivencia y reproducción (Zuk y Stoehr 2002). A pesar del amplio conocimiento que se ha generado en los últimos años dentro de la rama de la inmunoeología y que demuestra la existencia de disyuntivas, y su repercusión en la evolución de la respuesta inmune, existen tópicos poco examinados. Dentro de estos se encuentran los

efectos que tiene la constante presencia de un patógeno en la respuesta inmune y componentes de adecuación del hospedero; y la variación fenotípica de origen genético y no genético y a la sensibilidad de la respuesta inmune ante condiciones ambientales heterogéneas.

El objetivo general de esta tesis es incrementar el conocimiento de estos dos puntos antes mencionados. Esto mediante (1) la evaluación del efecto que tienen encuentros consecutivos con un patógeno sobre la respuesta inmune, supervivencia y reproducción del mosquito *Aedes aegypti*; y (2) cómo se afecta la respuesta inmune ante la heterogeneidad en los recursos alimenticios. Este último punto, abordando las diferencias que existen entre la hembra y macho del mosquito, y las determinantes genéticas y ambientales de la respuesta inmune. El conocimiento generado a partir de esta tesis podrá ser utilizado para aumentar la comprensión de la biología de esta especie de mosquito vector del virus Dengue y del de la Fiebre amarilla, y en un futuro proponer estrategias para su control. Esto con la finalidad de reducir el impacto que tiene sobre la salud de las poblaciones humanas.

La tesis comienza con un apartado (*capítulo II*) en el cual se revisan las disyuntivas generadas entre efectores de la respuesta inmune y otras características (morfológicas y conductuales). Se exponen las teorías de las razones de dichas disyuntivas. Se abordan temas que han sido poco estudiados como interacciones patógeno-hospedero y de competencia patógeno-patógeno y su influencia en la evolución del hospedero y del patógeno. Se contemplan las diferencias entre sexos, y su consecuencia evolutiva, que sería el dimorfismo sexual en caracteres morfológicos y de respuesta inmune. Brevemente, y para contemplar su relación con factores ecológicos, se mencionan los mecanismos de generación de la respuesta inmune, y se exploran algunas ideas sobre la capacidad que tiene el sistema inmune de insectos para responder de una manera eficaz a los ataques de patógenos. Por último, se exponen hipótesis que podrían explicar la evolución de

la respuesta inmune, esto considerando la coevolución entre el patógeno y su hospedero; a través de la correlación entre componentes del sistema inmune; y limitaciones generadas por estas asociaciones. Se contempla al estudio de la variación fenotípica de características de respuesta inmune para determinar cómo influye el ambiente sobre la expresión de caracteres cuantitativos (controlados por más de un gen). Se examinan los posibles efectos de la estocasticidad ambiental y efectos maternos sobre dichas características.

El grado de especificidad del sistema inmune de insectos no cuenta con un reconocimiento y respuesta contra agentes infecciosos después de un contacto inicial (como ocurre en vertebrados), es posible que tengan la capacidad de mostrar una mejora su respuesta inmune a lo largo de la ontogenia e inclusive que pueda ser heredable. Existe evidencia (ver Elliot et al, 2003; Kurtz y Franz, 2003; Moret y Siva-Jothy, 2003) que insectos, presentan un fenómeno análogo a la memoria de vertebrados, denominado prevención inmunológica (“immunological priming”; Little y Kraaijeveld, 2004; Little et al, 2005). Los organismos generan una posterior respuesta inmune más eficaz después de haber tenido una experiencia previa con agentes infecciosos. Sin embargo cabe aclarar que para invertebrados el uso del término memoria se limita a una mejora en la respuesta inmune en posteriores retos inmunes. En el *capítulo III* se evalúa la capacidad que tiene el sistema inmune del mosquito para incrementar su supervivencia al ser infectado con patógenos con los que previamente ya ha tenido contacto. De igual forma, se evalúan dos parámetros inmunes (actividad de fenoloxidasa y producción de óxido nítrico) que posiblemente están relacionados con la capacidad de desarrollar un tipo de memoria inmunológica.

Los organismos tienen que enfrentar el problema de cómo maximizar su adecuación (en términos de supervivencia y reproducción) en ambientes heterogéneos y/o estresantes, los individuos con

una constitución genética que permita variaciones fenotípicas para ajustarse a los diferentes cambios ambientales podrían tener una ventaja selectiva en dicho ambiente (Zhivotovsky *et al*, 1996). En algunas ocasiones un solo genotipo puede tener la capacidad de producir varios fenotipos alternativos como resultado de su interacción (y sensibilidad) con diferentes ambientes. A esto se le conoce como plasticidad fenotípica (Roff, 1997; Nylin y Gotthard, 1998). Existe evidencia de que la limitación de alimento afecta negativamente la respuesta inmune en insectos (Fellowes, 1998; Siva-Jothy y Thompson, 2002). Dado que los organismos tienen que enfrentar el problema de cómo maximizar su supervivencia y reproducción en ambientes que están en constante cambio, los individuos con una constitución genética que permita variaciones fenotípicas para ajustarse a los diferentes cambios ambientales podrían tener una ventaja selectiva en un ambiente cambiante (Via *et al*, 1995). Bajo este contexto, en el *capítulo IV* se estimó el componente genético y ambiental de la respuesta inmune, si existe plasticidad fenotípica y si existe variación genética para la plasticidad (componente ambiental e interacción GxA) de la respuesta inmune de mosquitos en condiciones de alimentación limitadas. Asociado a esto, se evalúa si existen diferencias entre los sexos en la forma de responder al ambiente. Es probable que, en caso de existir, las diferencias inmunológicas entre sexos sean consecuencia de la variación en la disponibilidad y tipo de recursos utilizados distintamente por hembras y machos.

En el capítulo IV se expone una breve discusión resaltando los principales resultados obtenidos en esta tesis. Además se proponen perspectivas de estudio y el potencial de aplicación en un futuro del conocimiento generado. Adicionalmente en el *apéndice I* se explican las metodologías comunmente usadas para la cuantificación de respuesta a parámetros inmunológicos. Se describe el contexto ecológico y evolutivo de cada uno de estos parámetros y se ofrecen hipótesis que explican la posible relación entre efectores de la respuesta, recursos necesarios para esto y la

consecuencia ecológica (disyuntivas) y evolutiva. Se ofrecen sugerencias para lograr una óptima medición de la respuesta inmune.

La especie de estudio: Aedes aegypti

Aedes aegypti (Diptera, Culicidae) tiene una amplia distribución entre los trópicos y zonas subtropicales llegando hasta los 40° Sur y 45° Norte, (Nelson, 1986; Badii et al., 2007). Por lo general habita en áreas geográficas con una temperatura media anual entre 17-30°C (Ibáñez-Bernal & Gómez-Dantés 1995). Su rango de distribución altitudinal llega a los 2400 m.s.n.m. (Badii et al 2007). En México el rango de distribución estimado abarca 29 estados (Ibáñez-Bernal y Gómez-Dantés 1995).

A. aegypti tiene cuatro estadios en su desarrollo postembrionario: huevo, larva, pupa y mosquito adulto (Fig. 1.). Los *huevos* son puestos en sustratos sólidos ubicados en la interface agua-tierra. Después de que ha ocurrido la oviposición, se da la melanización, lo que les confiere resistencia a la desecación, entrando en diapausa durante periodos secos, permaneciendo viables hasta 2 años (Christophers 1960). Cada hembra produce de 100 a 120 huevos por puesta (Apóstol et al 1994), con un éxito de eclosión muy alto. La *larva* eclosionada es acuática y pasa por cuatro fases. Las tres primeras fases tienen un desarrollo rápido, mientras que la última es más prolongada que es cuando la larva aumenta de tamaño y peso adquiriendo recursos para el periodo de metamorfosis (Nasci 1986). Por lo general, el tiempo de desarrollo de la etapa larval es de una semana (Christophers 1960), pero en condiciones de baja temperatura o escasez de alimento, la cuarta fase larval puede prolongarse por varias semanas antes de transformarse en pupa (Tun-Lin et al 2000). La mortalidad más alta ocurre durante las dos primeras fases larvales. Su fuente de alimentación se compone de microorganismos, particularmente hongos, algas, protozoarios y otros insectos (Merritt et al., 1992). La *pupa* es la fase de metamorfosis de larva a

adulto, la cual tiene la cualidad de desplazarse activamente en el medio acuático en respuesta a estímulos externos como vibraciones o cambios en intensidad lumínica. Esta etapa del ciclo de vida dura aproximadamente de dos a tres días (Christophers, 1960). Las larvas y las pupas de los machos se desarrollan más rápido que las de las hembras.

El *mosquito adulto* recién emergido pasa sus primeras 24 horas en reposo, periodo durante el cual se completa el desarrollo (Clements, 1999). El mosquito presenta dimorfismo sexual, las hembras son más grandes que los machos, sus antenas tienen vellos cortos y escasos, y los palpos son de un tercio o menos de longitud que la proboscis (Busvine, 1975), mientras que el macho tiene antenas plumosas con pelos largos y abundantes y palpos con un tamaño similar a la proboscis (Busvine, 1975). Los machos son poligínicos, y las hembras presentan un patrón de apareamiento monándrico. Un apareamiento es suficiente para fecundar sus huevos. La conducta reproductiva y de alimentación de sangre (en hembras) aparece entre las 24 y 72 horas post-emergencia adulta (Clements, 1999). Los machos se alimentan del néctar de las flores, las hembras de azúcares y de sangre; esta última le confiere nutrientes para la maduración de los huevos (Clements, 1999). Después de cada alimentación sanguínea se desarrolla un lote de huevos, aunque una alimentación reducida no conlleva a la producción de huevos. Después de 48 a 72 horas post alimentación, se realiza la oviposición (Carrada et al., 1984). Terminada la oviposición, la hembra reanuda la conducta de búsqueda de una nueva fuente de sangre para la producción del siguiente grupo de huevos. El adulto, en condiciones naturales sobrevive en promedio de 15 a 30 días (Badii et al., 2007).

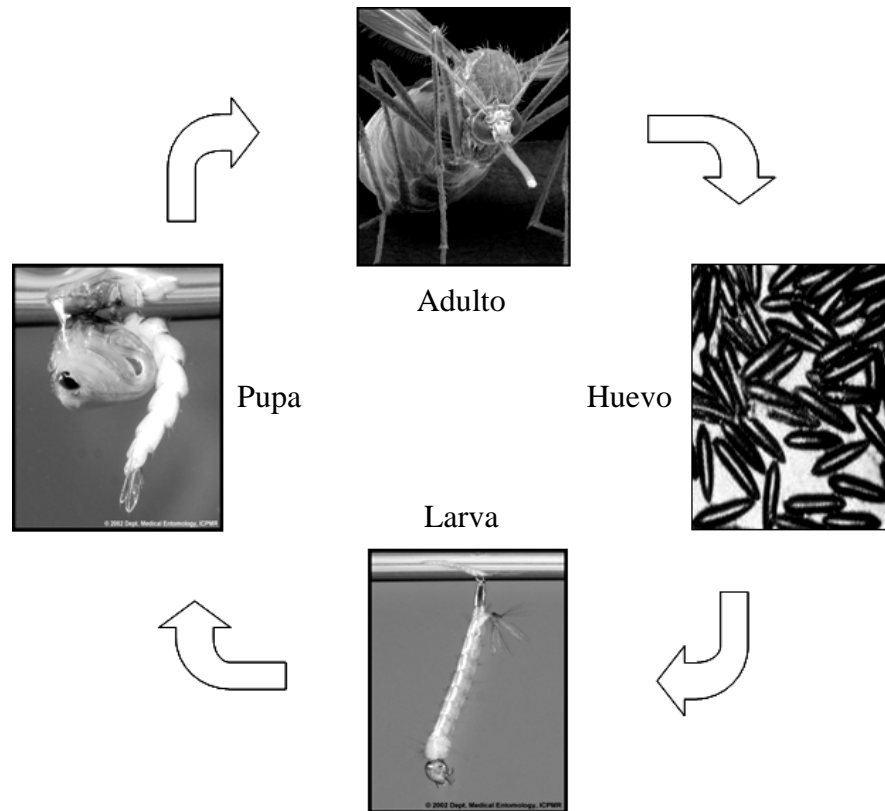


Figura 1. Ciclo de desarrollo de *Aedes aegypti*.

***Aedes aegypti* y el virus Dengue**

El mosquito *Aedes aegypti* es el principal vector del virus dengue, ocasionando fiebre del dengue y fiebre hemorrágica del dengue en humanos (Clarke, 2002). La invasión del virus al mosquito solo ocurre en los casos en que la hembra, al momento de la alimentación sanguínea, pica a un hospedero que porta al virus. Esta hembra ya infectada podrá transmitir el virus a hospederos sanos. La capacidad de diapausa que tienen los huevos del mosquito, rápido ciclo de vida y la preferencia de la hembra por ovipositar en cuerpos de agua relativamente pequeños, son factores que han contribuido en la expansión del rango de distribución del mosquito (Fig. 2). Unido a esto, la preferencia del mosquito a ambientes domésticos en su ciclo de vida y una aclimatación muy rápida a áreas urbanas rurales, y probablemente el cambio climático global, contribuye a que el virus dengue también pueda expandir su rango de distribución (Gubler, 2002), dado que la presencia del vector es indispensable para la transmisión del virus. El control de esta enfermedad

requiere del conocimiento integral del mosquito con una visión holística que genere información para el manejo racional de este vector.

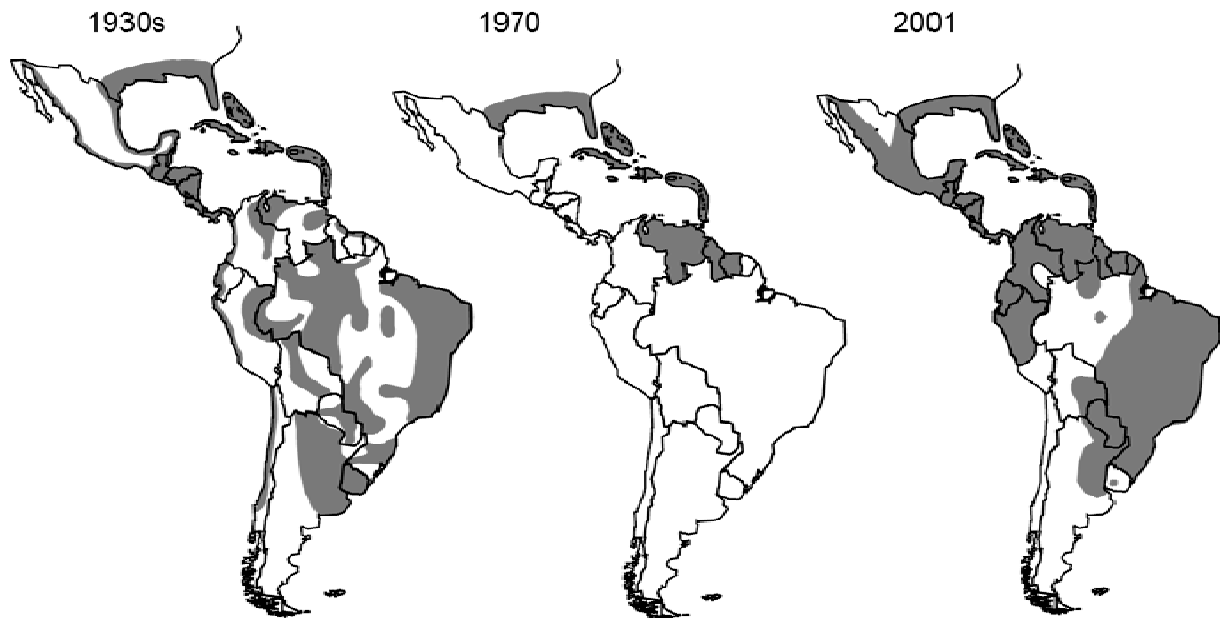


Figura 2. Expansión estimada del rango de distribución de *Aedes aegypti* en América hasta 2001 (tomado de Gubler, 2002).

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CAPÍTULO II

EVOLUTIONARY ECOLOGY OF INSECT IMMUNITY

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ABSTRACT

A large number of recent ecological studies have reported interactions between immunity and traits related to survival and reproduction (i.e. life history traits). In addition, recent molecular/physiological immune mechanisms studies have revealed a wide array of efficient immune responses, which can be enhanced through ontogeny and transferred to the offspring (immunological priming). Here, we review and identify areas where more investigation and integration of information from these different biological fields is needed. We also discuss possible explanations for variation in insect immune response highlighting the ecological and physiological contexts of trade-offs between immunity and other life history traits. We emphasize the analysis of the genetical and environmental bases of variation in immune response, as a way to expand our knowledge of insect immunity evolution and its mechanisms. An interdisciplinary approach will be a key to the advancement of the emerging discipline of immunoecology.

KEYWORDS

Immune response, insects, immunological priming, immunoecology, life history theory, quantitative genetics

INTRODUCTION

The understanding of insect immunity has considerably expanded in recent years. Ecological evidence demonstrates that it is a trait deeply involved in the evolutionary ecology of the reproductive and survival strategies of the organisms. The vast wealth of information on the area, demands integration in order to fully conceptualize the participation and consequences of immune response, and its genetic and phenotypic variation within and among individuals, populations and species. The primary goal of this review is to analyse the current knowledge and ideas related to the adaptive functionality and variation of insect immune response, considering: (1) the ecological context in which immune response is expressed in terms of required resources and related with other fitness-related traits, as well as the evolutionary consequences on life history strategies and the effect of pathogen virulence on the host; (2) the mechanisms responsible for immune response operation with a focus on the possible evolutionary outcome of maladaptive immune responses and individual-primed immune responses through ontogeny and between generations; and (3) the evolutionary features via individual phenotypic variation (quantitative genetics) of immune response, genetic correlations, and the production of different phenotypes according to distinct environmental conditions, as well as the causes and possible adaptive value of this variation. All these ideas are linked using a standpoint that integrates ecological, physiological and population genetics perspectives. It is hoped that this review will facilitate the reflection on encourage a more unified approach to the study of insect immune responses

LIFE HISTORY THEORY AND IMMUNITY

The majority of empirical research in the field of ecological immunity has been under the life history and trade-offs theory perspective. Life history theory seeks to explain the evolution of traits (e.g. size at birth, growth and mortality rates, size and age at maturity, clutch size and reproductive effort) (modified from Stearns, 1976; Roff, 1992; Stearns, 1992) all related with the

life history of an individual, and the changing environmental conditions to which organisms are exposed. These traits are commonly shaped by intrinsic (genetic, or physiological) restriction called “trade-offs” (Stearns, 2000), which affect both survival and reproduction. Trade-offs occur when natural and/or sexual selection is unable to maximize the expression of two or more traits at the same time (Núñez-Farfán, 1993), due to possible genetic or phenotypic correlation with other traits, and despite their potential contribution to fitness (Cheverud, Rutledge & Atchley, 1983). In insects there is growing empirical evidence of trade-offs between these traits and immune defense. Examples include the reduction in larval competitive ability for dietary resources (Kraaijeveld & Godfray, 1997), an increase in predation vulnerability (Rigby & Jokela, 2000), longer developmental time, reduction in egg viability, lower survival rates (Fellowes, Kraaijeveld & Godfray, 1998; Boots & Begon, 1993; Moret & Schmid-Hempel, 2000; Hoang, 2001), and a decrease in reproductive success (Sandland & Minchella, 2003a; Schmid-Hempel & Schmid-Hempel, 1998). Thus, the ability of an insect to minimize an infection (i.e. immunocompetence, Owens & Wilson, 1999) represents an evolutionary cost given its negative covariation with other fitness components (Schmid-Hempel, 2003).

Ecological basis of trade-offs

Infections affect the optimal allocation of host resources, changing the host’s “internal state” (Agnew, Koella & Michalakis, 2000), in terms of energy reserves and viability (Iwasa, Pomiankowski & Nee, 1991; Iwasa & Pomiankowski, 1994). In addition, the required resources used for the immune response are also demanded for the expression of other traits linked with survival and reproduction (Sheldon & Verhulst, 1996), resulting in a trade-off. Organisms have limited resources; as a result, they cannot maximize investment on all life history and/or intermediate traits (e.g. body size, colored traits, behavioral traits, etc.) or immunity. Therefore, organisms must physiologically “decide” how to allocate available resources to maximize fitness.

Optimal allocation depends on the strategies used to increase fitness in the particular habitat of the organism (Schlichting & Pigliucci, 1998).

Trade-offs between the immune response and other traits are usually detected in stressing environments (Sandland & Minchella 2003b), particularly with variables involved in the organism's internal state, or condition such as parasite load or prevalence/intensity of infection (McNamara & Houston, 1996), and external variables (e.g. food abundance, predators, population density, etc). Under favorable environmental conditions, trade-offs can be masked, since the value of life history traits are high (de Jong & Van Noordwijk, 1992) or the costs imposed by immune system activation and the consequent trade-offs are not evident (Moret & Schmid-Hempel, 2000). The occurrence of trade-offs may therefore depend on the amount of resources available, acquisition efficiency and their optimal allocation (Van Noordwijk & de Jong, 1986).

Survival, reproduction and immunity

Life history theory predicts that survival and fecundity are not usually maximized simultaneously (Roff, 1992). For example, current reproduction can negatively affect future reproduction due to the prevailing trade-off between reproduction (which uses resources for the expression of sexually selected traits, quantity and quality of eggs produced, etc.) and survival (i.e. resources used to escape from predators, competence or immunity; Reznick, Nunney & Tessier, 2000). The organism can also allocate its resources in relation to chances of future reproduction which is strongly related to life expectancy. Thus, when there is a high probability of future reproduction, an organism must allocate enough resources to current reproduction in such a way that this investment will not affect subsequent reproductive events (Williams, 1966). On the other hand, when the probability of future reproduction is low (i.e. lack of resources, environmental heterogeneity, incidence of infectious agents), the organism must allocate a large portion of its resources to current reproduction (Fessler

et al., 2005). Although this terminal investment theory was originally laid out for iteroparous organisms, it may be applied to semelparous organisms, as long as reproduction occurs in multiple egg laying events (as with insects) where there is sufficient temporal separation in which the organism is expected to vary in investment. This can be expected in social insects, given their generally long life expectancies and high risk of infection due to living in large colonies compared to non-social insects.

Survival and reproductive strategies adopted by an individual also depend on the force and course of infection. Van Baalen (1998) proposed a theoretical host recovery threshold, thus when the intensity of infection is below this threshold (i.e. no significant damage to the host), the organism should not make any investment to immune response at all. Above, but still close to the threshold, it then pays to allocate more into recovery and survival. However, when infection is high, without recovery possibilities, organism must allocate all resources to reproduction because a new infection may occur almost immediately leaving the animal without survival probabilities. Heterogeneity in food availability and kind and burden (and their interaction) of pathogens will have an effect on the evolution of the survival and reproductive strategies. Social insects seem very adequate to test these ideas as the risk of infection is higher in these animals compared to non-social insects.

IMMUNITY AND SEXUAL SELECTION

Sexually selected traits and immune response

The evolution of the immune response has been also examined under the context of sexual selection (competition to leave more offspring; Darwin, 1871), Recent evidence suggests that trade-offs between immunity and sexually selected traits (SST) favored during competition for mates, are common in nature (e.g. Siva-Jothy, 2000; Rantala *et al.*, 2003; Contreras-Garduño, Canales-Lazcano & Córdoba-Aguilar, 2006; Hosken, 2001; McKean & Nunney, 2001). Hamilton

and Zuk (1982) were the first to suggest that in vertebrates pathogen resistance is correlated with the expression of SST, presumably because, androgenic hormones necessary for the expression of sexual traits and behavior can act as immunosuppressors, giving rise to a physiological trade-off (Folstad & Karter, 1992), and rendering males more prone to infection. Thus, SST only could be expressed if males are able to fight-off infections in spite of the immunosuppressive action of hormones, so that the expression of SST becomes an honest indicator of male immunocompetence both to females (during intersexual selection) and/or males (during intrasexual selection) (Folstad & Karter, 1992).

Insects have fundamentally different hormones to those used by vertebrates (Sheridan *et al.*, 2000), but the rationale would be the same as vertebrates. Insects produce juvenile hormone, which has a physiological effect in several traits, for example: sexual maturation, pheromone production, ovaries development, courtship behavior, morphological polyphenisms (Flatt, Tu & Tatar, 2005) and immune function (Tu, Flatt & Tatar, 2005, Rolff & Siva-Jothy, 2002). This suggests that probable, JH induces or prevents allocation of resources to different functions. In beetles the trade-off between immune function and pheromone production seems physiologically mediated by JH (Rantala, Vainikka & Kortet, 2003). As in vertebrates, only males in good condition (i.e. males that can deal successfully with infections,) will be able to produce and maintain SST. These traits could be assessed by females when basing their mating decisions with the presumable indirect benefit of giving birth to resistant offspring.

One essential piece in the immunocompetence hypothesis is that female offspring would obtain disease resistance genes as signaled by SST. However, the relation between SST and the acquisition of genes related to disease resistance is not necessarily direct. Adamo and Spiteri (2005) recently proposed that females may accrue the direct benefit of avoiding infections when mating with males of current good health status (Able, 1996). Furthermore, selection for healthy

males may not only be maintained via female choice; intrasexual competition can also contribute to the maintenance of honest traits, as long as male-male competition implies energetically costly endurance. If males use signals to indicate their fighting ability, it is then likely that these signals may indirectly reveal immune condition. Contreras-Garduño *et al.* (2006) and Serrano-Meneses *et al.* (2007) found that in the damselfly *Hetaerina americana*, male wing coloration seemed to indicate territorial fighting ability, in terms of energy reserves. However, unlike other calopterygids of more recent origin (e.g. those of the genus *Calopteryx*), *H. americana* males do not court females but simply grab them and invariably mate with them. Wing coloration areas in species where males court females (e.g. *Calopteryx*) and *Hetaerina* males also positively correlate with immune ability. Given these relationships, in the *Hetaerina* case, it makes more sense to admit that SST indicates energetic condition, rather than immune ability to potential fighting contestants. However, even if courtship does not seem to occur in *Hetaerina* it does not reject the hypothesis that females still obtain direct benefits for their offspring if they mate with high fighting abilities (if it has a heritable basis) (Adamo & Spiteri, 2005; Contreras-Garduño, Lanz-Mendoza & Córdoba-Aguilar, 2007).

In some fly and cricket species, male seminal products bear an immunosuppressive effect on the female (McGraw, Gibson & Clark, 2004; Fedorka, Zuk & Mousseau, 2004) diminishing her fitness (Fedorka & Zuk, 2005). Males could increase their fertilization success by reducing female immunocompetence, increasing sperm survival, preventing sperm to be recognized as foreign invading bodies while avoiding female immune response (Fedorka & Zuk, 2005). This sexual conflict perspective for immune response evolution is unique and gives support to this hypothesis, and it should be considered as an alternative to female choice.

Mating systems and differences between the sexes

Sexual selection intensity differs among species depending on their mating system (monogamy vs. polygamy) (Andersson, 1994). This varying intensity should promote differences in the pattern of investment to immune defense by each sex. In polygynous species, males are expected to increase their fitness by reducing their investment on immune defense while investing resources on reproductive effort (for example SST) (Zuk & McKean, 1996; Sadd *et al.*, 2006). On the other hand, natural selection would favor an increase in resource investment to immunity in females, under the assumption that increased longevity could enhance fitness via egg production (Rolf, 2002). In *Acheta domesticus*, females increase egg production when a bacterial infection occurs, compensating the reduced life expectancy and future reproduction due to infection (Adamo 1999). Whereas in another cricket; *Gryllus texensis*, male immunocompetence decreased in the face of an infection, supporting the hypothesis that males trade-off immune response for reproduction (Adamo et al 2001).

Sex biases in immunity may be due to the different cost for SST compared to egg production and laying, and the presumed selection for longer life expectancy for females compared to males (Stoehr & Kokko 2006; Forbes, 2007). One missing piece in the presumed sex bias in immunity is the difference in kinds of pathogens attacking each sex. For example, in mosquitoes, adult females require carbohydrates (sugar) and proteins for vitellogenesis which usually comes in the blood meal. While males, feed on sugar solutions (nectar) (Clements, 1999). This difference does not mean that females are more prone to infections than males, but divergent host-pathogen interactions for each sex are possible and can be reflected in the course of action of individual immune responses. This disparity could lead to wrong conclusions when testing for sex immunity differences, where the same artificial infection (e.g. inoculation of bacteria, yeast, etc.) is employed equally in males as in females. Future studies must consider if males and females share the same pathogens before testing and assuming other reasons for sex immunity biases.

PATHOGEN VIRULENCE, MULTIPLE INFECTIONS AND WITHIN-HOST COMPETITION

In order to explain immunological survival or reproductive differences in the kind of pathogens and its virulence (i.e. harm imposed on a host measured in terms of a reduction in survival or fecundity due to pathogen growth or reproduction) must be considered. Pathogen fitness depends on transmission success to new hosts (May & Anderson, 1983). The transmission increases through enhanced growth or replication rates (Ebert, 1998), but with an associate cost: host fitness. Since this can lead to pathogen dead too, there will be a trade-off between fecundity and longevity for the pathogen (Frank, 1996). Therefore it is expected that pathogens change their virulence depending on the host's natural history. For example, if reproductive adults have a decreased immune response, pathogens should reduce their virulence, although high levels of virulence can be advantageous when infecting non-immune suppressed females (see Pfennig, 2001). Selection could be then favoring the maintenance of sex-or taxon-specific strategies, based on an optimal virulence level understood as that strategy that increases fecundity and longevity of pathogens with low host damage. Different immunological degree responses among hosts are also expected. In order to prove differential effects of pathogens on host, we need measures of differential virulence damage since, currently, variation in virulence is ignored.

An individual can be infected by more than one pathogen lineage (Read & Taylor, 2001). Higher levels of virulence could arise because within-host competition (Frank, 1996). In a mixed infection, low lethal pathogens can be eliminated through competitive exclusion by more virulent pathogens or strains, so selection will favor pathogens or strains less likely to be competitively suppressed (de Roode *et al.*, 2005). Different spatial and temporarily virulence heterogeneity in within-host competition becomes a fluctuating selective pressure altering the expression, variation and evolution of immune mechanisms of insects through generations. How the host's immune ability

varies, according to both virulence and within-host competition, in the context of studies of the evolution of immune response is still an open niche for further investigation.

MECHANISMS OF INSECT IMMUNE RESPONSE

The immune system seeks to maintain a relative homeostatic state under a wide variety of internal and external conditions (e.g. development, infectious agents). The immune responses in insects include cells and products which are interconnected (Hoffmann *et al.*, 1999) (Fig. 1). Here we briefly review general mechanism of insect immunity to build up frameworks of physiological trade-offs, linking immunopathology and immunological priming.

The first lines of defense include the exoskeleton cuticle as a physical barrier, and epidermis, gut epithelium, and male and female reproductive accessory glands (Gillespie *et al.*, 1997; Casteels, 1998). These tissues can secrete cytotoxic molecules like lysozymes, reactive oxygen species (ROS, e.g. superoxide anions, peroxides, hydroxyl radicals) (Schmid-Hempel, 2005a), which are transported to the wound or where infection take place (Nappi & Ottoviani, 2000).

Recongnition of pathogens is needed. The membrane of haemocytes comprises proteins called Pattern Recognition Receptors (PRRs). PRRs recognize conserved molecular features of pathogens called Pathogen-AssociatedMolecular Patterns (PAMPs). Lipopolysaccharides (LPS), mannoses, β -1, 3 glucans and peptidoglycans are considered the most common PAMPs (Gillespie *et al.*, 1997). Lectins are also involved in the recognition of oligosaccharides and polysaccharides present on the pathogen cell membrane (Wilson, Chen & Ratcliffe, 1999). PRRs include the gram-negative bacteria-binding protein (GNPB) and the peptidoglycan recognition proteins (PGRP); the latter includes molecules of long transmembranal form that are secreted into the haemolymph (Leclerc & Reichhart, 2004).

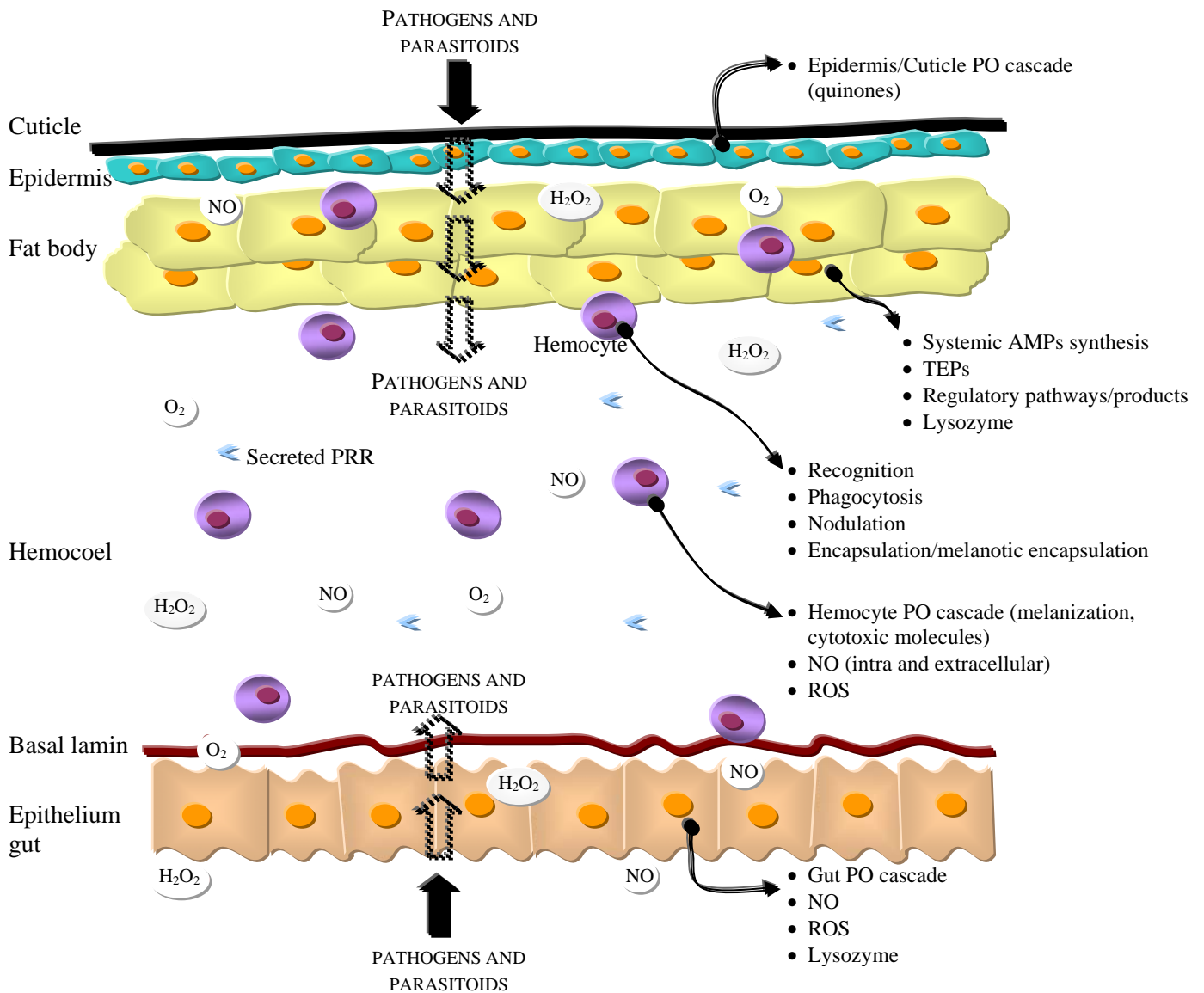


Figure 1. Components of insect immunity.

After recognition, coordinated responses of haemocytes begin. Phagocytosis is a response by which infectious agents become engulfed and destroyed (Gillespie *et al.*, 1997). It may also occur that a large number of haemocytes bind to bacterial aggregations to form a nodule (Gillespie *et al.*, 1997). When infectious agents are large in size (for example, parasitoid larvae), the process of encapsulation, which is similar to nodulation (the difference between both process depends on the size of infectious agents), is used. Occasionally, during encapsulation, a melanin layer is produced to cover parasites which die by anoxia, host generation of free radicals or starvation (Nappi *et al.*,

2000; Narayanan, 2004). Haemocytes also contribute to clot formation by aggregating at wound sites (Gregoire, 1974). Other immune responses are activated, which includes antimicrobial molecules produced within the reproductive accessory glands, gut cells, fat body and haemocytes (Manetti, Rosetto & Marchini, 1998 1998; Schmid-Hempel, 2005a) which are commonly referred as humoral response. Most of these molecules can be secreted into the haemolymph, epitheliums, Malpighian tubes or near cuticle. These molecules include lysozymes with depolymerizing bacteria (mainly Gram⁻) cell wall action (Gillespie *et al*, 1997); Thoiester-containing proteins (TEPs), with opsonization activity that enables phagocytosis (Tzou, De Gregorio & Lemaitre, 2002); nitric oxide (NO), a highly reactive and unstable free radical gas produced during the oxidation of L-arginine to L-citrulline by the nitric oxide synthase (NOS) (Müller, 1997) that crosses cell membranes to act in nearby targets (Müller, 1997), inhibits protein catalytic activity and has protein and pathogen DNA harming effects (Colasanti *et al.*, 2001; Rivero, 2006); ROS, which damages pathogen nucleic acids, proteins and cell membrane (Nappi *et al*, 2000); and antimicrobial peptides (AMPs; e.g. cecropins, atticins, dipterocins, drosomycins, metchnikowins, and defensins, all with isoforms; Lemaitre, Reichart & Hoffmann, 1997; Narayanan, 2004) which induce a collapse of the pathogen membrane and/or prevent the synthesis of molecules within the pathogen (Otvos Jr, 2000; Bulet, Charlet & Hetru, 2003).

The proteolytic pro-phenoloxidase (proPO) cascade is a key immune component for the synthesis of ROS, cytotoxic molecules and melanin (Iwanaga & Lee, 2005). After pathogen recognition, proPO system activation starts. As a first step, phenylalanine is hydroxylated and converted into tyrosine (Christensen *et al*, 2005). Then proPO is activated by a serin protease to its active form PO (an oxidoreductase enzyme) which catalyzes the reaction where tyrosine is transformed into dopa and then to dopaquinone (Söderhäll & Cerenius, 1998). After non-enzymatic polymerizations, dopaquinone is finally converted to melanin which will be used for wrapping up pathogens and

wound clotting (Nappi & Christensen, 2005). Opsonic factors, ROS, and cytotoxins such as quinones and semiquinones are important intermediate molecules, which are highly reactive and toxic to pathogens that greatly amplify immune response (Cerenius & Söderhäll, 2004; Nappi & Ottovianni, 2000).

In *Drosophila*, the immune system is regulated by the products of three signaling pathways: TOLL, Immune Deficiency (Imd) and the Janus kinase/Signal Transducers and Activators of Transcription (JAK/STAT). Regulation is ensured by transcriptional factors produced in the fat body, preventing self tissue damage and keeping haemocyte proliferation and differentiation controlled (Agaisse & Perriomon, 2004). The TOLL pathway bears a key function during antimicrobial peptide production after being activated by fungal and Gram⁺ bacterial infections (Janeway & Medzithov, 2002; Hoffmann & Ligoxygakis, 2004). The Cactus protein, a TOLL transduction pathway product, maintains haemocyte proliferation regulated (Qiu, Pan & Govind, 1998). The Imd pathway gives rise to antimicrobial peptides when activated by Gram⁻ bacteria (Khush, Leulier & Lemaitre, 2002; Hoffmann, 2003). Zaidman-Rémy *et al.* (2006) found that the Imd pathway can be regulated by PGRP-L, preventing host tissue damage as a consequence of extended immune activity. The JAK/STAT pathway regulates the secretion of humoral factors like TEPs (Tzou, De Gregorio & Lemaitre, 2002). This pathway also regulates haemocyte proliferation and differentiation (Agaisse & Perriomon, 2004). Interestingly, not all insects use this differential activation and regulation pathways. For example, in the mosquito *Aedes aegypti* phagocytic and melanization responses are independent of bacterial Gram type (see Hillyer, Schmidt & Christensen, 2004).

Physiology of immune responses and trade-offs

In ecological studies, trade-offs between immune responses and other traits are usually treated as a black box, with very little understanding of their underlying physiological mechanism. This kind of knowledge must be used as a starting point to understand some general aspects about the basis of the trade-offs generated and those that should be looked for. Tyrosine is responsible for coloring the egg chorion in some insects (Li & Christensen, 1993) and other metabolic pathways. Arginine, another essential amino acid, is necessary for NO generation used in immune response (Rivero, 2006), but it is also important for sperm maturation (Osanai & Chen, 1993), egg production (Uchida, 1993), long term memory, chemosensory (antennal lobes, olfaction) and visual information processing (Müller, 1997). Other effectors are the AMPs, which contains fewer than 150–200 amino acids (Bulet *et al.*, 1999). The overall AMPs concentration, in *Drosophila* haemolymph, can reach up to 200 μM (Otvos Jr, 2000), at this concentration, antimicrobial peptides seem costly to produce. Resources (proteins) gathered through ontogeny are indispensable for antimicrobial peptide generation as well as for the synthesis of other molecules. In addition most of the AMPs are cationic, due to their higher content in arginine (Bulet *et al.*, 1999). Consequently arginine could be a limited resource for NO production and AMPs synthesis leading to negative correlations between immune responses. During the PO cascade, tyrosine is used as substrate for the formation of melanin and highly reactive and toxic intermediate molecules. For these reasons, it is likely that tyrosine and arginine are restrictive resources in insects and may lead to trade-offs between immune response and other traits. Experimental manipulation of doses of these amino acids could be used to investigate the insect's response to these changes and trade offs.

Immunopathology: evolutionary and ecological implications

As mentioned before, several molecules (quinones, reactive oxygen and nitrogen molecules, peroxides, etc.) are produced during insect immune response which can be harmful to their own

tissues and cells, a phenomenon referred as immunopathology or autoreactivity. This process has been frequently ignored as a possible constraint for the insect immune system (Schmid-Hempel, 2005a).

It can be argued that natural selection may favor high responsiveness to ensure control of pathogens despite the risk of immunopathology (Graham, Allen & Read, 2005). Nonetheless, activation and upholding of immune response depend on the kind and burden of pathogens, and a novel pathogen could elicit responses beyond immunological control with potentially harmful effects. Mechanisms and molecules produced to kill pathogens could be over expressed or overproduced if monitoring, regulatory, and recognizing systems are not synchronized. On the other hand, if virulence of pathogens elicits autoreactive host responses, low virulence could be favored. Only in cases where competition among pathogens occurs, natural selection will favor high levels of virulence. In both cases there must be fitness advantages to the pathogen regardless of autoreactivity and decrease in host survival. In some occasions hosts could kill pathogens, but bears an extreme survival cost. Meanwhile other species may suffer the damage generated by pathogens but, nevertheless they could be able to survive and reproduce, so a large immunological response could not be the best strategy. The strategy adopted via immune mechanisms and their regulation will depend on pathogens (probably not all novel pathogens will elicit the same degree of autoreactivity), differences in host immune response of the hosts (i.e. unstressed individuals with much to invest in immunity) may be more prone to immunopathological costs (Graham *et al.*, 2005), and cost and benefits between autoreactivity and fitness of pathogens and hosts.

Furthermore, immune response suppression during stressful and energetically demanding situations (e.g. molting or during display of sexual traits) can be another strategy to avoid autoreactivity (Kotiaho, 2001 and references therein). In this sense, it is not resource limitation the factor that

produces the phenomenon, the assumed trade-off, but a type of “on-off” switch regulation of the immune system during ontogeny.

Immunological Priming

It has been argued that invertebrates are not capable of developing any kind of immunological memory (as vertebrates do). Nevertheless, recent experiments (Faulhaber & Karp, 1992; Moret & Schmid-Hempel, 2001; Kurtz & Franz, 2003; Moret & Siva-Jothy, 2003) have provided evidence of a phenomenon similar to that in with vertebrate acquired immune response which has been called immunological priming (Little & Kraaijeveld, 2004; Little, Hultmark & Read, 2005; Kurtz, 2005; Schmid-Hempel, 2005a, b). This process assumes that previous experience with infectious agents might enhance individual immunity which can be transmitted to subsequent generations (transgenerational priming) (Kurtz & Franz, 2003; Little *et al.*, 2003; Rahman *et al.*, 2004; Sadd *et al.*, 2005; Moret, 2006; Sadd & Schimid-Hempel, 2006; Pham *et al.*, 2007). This novel idea is nowadays being explored in detail by physiologists and immunoecologists, and, given its key importance, has been coined as prompting a new era in the evolutionary and ecological studies of the immune system (for experimental designs see Little & Kraaijeveld, 2004).

The immune invertebrate memory argument has been criticized (Klein, 1997). The basis of this is the lack of mechanisms for specific Ig production, rearrangement of somatical genes and clonal expansion. Then, if an immune response is not specific it makes no sense to refer it as “memory”. So, the term “anticipatory response” could be more suitable in a system with an enhanced but unspecific response after secondary immune challenge (Kurtz, 2004). The specificity of molecules with potential memory function, which enable the host to adapt to infectious agents during its lifetime, needs further investigation. Until then, the term memory must be used with caution. That is why we consider adequate to use the term “immunological priming” to refer to any anticipatory

response with or without specific memory in insects (and invertebrates) (see also Schmid-Hempel, 2005b). Below we put forward some general mechanisms proposed as for how this priming could operate, however, these possible mechanisms need further investigation.

Lectin bindings (Kurtz & Franz, 2003), AMPs (Little *et al.*, 2003, Schmid-Hempel, 2005b) and PRRs (Kurtz, 2005) could be the molecules involved in the priming process. Lectins, through their high structural diversity, highly specific microorganism recognition, agglutination action and opsonization role in phagocytosis (Marques & Barraco, 2000), could be mediators for the enhancing of secondary immune responses. AMPs show great structural diversity (more than 170 isoforms have been found in insects; Bulet *et al.*, 1999) and are produced soon after foreigner recognition (1-4 hours) with a pathogen-specific and efficient killing action (Bulet *et al.*, 1999, Otvos Jr., 2000, Schmid-Hempel, 2005b). Although transcription of AMPs became turned off, some AMPs may remain in haemolymph for up to three weeks (Schmid-Hempel, 2005b), which is convenient during subsequent pathogen encounters. PRRs are differentially induced depending on the infectious agents, and particularly the recognition proteins secreted into the haemolymph could be molecules conferring specificity to invertebrate immune response.

It cannot be excluded the possibility of somatic gene alterations of immune molecules to enhance the response to infectious agents (Flajnik, Miller & Du Pasquier, 2003). In fact, current knowledge shows that some invertebrate IgSF proteins called Down syndrome cell adhesion molecule (Dscam; Watson *et al.*, 2005). Dscam proteins are used as opsonization factors that enhance the phagocytic efficiency of haemocytes, increasing host survival (Dong *et al.* 2006). Also, the challenge with different pathogens induces specific Dscam repertoires with different affinity, showing specificity. In snails, fibrinogen related proteins (Freps) somatically diversify (like Ig of vertebrates) and are produced in the haemolymph in the presence of pathogens (Zhang *et al.*, 2004). The genetic and

molecular mechanism underlying these rearrangements also needs further investigations. However, given the diversity of insect species, it cannot be excluded the presence of a greater diversity of mechanisms behind immunological memory.

Enhanced immunity of offspring from mothers that were immune stimulated, has been also reported. Mother AMPs (or lectins and PRRs), and also mRNA (Huang & Song, 1999), in a transcription factor-like mode, could be promoting transcriptional initiation of immune response genes. So, there can be heritable changes in immune response among generations that cannot be due to DNA sequences changes or genetic variation, i.e. inherited epigenetic variation (Richards, 2006). DNA methylation can modify histones. These modifications alter affinity of proteins that mediate transcription and affect interaction between nucleosomes and chromatin. As a consequence gene expression and phenotype could be affected (Richards, 2006). As a result heritable changes in gene expression could affect immune response outcome. Therefore, these molecules could be working as elicitors that could interact with the offspring embryonic cells inducing defense; so when immature stages start fighting against infectious agents, they are already primed, increasing survival chances (Rahman *et al.*, 2004).

This idea is not so radical, in distant organisms such as plants. Molinier *et al.* (2006) found that immune elicitors increased epigenetic somatic homologous recombination of a transgenic reporter which can persist in subsequent generations, which could potentially enhance offspring defenses. Although we are not proposing homologous recombination epigenetic change as the general mechanism of immune transgenerational priming, the fact is that transgenerational memory is not an exclusive property of vertebrates and therefore, can take place in different plant and animal groups.

Other relevant issue of priming is the high energetic cost of prolonged activation or synthesis of immune molecules at a higher level. Since pathogens can act as a factor that changes the host's internal environment, an optimal resource allocation is likely to change over time, partly because infections are unpredictable and their impact is not usually immediate. One can expect that the first contact with a pathogen not only enhances immune response, but changes the host's life history strategies. For example, reproductive events, before a secondary infection occurs. It has been widely documented that survival is impaired when hosts devote more resources to immune defense (Schmid-Hempel, 2005a). However, the direction of change in reproductive strategies under priming is unclear. If immune priming occurs, not only survival or immune response need to be measured, but other important trait intimately linked with reproduction (e.g. egg production).

QUANTITATIVE GENETICS OF THE IMMUNE RESPONSE

Traits closely related to fitness are commonly under natural or sexual selective pressures. Traits in ecological and evolutionary analyses are continuous variables and their variation is thought to be polygenic and intimately affected by the biotic and abiotic environment. These traits must be analyzed using quantitative genetics, by partitioning and estimation of variances and covariances into causal components (Falconer & Mackay, 1996) to understand how they respond to selection. Immune response occurs via a set of physiological traits and is supposed to show low additive variance and hence low heritability values (like other physiological traits, see Mousseau & Roff, 1987). Nevertheless, recent studies have shown that different immune components such as haemocyte load (Ryder & Siva-Jothy, 2001; Cotter, Kruuk & Wilson, 2004a; Rolff, Armitage & Coltman, 2005; Simmons & Roberts, 2005), antibacterial activity (Kurtz & Sauer, 1999; Cotter *et al.*, 2004; Simmons & Roberts, 2005), PO activity (Hosken, 2001; Cotter & Wilson, 2002; Rolff *et al.*, 2005; Schwarzenbach, Hosken & Ward, 2005) and melanization (Fellowes *et al.*, 1998; Simmons & Roberts, 2005) tend to have high heritability (from 0.24 to 0.91), indicating high

genetic variance. There are at least three non-mutually excluding hypotheses to explain variance of immune response: host-pathogen co-adaptation cycles, genetic basis of trade-offs (genetic correlations), and phenotypic plasticity.

Host-pathogen co-adaptation cycles

Genes involved in immune resistance always show significant variance because of host-pathogen co-adaptation cycles (Hamilton & Zuk, 1982). The pathogen's short generational cycles may provide enough time to adapt to the host's immune response and given the number of infectious agents, there will be grounds for high genetic variability in immune responses. Variation will allow the host fighting against all varieties of infections produced by different pathogens. This has been used as the explanation for the maintenance of variation in SSTs which are under intense sexual selection (for example female choice; Kirkpatrick & Ryan, 1991). According to a perspective of sexual selection, if genetic variation of immune response is maintained, females (if female choice is the selective filter, but the same may apply for male-male competition and sexual conflict) will be able to continue exerting mate choice over SSTs (as these covariate with immune ability, reflecting male genetic quality), which will be passed on to offspring (Roff *et al.*, 2005).

Correlation among immune effectors

In an individual, different traits are frequently found to be genetic or phenotypically correlated. Genetic correlation arises because a single gene can influence multiple traits in a positive and negative fashion (pleiotropy and antagonistic pleiotropy, respectively) or because of linkage disequilibrium between genes affecting different characters (Falconer & Mackay, 1996). Meanwhile phenotypic correlations include the genetic causes and the positive or negative influences of environmental factors between traits (Roff, 1992). Negative correlations between immune responses and other life history traits (or intermediated traits) have been found and are

clearly detected under stressful conditions (usually resource limitation). However, trade-offs within immune system parameters are also possible. A number of studies have examined genetic and phenotypic correlations among encapsulation, lytic activity, cuticular darkness, PO activity and haemocyte load, showing an unclear pattern with positive and negative correlations (see Rantala & Kortet, 2003; Cotter *et al.*, 2004b; Fedorka *et al.*, 2004; Ryder & Siva-Jothy, 2004; Rantala & Roff, 2005; Roff *et al.*, 2005).

It must be considered that not all pathogens will induce the same reaction, or at least the magnitude of response can differ depending on the host-pathogen interaction type (coevolutionary history). So, genetic correlations, type and kind of pathogens could be closely related with the type and the intensity of correlations. The evolutionary trajectories of different immune responses can be difficult to predict when hosts are exposed to changing pathogens. Pathogens could differ spatially and temporarily in occurrence, burden and virulence so genetic/phenotypic correlations could change among generations. Therefore, pathogens to which different organisms have been exposed to may have played a role in the evolution of immune responses. Natural or sexual selection will be acting simultaneously on the different immune responses constraining their independent evolution, but probably with a fitness advantage for the host.

Phenotypic plasticity

As long as the organism has to deal with the problem of maximizing fitness in changing or stressful environments, a genetic background which can express changes in phenotype expression could have a selective advantage in that environment (Zhivotovsky, Feldman & Bergman, 1996). Occasionally, a single genotype can have the ability to produce distinct phenotypes when exposed to different environments, a phenomenon known as phenotypic plasticity (Schlichting, 1986; Roff, 1997; Nylin & Gotthard, 1998). Phenotypic plasticity could explain the maintenance of genetic

variance because plasticity uncouples the phenotype from genotype, buffering the impact of natural or sexual selection in the gene pool (Stearns, 1992), leading to a slow depletion of genetic variance. Phenotypic plasticity in quantitative traits therefore represents a genetic response to environmental heterogeneity. The quantitative and qualitative differences in immune response could be used as an indicator of the strategies followed by an organism in relation to environment stochasticity, and this is one reason why the evolution of immune response must be intimately related to the adaptive evolution to environmental heterogeneity. Despite the evidence that immune response can be strongly affected by the environment, really only a handful of studies have approached immune response using a phenotypic plasticity perspective. Related to this, Barnes & Siva-Jothy (2000) in beetles, Cotter *et al.* (2004b) in butterflies (reared at different population densities) and Mucklow & Ebert (2003) in water fleas, found phenotypic plasticity for mounting an immune response, concluding that investment in immunity depends on the infection probability based on the number of conspecifics, and that re-distribution of resources is adaptive. However, at the same time that some immune effectors are elevated at high population densities, while others are not. Since immune response is costly to express, this flexibility to cope with fluctuating environments could give fitness advantages.

The adaptive value of phenotypic plasticity depends on the ecological context in which it is expressed. Other ecological studies arguing the adaptive value of phenotypic plasticity (see Stirling, Roff & Fairbairn, 1999; Gebhardt & Stearns, 1993; David, Capy & Gauthier, 1990) have concluded that alternative strategies can be equally adaptive in some environments, but occasionally may not render advantages for the organism in other environments. Even the parental environment could have effects in the offspring behavioral phenotype (e.g. gregarious and solitary), related to immune response enhancing (Elliot, Blandford & Horton, 2003). It is therefore necessary to evaluate if different environments produce different phenotypes and if genotypes

respond differently to these environments (genotype-by-environment interaction, which represents genetic variation for phenotypic plasticity) and if these alternative phenotypes have fitness advantages.

Due to these reactions, phenotypic plasticity should be carefully considered in immunological studies. For example, it may explain biases in immune response between sexes (Joop & Rolff, 2004; McKean & Nunney, 2005), which may represent different male and female reproductive strategies. Sexual differences may not be genetic but simply a differential phenotypic expression under varying environment regimes. Phenotypic plasticity should be carefully analyzed in current immunocompetence hypotheses of male quality. It can be argued that a possible sexual selection cost of the expression of phenotypic plasticity is that the phenotypic traits may not be good indicators of male immunocompetence ability, because there is not a direct concordance between the phenotype, immune response and genotype (for analogous discussions see Adamo & Spiteri, 2005). Nevertheless in a broad sense immunocompetence could not only include the capacity of an individual to resist an infection but also the ability to produce distinct immune response phenotypes when exposed to different environments. This line of thought will enrich our comprehension of what male quality, in immunity terms, may mean and its interpretation in sexual selection studies.

Maternal Effects and Immunological Priming

Maternal effects arise when a mother's phenotype, which is intimately correlated with the environment she experiences, has a phenotypic effect on her offspring (Mousseau and Fox 1998). The local maternal environment might influence offspring phenotype, as maternal experiences may provide an indicator of the environmental conditions that offspring will face (Fox & Mousseau, 1998; Rossiter; 1996). If mothers have been exposed to infectious agents, information could be inherited (via maternal lectins, AMPs, PRRs or mRNA; already discussed previously), inducing

offspring phenotypic variation in immune defense (a kind of transgenerational phenotypic plasticity). There are four ecological conditions for transgenerational phenotypic plasticity in immune response to evolve: (1) infectious agents are variable and unpredictable on time, (2) a direct relation between cue and response exists, (3) induced defense is effective and (4) immune response is costly to express (adapted from Harvell, 1990; Harvell & Tollrain, 1999). For these reasons, maternally inherited effects have been proposed as the main explanation for transgenerational priming (Little *et al.*, 2003; Grindstaff, Brodie III & Ketterson, 2003; Sadd *et al.*, 2005; Moret, 2006). Again little evidence has been forwarded to support this.

Little & Kraaijeveld (2004) have proposed immunological priming is expected to evolve under high longevity and clonal reproduction. However, nearly any organism (clonal or sexual, short or long-lived) may be attacked repeatedly by a disease as this is temporally and spatially unpredictable. If, an infectious agent appears in the population, there is a high probability of being reinfected by the same agent and it is the same for the offspring; this is because as prevalence increases, the infectious agent also can increase in frequency (Moret & Siva-Jothy, 2003). Furthermore, immunological (transgenerational) priming can evolve by natural selection in gregarious or social organisms with low mobility and overlapping generations. Additionally, maternal inheritance can produce time lags in the response to selection, hence low depletion of genetic variance. The response to selection will depend on the kind of pathogens (and their virulence) and the evolutionary response for those in the previous generation, as well as the co-occurrence of the same pathogens in the current generation.

Heritability measures and artificial selection experiments (see Kraaijeveld & Godfray, 1997; Fellowes *et al.*, 1998; Rahman *et al.*, 2004), have shown that immune response can evolve. The phenotypic differences between individuals are related with differences in survival and

reproductive success. Natural or sexual selection will act on phenotypic variation, favoring some phenotypes. As long as the phenotypic variation has a genetic background, the population can evolve. The fact that the immune response is enhanced across generations necessarily implies that transgenerational immunological priming not only depends on the offspring's phenotype but on the mother's phenotype too (Kirkpatrick & Lande, 1989). So, variation among mothers in the molecule transmission capacity can also be under selective pressure.

Future studies are essential to understand the mechanisms of intra and intergenerational priming. Until these results are available and similar to what we discussed before with phenotypic plasticity, it is necessary to be careful when interpreting enhanced immunity ability during ontogeny and its heritability basis. As mentioned before, organisms and their offspring are under constant attack by infectious agents, which occasionally lead to infections. However, at other times, the host cannot totally eliminate the pathogen but, just keeps it "under control". This control can be due to the organism's ability to express different survival and reproductive strategies in response to stressful environmental conditions to which is exposed during its lifetime. Previous experience with infectious agents could also generate the expression of behaviors aimed to avoid negative effects of infections (see Loehle, 1995; Moore, 2002). For example behaviors to avoid areas with parasites (Hart, 1990), foraging behaviors for increase resource intake, and also physiological changes for the optimization of resource allocation.

CONCLUSIONS

This review illustrates some evolutionary and ecological principles that can direct the study and understand how the immune system fights infectious agents and how far we are from merging the evolutionary, ecological and physiological disciplines to understand the evolution of immune response. Evolutionary biologists have focused on quantifying phenotypic variation rather than on

understanding its nature this by means, the immune mechanisms, the underlying trait architecture and trade-offs. Ecological insights can be very relevant to understanding immune system dynamics and vice versa. Since immune system is highly complex, several aspects of immune response must be therefore integrated (e.g. immunopathology, immunological priming) to understand the evolutionary ecology of insect immunity. We have attempted to look at the available data on insect immune response pointing out some research avenues and careful considerations. The starting line in insect immunoecology must incorporate the great range of environmental factors involved using a trade-off framework. Resource allocation and investment strategies should be studied by taking into account differences between the sexes, evidence of the real cost in natural condition and their effects in the expression of immune response and generational changes. It must be also considered that host life history traits are evolving in response to selection pressures imposed by the type of pathogens and the interactions among them. Also the mechanisms, the metabolic pathways, synthesis and molecular properties of the products of the immune system cannot be ignored.

It is necessary to include a framework that integrates the host's strategies and the process of coevolution with parasites to understand the host immune response. Finally, it is highly desirable that ecologist and evolutionary biologist alike must incorporate useful molecular techniques (e.g. microassays, proteomics, quantitative Reverse Transcriptase-PCR, etc.) to study individual variation within a framework of population genomics and genetics to identify the mechanisms and the genetic basis of trade-offs between immunity and other life history traits.

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1 **Running head: Anticipatory immune response in *Aedes aegypti***

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8 **Anticipatory immune response in *Aedes aegypti* against bacterial challenges and**
9 **its effects on female reproduction**

10

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24

25

26 **Abstract**

27 In insects exposure, to pathogens induces an anticipatory immune response, a phenomenon
28 called immune priming. However, little is known of whether immune effectors improve their
29 efficiency with time; the temporal dynamics of immune enhancement; and, its reproduction
30 costs. In this study, we demonstrate that priming (using live bacteria) protects the mosquito
31 *Aedes aegypti* against a lethal second challenge. We experimentally reveal the temporal
32 dynamics of phenoloxidase (a key immune effector) measurement which suggests a
33 deactivation rather than activation; this may be due to host use of the enzyme to control
34 recurrent infection with live bacteria. Meanwhile, nitric oxide (another key immune effector)
35 increased to some extent after second challenge. Finally, that egg production was not affected
36 during the priming dose and second challenge; however, the number of egg laying females
37 with two recurrent infections (i.e. priming dose+second challenge) was lower compared to the
38 unprimed groups. Our study not only corroborates the presence of insect immune priming, but
39 also documents the point that not all immune parameters can be enhanced when hosts have had
40 previous contact with pathogens. Immune priming may be considered not only as an enhanced
41 response, but an optimal response to protect the organisms after a previous experience with an
42 elicitor.

43

44 **Keywords:** immunological priming, phenoloxidase, nitric oxide, *Aedes aegypti*

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52 **Introduction**

53 It is widely acknowledged that, unlike vertebrates, insects do not have mechanisms for specific
54 Ig production, rearrangement of somatic genes and clonal expansion (Janeway, 2005). Also,
55 there are no molecules with potential specific immune memory function (but see Zhang *et al.*,
56 2004; Watson *et al.*, 2005), which enable the host to adapt to infectious agents during its
57 lifetime. However, it can be safely assumed that a vast majority of insects are attacked
58 repeatedly by the same pathogen because the incidence of the latter is temporally and spatially
59 unpredictable. If, for example, an infectious agent appears in the population, there is a high
60 probability of a host being re-infected by the same agent. This is because prevalence of the
61 infectious agent increases in the host population (Moret & Siva-Jothy, 2003). Thus, the ability
62 of an anticipatory immune response in non-vertebrate organisms could represent an adaptive
63 characteristic to such environmental selective pressure.

64
65 Despite the lack of the physiological and biochemical machinery that might allow for an
66 adaptive immune response in insects, it has been long recognized that immunization with
67 killed or living bacteria can induce an anticipatory immune response in insects (Boman *et al.*,
68 1972). This fact has been recently supported by experimental observations (Faulhaber & Karp,
69 1992; Moret & Schmid-Hempel, 2001; Kurtz & Franz, 2003; Moret & Siva-Jothy, 2003). The
70 phenomenon has been coined as immune priming (Little & Kraaijeveld, 2004; Little, Hultmark
71 & Read, 2005; Kurtz, 2004; Schmid-Hempel, 2005a, b) and assumes that a previous
72 experience with an elicitor might raise an enhance immune response when re-exposed to the
73 same agent (Pham & Schneider, 2008).

74
75 Protective enhancement of immune response due to priming (i.e. prolonged activation or
76 increased levels of previously immune stimulated organisms) has been assessed using different
77 response variable such as survival, behavior, reproductive capacity, and different immune

78 effectors (phagocytosis, antimicrobial peptide load and synthesis, and antimicrobial and
79 phenoloxidase activity) (Deverno *et al* 1983, Adamo, 1998, Rosengaus *et al* 1999, Brown *et al*
80 2003, Little *et al* 2003, Sadd & Schmid-Hempel 2006, Pham *et al* 2007). In these studies,
81 immune response became more effective upon repeated exposure to the elicitor, an effect that
82 probably persists during the host's lifetime (see model in Fig. 1). However, there is
83 controversy with respect to whether such efficiency applies to all immune effectors. For
84 example, Pham *et al* (2007) found that hemocytes, through phagocytosis, are responsible for
85 an enhancement effect, which was not the case for antimicrobial peptides or the
86 prophenoloxidase (proPO) system. One potential reason as for why immune effectors
87 apparently vary in effectiveness is that they may be better explained by methodological rather
88 than physiological factors. In particular, the moment of quantification of immune parameters
89 should be controlled, while characterizing the dynamics of immune response after a priming
90 dose. This will detect how fast improvement (if it occurs) is acquired. To our knowledge this
91 information has not been gathered by previous studies.

92
93 Priming has been referred to as either as “short” (hours, or few days) or “long” (weeks or
94 lifetime) in terms of immune protection persistence (Pham & Schneider, 2008). However these
95 definitions have yet to be concretely established, as several studies have reported different
96 results. One explanation for these inconsistencies is the variable period of time elapsed
97 between challenges, as well as the nature of the immune response being evaluated in these
98 cases. For example, in the lepidopteran *Heliothis virescens*, a second re-infection was
99 performed at 48 hrs after a priming dose, but immune enhancement reached only 3% after 12
100 hrs (Ourth & Parker, 2006); thus, these authors concluded that in this species priming is not
101 elicited. In this specific case, however, it is likely that the second challenge overlapped with
102 the response generated by the priming dose, avoiding the results of the prime effect. Related to
103 this explanation, in the moth *Galleria mellonella*, humoral antibacterial activity was noted 5 hr

104 after infection, reaching a maximum level at 18-48 hrs post infection, and decreasing over a
105 few days (Jarosz, 1993; Andrejko et al, 2009). Conversely, in other species, immune
106 parameters such as hemocytes and the PO system are very rapidly activated but quickly turned
107 down (Schmid-Hempel & Ebert 2003). Therefore, if priming occurs in these other species, the
108 second challenge and quantification of immune parameters should be performed after the
109 overall immune recovery period (Fig. 1). In general, all these results mean that the nature of
110 immune response time dynamics needs to be taken into account when looking for immune
111 priming effects.

112
113 Another important, yet unexplored, aspect relevant to the time of inducible protection is the
114 high energetic cost of prolonged activation or synthesis of immune molecules. A high
115 energetic cost is expected, as the required resources used for immune response can also be
116 demanded for the expression of other traits linked with survival and reproduction (Sheldon &
117 Verhulst, 1996, Roff, 1992), as predicted by the resource allocation theory (Stearns, 1992).
118 Modulation of the immune response can be expected after previous challenge in order to an
119 optimal adjustment of resources used in immunity and other traits (see Pham & Schneider
120 2008 for a similar claim) (Fig. 1). Then, a recovery period can be expected for resource
121 acquisition and a suitable enhanced immune response.

122
123 Organisms must physiologically “decide” how to allocate available resources to maximize
124 fitness. An optimal allocation depends on the strategies used to increase fitness in the habitat
125 of the organism (Schlichting & Pigluicci, 1998). Since pathogens can act as a factor that
126 changes the host’s internal environment, an optimal resource allocation is likely to change
127 over time, partly because infections are unpredictable and their impact is not usually
128 immediate. One can expect that the first contact with a pathogen not only activates immune
129 response, but changes the host’s life history strategies, for example, reproductive events,

130 before a secondary infection occurs. It has been widely documented that survival is impaired
131 when hosts devote more resources to immune defense (reviewed by Schmid-Hempel, 2005a).
132 However, the direction of changes in reproductive strategies under recurrent infections is
133 unclear.

134
135 Using *Aedes aegypti* Linnaeus (Diptera: Culicidae), the principal vector of Dengue and Yellow
136 Fever virus, as our study subject, we have first documented its survival following a challenge
137 with a lethal dose of the bacterium 7 days after being previously primed with a non lethal dose.
138 We consider this period of time long enough to detect an effect (if it exists) between the
139 priming dose and the second challenge. This period was based on information about the
140 activation and synthesis of immune molecules in insects (for example, antimicrobial peptide
141 transcription has a range from 6 hrs to 72 hrs after challenge; PO activation ; (Jarosz, 1993;
142 Lemaitre et al 1997; Uttenweiler-Joseph et al 1998; Schmid-Hempel & Ebert 2003; Andrejko
143 et al, 2009). Second, we have documented the temporal dynamics of two immune parameters
144 during the priming dose period and also after a second challenge. The immune parameters
145 used were phenoloxidase activity (PO) and nitric oxide production (NO), two key components
146 during insect immune defense. In insects, PO (an oxidoreductase enzyme), catalyzes the
147 transformation of tyrosine into dopa and then to dopaquinone (Söderhäll & Cerenius, 1998).
148 After a number of non-enzymatic polymerizations, dopaquinone is finally converted to
149 melanin which is used for wrapping up pathogens and wound clotting (Nappi & Christensen,
150 2005). Also, during the PO cascade highly reactive and toxic molecules are produced (opsonic
151 factors, reactive oxygen species, quinones and semiquinones) (Cerenius & Söderhäll, 2004;
152 Nappi & Ottovianni, 2000). NO is a highly reactive and unstable free radical gas produced
153 during the oxidation of L-arginine to L-citrulline by the nitric oxide synthase (NOS) (Müller,
154 1997). NO crosses cell membranes to act on nearby targets (Müller, 1997), inhibit protein
155 catalytic activity and harms protein and pathogen DNA (Colasanti *et al.*, 2001; Rivero, 2006).

156 Finally, and in terms of changes in reproductive strategies and resource allocation theory, we
157 have documented the number of eggs produced and the proportion of egg laying females with
158 and without a priming dose and with recurrent immune challenges

159

160

161 **Materials and Methods**

162 Mosquitoes were reared under insectary conditions at the Instituto Nacional de Salud
163 Pública (INSP), Cuernavaca, Mexico (the colony stock has been maintained at over 2000
164 individuals per generation with random mating). Three day-old adult female mosquitoes were
165 used at the beginning of each experiment. We used live or heat killed *Serratia marcescens* and
166 *Escherichia coli* (both Gram-negative bacteria). Both bacteria were kindly donated by Lilia
167 González Cerón (Centro de Investigaciones en Paludismo, INSP) and Jesús Silva (Centro de
168 Investigaciones Sobre Enfermedades Infecciosas, INSP) respectively. Bacteria were grown
169 overnight in LB-broth at 37°C while shaking at 200 RPM until they reached stationary phase.
170 To determine bacterial concentrations, 100 µl of a 1/1000, 1/10,000 or 1/100,000 bacterial
171 culture was spread on LB agar plates. Plates were grown overnight at 37°C and the colony
172 forming units (CFU) were counted. LD doses were previously determined by inoculating
173 separate groups of mosquitoes with serial bacterial dilutions and selecting the dose closest to
174 killing 100%, 50% , <10% of the animals. We recorded mosquito mortality during the next 24
175 hours after bacterial injection.

176

177 *Survival experiment*

178 For the survival analyses, two experiments were carried out. In the first experiment, three days
179 old adult female mosquitoes were randomly divided into four groups of 100 individuals each:
180 control and three experimental groups. For the priming dose, mosquitoes of group one (RPMI-
181 G) were refrigerated and then inoculated with RPMI directly into the hemocele. The second

182 group (Live-B) was inoculated with live *S. marcescens* ($LD_{<10}$; 74×10^4 CFU) suspended in
183 RPMI. The third group (Dead-B) was inoculated with dead *S. marcescens* (at 74×10^4 CFU)
184 suspended in RPMI. The Control group (C) was only refrigerated at 4°C for about 10 minutes.
185 At day 7 post priming dose, mosquitoes were re-infected via inoculation in the abdomen (see
186 below). For the second challenge RPMI-G, Live-B and Dead-B, were inoculated with live *S.*
187 *marcescens* at a higher concentration ($LD_{>50}$; 12.3×10^5 CFU). C group was again refrigerated.
188 Survival was quantified until the live mosquito proportion ceased to change.

189
190 For the second experiment we used *E. coli*. Three groups (Live-B, RPMI-G and C) of 100
191 females each were used. Dead-B was not included given that previous observations showed
192 that dead bacteria do not elicit priming condition (unpublished data). For the priming dose,
193 Live-B was inoculated with bacteria at 92.5×10^3 CFU ($LD_{<10}$), RPMI-G was inoculated with
194 RPMI medium, and C (negative control) group was only refrigerated. At day 7 post priming
195 dose, mosquitoes were re-infected via inoculation in the abdomen (see below). RPMI-G and
196 Live-B were inoculated with a higher dose of bacteria ($LD_{>50}$; 10.6×10^4 CFU). For this
197 experiment we included a primed RPMI-G challenged again with RPMI medium as positive
198 control. Survival was quantified until live mosquitoes proportion ceased to change.

199
200 All inoculations were made using a pulled glass needle attached to a Drumond microinjector.
201 For manipulation, mosquitoes were previously refrigerated at 4°C for 10 min. Organisms were
202 always injected in the abdomen close to the junction between the ventral and dorsal cuticles.
203 The volume of inoculate was $\sim 1\mu\text{l}$. After inoculation, all groups were transferred to an
204 insectary and maintained on a 12:12h light:dark cycle at $25\text{-}26^\circ\text{C}$ and allowed to feed ad
205 libitum on cotton soaked with sugar solution.

206

207 *Kinetics of immune response, hemolymph collection and immune measurements*

208 Three groups were formed (Live-B, RPMI-G and C). Live-B group was inoculated with *E. coli*
209 in RPMI (87×10^3 CFU; non lethal dose) directly to hemocele. RPMI-G was inoculated with
210 RPMI medium only. At day 7 post priming dose, mosquitoes were re-infected. One half of the
211 individuals in RPMI-G was inoculated with RPMI, while the other RPMI-G half and Live-B
212 groups were inoculated with live *E. coli* at 99×10^3 CFU ($LD_{<5}$), a higher but minimal lethal
213 dose after priming. A minimal but lethal dose was used because mosquitoes needed to survive
214 in order to be enough individuals for daily hemolymph collection. C group was never
215 manipulated. Kinetics experiments were repeated twice. C group was included only in the
216 second repetition. Mosquitoes were inoculated as above. After inoculation, all groups were
217 transferred to an insectary and maintained as above.

218
219 Daily, after the priming dose, mosquitoes were macerated with a biovortexer in 130 μ l of PBS
220 buffer and each sample was centrifuged for 10 min at 10,000 rpm (4°C). Supernatant was used
221 to record protein load concentration, PO activity and NO production. Prior to recording PO
222 activity, protein concentration was determinate to control for differences in protein content
223 among samples (see Contreras-Garduño *et al.* 2007). The BCATM (Pierce) assay kit was used
224 to determine protein concentration for each sample. Briefly, sample supernatant, 40 μ l of PBS
225 and 150 μ l of BCATM kit reagents mix were added to a 96 microwell plate and incubated for 10
226 min. at 37°C . A known concentration of albumin (5-60 μ gr) was used as a standard reference
227 curve. The absorbance was recorded at 562nm on a plate reader. After protein load adjustment
228 and to measure PO activity, the sample supernatant plus PBS gauged at 50 μ l was mixed on a
229 96 microwell plate with 50 μ l L-DOPA (L-dihydroxyphenylalanine; 4mg/ml) as substrate and
230 incubated for 10 min at room temperature (24°C). 50 μ l of buffer mixed with 50 μ l of L-DOPA
231 was used as blank. Absorbance was recorded at 490nm on a plate reader. An increment in OD
232 after 30 minutes was defined as PO activity. Three mosquitoes were required for a single

233 sample since one individual does not provide enough material for spectrophotometer readings.
234 Three samples per group were quantified every day.

235
236 The Griess reaction was used to determine NO concentration (Eckmann et al. 2000). 50 µl of
237 each sample supernatant were mixed with 50µl of 1% sulfanilamide and 50 µl of 0.1%
238 naphthylethylenediamine on a 96 microwell plate and incubated for 10 min at room
239 temperature (24°C). NO was quantified using a NaNO₂ (1-100 µM) standard reference curve
240 for each assay. Absorbance was recorded at 540nm on a plate reader. The highest readings
241 obtained in an interval of 30 min (with measurements every 5 min.) were defined as NO
242 production (expressed as µM).

243
244 Maceration and plate filling (for PO and NO) were done in a cold room (4°C) to exclude room
245 temperature changes that could affect quantifications.

246

247 *Egg production and proportion of egg laying females*

248 To record the number of eggs produced and egg laying females, three groups were used (C,
249 RPMI-G, and Live-B). Live-B was inoculated with *E. coli* in RPMI (87x10³ CFU) directly to
250 hemocele. RPMI-G was inoculated with RPMI medium. After inoculation, all groups were
251 maintained on a 12:12h light: dark cycle at 25-26°C and allowed to feed ad libitum on cotton
252 soaked with sugar solution. At day 7 post priming dose, half of the individuals of each
253 treatment were allowed to artificially feed on male sheep blood. In order to ensure that only
254 blood-fed females were used, all mosquitoes were chilled on ice and non-fed-females were
255 removed after blood feeding. For egg laying, single females isolated in plastic glass containers
256 with humid filter paper as the egg laying substrate. Half of mosquitoes were re-infected for
257 which half of which were inoculated with RPMI, and the other half with RPMI-G and Live-B
258 groups were inoculated with live *E. coli* at 99x10³ CFU. C group was never manipulated. After

259 inoculation mosquitoes were maintained as above. At day 7 post second challenge, females of
260 each treatment were allowed to artificially feed and lay eggs as above. The number of eggs
261 produced by each female and the number of egg laying females were recorded.

262

263 *Statistical analysis*

264 We performed a survival analysis using a Log-rank χ^2 to detect differences in survival curves.
265 Analyses were performed using JMP 7.0 (SAS Institute, 2004). For the kinetics of PO and NO,
266 we performed repeated measures ANOVAs with time (time period elapsed since the priming
267 dose) and treatment (C, RPMI-G+RPMI-G, RPMI-G+ Live-B, and Live-B+Live-B) as fixed
268 factors. For kinetics, various transformations did not lead to normal distribution of data.
269 However we still used the repeated measures ANOVA since this test is still appropriate for
270 finding significance when they exists even with non-transformed data (Zar, 1999). Analyses
271 were performed using STATISTICA 7.0 (Statsoft, 2004).

272

273 A one-way ANOVA was conducted to test whether the number of eggs produced at day 7
274 (priming dose effect) was significantly different among treatments (C, RPMI-G, Live-B). To
275 achieve normal error distribution, data were transformed using a Box-Cox transformation (Fit
276 Model module of JMP Ver. 7.0). Since the absolute values of egg production at day 14 (second
277 challenge effect) were not normally distributed, a non-parametric Kruskal-Wallis test was
278 performed to compare among groups. A χ^2 test analysis was performed to detect differences in
279 the proportion of egg laying females among groups (C, RPMI+RPMI, RPMI+ Live-B, and
280 Live-B+Live-B) at day 7 (priming dose effect) and at day 14 (second challenge effect)

281

282

283 **Results**

284 *Survival*

285 Previous exposure to a minimal lethal dose of *S. marcescens* provided protection to
286 mosquitoes against a lethal challenge administered 7 days later (Log-Rank $\chi^2=13.72$, $df=3$, $P=$
287 0.0033 ; notice that removal of C group from the survival model did not change the significant
288 differences, Log-Rank $\chi^2=12.30$, $df=2$, $P=0.0021$) (Fig. 2). Thus, individuals primed with live
289 bacteria died at a slower rate than the RPMI-G+RPMI-G groups. Dead bacteria did not induce
290 the same magnitude of protection against the second lethal challenge (Fig. 2). At day 4 post
291 second challenge, survival curves between RPMI-G+Live-B and Dead-B+Live-B achieved the
292 same survival probability (Fig. 2).

293
294 In groups inoculated with *E. coli*, similar results were observed. Inoculation with a low dose of
295 bacteria affected mosquito survival. After the second inoculation with a high bacterial dose,
296 the Live-B+Live-B group died slower than the RPMI-G+Live-B group, meanwhile C and
297 RPMI-G+RPMI-G groups exhibited a reduced mortality rate (overall model Log-rank $\chi^2=$
298 18.61 , $df=3$, $P=0.0003$). When C and RPMI-G+RPMI-G groups were excluded from the
299 survival model, the differences between the curves of RPMI-G+Live-B and Live-B+Live-B
300 groups were still statistically significant (Log-rank $\chi^2=11.16$, $df=1$, $P=0.008$) until day
301 13, which no longer occurred at day 14 (Log-rank $\chi^2=1.74$, $df=1$, $P=0.18$).

302

303 *Kinetics of PO and NO*

304 For PO activity, there were differences among treatments (Table 1a) and a significant
305 interaction Time by Treatment interaction (only for the second repetition; Table 1b) which
306 indicates that the PO activity changed over time but according to treatment. PO values, in
307 both repetitions, for the Live-B+Live-B group were always below the mean PO of the other
308 groups (Fig. 3a, b). However, on some days, mostly after the second challenge, the RPMI-
309 G+Live-B group showed lower PO values (Fig. 3a: day 7 and 9; Fig. 3b: day 11 and 10).
310 Interestingly, the PO dynamics of this group appears less complex in comparison to those that

311 occurred within the RPMI-G+RPMI-G and RPMI-G+Live-B groups. PO mean peaks and falls
312 for the Live-B+Live-B group were not as intense as compared to the other experimental
313 groups in different days. Bacteria induced differences in the dynamics among groups, this
314 because the groups inoculated with bacteria showed an overall mean PO activity lower than
315 the RPMI-G+RPMI-G group.

316
317 NO production, by contrast, showed a more complex dynamics than PO activity (Fig. 4a, b).
318 There were differences among treatments (Table 2a, b) and a significant interaction Time by
319 Treatment interaction (only for first the repetition; Table 2a) which indicates that NO
320 production changed over time but differential for each treatment. During the first repetition,
321 NO values of Live-B + Live-B were always below the mean NO values of the other
322 treatments. Interestingly, in both repetitions, after the second challenge, mean NO production
323 for priming mosquitoes was above that of the other groups (at day 9 and 10 for the first
324 repetition, and markedly after day 8 for the second repetition). For this immune parameter we
325 also noticed an effect of the inoculation with live bacteria: mean NO values of the bacteria
326 challenged groups were constantly below the mean values of the group inoculated only with
327 RPMI. Intriguingly, in the second repetition, C group after day 7 (without any manipulation)
328 showed lower values compared to the manipulated groups (Fig. 4b).

329
330 *Effects on number of eggs produced and egg laying females*
331 We found differences in eggs production among groups ($F_{2,73} = 3.58$, $P = 0.032$): the Live-B
332 group fed at day seven (priming dose effect) was significantly larger in comparison to egg
333 production of females injected with RPMI-G (Fig. 5A). Egg production between Live-B and C
334 groups was similar. Females in the Live-B+Live-B group produced more eggs at day 14
335 (second challenge effect) compared with the RPMI-G+RPMI-G and RPMI-G+Live-B groups;
336 however, this difference was not significant (Fig. 5B).

337
338 We did not find differences in the proportion of females (fed at day 7; priming dose effect) that
339 laid eggs among C group (96%), RPMI (80%) and Live-B (80%) (Fig. 6A). After the second
340 challenge, the proportion of egg laying females in the Live-B+Live-B group was smaller
341 (45%) compared with the other three groups (C= 66%; RPMI-G+RPMI-G= 61%; RPMI-
342 G+Live-B=65%) (Fig. 6B), however, the overall model was not statistically significant
343 ($\chi^2=2.88$; $P=0.40$).

344

345 **Discussion**

346 Our results demonstrate that *Ae. aegypti* mosquitoes that were previously exposed to live
347 bacteria are more likely to survive a re-exposure 7 days later to the same bacteria at a higher
348 doses. This supports the idea that insects are capable of developing resistance upon a
349 secondary exposure to a pathogen (Little & Kraaijeveld, 2004; Little, Hultmark & Read, 2005;
350 Kurtz, 2005; Schmid-Hempel, 2005a, b; Moret & Schmid-Hempel, 2001; Kurtz & Franz,
351 2003). However, a previous exposure to *E. coli* does not permanently alter the mosquito's
352 response, as it does with *S. marcescens*. The total experimental time span was 23 days (3 days
353 given age of the mosquitoes + 7 days after priming dose +13 days after the second challenge),
354 which is a considerably long time taking into account the maximum lifespan of 21 days
355 reported in the wild for this animal (Clements, 1999). Despite the fact that *E. coli* does not
356 permanently change mosquito resistance, the time of protection was still long enough to
357 ameliorate the negative impact caused by a second infection. The crucial point in this case is
358 the absolute life span relative to time elapsed between exposures.

359

360 While inoculation with dead bacteria did not induce “long” protection (just a weak 3-4 days
361 potentiation), live bacteria *S. marcescens* did enhance protection. Immunization with *E. coli*
362 provided protection for 13 days. The reasons for these differences remain unclear. It is

363 possible that dead bacteria, used as an elicitor, do not release molecules that act as the
364 inducing agents. A similar phenomenon has been observed in the tsetse fly, *Glossina*
365 *morsitans* (Kaaya and Darji, 1988): inoculation with dead bacteria did not stimulate
366 antibacterial activity; however lysozyme response was weaker in comparison to the response
367 of live bacteria. The use of non-living molecules (e.g. sephadex beads, LPS vs. bacteria) could
368 be useful to prove if soluble substances from the elicitors are responsible for induction, as
369 occurs in other insects (Söderhäll 1982; Lemaitre *et al.* 1997; Moret & Siva-Jothy, 2003; Pham
370 *et al.*, 2007) (see Wiesner, 1991, for similar discussion).

371
372 After the priming dose, we found overall lower PO values for immune-stimulated mosquitoes
373 when compared to RPMI-G+RPMI-G and RPMI-G+Live-B groups. These results suggest a PO
374 deactivation rather than activation possibly because the organism is using it to control live
375 bacteria. Since bacteria used for the priming dose may cause the conversion of proPO to PO,
376 post- priming dose, it is possible that when we collect hemolymph, we are just detected the
377 remaining PO after being used to defend against bacteria. Groups RPMI-G+RPMI-G and C (in
378 second repetition) were always the groups with highest PO activity (Fig 3a, b). This result
379 indicates the natural dynamics of PO in the mosquito and the effect of the wound post
380 manipulation. Contrary to what it was expected with our model (Fig. 1), no enhancement of PO
381 activity was noticed after a second challenge. Immediately after the second challenge, a
382 decrement in the PO activity in the Live-B+Live-B group was observed, however, at day 10 a
383 slight recovery was detected. The effect of the bacteria used for the second challenge was also
384 detected. As mentioned, the dynamics of PO activity for Live-B+Live-B is not as complex as
385 those in RPMI-G+Live-B. This is probably due to previous contact with bacteria, which induced
386 long-term changes, not enhancing PO activity, but giving place to an efficient use of resources.
387 For example, tyrosine is a PO substrate, and is used for the expression and manufacture of other
388 traits. One of this is coloring the egg chorion in some insects (Li & Christensen 1993). This is

389 why in mosquitoes, melanization responses against worms generate a delay in TYR
390 accumulation in the ovaries, and therefore a decrease in the number of eggs produced after an
391 immune challenge (Li & Christensen, 1993) and a delay in oviposition (Ferdig et al. 1993).
392 Examples like this are suggestive that the substrates of PO and melanin substrates - PHE and
393 TYR - are restrictive resources that may lead to trade-offs among immune response and other
394 key functions including molting, basal metabolic rates and protein synthesis. Also, the PO
395 enzyme is used for different physiological processes (wounding, clotting, cuticle composition,
396 melanotic encapsulation, and probably spermatheca formation (Ilango 2005). Due to these
397 different functions, PO is constantly synthesized; therefore it could be costly to produce for the
398 organism. Also, it must be considered that intermediate molecules produced during the PO
399 cascade serve to amplify immune response (Cerenius & Söderhäll 2004; Nappi & Ottovianni
400 2000).

401
402 We have considered two alternative explanations for the low levels of PO. One is based on
403 resource allocation theory. PO enzyme is used for different immune responses (wounding,
404 clotting, cuticle composition, melanotic encapsulation, production of cytotoxic molecules;
405 (Christensen et al. 2005; Nappi & Christensen 2005) and therefore seems costly for the
406 organisms to produce (Schmid-Hempel, 2005b). It is possible that proPO gets activated in the
407 hemocele, consequently leaving other target tissues with a proPO deficiency. This means that a
408 trade off may arise because of a proPO shortage. Other negative effects may arise when PO
409 cascade is continuously activated; this leads to our second explanation of the low PO levels
410 observed. Several molecules (e.g. quinones, reactive oxygen molecules) are produced during
411 the PO cascade (Nappi & Christensen, 2005). These molecules can be harmful to pathogens
412 but also to host tissues and cells, a phenomenon referred to as immunopathology (Graham,
413 Allen & Read, 2005). It is possible that a controlled activation of PO could be a strategy to

414 avoid this negative effect. In this sense, low PO levels would not be the result of resource
415 limitation but a type of “on-off” switch regulation to avoid self harm.
416
417 NO production was similar to that of PO. During first repetition, after a priming dose, NO
418 levels for Live-B+Live-B were below the other groups. This reveal that the production of NO
419 after an injury induced by manipulation is not similar to the NO response to bacteria.
420 Additionally, an interesting NO production dynamics was observed after the second challenge.
421 NO production for Live-B+Live-B was higher than RPMI-G+Live-B, RPMI-B+RPMI-B and
422 C groups after the second challenge. This result was anticipated by our model (Fig. 1). In *Ae.*
423 *egypti* NO participates in the control of the dengue virus load (Ramos-Castañeda et al. 2008).
424 It is known that during malaria parasite infection, *Anopheles spp* generate NO to an extent that
425 limits parasite development (Herrera-Ortiz *et al.*, 2004; Peterson, Gow & Luckhart, 2007). In
426 *Drosophila*, NO is key for activating both the Immune Deficiency (Imd) (Foley & O’Farrel
427 2003) and upstream Imd pathways. It is also a signaling molecule, so a constant production of
428 NOS is necessary for homeostatic purposes, while inducible NOS is only synthesized after an
429 immune challenge (Nappi et al. 2000). Given the importance of NO for control pathogens, it is
430 therefore expected that production increases after an immune challenge. Despite this,
431 Krishnan, Hyršl & Šimek (2006) found that NO production was similar in non-stimulated and
432 stimulated hemocytes in lepidopteran larvae. In this particular case, the authors call attention
433 to the fact that NO synthesized by an inducible NOS, can persist for long periods of time
434 (hours to days). This means that efficacy of NO in killing pathogens may not only reside on
435 NO concentration, but on the duration of NO activity (see Laurent et al. 1996). However our
436 results show that NO production dynamics were affected differently by the presence of
437 bacteria compared to the effect of inoculation. The higher production of NO in the group Live-
438 B+Live-B after the second challenge can be explained by the previous contact with bacteria
439 inducing priming, causing enhanced NO production.

440
441 We detected differences in the enhancement levels of NO production between repetitions. NO
442 production is probably dependent on animal condition. In a previous study there was no
443 environmental effect on basal (without immune challenge) NO production and PO activity,
444 due to food quality and quantity limitation (Moreno-García *et al*, 2010). However, the
445 presence of recurrent infections could lead to different levels of immune parameters activation.
446 If there is a difference in nutrition among cohorts, possibly priming may be detected only
447 when the organisms had enough resources to mount an enhanced immune response after
448 continuous infections. Related to this, NO is produced during the oxidation of L-Arginine
449 (Müller 1997). Arginine is an amino acid that must be obtained from the diet (Rivero 2006).
450 Arginine is also important in sperm maturation (Osanai & Chen 1993), egg production
451 (Uchida 1993), long term memory, chemosensory (antennal lobes, olfaction), and visual
452 information processing (Müller 1997).
453
454
455 There is currently a gap in terms of the life history consequences of immune priming. Our data
456 demonstrate that a first immune stimulation results in an increase in egg production. It is
457 probable that when this first infection occurs, the organism is allocating resources to
458 reproduction. It is possible that the egg laying strategy depends on the force and course of
459 infection. After the first challenge, females could be allocating their resources to lay eggs, in
460 an effort to compensate for the reduced life expectancy that would diminish their chances of
461 future reproduction. Related to costs, in *Ae. aegypti* and *D. melanogaster*, up regulation of
462 immune genes reduce the life span of the organism (Libert *et al*, 2006; Kambris *et al*, 2009;
463 McMeniman *et al*, 2009), so, investment in early reproduction could be adaptive. After the
464 second challenge there was no difference in egg production among individuals in the different
465 treatments, however, survival and NO production for Live-B+Live-B was higher than the other

466 groups. It can be argued that survival (and related immune traits) and fecundity are being
467 maximized simultaneously. On the other hand, our results also indicate that the proportion of
468 egg laying females in the Live-B+Live-B group was lower than the other groups, indicating
469 that not all females are disposed to produce eggs, increment survival and enhance immune
470 response. It is possible that the egg laying strategy depends on the force and course of
471 infection, and that the organisms that recover from infection and reduce immune activation
472 costs use its resources in relation to chances of future reproduction which is strongly related to
473 life expectancy.

474
475 In summary, we have provided evidence that PO activity and NO production do not show an
476 overall increased level. However, NO production may be enhanced when the host has had a
477 previous contact with live bacteria. Combined with survival results, the immune levels found
478 here can be explained as the efficacy of organisms in fighting bacteria. Depending on the
479 immune markers measured, immune priming can be considered not only an enhanced
480 response, but an optimal response (in terms of survival and reproduction) to protect the
481 organisms after a previous exposure to an elicitor. Reactions and products of immune response
482 are interconnected and the kind of response is related to the pathogens virulence. It is possible
483 that not all pathogens will induce the same reaction, or at least the magnitude of response can
484 differ depending on the host-pathogen interaction type. Therefore, it is recommended to use
485 more than one immune marker when possible. The mechanisms of immune response in insects
486 include reactions and molecules that are interconnected. It is also important to ascertain the
487 contribution of other immune parameters such as hemocyte activation or antimicrobial
488 synthesis whose activities are enhanced after a second pathogen exposure and have an
489 integrated system of dynamics within immune response. Finally, in terms of reproduction
490 effects, we did not detect an obvious negative result on egg production and number of egg
491 laying females. Reasons for this result granted further studies.

492

493

494 **Acknowledgements**

495 We thank L. González-Cerón and J. Silva for kindly donating bacterial strains. We also thank
496 the Malaria-Dengue Group at the INSP for useful comments. We thank Gloria Tavera for
497 revising this paper in English. MM-G thanks the Posgrado en Ciencias Biomédicas
498 (CONACYT grant No. 172947), Instituto de Ecología, Universidad Nacional Autónoma de
499 México, and Centro de Investigaciones Sobre Enfermedades Infecciosas, INSP, México. AC-
500 A was supported by a PAPIIT-UNAM grant (Project No. 211506). HL-M thanks the Bill and
501 Miranda Gates Foundation for the Grand Challenges Exploration Grant (Grant ID 51996).

502

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- 669

670 Table 1. Results of repeated measures ANOVA of immune kinetics for (a) NO production and
 671 (b) PO activity (first repetition).

<i>NO production</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P</i>
Treatment	363.3	2	181.6	122.7	0.008
Error	2.9	2	1.4		
Time	1846	10	184.6	5.4	0.000
Time*Treatment	1597.4	20	79.8	2.3	0.032
Error	683.2	20	34.1		

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<i>PO activity</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P</i>
Treatment	0.205	2	0.102	44.6	0.021
Error	0.004	2	0.002		
Time	0.309	10	0.030	3.5	0.007
Time*Treatment	0.225	20	0.011	1.2	0.289
Error	0.175	20	0.008		

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689 Table 2. Results of repeated measures ANOVA of immune kinetics for (a) NO production and
 690 (b) PO activity (second repetition).

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<i>NO production</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P</i>
Treatment	265.2	3	88.4	4.43	0.047
Error	139.5	7	19.9		
Time	1811.2	11	164.6	5.48	0.000
Time*Treatment	1251.7	33	37.9	1.26	0.200
Error	2312.1	77	30		

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<i>PO activity</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P</i>
Treatment	0.573	3	0.191	50.79	0.000
Error	0.022	6	0.003		
Time	0.085	11	0.007	2.17	0.026
Time*Treatment	0.256	33	0.007	2.18	0.003
Error	0.234	66	0.003		

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708 **Figure 1.** A hypothetical model of the temporal dynamics of the immune response after a
709 priming dose and a second challenge. Dashed line= Prolonged activation.

710 **Figure 2.** Survival curves of mosquitoes after a second exposure with A) *S. marcescens*, and
711 B) *E. coli*.

712 **Figure 3.** In vivo dynamics of (a) PO activity and (b) NO production (first repetition).

713 **Figure 4.** In vivo dynamics of (a) PO activity and (b) NO production (second repetition).

714 **Figure 5.** Egg production in females fed at day 7 after priming dose (a), and in females fed at
715 day 14, after second challenge (b).

716 **Figure 6.** Proportion of egg laying in females fed at day 7 after priming dose (a), and in
717 females fed at day 14, after second challenge (b).

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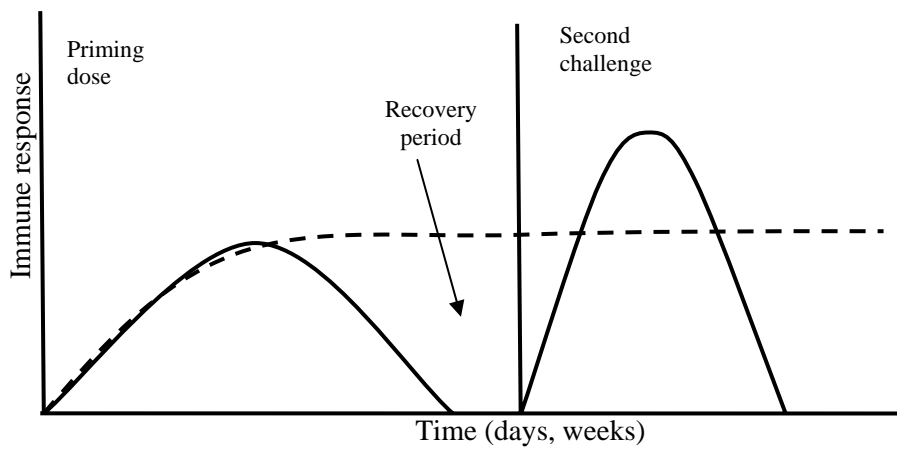
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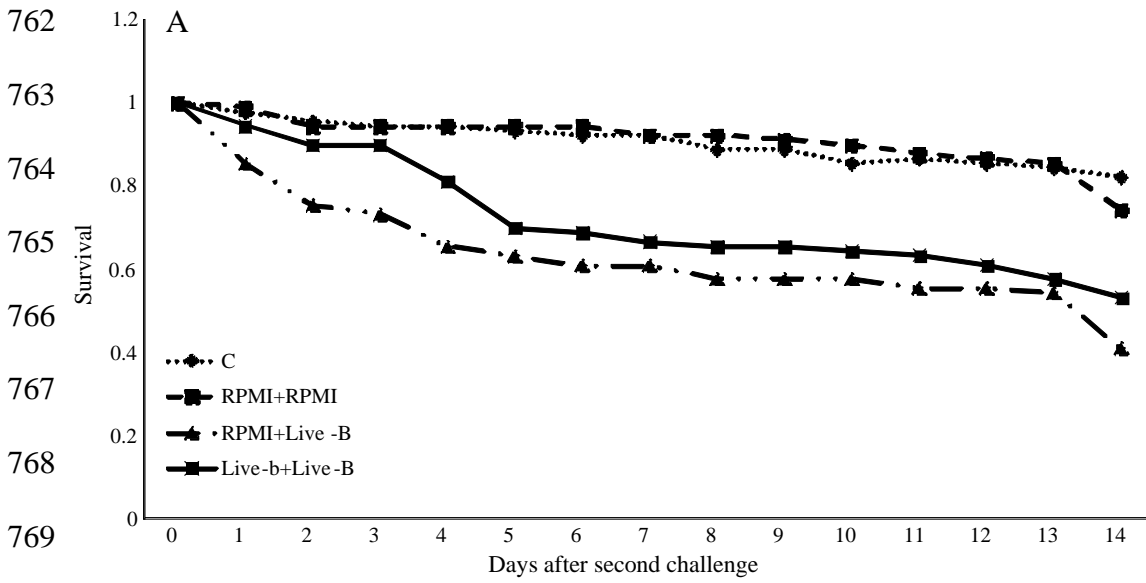
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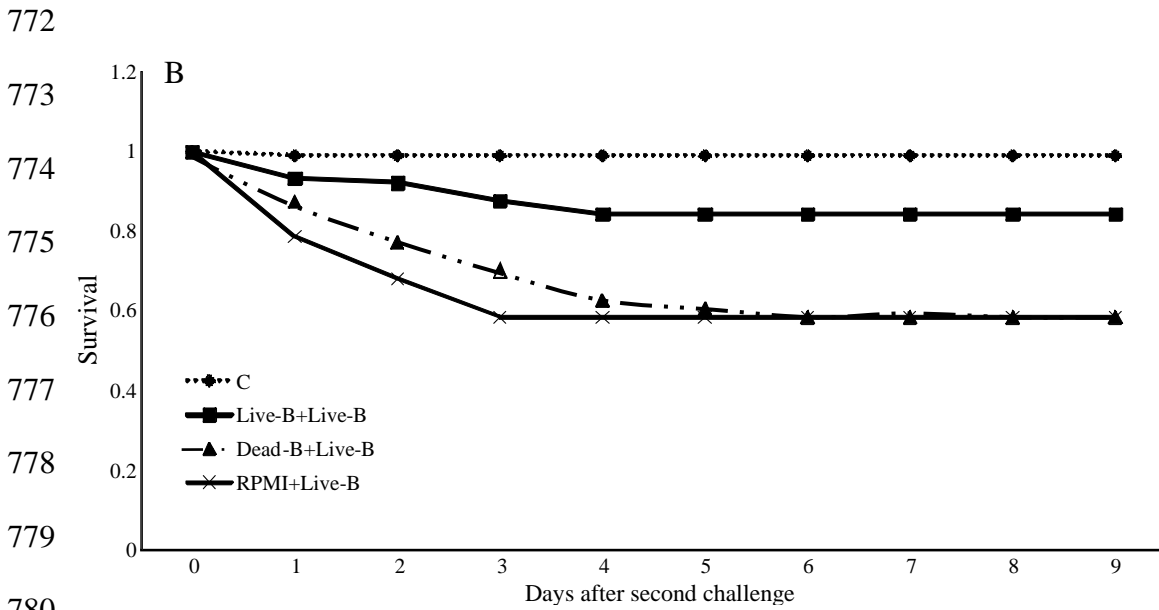
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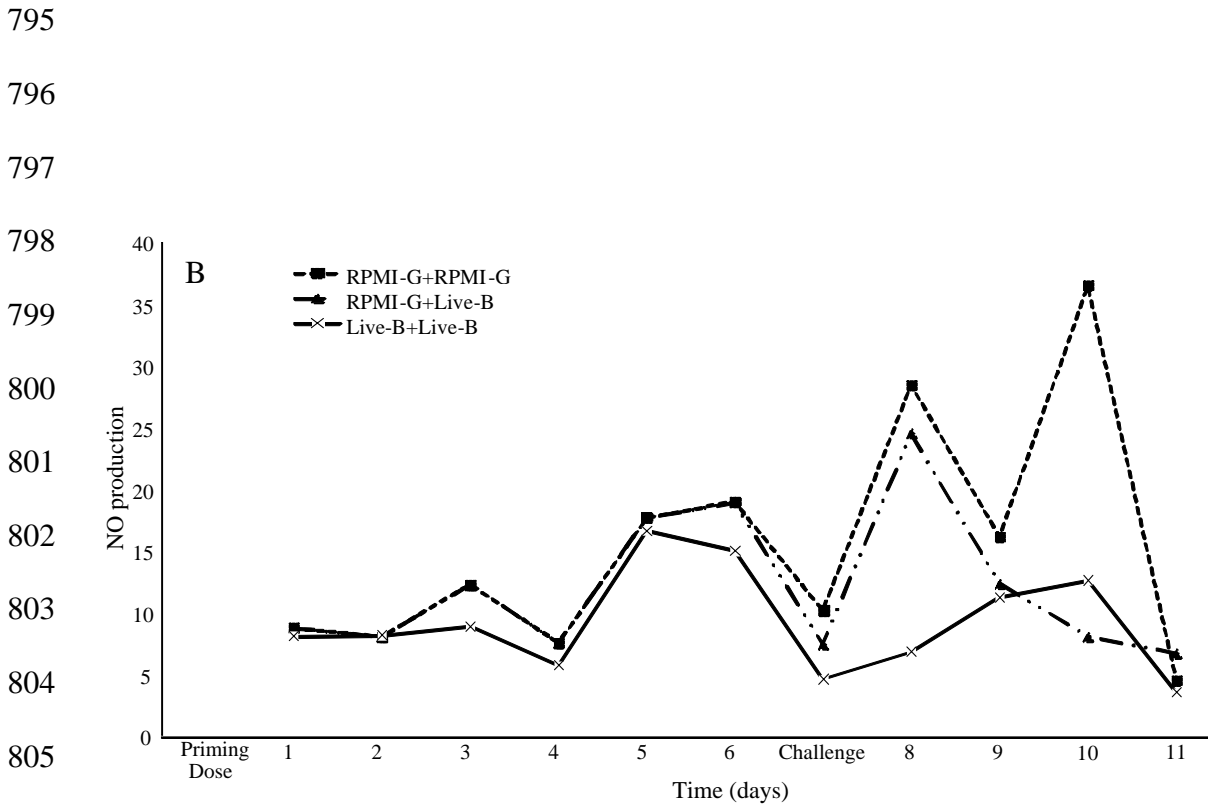
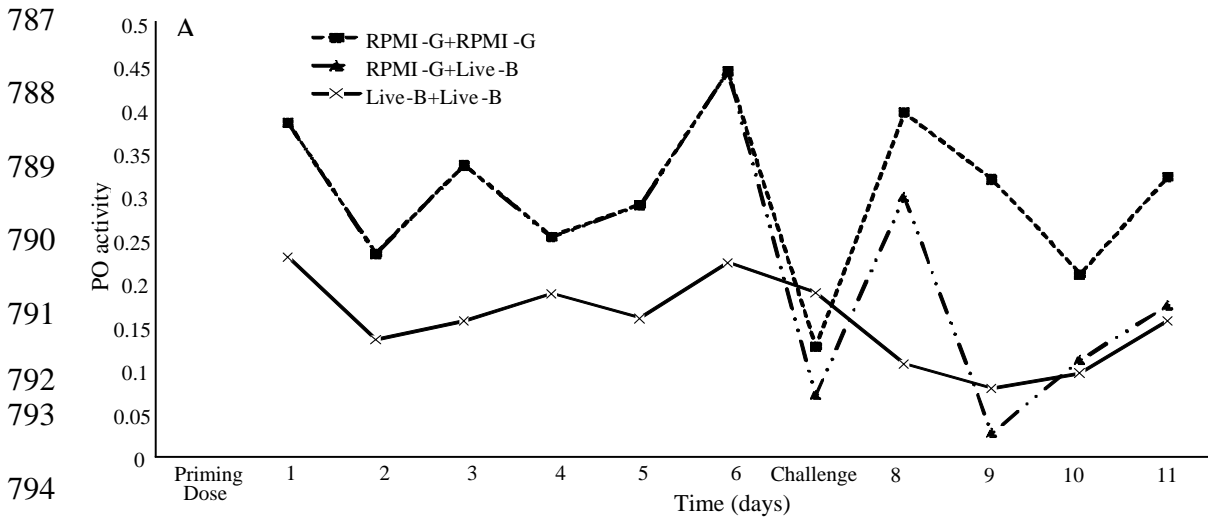
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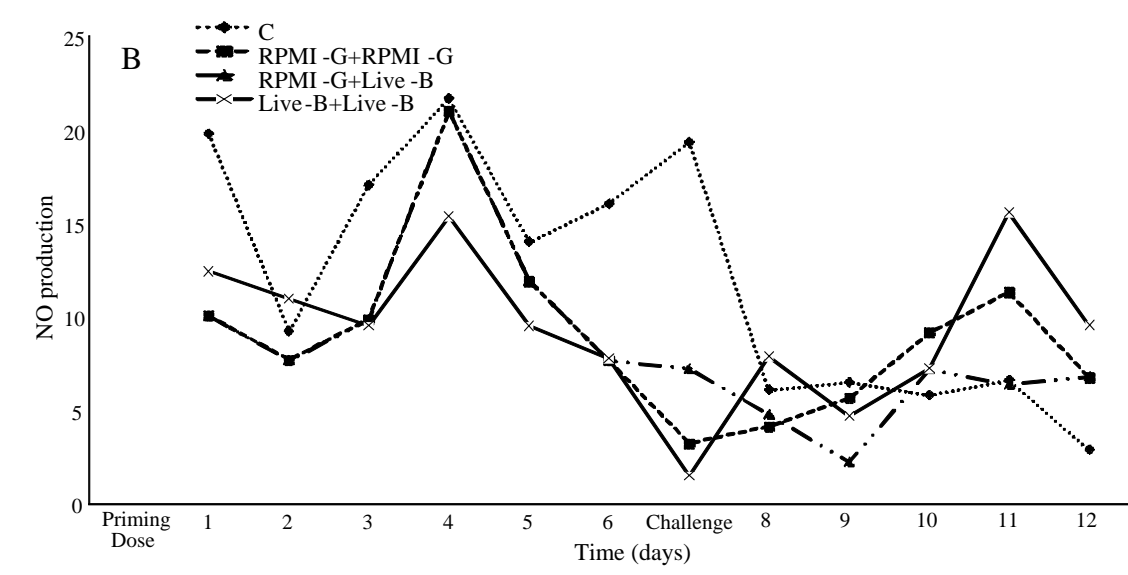
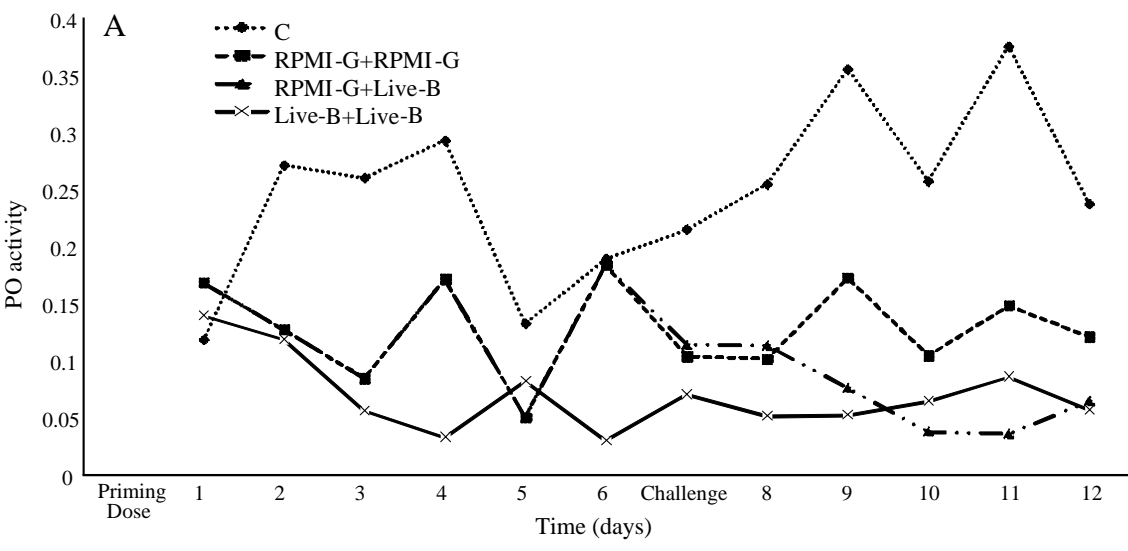
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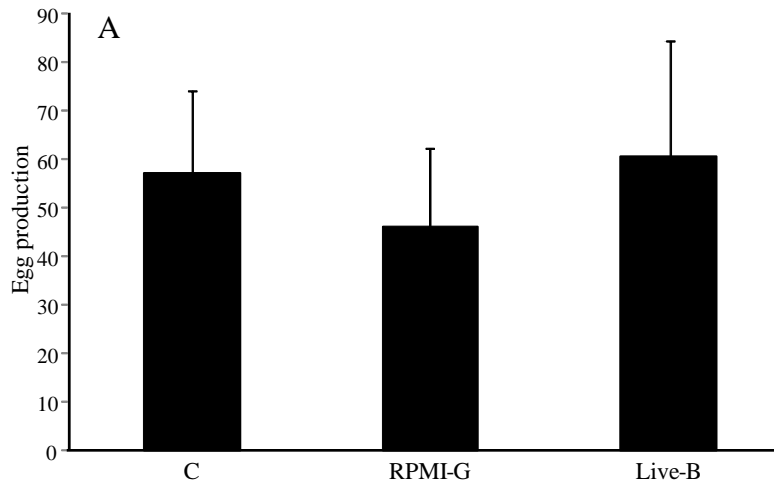
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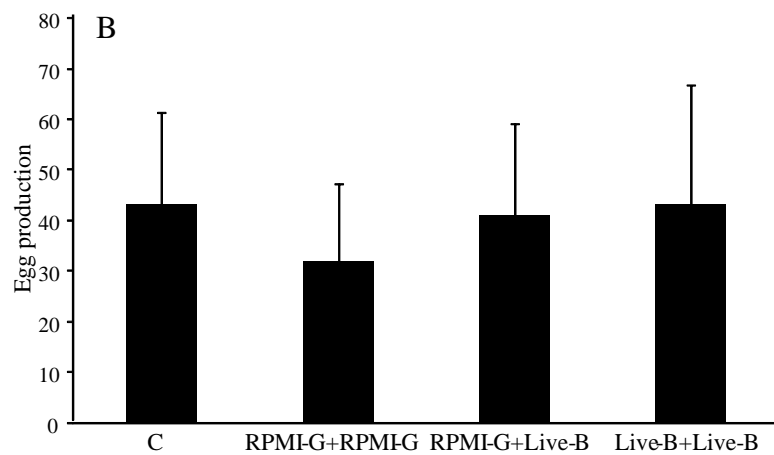
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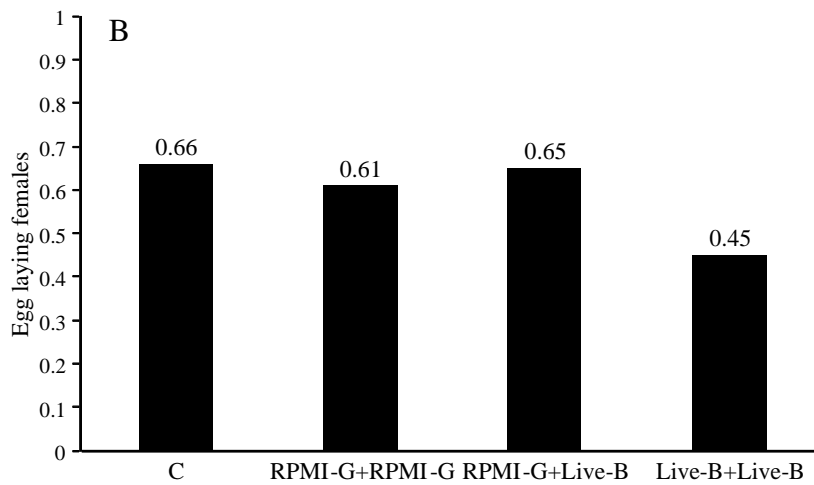
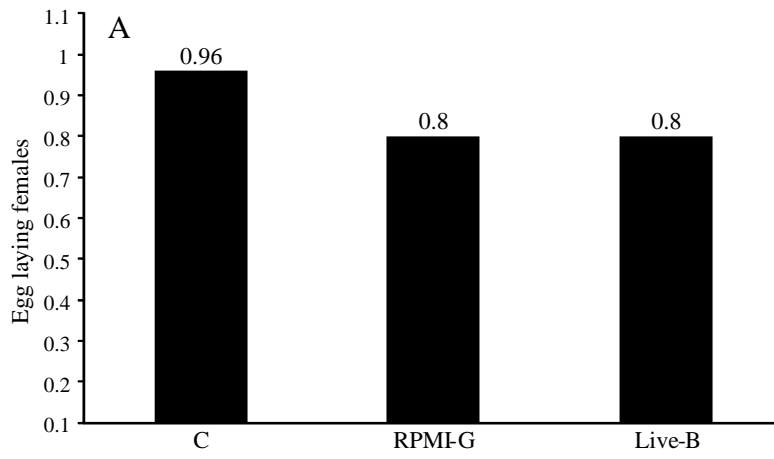
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Genetic Variance and Genotype-by-Environment Interaction of Immune Response in *Aedes aegypti* (Diptera: Culicidae)

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J. Med. Entomol. 47(2): 111–120 (2010); DOI: 10.1603/ME08267

ABSTRACT Immune response can be negatively affected by resource limitation, so it is expected that organisms evolve strategies to minimize the impact of this environmental outcome. Phenotypic plasticity in immune response could represent a genetic response to face such situations. We investigated the effects of high and low quality and quantity of food at the larval stage on two important immune components, phenoloxidase activity (PO) and nitric oxide production (NO) measured in adults of the Dengue vector, *Aedes aegypti*. We reared families to determine the magnitude and pattern of expression of genetic variance, environmental variance and genotype-by-environment interaction (GEI). In addition, we quantified whether there were differences in plastic immune responses in both sexes. Our results indicated additive variance for PO and NO, but rearing environment did not produce differences among individuals. For NO and PO in males, there were large differences among families in plasticity, as indicated by the different slopes produced by each reaction norm. Therefore, there is additive genetic variation in plasticity for NO production and PO activity. One possible interpretation of these results is that different genotypes may be favored to fight pathogens under the different food quality situations. Males and females showed similar overall GEI strategies but there were differences in PO and NO. Males showed a phenotypic correlation between PO and NO, but we did not find genetic correlations between immune parameters in both sexes.

KEY WORDS immune response, quantitative genetics, phenotypic plasticity, *Aedes aegypti*

Immune response is a trait closely linked to survival and reproduction (Schmid-Hempel 2003, Schulenberg et al. 2009). Despite the fact that investment to immunity is adaptive, immune response is strongly impacted by environmental conditions. This occurs in situations of environmental heterogeneity (such as variation in food abundance), in which an overall fitness decrement for a given genotype is observed (e.g., Leclaire and Brandl 1994, Fellowes et al. 1998, Metcalf and Monaghan 2001, Siva-Jothy and Thompson 2002). In mosquitoes, for example, larvae that have been reared in crowded or undernourished conditions, give rise to adults with a weak cellular encapsulation (Suwanchaichinda and Paskewitz 1998). On the same line of research, it has been also found that when mosquito adults are stressed with food shortage, encapsulation immune response decreases (Chun et al. 1995, Schwartz and Koella 2002). Therefore, the organisms are expected to deal with the problem of

maximizing fitness in changing or stressful environments.

A genetic background that expresses changes in phenotype expression could have a selective advantage in a given environment (Zhivotovsky et al. 1996). In some cases, a single genotype can have the ability to produce distinct phenotypes when exposed to different environments, a phenomenon known as phenotypic plasticity (Roff 1997, Nylin and Gotthard 1998, Schlichting and Pigliucci, 1998). Phenotypic plasticity is shown by many traits and immune response is not an exception (Fordyce 2006). In fact, such immune plasticity could be used as an indicator of the strategies that an organism has followed when dealing with environmental heterogeneity. Paradoxically, given the recent explosion in ecological and evolutionary studies of immunity, only a handful of studies have approached immune response using a phenotypic plasticity perspective (but see Barnes and Siva-Jothy 2000, Mucklow and Ebert 2003, Cotter et al. 2004a, Lazzaro et al. 2008, McKean et al. 2008). To have a better understanding of the evolution of immune responses, more studies of phenotypic plasticity are therefore needed.

To afford the problem of whether plastic immune responses could have an impact in population evolu-

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tion, it is necessary to evaluate if different environments produce different phenotypes and if genotypes respond differently to these environments (genotype-by-environment interaction GEI, which represents genetic variation for phenotypic plasticity). It has been proposed that GEI can allow populations to evolve to an optimum phenotypic mean in different environments, promoting adaptation to heterogeneous environments (Via and Lande 1985). As a result, the amount of genetic variance, as assessed by the slope of reaction norms (i.e., a graphical description of the GEI), may be a strong determinant of the population fitness (Fry 1996). Notwithstanding, a lack of variation may lead to a failure to respond adaptively to environmental changes. There are theoretical advantages of using this approach. For example, phenotypic plasticity could explain the maintenance of genetic variance recently found in different immune components (e.g., see Ryder and Siva-Jothy 2001, Simmons and Roberts 2005, Cotter et al. 2004b, Rolff et al. 2005, Schwarzenbach et al. 2005, Fellowes et al. 1998). The fact that plasticity uncouples the phenotype from genotype and thus releases the gene pool from the immediate impact of natural or sexual selection (Stearns 1992), will lead to slow depletion of genetic variance. However, some reaction norms will not produce the optimal mean trait value in a given environment (Schlichting and Pigliucci 1998, Roff 1997), slowing down the rate of adaptation for immediate generations.

The conventional approach to see how a trait responds to selective pressures is to analyze the trait in question using quantitative genetics by partitioning and estimating variances and covariances into causal components (Falconer and Mackay 1996). In this study we have explored the environmental heterogeneity affecting immune components in the mosquito *Aedes aegypti* Linnaeus (Diptera: Culicidae), the principal vector of Dengue and Yellow Fever virus. We varied levels of food quality and quantity in the larval stage, and as response variables two immune markers were measured in the adult: the basal levels of phenoloxidase (PO) activity and nitric oxide (NO) production. PO is an oxidoreductase enzyme used in insect cellular and humoral response such as cuticle melanization, wound repair, cytotoxin production, and melanotic encapsulation (Söderhäll and Cerenius 1998). NO is a highly reactive and unstable free radical gas that inhibits protein catalytic activity and produces protein and harming effects on pathogens' DNA (Rivero 2006). Recent studies have shown that PO activity is costly to produce (reviewed by Kanost and Gorman 2008). Although there is no direct evidence for such costs for NO, their physiological pathways strongly indicate that it is potentially costly (Rivero 2006, Carton et al. 2008). We evaluated genetic variation, phenotypic plasticity and tested for GEIs of PO and NO using a split family design. We predicted the existence of genetic differences and environmentally sensitive production of alternative phenotypes by given genotypes on these immune markers and a negative effect because of limitation in food quality: mosquito adults

whose larval stage had poor quality food would show decreased basal PO activity and NO production while the opposite would be found for mosquito adults whose larval stage had access to high quality food.

Reproductive strategies could promote differences in the pattern of investment to immune defense by each sex. Males are expected to increase their fitness by reducing their investment to immune defense while investing resources to reproductive effort (Zuk and McKean 1996, Sadd et al. 2006). Meanwhile, in females natural selection would favor an increase in resource investment to immunity, under the assumption that increased longevity could enhance fitness via egg production (Rolff 2002). How this presumable sexual dimorphism translates into plastic response differences in both sexes has been little explored. McKean and Nunney (2005) found sex specific plastic responses in relation to the availability of limiting resources. Given this background information, we thus evaluated if males and females express different strategies in their plastic responses in PO activity, NO production, and GEIs strategies.

Materials and Methods

Mosquitoes were obtained from an insectary at the Instituto Nacional de Salud Pública, Cuernavaca, Mexico. The stock population had been held in this place for at least 130 generations. The colony was maintained with a protocol that minimizes inbreeding: fairly high number of individuals per generation (over 2,000 individuals per generation) and random mating. During the study, the colony was kept on a 12L:12D cycle at 25–28°C.

Experimental Design. For PO activity and NO production, we had 44 families by randomly choosing 44 mated and blood fed females. We then split their F1 hatched larvae into one of two rearing environments that differed in food quality and quantity: the high quality food (HQF) environment was based on rat chow, yeast extract and lactoalbumin hydrolysate (1:1:1 mix; 25 g/200 ml), while the low quality food (LQF) environment was based on rat chow only (25 g/200 ml). Preliminary observations have shown these dietary requirements are effective enough to change the individual condition, as reflected by adult size (S. Hernandez-Martinez personal communication), lifespan, and egg clutch and size (see Nasci 1986, Packer and Corbet 1989). Larvae were fed according to the schedule shown in Table 1. Each family of larvae was reared in plastic glasses containing 100 ml of water. Adult females and males of 3 d postemergence from these larvae were used to obtain PO and NO readings.

To minimize degradation of molecules, individuals were collected and frozen at -70°C until they were processed. Given the final large number of individuals and with the aim to avoid sample degradation, maceration, supernatant collection and readings were only done for the exact number of tests supported in the microwell plate (80 wells for test samples plus 16 for standard reference curves, see below) that can be processed and quantified in a single day. A group of

Table 1. Rearing schedule according to food regimes

Day posteclosion	HQF	LQF
1	30 μ l	15 μ l
2	0 μ l	0 μ l
3	30 μ l	15 μ l
4	70 μ l	35 μ l
5	110 μ l	55 μ l
6	50 μ l	25 μ l
7	50 μ l	25 μ l
8	50 μ l	25 μ l
9	50 μ l	25 μ l

HQF, high quality food; LQF, low quality food.

three mosquitoes was required for a single sample because one individual cannot provide enough sample for spectrophotometer readings (MM-G, unpublished data).

All organisms were macerated with a biovortexer in 120 μ l of PBS buffer (4°C) and each sample was centrifuged for 10 min at 10,000 rpm (4°C). Supernatant was used to record protein load concentration and PO activity. Before recording PO activity, we determined protein load concentration to control for individual differences in protein content among samples that may bias PO readings (see Contreras-Garduño et al. 2007). The BCA (Pierce) assay kit was used to determine protein concentration for each sample. There were 10 μ l of sample supernatant, 40 μ l of PBS, and 150 μ l of BCA kit reagents mix added to a 96 microwell plate and incubated for 10 min at 37°C. A known concentration of albumin (5–60 μ g) was used as a standard reference curve. The absorbance was recorded at 562 nm in a plate reader. For PO activity, the sample supernatant plus PBS gauged at 50 μ l (after protein load adjustment) was mixed on a 96 microwell plate with 50 μ l L-DOPA (L-dihydroxyphenylalanine; 4 mg/ml) as substrate and incubated for 10 min at room temperature, 50 μ l of buffer mixed with 50 μ l of L-DOPA was used as blank. The absorbance was recorded at 490 nm in a plate reader. An increment in OD after 30 min was defined as PO activity. PO readings of each microwell plate were obtained as proportional values (highest reading = 100%; lowest reading = 0%).

The Griess reaction was used to determine NO concentration (Eckmann et al. 2000). There were 50 μ l of each sample supernatant mixed with 50 μ l of 1% sulfanilamide and 50 μ l of 0.1% naphthylethylenediamine on a 96 microwell plate and incubated for 10 min at room temperature. NO was quantified using a NaNO₂ (1–100 μ M) standard reference curve for each assay. Absorbance was recorded at 540 nm in a plate reader. The highest readings obtained in an interval of 30 min (with readings every 5 min) were defined as NO production (expresses as micrometers). NO readings of each microwell plate were obtained as proportional values (highest reading = 100%; lowest reading = 0%).

Plasticity and Genotype-by-Environment Interaction Data Analysis. We first investigated whether there were differences in wing length according to

food treatments. For this, a *t*-test was used to compare wing length between adult females raised when larvae in LQF and HQF. We investigated the effects of family, food quality and quantity environment (HQF-LQF) and the interaction (GEI) of PO activity and NO production using a mixed-model ANOVA (type III SS) for unbalanced data using the Variance Components module of STATISTICA 7.0 (StatSoft 2004). Family was entered as a random effect while environment was entered as a fixed effect. This method corresponds to the Scheffé model of Fry's (1992), in which the effect of family is tested using the formula $F = MS_{family} / MS_{error}$. The MS of the interaction was used as denominator for the fixed effect. To estimate GEIs we first tested for family by rearing environment interactions using the mixed model ANOVA. If a significant interaction was detected, we then evaluated GEI by calculating the cross-environment genetic correlation (r_g ; see Fry 1992). r_g is defined as the correlation between the mean of a trait of a genetic group in one environment and the group's mean in another environment. Thus, r_g assumes that a single trait expressed in different environments represents two separate traits (Falconer and Mackay 1996). We used the SAS model of Fry (1992) where $r_g = Cov(M_{1j}, M_{2j}) / \sqrt{[Var(M_{1j}) \times Var(M_{2j})]}$, where M_{1j} and M_{2j} are the mean trait values of genetic group *j* under environmental conditions 1(HQF) and 2(LQF) where $Cov(M_{1j}, M_{2j})$ is the covariance of the mean trait values between conditions HQF and LQF, and where $Var(M_{1j})$ and $Var(M_{2j})$ are the variances of the mean trait values under conditions HQF and LQF. Cross-over interactions in reaction norms are more feasible when r_g values are $\ll 1$ (Fry 1992). We used families with at least one sample in both rearing environments. ANOVA were performed on untransformed data when non-normal distribution was present. Various transformations did not lead to a normal distribution of data. However, we still performed the ANOVA using the nontransformed as this test is still appropriate as it gives a better chance of finding significances when they exist (Zar 1999). Also, it was necessary to perform the parametric ANOVA to obtain the MS given by STATISTICA 7.0 Variance Component module.

Sexual Dimorphism in PO Activity and NO Production. For PO a *t*-test was used to compare the average reaction norm of each sex, using the GEI resultant least-square means of each family reared at HQF and LQF environment. For NO, data has non-normal distribution, so a Mann-Whitney *U* test was used to compare the average reaction norm of each sex, using the GEI resultant least-square means of each family reared at HQF and LQF environment. We also evaluated if rearing environment induced different plastic responses for PO activity and NO production between the sexes using a Mann-Whitney *U* test for which we used nontransformed data.

Genetic and Phenotypic Correlations. We examined phenotypic and genetic correlations separately for each rearing environment and sex. We estimated genetic correlations between PO activity and NO production by calculating Spearman correlation coeffi-

Table 2. Mixed model ANOVA testing for the effect of family, rearing environment, and GEI on PO activity in adult female and male mosquitoes

Adult mosquitos	df	SS	MS	F	P
A. Female					
Family	43	2.8752	0.0668	2.0651	<0.0005
Rearing environment (HQF-LQF)	1	0.0093	0.0093	0.2405	>0.25
Family × environment	43	1.666	0.0387	1.1970	0.19
Error	337	10.911	0.0323		
B. Male					
Family	43	3.704	0.0861	3.3089	<0.0005
Rearing environment (HQF-LQF)	1	0.065	0.0657	1.3674	>0.25
Family × environment	43	2.066	0.0480	1.8459	0.0014
Error	396	10.309	0.0260		

HQF, high quality food; LQF, low quality food.

cient for mean trait values for each family. We estimated phenotypic correlations between traits with Spearman correlation coefficient for individual trait values for which we used nontransformed data.

Results

Plasticity and Genotype-by-Environment Interaction. No differences in wing length were observed between diet regimes ($P > 0.05$). The ANOVA revealed a strong family effect on female (Table 2; Fig. 1) and male (Table 3; Fig. 2A) PO activity, which means that there was genetic variation (Family source) for this immune response. There was no significant effect for rearing environment for males and females and no interaction between the effects of family and rearing environment in females. However, our analysis revealed strong effects of family-by-rearing environment on PO activity in males (Table 3; Fig. 2B).

NO production in females had a significant family effect (Table 3A; Fig. 3A). Females reared in the LQF regime produced more NO than females reared in the HQF regime (Fig. 3B) although such differences were not statistically significant (Table 3A.). For males, rearing environment did not have a significant effect, family revealed a strong effect, and the family-by-rearing environment was marginally nonsignificant

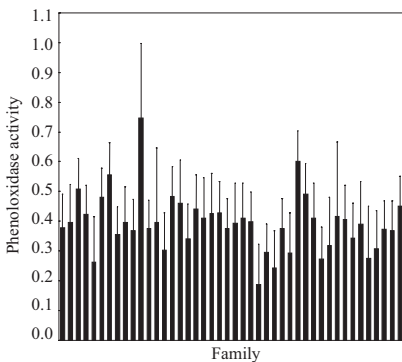


Fig. 1. Genetic differences (means ± SE) of 44 half-sib families for PO activity in adult female mosquitoes.

Table 3. Mixed model ANOVA testing for the effect of family, rearing environment, and GEI on NO production activity in adult female and male mosquitoes

Adult mosquitos	df	SS	MS	F	P
A. Female					
Family	42	2.4284	0.0578	1.6114	<0.025
Rearing environment (HQF-LQF)	1	0.0098	0.0098	0.2751	>0.25
Family × environment	42	1.5019	0.0357	0.9966	0.48
Error	337	12.0920	0.0358		
B. Male					
Family	43	1.0886	0.0253	1.6180	<0.025
Rearing environment (HQF-LQF)	1	0.0024	0.0024	0.1120	>0.25
Family × environment	43	0.9446	0.0219	1.4039	0.052
Error	396	6.1963	0.0156		

HQF, high quality food; LQF, low quality food.

(Table 3B; Fig. 4A and B). It is possible that genetic variation exists in NO production and genotypes differ in the level or direction of plasticity in this immune component.

The family-by-rearing environment interaction detected for PO activity and NO production in males suggested GEIs. We thus calculated r_g from the variance components of the mixed-model ANOVA (Table 2B; 3B): for PO there was a $r_g = 0.641$; for NO there was a $r_g = 0.535$. These values are consistent with the crossover interactions in reaction norms. Visual inspection of Fig. 2B revealed 19 families (out of 44) that showed higher PO values for individuals reared at LQF compared with his brothers reared at HQF. For NO (Fig. 4B), 16 families (out of 43) showed higher NO production when individuals were reared at LQF contrasting with their brothers reared at HQF. These cross-environmental genetic correlations showed that the PO production and NO activity can be viewed as a different trait whose production depends on the environment.

Sexual Dimorphism in PO Activity and NO Production. PO GEI least-square average means (the average slopes for the reaction norms of each family) produced a sexual difference ($t = 8.862$; $df = 1, 172$; $P < 0.0001$; Fig. 5A): males showed higher average reaction norm than females. Meanwhile, NO average reaction norm was higher in females ($U = 327$; $P < 0.0001$; Fig. 5B). Concordant with the average reaction norms, males of both rearing environments had higher PO activity than females ($U = 55146$; $P < 0.0001$; Fig. 6A). Females, compared with males, showed higher NO production in both rearing environments ($U = 29392$; $P < 0.0001$; Fig. 6B).

Genetic and Phenotypic Correlations. We only detected a phenotypic correlation between PO and NO in males in both environments (Table 4). This result indicates that variance in PO is related to variance of NO, variation in this immune parameters is not independent. We did not detect any correlation in females.

Discussion

We explored whether there was variance in PO activity and NO production that can be ascribed to

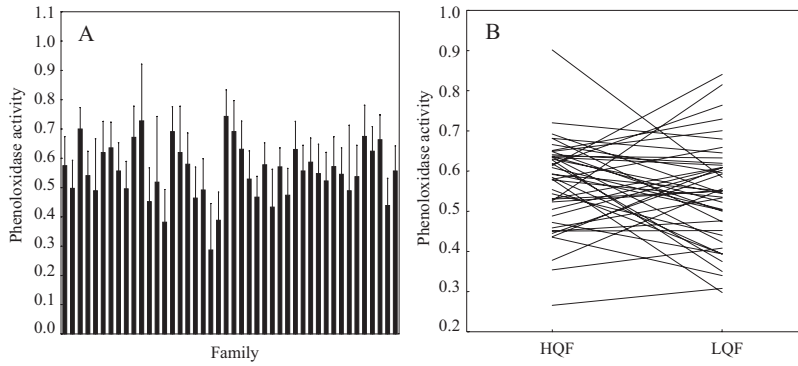


Fig. 2. (A) Genetic differences (means \pm SE), and (B) reaction norms of PO according to different food treatments for males of 44 half-sib families. HQF, high quality food; LQF, low quality food.

genetic differences between individuals, as well as the presence of additive genetic variation for plasticity (GEI). PO activity and NO production show genetic variation for both sexes, this fact has been already documented in other insects (e.g., Cotter et al. 2003b; Schwarzenbach and Ward 2006). In these cases, genetic variation has been interpreted as an adaptive response to face pathogen attack: given the large number of infectious agents, there will be ground for high genetic variability in immune response. In the case of our study subject, such variation could allow the male and female mosquitoes to fight against all varieties of infections produced by different pathogens (e.g., the coexistence of different dengue serotypes; Thavara et al. 2006).

Interestingly, no environmental variance was detected for both overall PO activity and NO production despite contrary evidence suggesting that immunity is affected by, for example, dietary restrictions (e.g., Siva-Jothy and Thompson 2002) among other environmental factors (reviewed by Schmid-Hempel 2003). It may be that the specific dietary components that we varied may not be directly involved in PO activity. One component ascribed to be key as PO substrate is tyrosine, which is derived from phenylalanine hydroxylation (Christensen et al. 2005). Phenylalanine (thus tyrosine) is gathered through larval

development in many insects (Kramer and Hopkins 1987). Furthermore, melanin, the final product of the phenoloxidase cascade, is a nitrogen-rich compound. Nitrogen or protein investment is expected to be required for its production (Blois 1978). Poor diet could reduce these compounds destined for melanin production, consequently the production of PO could be synthesized in low quantities. Meanwhile, NO is produced during the oxidation of L-arginine to L-citrulline (Müller 1997); arginine is an essential amino acid which must be obtained from diet (Rivero 2002) mainly through ontogeny. Therefore, suboptimal food quality conditions for larvae could result in a high genetic variability in adult immunocompetence. A situation like this can be found in the wild. It was expected that the beneficial effect of supplementing lactoalbumin and yeast extract (as amino acids donors; see Davis, 1975) on larvae diet may render differences in PO activity and NO production between HQF and LQF. However, rearing effects were not detected in the Mixed ANOVA. It may be that the concentration of proteins mixture in the LQF in this study may not be suboptimal for these immune parameters. However, such explanation does not seem to be entirely correct as the presence of phenotypic correlations in males indicate a degree to which PO

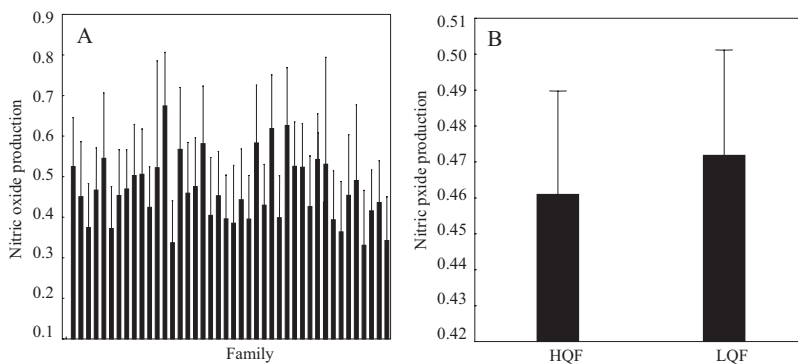


Fig. 3. (A) Genetic differences (means \pm SE), and (B) rearing environment effects in NO for adult female mosquitoes (43 families). HQF, high quality food; LQF, low quality food.

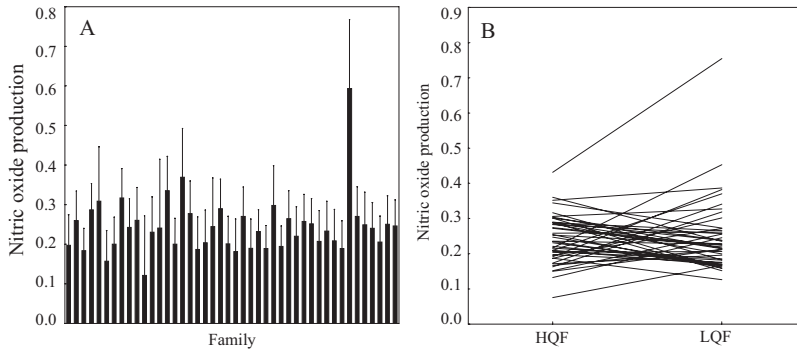


Fig. 4. (A) Genetic differences (means \pm SE), and (B) rearing environment effects in NO for adult male mosquitoes (44 families). HQF, high quality food; LQF, low quality food.

and NO respond to variation in the same environmental factors (HQF or LQF).

GEI was not present in females. However, in males there exists some genetic variation in the sensitivity to environmental effects among different mosquito families. The presumed sex-biased genetic-by-environment difference in immunity could be a consequence of the difference in pathogens, quantity of resource acquired during ontogeny, and differences in survival and reproductive strategies. It is possible that females are under the selective pressures of a large variety of pathogens for which, consequently, genetic variance should be expected. However, a female strategy could be to buffer the variation caused by stressing environmental factors (i.e., limited resources), ensuring phenotypic expression within individuals given a specific genotype and environment. The absence of GEI in females for NO and PO could be the result of the ability of females to withstand environmental perturbations and, in combination with the presence of genetic variance, to respond adaptively to changes in the environment and pathogen challenges as a result of stabilizing selection. However, the intrinsic physiological differences between males and females (e.g., fat body reserves, life span and reproductive effort) might also preclude the GEI in females (see below for discussion about sexual dimorphism).

For males, the cross-environmental genetic correlations (r_g) for both immune parameters suggest differences in genetic architecture across the environmental food conditions. It is possible that differences in the amount of reserves accumulated during the larval life affect the immune response of adult mosquitoes that would depend on their genotype. It has been suggested that the presence of GEI may allow adjustment of development that maximizes fitness in a particular environment (Stearns 1992, Fry 1996). For example, under poor food conditions, it may be advantageous to cease growth and allocate the available energy to immunity. However a similar phenotypic response may also occur as consequence of stress, lacking any fitness benefits (Kearsey and Pooni 1996). Although we cannot address the possible adaptive significance of the observed GEI in this study, the genotypes differing in their reaction norms have the potential to ensure immunocompetence during periods of food quality stress and influence the course and rate of evolution in heterogeneous environments (e.g., Gillespie and Turelli 1989). Nevertheless, to see how realistic this situation is in periods of food stress and test that our results are adaptive, further experimental approaches need to be done (for a similar claim see Siva-Jothy et al. 2005).

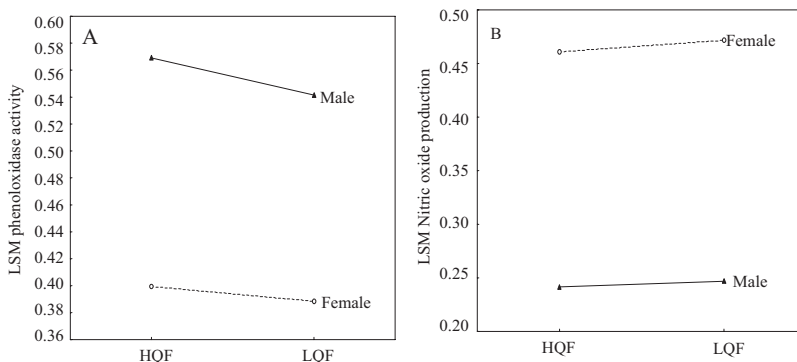


Fig. 5. Means of slopes of each sex's reaction norm for (A) PO activity and (B) NO production based on least-square means. HQF, high quality food; LQF, low quality food.

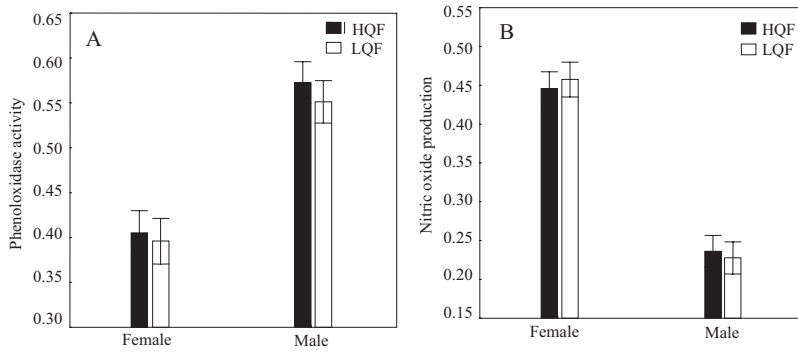


Fig. 6. Sexual dimorphism in (A) PO activity and (B) NO production for adult mosquitoes reared in varying regimes of food quality and quantity. HQF, high quality food; LQF, low quality food.

There were differences between male and female overall values of GEI (Fig. 4A and B), and PO activity and NO production (Fig. 5A and B), which suggests that the two sexes exhibit dissimilar immune strategies. This difference is consistent with the idea that traits involved in immune response do not have the same importance for the adult fitness of the two sexes in terms of condition (McKean and Nunney 2005). PO activity was larger in males than females that can be explained as a consequence of resources being allocated to reproductive activity (host search and oogenesis) rather than to the maintenance of PO cascade. During this cascade, phenylalanine and tyrosine are used as substrates for the formation of melanin and highly reactive and toxic intermediate molecules. Phenylalanine is also involved in cuticle formation and pigmentation, relevant in natural selection functions such as aposematism and crypsis (to avoid predation), thermoregulation, resistance to UV radiation and colored traits that can be under sexual selection (True 2003). Meanwhile, tyrosine is also responsible for coloring the egg chorion in some insects (Li and Christensen, 1993) and other metabolic pathways. It has been shown that mosquitoes undergoing melanization responses against filarial worms after blood-feeding exhibit a delay in tyrosine accumulation in the ovaries, and therefore a delay in egg production and oviposition (Ferdig et al. 1993). For these reasons, it is likely that phenylalanine and tyrosine are restrictive resources in females mosquitoes leading to trade-offs between immune response and other life history traits.

In the absence of an immune challenge, females would favor an increase in resource investment to longevity and egg production rather to immunity (see also Rolff 2002).

NO was produced by females in a twofold relation compared with males. It could be argued that a trade-off is also present. Arginine is an important immune molecule, but it is also essential for sperm maturation (Osanai and Chen 1993), egg production (Uchida 1993), long-term memory, chemosensory (antennal lobes, olfaction) and visual information processing (Müller, 1997). However, in *Anopheles stephensi*, NO limits parasite development (Tina et al. 2007), and in *Ae. aegypti* NO participates in the control of the dengue virus load (Ramos-Castañeda et al. 2008). This presumed sex bias in NO production could result from different kinds of pathogens attacking each sex. For example in mosquitoes, adult females require carbohydrates (sugar) and proteins for vitellogenesis that are usually present in the blood meal. Meanwhile, males need sugar solutions (nectar) (Clements 1999). If some pathogens are acquired when feeding (as it is the case of the dengue virus) when each sex occupies or uses different habitats, then both sexes should show extremely different pathogens and, therefore, distinct immune adaptations against pathogens. Also, if adult female mosquitoes that have inherited pathogens from their parents, or become contaminated when mating with infected males (males that have inherited pathogens too) (e.g., Diallo et al. 2000) are expected to enhance survival. Thus, not only there are differences

Table 4. Genetic and phenotypic correlations between PO activity and NO production in adult male and female mosquitoes reared in two different food regimes during larval development

	Males		Females	
	At HQF PO activity	At LQF PO activity	At HQF PO activity	At LQF PO activity
Phenotypic correlations				
NO production	0.191 (**) N = 244	0.144 (*) N = 242	0.082 (NS) N = 221	0.004 (NS) N = 207
Genetic correlations				
NO production	0.252 (NS) N = 44	0.031 (NS) N = 44	0.172 (NS) N = 44	0.218 (NS) N = 44

Spearman correlation coefficients calculated with untransformed data. P values are indicated between parentheses (*, $P < 0.05$; **, $P < 0.005$). HQF, high quality food; LQF, low quality food; NS, nonsignificant.

in the incidence of different pathogens affecting males and females that explain the sexual dimorphism observed for NO and PO, but different reproductive strategies followed by each sex that could be important. This fact could be reflected in the course of action of individual immune response.

Immune response occurs via a set of physiological traits and is supposed to show low additive variance and hence low heritability values (like other physiological traits, see Mousseau and Roff 1987). Our breeding results of PO activity and NO production confirmed the (additive) genetic variance previously reported for other immune parameters in other insects (e.g., Fellowes et al. 1998, Kurtz and Sauer 1999, Hosken 2001, Rolff et al. 2005, Simmons and Roberts 2005). In male mosquitoes, genetic variation can be maintained if environmental conditions vary and if distinct alleles maximize fitness under each environment. GEIs may contribute to genetic variability in immunocompetence although this consideration has been rarely investigated. Natural or sexual selection will act on phenotypic individual variance, favoring some phenotypes, and as long as the phenotypic variance has a genetic background, the population can evolve. The lack of genetic correlations (pleiotropy) (in males and females) and r_g (for males) indicate that if some genotypes are favored under different environmental conditions, there should be no strong genetic constraints for adaptation to each of the environments for independent PO and NO responses. However, the presence of additive genetic variation could be the result not only of direct selection on the mosquito's immune response, but also because of differences in immune strategies between the sexes, and indirect selection acting on genetically correlated traits (see Koella and Boëte 2002).

It has to be mentioned that our study was conducted using basal immune components (i.e., in the absence of an immune challenge). Moreover, this basal immune response could be adaptive in cases where females have inherited pathogens from their parents, or progeny that do not carry the virus at their emergence but become contaminated while mating with infected partners. Despite our methodological approach, it is likely that the interaction between *Ae. aegypti* genotypes and abiotic environment can affect the evolution of resistance to infection because the performance of different genotypes changed across food quality and quantity environments. In view of this, the effects of environmental variation must not be ignored in future studies of immunocompetence of mosquitoes and other insects. These results may also be important to understand the colonizing success of *Ae. aegypti* to new habitats (in natural or urbanized areas) and/or habitats where resources and pathogens vary seasonally and geographically (Kittayapong et al. 1999, Wearing and Rohani 2006).

Acknowledgments

We thank Salvador Hernández-Martínez for allowing access to protocols and helpful suggestions. We are deeply

grateful with Priscila Bascuñan, Guadalupe Hernández, Jimena Patiño, and Valeria Vargas for their help with larval maintenance and laboratory work. To Betsabé Ruiz for helpful suggestions on an early version of the manuscript. To Kim Van Ryzin for invaluable help. M.M-G. thanks the Posgrado en Ciencias Biomédicas (CONACYT Grant No. 172947), Instituto de Ecología, Universidad Nacional Autónoma de México, and Centro de Investigaciones Sobre Enfermedades Infecciosas, Instituto Nacional de Salud Pública, México. AC-A was supported from a PAPIIT-UNAM grant (Project No. 211506). Two anonymous reviewers provided key comments that greatly enriched this work.

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Received 6 November 2008; accepted 3 November 2009.

CAPÍTULO V. DISCUSIÓN Y CONCLUSIÓN GENERAL

En este trabajo se contribuyó a entender cómo la respuesta inmune afecta y se ve afectada por estrategias y características de historia de vida. En específico, cómo se ve afectada la respuesta inmune ante factores bióticos y abióticos como lo son la presencia recurrente de patógenos, y la heterogeneidad en los recursos alimenticios. Para esto se evaluaron dos aspectos: (1) el efecto de la presencia recurrente de un mismo patógeno en hembras del mosquito *Ae. aegypti*, la forma en que responde inmunológicamente y su consecuencia en la supervivencia y reproducción. (2) El efecto que tiene la variación en la calidad y cantidad de la dieta a lo largo del desarrollo larval en la actividad de fenoloxidasa y producción de óxido nítrico de hembras y machos adultos de *Aedes aegypti*. A continuación se discuten los puntos esenciales de cada una de estas observaciones, y se plantean perspectivas de estudio.

Factores que moldean la respuesta inmune de insectos

A pesar de que los insectos cuentan con un sistema inmune que consta de células especializadas, moléculas efectoras, y mecanismos de reconocimiento y regulación muy eficaces, constantemente sufren infecciones. Además la respuesta inmune varía entre especies. La amplia evidencia generada en diversos estudios de insectos indica que fisiológicamente el individuo tiene que decidir, en base a los recursos energéticos disponibles, la asignación de estos hacia diversas actividades, unas involucradas con la supervivencia y otras con la reproducción (Moret & Schmid-Hempel 2002). La inversión dirigida a montar y mantener la respuesta inmune para combatir patógenos infecciosos en muchas ocasiones limita la reproducción de los organismos (ej. Adamo et al, 2002). El individuo se verá favorecido con la inversión que al final le genere el dejar la mayor descendencia viable. La inversión diferencial de los recursos hacia distintas estrategias afectará directamente la evolución de de la respuesta inmune y otras características (ej. caracteres sexuales secundarios). Además hay que considerar la patogenicidad y virulencia del agente infeccioso (Pfennig, 2001). El daño que un patógeno puede infringir a su hospedero

está en función de estos dos aspectos. El hospedero también tendrá que “decidir” la inversión en energía en función del potencial daño.

Otro aspecto que se considera es que el sistema inmune no es simple, distintos parámetros que lo comprenden pueden estar correlacionados positiva o negativamente. El sistema inmune se activa cuando el organismo reconoce moléculas no propias, sin embargo, se sabe que en ocasiones el tipo de patógeno determinará la activación diferencial de los parámetros inmunes (Janeway & Medzhitov, 2002). Las correlaciones negativas podrán verse favorecidas en el caso en esta correlación tenga como consecuencia la eficaz eliminación del patógeno. Aunque puede darse el caso en que la correlación negativa entre parámetros inmunes sea consecuencia de la limitante en los recursos (de Jong & Van Noordwijk, 1992).

Se debe tener siempre en consideración que la disponibilidad de los recursos y la presencia (y recurrencia) de patógenos son temporal y espacialmente estocásticos. Ambos factores son causas próximas que actúan sobre el sistema inmune de insectos. La expresión de estrategias inmunes estará íntimamente relacionada con estos dos factores ecológicos.

*Encuentros recurrentes con patógenos y su efecto en *Aedes aegypti**

En esta tesis (Capítulo III) se comprobó que el mosquito puede resistir dosis letales de bacterias después de haber tenido un contacto previo con la misma bacteria (viva y a bajas concentraciones). Los datos de supervivencia a constantes retos con patógenos encontrados en este trabajo se une a la evidencia previa que hace constar que mosquitos tengan la capacidad de mostrar un tipo de memoria lo largo de la vida del individuo (i.e. priming inmunológico) (Little & Kraaijeveld, 2004; Little, Hultmark & Read, 2005, Pham & Schneider, 2009). Por primera vez se realizó una cinética midiendo dos parámetros en los que se esperaba que mostraran una mejora en su producción y activación. Sin embargo, ninguno de los efectores (fenoloxidasa y óxido nítrico) cuantificados mostró incremento significativo durante el segundo encuentro con el mismo patógeno. La actividad de fenoloxidasa muestra una continua deactivación. Es

posible que el que la continua activación de la fenoloxidasa evite daño (debido a las moléculas tóxicas producidas durante la cascada) a tejidos propios del mosquito. Lo que indica que el incremento en la supervivencia de los individuos esta relacionada con una regulación de los efectores inmunes, lo que sería ventajoso en caso que se incremente la eficiencia para controlar patógenos. Es posible que la actividad de fenoloxidasa, se vea comprometida con otras características, por ejemplo la producción de huevos. Se necesitan más estudios para poner en prueba esta idea. Así, se tendría un excelente ejemplo de la disyuntiva entre supervivencia y reproducción, y que la memoria inmunológica en el mosquito, a pesar de ser una estrategia adaptativa, también ha sido y es moldeada por factores ecológicos como es la limitante de recursos energéticos.

Como ocurre con frecuencia una pregunta biológica, a parte de responderse, genera más preguntas. En el caso del priming inmune encontrada en *Ae. aegypti* ocurre lo mismo y generan perspectivas de estudio. Por ejemplo, el priming inmune en otros organismos muestra especificidad hacia el patógeno. En insectos es probable que también exista este fenómeno, lo cual sería ventajoso si es que existen vías moleculares inmunes específicas para los distintos tipos de patógenos (bacterias Gram-, Gram+, hongos, protistas, parasitoides o virus). En el mosquito podrían hacerse infecciones cruzadas (ej. Gram- vs Gram+ o viceversa) para establecer si la especificidad también esta presente. Otro aspecto sería evaluar la presencia de un efecto transgeneracional del fenómeno de priming inmune. Este aspecto fue contemplado teóricamente en el Capítulo II. Se presentó evidencia, a partir de diversos estudios llevados a cabo por distintos grupos de investigación, de que padres inmunizados pueden heredar la información a sus hijos, haciéndolos menos susceptibles al ataque de patógenos. Aunque temporalmente es difícil que un mosquito adulto tenga contacto con sus hijos, el ambiente en el que viven sí puede ser compartido. La existencia de priming transgeneracional podría ser una estrategia adaptativa en este grupo de insectos. Por último, no hay que dejar el lado el necesario estudio de los mecanismos genéticos, moleculares y celulares que se dan para que los insectos puedan presentar el fenómeno de priming inmune. En esta tesis se observó la existencia del priming en el mosquito, sin embargo se desconoce el mecanismos de cómo

es que esto ocurre. Actualmente el estudio de los mecanismos inmunes es llevado a cabo por un gran número de grupos de investigación, diversas herramientas están siendo utilizadas para investigar cuáles son los parámetros inmunes que se activan y cómo es que están regulados.

Efecto de la variación en la calidad y cantidad de la dieta en la respuesta inmune del mosquito

Los resultados obtenidos en esta tesis (Capítulo IV) muestran que los individuos tienen una constitución genética que permite variaciones fenotípicas para ajustarse a la heterogeneidad ambiental. En específico, en ocasiones un solo genotipo puede tener la capacidad de producir dos fenotipos alternativos como resultado de su interacción con la limitación de alimento durante el desarrollo (Schlichting 1986). La respuesta inmune se puede ver como la suma de los efectos genéticos y los efectos ambientales que se dan durante el desarrollo larval, dando como resultado variación en la respuesta inmune del adulto. Es posible que la plasticidad pueda estar manteniendo la variación en la población. Las diferencias fenotípicas entre los individuos son, generalmente, las responsables de las diferencias en la adecuación, por lo tanto el cambio evolutivo en la respuesta inmune, debido a selección natural, estará determinado por dicha variación. Relacionado con lo anterior, las diferencias en la respuesta inmune entre hembras y machos pueden ser utilizadas como un indicador de las estrategias que cada sexo sigue dependiendo del ambiente y la forma de respuesta ante éste. Lo que podría estar cambiando entre generaciones no son los genes sino las normas de reacción de los genotipos, los beneficios o costos que la plasticidad traiga va a dar como resultado la evolución de las normas de reacción del fenotipo inmune que se está expresando.

Al igual que para el priming, los mecanismos moleculares de plasticidad fenotípica son desconocidos. Sin embargo, es un hecho que las señales ambientales son moduladores de la actividad transcripcional de genes, alterando su expresión (Kent et al. 2009). La base molecular se desconoce, sin embargo, se ha propuesto que genes con estructura TATA box, son capaces de de respuesta rápidas y variables (Richards et al. 2006; Liefting et al. 2009). Procesos epigenéticos, como la metilación de ADN, puede modificar el nivel de expresión de caracteres (Bender 2004). Es posible que estos mecanismos sean los

responsables de las diferencias fenotípicas observadas dentro de los genotipos. Estas diferencias estarían dadas como respuesta a la variación ambiental. El priming inmune también podría estar siendo modulado por fenómenos epigenéticos. La presencia de patógenos puede ser considerada como una señal tanto de ausencia y presencia, como de intensidad (virulencia). La modulación transcripcional a través de epigenesis necesita ser evaluada para lograr entender los mecanismos detrás de la expresión de la respuesta inmune en insectos.

Implicaciones del estudio de la ecología evolutiva del mosquito en salud pública

El mosquito actualmente en zonas en las cuales comúnmente no se encontraba (ver Fig. 2, Capítulo I). Este hecho no solo se debe al incremento de temperatura, que posiblemente contribuya a ampliar el rango de distribución del mosquito, sino a la capacidad del mosquito de colonizar y mantenerse en nuevos nichos ecológicos. Esta capacidad inherente del mosquito muy posiblemente está relacionada a la plasticidad del organismo para adaptarse al ambiente. El estudio de la existencia de variación genética, efecto del ambiente y la interacción Genotipo x Ambiente en diversas poblaciones de mosquitos de campo sería valioso para poder concluir si en realidad la plasticidad fenotípica es realmente una característica esencial para el éxito del mosquito.

Todos los organismos están bajo en continuo contacto con agentes infecciosos. Debido a esto es necesario un adecuado y eficiente sistema de defensa para proteger la integridad, y asegurar la supervivencia y reproducción, del organismo. Barreras físicas, células y sobre todo moléculas son comunes entre los distintos taxones (incluyendo plantas y probablemente hongos) (Heine, 2008). En los últimos años la concepción del sistema innato de defensa ha dejado de considerarse “sencillo” para ahora tener el lugar que desde un principio le correspondía que es el de “esencial e imprescindible”. El sistema inmune de insectos difiere con mamíferos por la ausencia del llamado sistema adaptativo (linfocitos y anticuerpos). Sin embargo cuentan con un sistema eficaz de defensa. Este sistema, tiene como finalidad conservar un estado de homeostasis entre el organismo y el medio que lo rodea. El estudio de fenómenos

como la memoria inmunológica y aspectos genéticos cuantitativos de la respuesta inmune de los mosquitos son útiles para el entendimiento de los procesos evolutivos y ecológicos que la están moldeando. Este conocimiento puede y debe ser aplicado, por ejemplo la memoria inmune de mosquitos puede ser útil para limitar la transmisión del virus Dengue. El esparcir virus Dengue inactivo (inclusive mezclado con insecticida) podría hacer que los mosquitos que sobreviven puedan generar memoria, para que cuando estén en contacto con el virus activo, este tenga una nula o menor tasa de replicación dentro del mosquito y en consecuencia disminuir la incidencia de casos de dengue clásico o hemorrágico en las poblaciones humanas.

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Current immunity markers in insect ecological immunology: assumed trade-offs and methodological issues.



Journal:	<i>Methods in Ecology and Evolution</i>
Manuscript ID:	Draft
Manuscript type:	Review
Date Submitted by the Author:	n/a
Complete List of Authors:	Moreno-García, Miguel; Instituto Nacional de Salud Pública, Centro de Investigaciones Sobre Enfermedades Infecciosas; Universidad Nacional Autónoma de México, Instituto de Ecología Córdoba-Aguilar, Alex; Universidad Nacional Autónoma de México, Ecología Evolutiva Lanz-Mendoza, Humberto; Instituto Nacional de Salud Pública, Centro de Investigaciones Sobre Enfermedades Infecciosas
Keywords:	Evolutionary Biology, Population Genetics
Abstract:	The field of ecological immunology currently relies on using a number of immune effectors or markers. These markers are usually used to infer ecological trade-offs (via conflicts in resource allocation), though physiological nature of these markers remains elusive. Here, we review markers frequently used in insect evolutionary ecology research: cuticle darkening, haemocyte density, nodule/capsule formation, phagocytosis and encapsulation/melanization via use of nylon filaments and beads, phenoloxidase activity, nitric oxide production, lysozyme and antimicrobial peptide production. We also provide physiologically based information that may shed light on the probable trade-offs inferred when these markers are used. In addition, we provide a number of methodological suggestions to improve immune marker assessment.

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3 **Current immunity markers in insect ecological immunology: assumed**
4 **trade-offs and methodological issues**

5

6 RUNNING TITLE: INSECT IMMUNO-ECOLOGY MARKERS, MORENO-GARCÍA

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20 **Abstract**

21 THE FIELD OF ECOLOGICAL IMMUNOLOGY CURRENTLY RELIES ON USING A NUMBER OF IMMUNE
22 EFFECTORS OR MARKERS. THESE MARKERS ARE USUALLY USED TO INFER ECOLOGICAL TRENDS (E.G. RESOURCE
23 CONFLICTS IN RESOURCE ALLOCATION), THOUGH PHYSIOLOGICAL NATURE OF THESE MARKERS IS OFTEN
24 ELUSIVE. HERE, WE REVIEW MARKERS FREQUENTLY USED IN INSECT EVOLUTIONARY ECOLOGY
25 RESEARCH: CUTICLE DARKENING, HAEMOCYTE DENSITY, NODULE/CAPSULE FORMATION, PHENOLASE
26 AND ENCAPSULATION/MELANIZATION VIA USE OF NYLON FILAMENTS AND BEADS, PHENOLASE
27 ACTIVITY, NITRIC OXIDE PRODUCTION, LYSOZYME AND ANTIMICROBIAL PEPTIDE PRODUCTION.
28 PROVIDE PHYSIOLOGICALLY BASED INFORMATION THAT MAY SHED LIGHT ON THE PROBLEMS
29 INFERRED WHEN THESE MARKERS ARE USED. IN ADDITION, WE PROVIDE A NUMBER OF
30 METHODOLOGICAL SUGGESTIONS TO IMPROVE IMMUNE MARKER ASSESSMENT.

31

32 **Keywords:** EVOLUTIONARY ECOLOGY, INSECT IMMUNITY, IMMUNE MARKERS

33 Introduction

34

35 ECOLOGICAL IMMUNOLOGY IS A RAPIDLY EXPANDING YOUNG DISCIPLINE (ROLFF & REYNOLDS
36 2009). WITHIN THIS FIELD, IMMUNE RESPONSE IS SEEN AS A SET OF TRAITS THAT INVOLVE
37 (SHELDON & VERHULST 1996). RESOURCES MUST BE ALLOCATED BETWEEN AN ORGANISM'S
38 MAINTENANCE AND PROPER FUNCTIONING AND THE EXPRESSION OF OTHER TRAITS LINKED TO
39 SURVIVAL AND REPRODUCTION, RESULTING IN ECOLOGICAL TRADE-OFFS AMONG TRAIT
40 MOMENT, THERE IS GROWING EMPIRICAL EVIDENCE OF TRADE-OFFS ASSOCIATED BETWEEN
41 TRAITS (SIZE AT BIRTH, GROWTH AND MORTALITY RATES, SIZE AND AGE AT MATURITY, LIFETIME
42 SURVIVAL, CLUTCH SIZE AND REPRODUCTIVE EFFORT; REVIEWED IN STEARNS 1992) AND
43 DEFENSE (E.G. KRAAIJEVELD & GODFRAY 1997, RIGBY & JOKELA 2000, FELLOWS
44 BOOTS & BEGON 1993, MORET & SCHMID-HEMPEL 2000, HOANG 2001).

45

46 THE ECOLOGICAL IMMUNITY FRAMEWORK HAS BEEN APPLIED TO A NUMBER OF DISCIPLINES
47 INCLUDING SEXUAL SELECTION (I.E. DIFFERENTIAL MATING AND FERTILIZATION SUCCESS
48 1871) (E.G. SIVA-JOTHY 2000; MCKEAN & NUNNEY 2001, ROBERTSON *et al.* 2007), TO A
49 LESSER EXTENT, SOCIAL EVOLUTION (E.G. BOEDERIE *et al.* 2008), PREDATOR-PREY
50 RELATIONSHIPS (E.G. PACKER & ROY & HOLT 2008) AND PARENTAL INVESTMENT (E.G.
51 HOI-LEITNER *et al.* 2001; BRZEK & KONARZEWSKI 2007). THE INFLUENCE OF ECOLOGICAL
52 IMMUNITY HAS ALSO EXTENDED TO BROADER FIELDS SUCH AS LIFE-HISTORY THEORY (SHELDON
53 2005A), PARASITE-HOST COEVOLUTION (E.G. DUFFY *et al.* 2007), CONSERVATION
54 BIOLOGY (E.G. STEVENSON 2006, TARLOW & BLUMSTEIN 2007) AND LEARNING (E.G. BARNARD
55 *et al.* 2006), AMONG OTHERS.

56

57 ONE ANIMAL GROUP WHERE ECOLOGICAL IMMUNOLOGY HAS BEEN EXTENSIVELY INVESTIGATED
58 INSECTS (SEE, FOR EXAMPLE, LAWRENCE & KROMBEIN 2006). INSECTS HAVE PROPERTIES (E.G. SMALL SIZE,
59 LARGE SAMPLE SIZES, SHORT DEVELOPMENT TIMES) THAT ALLOW A WIDE RANGE OF EXPERIMENTAL
60 INSECT STUDIES HAVE MEASURED A NUMBER OF IMMUNE EFFECTORS RELATED TO BOTH
61 HUMORAL DEFENCE REACTIONS (HEREON REFERRED TO AS IMMUNE MARKERS), IN WHICH
62 SUCH IMMUNE RESPONSE, MEASURED THROUGH THESE MARKERS, IS ASSUMED (SEE ALSO
63 2004). THE LOGIC OF CHOOSING SUCH MARKERS HAS BEEN BASED ON THE ASSUMPTION THAT THEY
64 PROTECT THE INSECT, BUT THEIR PRODUCTION NEGATIVELY CORRELATES WITH OTHER LIFE-HISTORY
65 INTERMEDIATE TRAITS (E.G. COLORED TRAITS, BEHAVIORAL TRAITS, ETC.). MANY FEASIBLE
66 SENSITIVE TECHNIQUES IN INSECT IMMUNOLOGY (E.G. PROTEOMIC, TRANSCRIPTOMIC, LIQUID CHROMATOGRAPHY /MASS SPECTROMETRY) HAVE BEEN EMPLOYED AT THE INDIVIDUAL-LEVEL. HOWEVER, HIGH COSTS CONFINE THESE APPROACHES TO EXPLORATORY OR QUALITATIVE STUDIES AT THE POPULATION-LEVEL. THIS TURNS INTO A RESTRICTION WHEN THE OBJECTS OF STUDY ARE PRIMARILY INDIVIDUALS IN NATURAL POPULATIONS; BECAUSE DIFFERENCES AMONG ORGANISMS IN IMMUNE RESPONSES (AND THEIR TRADE-OFFS AMONG OTHER BEHAVIOURAL OR REPRODUCTIVE TRAITS) ARE AN IMPORTANT CAUSE OF ADAPTIVE CHANGE OF THE POPULATION.

73

74 HERE WE REVIEW EACH OF THE MOST COMMON IMMUNE MARKERS USED IN ECOLOGICAL AND EVOLUTIONARY STUDIES THAT CAN BE UTILIZED ACROSS LARGE SAMPLE SIZES. OUR FIRST AIM IS TO IDENTIFY, WHEN POSSIBLE, THE PHYSIOLOGICAL PROCESSES BEHIND EACH MARKER. THIS IS FOLLOWED BY THE INFERENCE OF TRADE-OFFS WITH OTHER FUNCTIONS THROUGH INDIVIDUAL DATA. SECOND, WE BRIEFLY OUTLINE THE METHODOLOGICAL INCONVENIENCES IN THE ASSESSMENT OF SUCH TRADE-OFFS AND SUGGEST SOLUTIONS. OUR AIM IS TO PROVIDE A SET OF MORE ROBUST METHODOLOGICAL APPROACHES. AN OUTLINE OF THESE TWO AIMS APPEARS IN TABLE 1, WE PRESENT EACH IN GREATER DETAIL IN THE

81 BELOW. FIRST, HOWEVER, WE PRESENT A BASIC SHORT REVIEW OF HOW INSECT IMMUNITY
82 WHICH SERVES AS THE FRAMEWORK FOR DISCUSSING THE TRADE-OFFS AND METHODOLOGY
83

84 **Insect Immune Mechanisms**

85 THE INSECT IMMUNE SYSTEM IS FORMED BY A SET OF CELLS, MOLECULES AND REACTIONS
86 THESE FEATURES ARE CONTINUOUSLY EVOLVING TO RESIST (ATTACK AND ELIMINATE) PATHOGENS
87 INVASION AND TO LIMIT THE NEGATIVE CONSEQUENCES OF THE INFECTION (HOFFMAN &
88 2002, SCHMID-HEMPEL 2005A, SCHNEIDER 2009). THE FIRST LINES OF DEFENCE INCLUDE THE
89 EXOSKELETON CUTICLE, PHYSICAL AND CHEMICAL PROPERTIES (E.G. PH) OF THE EPIDERMAL
90 EPITHELIUM, AND MALE AND FEMALE REPRODUCTIVE ACCESSORIES (GILLESPIE
91 CASTEELS 1998). THESE TISSUES ALSO SECRETE CYTOTOXIC MOLECULES LIKE LYSOZYME
92 REACTIVE OXYGEN SPECIES (ROS: E.G. SUPEROXIDE ANIONS, PEROXIDES, HYDROXYL RADICALS)
93 (SCHMID-HEMPEL 2005A). PATHOGENS ARE MAINLY RECOGNIZED BY THE MEMBRANE OF
94 HAEMOCYTES (FREE IN THE HAEMOLYMPH) OR BY THE MEMBRANES OF EPITHELIAL CELL
95 MEMBRANES BEAR PROTEINS CALLED PATTERN RECOGNITION RECEPTORS (PRRS) THAT RECOGNIZE
96 CONSERVED MOLECULAR FEATURES OF PATHOGENS CALLED MICELLAR-ASSOCIATED
97 PATTERNS (PAMPS: LPS, MANNAN, CHITIN, GLUCANS AND PEPTIDOGLYCANS) (GILLESPIE
98 1997). ONCE PATHOGENS ARE RECOGNIZED AS NON-SELF, CELLULAR AND HUMORAL IMMUNE
99 MECHANISMS ARE ACTIVATED. CELLULAR RESPONSES INCLUDE PHAGOCYTOSIS, NODULATION
100 ENCAPSULATION. DURING ENCAPSULATION, SMALL AND LARGE PATHOGENS ARE SURROUNDED
101 BOUND BY HAEMOCYTES (GILLESPIE *et al.* 1997). DURING NODULATION AND ENCAPSULATION
102 MELANIN LAYER IS CONSTRUCTED (FREQUENTLY REFERRED TO AS MELANOTIC
103 NODULATION/ENCAPSULATION) TO COVER FOREIGN AGENTS, WHICH ULTIMATELY DIE BY
104 ROS OR STARVATION (NOTTOVIANI 2000, NARAYANAN 2004).

105

106 MELANIN PRODUCTION IS A CONSEQUENCE OF THE PHENOLOXIDASE (PO) CASCADE (SÖDERHÄLL & CERENIUS 1998). DURING THE CASCADE, OPSONIC FACTORS, ROS, AND CYTOTOXINS SUCH AS QUINONES AND SEMIQUINONES ARE PRODUCED. THESE IMPORTANT INTERMEDIATE MOLECULES ARE HIGHLY REACTIVE AND TOXIC TO PATHOGENS, AND SERVE TO AMPLIFY THE IMMUNE RESPONSE (CERENIUS & SÖDERHÄLL 2004, NAPPI & OTTOVIANNI 2000). PO IS INVOLVED IN VARIOUS PHYSIOLOGICAL PROCESSES INCLUDING: CUTICULAR SCLEROTIZATION, IMMUNE DEFENSE (MELANOTIC ENCAPSULATION AND WOUND HEALING). THREE KINDS OF PO EXIST (SUGUMARAN & KANOST 1993): MONOPHENOL MONOOXIGENASE (ALSO REFERRED TO AS TYROSINASE-TYROSINASE (ASHIDA & BREY 1997), *o*-DIPHENOLOXIDASE (ALSO REFERRED TO AS CATHECOLOXIDASE-TYROSINASE (DECKER & JAENICKE 2004) AND *p*-DIPHENOLOXIDASE (ALSO REFERRED TO AS LACCASE-TYROSINASE (SUGUMARAN & KANOST 1993). THESE THREE KINDS OF PO AND THEIR ACTIVATING SYSTEMS ARE STRUCTURALLY ALMOST INDISTINGUISHABLE; HENCE, THE TERM PHENOLOXIDASE IS OFTEN USED IN THE LITERATURE WITHOUT DISTINGUISHING AMONG THEM. PO ASSOCIATED WITH SCLEROTIZATION AND PIGMENTATION OF CUTICLE (MONOPHENOL MONOOXIGENASE AND *p*-DIPHENOLOXIDASE) CAN BE UNDER HORMONAL CONTROL (ASHIDA & BREY 1997), WHILE *o*-DIPHENOLOXIDASE IS TRIGGERED BY RECOGNITION OF NON-SELF PARTICLES (ASHIDA & BREY 1997). HAEMOLYMPH PO IS MAINLY KEPT INSIDE OR NEAR HAEMOCYTES; NEVERTHELESS, IT CAN BE A FEATURE OF THE HUMORAL IMMUNE RESPONSE. THE PRECURSOR OF PO, PRO-PHENOLOXIDASE (PROPO) IS PRIMARILY RELEASED FROM HAEMOCYTES INTO THE HAEMOLYMPH AFTER CONTACT WITH NON-SELF PARTICLES (LING & YU 2006). IT CONTRIBUTES TO HUMORAL MELANIZATION OF PATHOGENS. THE GENERAL PO ACTIVATION IS INITIATED WHEN PHENYLALANINE IS HYDROXYLATED AND CONVERTED INTO TYROSINE. TYROSINE IS THEN INDUCED INTO ACTIVE FORM PO. PO CATALYZES BOTH THE HYDROXYLATION OF TYROSINE AND THE OXIDATION OF DOPA TO DOPAQUINONE. FINALLY, DOPAQUINONE IS CONVERTED TO MELANIN, WHICH IS USED FOR WRAPPING PATHOGENS AND WOUND CLOSURE (CHRISTENSEN *et al* 2005, SÖDERHÄLL & CERENIUS 1998, NAPPI & CHRISTENSEN 2005).

131

132 OTHER HUMORAL RESPONSES INCLUDE ANTIMICROBIAL SYSTEMIC MOLECULES, WHICH A
133 SYNTHESIZED MAINLY IN THE FAT BODY AND REPRODUCTIVE ACCESSORY GLANDS, GUT
134 HAEMOCYTES (MANETTI 1998, SCHMID-HEMPEL 2005A). MOST OF THESE MOLECULES ARE
135 SECRETED CLOSE TO THE CUTICLE OR INTO THE HAEMOLYMPH, GUT TRACT, OR MALPIGHI
136 ANTIMICROBIAL PEPTIDE (AMP) MOLECULES INDUCE A NUMBER OF NEGATIVE EFFECTS ON
137 MEMBRANES INCLUDING MEMBRANE COLLAPSE, PREVENTION OF CELL DIVISION, AND PE
138 DISRUPTION (OTVOS JR 2000, BUDDE 2003). NITRIC OXIDE (NO) IS ANOTHER IMMUNE-
139 RELEVANT MOLECULE: IT IS A HIGHLY REACTIVE AND UNSTABLE FREE RADICAL GAS THAT
140 MEMBRANES TO ACT ON NEARBY TARGETS (MÜLLER 1997). NO ALSO INHIBITS PROTEIN CA
141 ACTIVITY AND HAS DAMAGING EFFECTS ON PATHOGEN PROTEIN AND DNA (REVIEWED IN
142 2006). MOLECULES SUCH AS ROS CAN DAMAGE PATHOGEN NUCLEIC ACIDS, PROTEINS AND
143 MEMBRANE (NAPPI 2000, HERRERA-ORTIZ 2004). LYSOZYMES ARE HYDROLYTIC
144 ENZYMES THAT CLEAVE THE GLYCOSIDIC BOND BETWEEN N-ACETYLMURAMIC ACID AND
145 ACETYLGLUCOSAMINE IN PEPTIDOGLYCAN, A MAJOR COMPONENT OF THE GRAM
146 (JOLLÈS 1996).

147

148 *Trade-offs among immune responses*

149 TRADE-OFFS WITHIN IMMUNE SYSTEM EFFECTORS ARE ALSO POSSIBLE. IN AN INDIVIDUAL
150 TRAITS ARE FREQUENTLY FOUND TO BE GENETICALLY OR PHENOTYPICALLY CORRELATE
151 CORRELATION ARISES BECAUSE A SINGLE GENE CAN INFLUENCE MULTIPLE TRAITS IN PO
152 NEGATIVE FASHIONS (PLEIOTROPY AND ANTAGONISTIC PLEIOTROPY, RESPECTIVELY) OR
153 LINKAGE DISEQUILIBRIUM BETWEEN GENES AFFECTING DIFFERENT CHARACTERS (FALCO
154 1996). PHENOTYPIC CORRELATIONS INCLUDE THE POSITIVE OR NEGATIVE INFLUENCES OF
155 ENVIRONMENTAL FACTORS ON TRAITS (ROFF 1992). REACTIONS AND PRODUCTS OF IMMUN

156 ARE INTERCONNECTED AND THE KIND OF RESPONSE COULD BE RELATED TO THE TYPE OF
157 PATHOGEN. A NUMBER OF STUDIES HAVE EXAMINED GENETIC AND PHENOTYPIC CORRELATIONS
158 AMONG ENCAPSULATION, LYTIC ACTIVITY, CUTICULAR DARKNESS, PO ACTIVITY AND HAIR PRESENCE
159 IN SUMMARY, THESE STUDIES REPORT POSITIVE AND NEGATIVE CORRELATIONS RESULTING IN NO
160 CLEAR PATTERN ACROSS SPECIES OR EVEN POPULATIONS (SEE RANTALA & KORTET 2003,
161 2004, FEDORKA *et al.* 2004, RYDER & SIVA-JOTHY 2001, RANTALA & ROFF 2005, ROFF *et al.*
162 2005, MORENO-GARCÍA 2010). THIS POINT WILL BE RECONSIDERED LATER.

163

164

165 **Immune Markers Used in Evolutionary Ecology Research**

166

167 *Cuticle darkness*

168 THE MAIN COMPONENTS OF INSECT CUTICLE –QUINONES AND MELANIN (SEE PHENOLOXIDASE
169 ACTIVITY) – ARE ALSO USED FOR OTHER FUNCTIONS, THUS STRONGLY SUGGESTING A TRADE-OFF
170 (BARNES & SIVA-JOTHY 2000, ARMITAGE & SIVA-JOTHY 2005). QUINONE COMPOUNDS
171 DETERMINE CUTICLE PIGMENTATION (VIA SCLEROTIZATION) BY MAKING COVALENT LINKS WITH
172 PROTEINS, RESULTING IN COLOURED PRODUCTS, A PROCESS CALLED QUINONE TANNING (HIRUMA
173 1988). QUINONES ALSO FORM METHIDE DERIVATIVES THAT CROSSLINK CUTICLE
174 THROUGH THE QUINONE SCLEROTIZATION REACTION) AND ARE RESPONSIBLE FOR CUTICLE
175 HARDENING (SAUL & SUGUMARAN 1988). CUTICLE DARKNESS IS THUS RELATED TO HOW MUCH
176 QUINONE IS DEPOSITED IN EVERY SHED OR WOUND REPAIR SITE.

177

178 *Methodology*

179

180 CUTICULAR COLOUR IS FREQUENTLY ASSESSED UNDER A FLUORESCENT WHITE LIGHT AND
181 COLOUR CLASSES ARE ANALYZED USING A GREYSCALE VALUE (E.G. BARNES & SIVA-JOTHY
182 THOMPSON *et al.* 2002, ROLFF *et al.* 2005). HOWEVER, SINCE THE CUTICLE STRUCTURE IS
183 COMPOSED OF HUNDREDS OF PROTEINS (ANDERSEN *et al.* 2007) AND LIPIDS (CHAPMAN
184 1998), AND SUCH PROTEIN COMPOSITION GREATLY INFLUENCES CUTICLE MECHANICAL PROPERTIES
185 (E.G. FLEXIBILITY; GOSLING 2002, HAAS *et al.* 2000), THERE COULD BE A POSSIBLE
186 OVERESTIMATION OF QUININE AND MELANIN TANNING. FURTHERMORE, SINCE CUTICLES OF
187 TAXA VARY CONSIDERABLY IN STIFFNESS, HARDNESS, AND PIGMENTATION, IT IS POSSIBLE THAT
188 ASPECT OF A PARTICULAR TISSUE MAY NOT NECESSARILY REFLECT OR BE RELATED TO A
189 HEIGHTENED IMMUNE ABILITY. IN COMBINATION WITH CUTICULAR COLOUR ANALYSIS, WITH
190 THE USE OF TRANSVERSE SECTION PREPARATIONS (USING A MICROTOME) FROM DIFFERENT
191 REPRESENTATIVE REGIONS OF THE INSECT BODY AND CUTICLE WIDTH MEASUREMENT. THIS
192 REFLECT HOW MUCH THE INSECT HAS INVESTED IN PRODUCING A THICK CUTICLE AND, PRODUCTION OF
193 QUININES AND MELANIN. NOTE, HOWEVER, THAT CUTICULAR MELANIZATION IS NOT ALWAYS
194 WITH GREATER IMMUNOGENIC RESPONSE (SEE ROBB *et al.* 2003).

195

196

197 *Cellular Responses*

198

199 *Total number of haemocytes (haemocyte density)*

200

201 HAEMOCYTE COUNTING AND/OR ACTIVITY IS A STRAIGHTFORWARD WAY OF ASSESSING
202 IMMUNE ABILITY (E.G. RANTALA *et al.* 2000, KRAAIJEVELD *et al.* 2001; COTTER *et al.* 2004).
203 HAEMOCYTES ARE GENERALLY RESPONSIBLE FOR IMMUNE ACTIVITIES SUCH AS ENGULFING
204 SURROUNDING AND DESTROYING INFECTIOUS AGENTS. AFTER BACTERIA HAVE PENETRATED

205 HAEMOCOEL, HAEMOCYTES ARE THE FIRST AND MOST EFFICIENT BACTERIA-CLEARING M
206 (HILLYER *et al.* 2003). HOWEVER, THERE ARE DIFFERENT HAEMOCYTE TYPES, EACH PLAYING A
207 DIFFERENT FUNCTION (RECOGNITION, PHAGOCYTOSIS, NODULATION, ENCAPSULATION OF
208 (SEE REVIEWS BY LAVINE & STRAND 2002, RIBEIRO & BREHELÍN 2006). DIFFERENCES IN
209 PATHOGEN TYPE (E.G. BACTERIA, VIRUS, NEMATODES, ETC.) AMONG INDIVIDUALS, SEXES
210 MUST BE CONSIDERED. FOR EXAMPLE, IN *Aedes aegypti* *Escherichia coli* IS MAINLY
211 PHAGOCYTOSED, MEANWHILE *Micrococcus melanos* IS MELANIZED (HILLYER 2003).
212 HOWEVER, THE FACTORS ELICITING PHAGOCYTTIC VS. MELANIZATION RESPONSES AGAINST
213 INDEPENDENT OF GRAM TYPE (HILL 2004)

214

215 *Methodology*

216

217 FOR TOTAL AND VIABLE COUNTS, HAEMOLYMPH IS COLLECTED AND INCUBATED IN HUMIDITY
218 THEN COUNTED USING AN HEMOCYTOMETER USING (PHASE CONTRAST) MICROSCOPY (E.
219 *al.* 2000; KRAAIJEVELD. 2001). FOR COLLECTION AND INCUBATION OF HAEMOLYMPH, WE
220 SUGGEST THE USE OF CELL CULTURE MEDIA SUCH AS RPMI, SCHNEIDER OR GRACE RATHER THAN
221 BUFFERS. THIS MEDIA MAY ALLOW A BETTER VIABILITY OF CELLS. TRYPAN BLUE CAN BE USED TO
222 EXCLUDE BETWEEN DEAD AND VIABLE CELLS. HAEMOLYMPH EXTRACTION MAY BE ACCOMPLISHED BY
223 PERFUSION-BLEED; A DROPLET CAN BE COLLECTED FROM A PUNCTURE/CUT ON THE THORAX OR
224 REMOVED LEG. ALTHOUGH THIS MAY APPEAR AN EASY TASK, HAEMOCYTES CAN ADHERE TO SURFACES
225 SUCH AS THE FAT BODY. ONE SOLUTION TO THIS PROBLEM IS TO USE ANTICOAGULANT BLEND
226 PH (PECH *et al.* 1994). AN ACCURATE EXTRACTION OF THESE CELLS CAN ALSO BE CARRIED OUT BY
227 INJECTING A PROTEASE INHIBITOR BLEND (E.G. PMSF, TLCK, LEUPEPTINE, EDTA). HOWEVER, THE
228 OPTIMAL CONDITIONS FOR OBTAINING INSECT HAEMOCYTES CANNOT BE GENERALIZED, AND THE
229 PROCEDURE OF EXTRACTION THE USE (OR NON-USE) OF ANTICOAGULANT MUST BE ADJUSTED

230

231

232 *Phagocytosis*

233

234 PHAGOCYTTIC ACTIVITIES REQUIRE THE INTERNALIZATION OF THE SURFACE MEMBRANE AND

235 SEVERAL ORGANELLES, INCLUDING ENDOSOMES AND LYSOSOMES, WHICH FUSE WITH THE

236 MEMBRANE AND PROVIDE THE MEMBRANE FOR PHAGOSOME FORMATION (DESJARDINS *et al.*

237 IN MOSQUITOES, THE SYNTHESIS OF LIPIDS FOR THE MEMBRANES OF THE PHAGOSOME, PHAGOCYTTIC

238 MEMBRANE AND FOR THE PRODUCTION OF CYTOTOXIC PROTEINS FOR THE DEGRADATION OF

239 PHAGOCYTTISED MATERIAL DECREASES THEIR REPRODUCTIVE OUTPUT (FERDIG

240 HURD 1995, HILLYER 2003). FOR THESE REASONS, TRADE-OFFS ARE EXPECTED AMONG

241 PHAGOCYTTIC ACTIVITIES AND FITNESS TRAITS. FURTHERMORE, THAT HAEMOCYTES

242 UNDERGO APOPTOSIS AFTER PHAGOCYTTOSIS, WHICH REDUCES THE NUMBER OF CIRCULATING

243 HAEMOCYTES (AS OCCURS IN *Aedes triseriatus* HILLYER *et al.* 2005). THIS WOULD LEAD TO ANOTHER

244 TRADE-OFF, AS IT WOULD GENERATE A DECREASE IN THE HOMEOSTASIS AND REMODELLING

245 AS WELL AS THE CAPACITY FOR FIGHTING PATHOGENS. PHAGOCYTTOSIS IS AN IMPORTANT MEANS TO

246 REMOVE DEAD OR DAMAGED CELLS, AS WELL AS TO REMODEL ORGANS DURING METAMORPHOSIS

247 (ABRAMS *et al.* 1993, FRANC ET AL, 1996). THUS, A DELAY IN GROWTH SEEMS NECESSARY FOR

248 PATHOGEN CLEARING, VIA RELEASING HAEMOCYTES FROM THEIR IMMUNE ACTIVITY AND

249 THE WOUND REPAIRING PROCESS BEFORE CUTICLE SHEDDING (LAVINE & STRAND 2002).

250

251 *Methodology*

252

253 PHAGOCYTTIC ACTIVITY HAS BEEN MEASURED BY EXPOSING THE HOST TO MICROBIAL PATHOGENS

254 AS BACTERIA AND YEAST CELLS OR POLYSTYRENE BEADS, AND ASSESSING *in vitro* PHAGOCYTTIC

255 (KURTZ & SAUER 1999, KURTZ ~~2000~~ KURTZ 2002). ONE WAY TO ASSESS THIS PROCESS IS BY
256 INOCULATING FLUORESCENT, LABELLED BEADS ~~(CORRECTOR)~~ IN EITHER PROCESS,
257 THE PHAGOCYTISED INTRUDERS ARE COUNTED MANUALLY OR BY USING IMAGE ANALYSIS
258 CARE MUST BE TAKEN TO ENSURE THE BEADS DO NOT BECOME OBSTRUCTED BY MELANIN
259 OF REMAINING BEADS (IF POSSIBLE) FROM THE HAEMOLYMPH IS A USEFUL METHOD TO EVALUATE
260 PHAGOCYTOSIS ACTIVITY. TRYPAN BLUE CAN ALSO FURTHER ASSESS THE VIABILITY OF INGESTED
261 MOLECULES. THE EFFECT OF THIS COLORANT IS THAT ONLY THE INGESTED PARTICLES REVEAL
262 FLUORESCENCE, WHILE NON-INGESTED PARTICLES BECOME COATED WITH TRYPAN BLUE
263 *et al.* 2000). ANOTHER METHOD IS TO USE PHRODO DYE CONJUGATED BACTERIA. BACTERIA
264 FLUORESCENT ONLY IN THE ACIDIC ENVIRONMENT OF THE HAEMOCYTE PHAGOSOME (SEE
265 FEDROW *et al.* 2009); WHILE EXTRACELLULAR NON-PHAGOCYTOSED BACTERIA DO NOT FLUORESCENE
266 WHEN USING LIVE BACTERIA TO EVALUATE THE EFFICIENCY OF PHAGOCYTOSIS, OR ANY OTHER
267 PARAMETERS REFERRED TO IN THIS PAPER, WE RECOMMEND FIRST TO ASSESS THE BACTERIAL
268 OF THE INFECTED HOST. TO DO THIS, SUPERNATANT OF HAEMOLYMPH OR HOMOGENIZED HAEMOLYMPH
269 CAN BE SPOTTED IN APPROPRIATE PLATES AND COLONY FORMING UNITS COUNTED (SEE
270 2007).

271

272

273 *Capacity to attach to non-self surfaces (Nodule/Capsule Formation)*

274

275 QUITE FREQUENTLY, RESEARCHERS HAVE INTRODUCED PATHOGENS OR MICROBIAL CELL
276 COMPONENTS TO INSECT HOSTS TO ASSESS CELLULAR IMMUNE ~~RESPONSE~~ (E.G HOWARD
277 THIS INDUCES THE PRODUCTION OF NODULES/CAPSULES, WHICH ARE THEN COUNTED WHEN
278 HAEMOLYMPH IS EXTRACTED AFTER THE CHALLENGE OR DIRECTLY IN THE HAEMOCOELM
279 *et al.* 1996, GOLDSWORTHY ~~2007~~). NODULES CAN PERSIST IN INSECTS FOR LONG PERIODS OF TIME

280 (CARTON & NAPPI 1997), WHICH MEANS THAT THE HAEMOCYTES USED FOR NODULE FORM
281 REMAIN ATTACHED, THEREBY DECREASING HAEMOCYTE DENSITY DURING INSECT LIFET
282 REDUCTION IN HAEMOCYTE NUMBER COULD IN TURN LIMIT THE AVAILABILITY OF HAEM
283 BECOME RELEVANT DURING MOULTING AND/OR METAMORPHOSIS. NOT ALL ADULT INSE
284 HAEMOCYTES SINCE HAEMATOPOIETIC ORGANS ARE NOT ALWAYS PRESENT AT THIS STA
285 1998). IN LEPIDOPTERANS AND DIPTERANS, HEMATOPOIETIC ORGANS HAVE BEEN IDENTIF
286 EXCLUSIVELY IN THE EMBRYONIC, LARVAL AND PUPAL STAGES (GARDINER & STRAND 20
287 *al.* 2003, HOLZ *et al.* 2003, WILLIAMS 2007). THIS MEANS THAT ANY HAEMOCYTES PRESENT IN
288 THE ADULT STAGE WERE PRODUCED DURING EMBRYOGENESIS, LARVAL STAGES AND ME
289 HOWEVER, THE PRESENCE OF HEMATOPOIETIC ORGANS IN ADULT GRASSHOPPERS HAS BE
290 (HOFFMAN 1973, HOFFMAN *et al.* 1974). NATURAL HISTORY DIFFERENCES AMONG INSECT GROUP
291 (E.G. THE PRESENCE OF METAMORPHOSIS) LIKELY LEADS TO VARIATION IN HAEMOCYTE I
292 (CONSEQUENTLY CELL NUMBER).

293

294 *Methodology*

295 PATHOGENS SUCH AS BACTERIA, YEAST CELLS ARE INOCULATED DIRECTLY TO HAEMOCO
296 *al.* 1989). ALSO, INJECTED PARASITES CAN BE USED TO ASSESS NODULE/ENCAPSULATION F
297 (SEE ELEFThERIANOS 2008). HAEMOLYMPH IS COLLECTED AND INCUBATED IN HUMIDITY
298 CHAMBER, NODULES ARE OBSERVED USING A MICROSCOPE AND COUNTED. CARE MUST B
299 PATHOGENS CAN BE DESTROYED BEFORE NODULATION TAKES PLACE, FOR EXAMPLE, IF F
300 THE MIDGUT (LUCKHART 1998) *al.* FURTHERMORE, NODULES MUST BE COUNTED EVEN WHEN TI
301 SIZE VARIES, WHICH CAN OCCUR EVEN WITHIN THE SAME INDIVIDUAL (SEE HOWARD
302 ENUMERATION OF DIFFERENT SIZES CAN PROVIDE FURTHER INFORMATION ON DIFFEREN
303 ABILITY AMONG HOSTS. WE RECOMMEND TO DIRECTLY ASSESS THE SIZE OF NODULES EIT

304 QUALITATIVELY (BY CATEGORIZING THE DIFFERENT SIZES AND COUNTING HOW MANY OF
305 THE DISTINCT CATEGORIES) OR QUANTITATIVELY WITH THE USE OF IMAGE ANALYSIS SOFTWARE
306

307

308 *Melanotic nodulation/encapsulation (Nylon Monofilament and Beads Insertion)*

309

310 FOR ENCAPSULATION/MELANOTIC ENCAPSULATION ASSESSMENT, INOCULATION WITH SPERMATHECA
311 POLYSTYRENE BEADS OR INSERTION OF A NYLON MONOFILAMENT INTO THE HAEMOCOEL
312 USED. IN THIS METHOD, NYLON AND BEADS ARE THOUGHT TO MIMIC NATURAL INFECTION BY
313 A PARASITE/PARASITOID) WHILE AVOIDING THE DAMAGING EFFECTS OF PATHOGENS (E.G. WANG
314 2000, KOSKIMÄKI *et al.* 2004, SIMMONS *et al.* 2005, RANTALA & ROFF 2007). CELLULAR AND
315 HUMORAL RESPONSES ARE ACTIVATED, WHICH COULD LEAD TO TRADE-OFFS SIMILAR TO THOSE
316 FORMATION AND PHAGOCYTOSIS ARE ELICITED. IN THE MELANIN LAYER – WHICH IS FREQUENTLY
317 PRODUCED DURING NODULATION/ENCAPSULATION -- SYNTHESIS OF MELANIN IS REQUIRED
318 COSTLY TO PRODUCE, NOT ONLY BECAUSE OF THE RESOURCES USED (SEE *Phenoloxidase Activity*
319 BELOW), BUT BECAUSE MELANIN IS REQUIRED FOR CUTICLE FORMATION, HUMORAL TOXIN
320 PRODUCTION, AND COLOURING AND HARDENING OF OTHER STRUCTURAL TRAITS (E.G. WANG
321 SPERMATHECA; LI & CHRISTENSEN 1993, ILANGO 2005).

322

323 *Methodology*

324

325 AFTER INSERTION, FILAMENT/BEADS ARE THEN RETRIEVED AND THE VOLUME OF MELANIN
326 MELANIZED CELL MASSES THAT ENCAPSULATE THE FILAMENT ARE MEASURED (SIVA-JOYNT
327 *in vitro* ENCAPSULATION ASSAYS CAN BE ALSO BE PERFORMED. FOR THIS, HAEMOLYMPH IS COLLECTED
328 FROM EACH INDIVIDUAL AND MIXED WITH BEADS. ENCAPSULATION AND MELANIZATION

329 AFTER INCUBATION BY MICROSCOPY. THREE METHODS HAVE BEEN DEVELOPED FOR MEASURING
330 MASSES. ONE METHOD OBSERVES THE MEAN GRAY SCALE OF THE COVERED AREA USING
331 ANALYSIS SOFTWARE (SIMMONS) WHILE A SECOND USES IMAGES SOFTWARE FILTERS TO
332 MEASURE THE DENSITY OF THE COVERED AREA (KIM). ANOTHER APPROACH
333 MEASURES THE ENCAPSULATED/MELANIZED AREA (CONTRERAS & GARDUÑO *et al*)
334 MEASURES, HOWEVER, ARE UNABLE TO DIFFERENTIATE THE FUNCTIONAL ASPECTS OF HAEMOLYMPH
335 CELLULAR NODULATION/ENCAPSULATION AND HUMORAL MELANIZATION ARE TWO SEPARATE
336 PROCESSES (LING & YU 2006). HAEMOCYTES RELEASE OR CONTAIN SURFACE PROTEINS THAT
337 ACTIVATED TO PO, CAUSES MELANIZATION. THIS CELLULAR ENCAPSULATION ALSO ENHANCES
338 INTERACTIONS BETWEEN HAEMOLYMPH AND PO. LING & YU (2006) NOTICED THAT MELANIZATION
339 DID NOT OCCUR WHEN ISOLATED HAEMOCYTES WERE USED, IT OCCURRED ONLY IN THE PRESENCE OF
340 PLASMA. THIS SUGGESTS THAT FACTORS IN HAEMOLYMPH ARE REQUIRED FOR PO CASCAID AND
341 AND HENCE MELANIZATION. THEREFORE, CELLULAR ENCAPSULATION COULD BE IMPLICATED IN
342 CELLULAR AND HUMORAL IMMUNE RESPONSES.

343

344 TRADE-OFFS AMONG IMMUNITY AND OTHER TRAITS (E.G. DEVELOPMENTAL TIME) USING
345 HAVE BEEN FOUND (SEE RANTALA & ROLFF 2005). THE USE OF NON-NATURAL INFECTIONS CAN
346 BE A SUITABLE INDEX OF HOST IMMUNE ACTIVATION-RECOGNITION AND ITS ECOLOGICAL
347 RANTALA & ROLFF 2007). FOR THE METHODS ABOVE DISCUSSED, UNFORTUNATELY, THE DENSITY
348 OF MELANIZED/ENCAPSULATED AREAS IS OFTEN SO IRREGULAR THAT IT IMPEDES THOROUGH
349 (I.E. IT DEPENDS ON THE POSITION OF THE ARTIFICIAL OBJECT DURING AREA MEASUREMENTS).
350 RECOMMEND SEVERAL MEASURES OF THE SAME IMPLANT. ADDITIONALLY, INSERTING AN IMPLANT
351 A DIFFICULT TASK FOR SMALL INSECTS. IN THIS CASE, WE RECOMMEND THE USE OF INOCULATED
352 WE ALSO RECOMMEND USING LIVE PATHOGENS THAT INDUCE MELANOTIC ENCAPSULATION AND
353 IMMUNE RESPONSES. THIS CAN BE USEFUL, FOR EXAMPLE, WHEN TESTING FOR IMMUNE T

354 AND BEHAVIOUR (SEE AYRES & SCHNEIDER 2009). TOLERANCE THEORY PREDICTS A POSSI
355 INCREASE IN THE PATHOGEN BURDEN, BUT THE IMMUNE RESPONSE WILL LIMIT THE HEAL
356 CONSEQUENCES OF THIS PATHOGEN BURDEN (SCHNEIDER & AYRES 2008, READ 2008).
357 ONLY LIVE PATHOGENS CAN REPLICATE AND CONTINUOUSLY DAMAGE A HOST. IT IS ALS
358 LIVE PATHOGENS ACTIVATE MORE IMMUNE PATHWAYS (COMPARED WITH INERT IMPLAN
359 THEREFORE GIVE A MORE COMPLETE PICTURE OF AN IMMUNE RESPONSE. HOWEVER, WHI
360 LIVE PATHOGENS, IT IS IMPORTANT TO FIRST ESTIMATE THE DOSAGE OF PATHOGEN NEED
361 MEASUREMENT OF A PARTICULAR IMMUNE RESPONSE. IT HAS BEEN OBSERVED THAT IN S
362 LOW TO MEDIUM DOSES ELICIT PHAGOCYTOSIS, MODERATE TO HIGH DOSES ELICIT NODU
363 OVERDOSES CAN PROMOTE SEPTIC SHOCK (SALVADOR HERNÁNDEZ-MARTÍNEZ, *personal*
364 *communication*). BEFORE A CHALLENGE, WE RECOMMEND TO DISCOVER THE IMMUNE RESP
365 THAT IS ELICITED IN RELATION TO THE TYPE AND DOSE OF INFECTIOUS AGENT (NATURAL
366 MODEL.

367

368

369 *Humoral responses*

370

371 *Phenoloxidase Activity*

372

373 PO IS THE MOST WIDELY USED IMMUNE MARKER IN ECOLOGICAL IMMUNOLOGY STUDIES
374 GOLDSWORTHY 2003, WILSON *et al.* 2003, JACOB *et al.* 2005). THE PO ENZYME IS USED
375 FOR DIFFERENT PHYSIOLOGICAL PROCESSES (WOUNDING, CLOTTING, CUTICLE COMPOSIT
376 ENCAPSULATION, PRODUCTION OF CYTOTOXIC MOLECULES), AND PROBABLY SPERMATH
377 (ILANGO 2005). DUE TO THESE DIFFERENT FUNCTIONS, PO IS CONSTANTLY SYNTHESIZED,
378 COULD BE COSTLY TO PRODUCE FOR THE ORGANISM. RECALL THAT PROPO MOLECULES A

379 MAINLY BY HAEMOCYTES (OTHER TISSUES, SUCH AS FAT BODY AND MIDGUT EPITHELIUM
380 PRODUCE PROPO IN SMALLER QUANTITIES), AND THAT SOME OF THESE MOLECULES REMA
381 TO HAEMOCYTES, WHILE SOME PROPO MIGHT END UP IN SEVERAL OTHER TISSUES (I.E. CU
382 MIDGUT, SALIVARY GLANDS, EGGS; ASHIDA & BREY 1997). IF PATHOGENS ARE DETECTED
383 PROPO TRANSPORTATION FROM HAEMOCYTES TO OTHER TISSUES, IT IS POSSIBLE THAT T
384 ACTIVATED IN THE HAEMOCELE FOR CLEARING PATHOGENS, CONSEQUENTLY LEAVING T
385 WITH A PROPO DEFICIENCY. THIS MEANS THAT A TRADE-OFF MAY ARISE BECAUSE OF A P
386 DEFICIENCY (BUT NOT OF OTHER COMPONENTS, SUCH AS PO SUBSTRATES; SEE BELOW), A
387 ASSUMED IN GENERAL STUDIES WHERE PO AND MELANIN HAVE BEEN INVOLVED (SIVA-JO
388 THIS DIFFERENT PERSPECTIVE OF A POSSIBLE TRADE-OFF WHEN MEASURING PO ACTIVITY
389 BEEN PUT FORWARD NOR EXAMINED PREVIOUSLY AND NEEDS FURTHER EMPIRICAL STUD
390
391 THE PO SUBSTRATE TYROSINE (TYR) AND TYROSINE SUBPRODUCTS ARE USED FOR THE EX
392 AND MANUFACTURE OF OTHER TRAITS. TYR IS OBTAINED THROUGH FOOD AND IS NATUR
393 SYNTHESIZED BY ORGANISMS OR BY THE HYDROXYLATION OF PHENYLALANINE (PHE) (C
394 *al.* 2005). SINCE PHE IS AN ESSENTIAL AMINO ACID FOR PROTEIN SYNTHESIS, ITS IMPORTAN
395 IN THE COST OF ACQUIRING THIS RESOURCE. TYR IS ALSO INVOLVED IN A VARIETY OF FU
396 OF THESE, FOR EXAMPLE, IS THE COLOURING OF THE EGG CHORION IN SOME INSECTS (LI &
397 CHRISTENSEN 1993). IN MOSQUITOES, MELANIZATION RESPONSES AGAINST WORMS GENER
398 DELAY IN TYR ACCUMULATION IN THE OVARIES, AND THEREFORE A DECREASE IN THE NU
399 PRODUCED AFTER AN IMMUNE CHALLENGE (LI & CHRISTENSEN, 1993) AND A DELAY IN OV
400 (FERDIO *al.* 1993). EXAMPLES LIKE THIS ARE SUGGESTIVE OF SUBSTRATES OF PO AND MEL
401 PHE AND TYR - BEING RESTRICTIVE RESOURCES THAT MAY LEAD TO TRADE-OFFS BETWEEN
402 IMMUNE RESPONSE AND OTHER KEY FUNCTIONS INCLUDING MOULTING, BASAL METABO
403 PROTEIN SYNTHESIS. ALTHOUGH TYR CAN BE STORED (AS TYROSINE-O-PHOSPHATE, BET

404 TYROSINE, TYROSINE GLUCOSIDE; MITCHELL & LUNAN 1964, LEVENBOOK *et al.* 1969, CHEN
405 1978), TRADE-OFFS COULD BE DETECTED IF NOT ENOUGH TYR IS GATHERED IN ORDER TO
406 AMINO ACID TO ALL THE METABOLIC REQUIREMENTS THROUGH THE INSECT LIFETIME. H
407 INSECTS NECESSARILY USE TYR OR PHE IN THE SAME WAY, PO MEASURES NEED TO BE CA
408 INTERPRETED IN EVERY INSECT MODEL. MELANIN, THE FINAL PRODUCT OF THE PO CASCA
409 INVOLVED IN OTHER MORPHOLOGICAL TRAITS SUCH AS CUTICLE FORMATION AND PIGME
410 MAY AFFECT FUNCTIONS RELATED TO APOSEMATISM AND CRYPISIS, THERMOREGULATION
411 UV RADIATION AND COLOURED SEXUAL TRAITS. MANY OF THESE FUNCTIONS AND THEIR
412 TRADE-OFFS HAVE NOT BEEN ADEQUATELY INVESTIGATED.

413

414 *Methodology*

415

416 PO ACTIVITY IS ASSAYED BY THAWING HAEMOLYMPH AND *o*-DIPHENOLS (DOPA OR DOPA
417 SUBSTRATE INTO A MICROPLATE WELL. THIS ASSAY WILL NOT DISTINGUISH THE THREE D
418 ACTIVITIES, SINCE IT IS INFERRABLE. PHENOL PEROXIDASE ACTIVITY OF ALL THREE ENZYMES W
419 BE DETECTED (SUGUMARAN & KANOST 1993). CARE MUST BE TAKEN WHEN USING
420 SINCE THERE COULD ALSO BE A PEROXIDASE-MEDIATED MELANIN FORMATION (CHRISTE
421 2005), WHICH CAN RESULT IN AN OVERESTIMATION OF PO ACTIVITY. WE RECOMMEND TH
422 PEROXIDASE SUPPRESSOR (E.G. HYDROGEN PEROXIDE IN METHANOL OR COMMERCIALY
423 ANOTHER CURRENT PROBLEM WITH ASSESSING PO IS THAT PROTEIN LOAD STANDARDIZA
424 BEFORE MEASUREMENT, USING A STANDARD PROTEIN ASSAY (CONTRERAS-GARDUÑO *et al.*
425 PRACTICE THAT IS RARELY CARRIED OUT AND WITHOUT WHICH MAY RESULT IN DISPARA
426 PO READINGS. SINCE OCCASIONALLY REAL PROTEIN LOAD CAN BE MASKED IF DIFFERENC
427 INDIVIDUALS, (E.G. BIGGER SIZE OF FEMALES THAN MALES; INACCURATE EXTRACT OF TO
428 HAEMOLYMPH); WE RECOMMEND THE PROTEIN LOAD PRACTICE MENTIONED ABOVE. FUR

429 THERE MIGHT BE A RAPID DEGRADATION OF PO, IF STORED FOR RELATIVELY LONG TIME.
430 BE RESOLVED BY USING A PROTEASE INHIBITOR COCKTAIL (E.G. PMSF, LEUPEPTIN). ANOT
431 PROBLEM IS THAT SMALL-SIZED INSECTS MAY NOT PROVIDE PO READINGS, SINCE PO QU
432 SO SMALL. THIS PROBLEM CAN BE SOLVED BY USING MORE THAN ONE INDIVIDUAL FOR A
433 SAMPLE TO OBTAIN PO READINGS. AS WELL, THE TOTAL PO ACTIVITY CAN BE ESTIMATED
434 ENZYME CHYMOTRYPSIN TO ACTIVATE ALL PRO-PO PRESENT IN THE HAEMOLYMPH (SEE I
435 ZUK 2010).

436

437

438 *Nitric Oxide*

439

440 NO QUANTIFICATION IS A RELATIVELY NEW IMMUNE MARKER IN ECOLOGICAL IMMUNITY
441 FOR EXAMPLE, MORENO-GARCÍA *et al.* (2010) NO DAMAGES PATHOGENS AND IS ALSO USED AS A
442 SIGNALLING MOLECULE IN *Drosophila*, NO IS INDISPENSABLE FOR ACTIVATION OF THE IMMUNE
443 DEFICIENCY (IMD) PATHWAY (FOLEY & O'FARRELL 2003). THIS IS ONE OF THE THREE PATHW
444 (THE OTHER TWO BEING TOLL AND JAK/STAT SEE FERRANDON & PERRIOMON 2004) TO PRODUCE
445 TRANSCRIPTIONAL FACTORS IN THE FAT BODY, LEADING TO THE SYNTHESIS OF ANTIMICR
446 AND OTHER FACTORS THAT PREVENT SELF TISSUE DAMAGE AND CONTROL HAEMOCYTE
447 DIFFERENTIATION (FOLEY & O'FARRELL 2003; AGAISSE & PERRIOMON 2004). ANOTHER INT
448 POINT IS THE FACT THAT THE NO SYNTHASE (NOS) (AN ENZYME THAT CONVERTS L-ARGIN
449 CITRULLINE AND GENERATES NO) IS CONSTITUTIVE AND INDUCIBLE (MÜLLER 1997). NO IS
450 SIGNALLING MOLECULE, SO A CONSTANT PRODUCTION OF NOS IS NECESSARY FOR HOME
451 PURPOSES, WHILE INDUCIBLE NOS IS ONLY SYNTHESIZED AFTER AN IMMUNE CHALLENGE
452 *al.* 2000). IT IS THEREFORE EXPECTED THAT NO PRODUCTION INCREASES AFTER AN IMMUN
453 CHALLENGE. NEVERTHELESS, KRISHNANA, HYRŠL & ŠIMEK (2006) FOUND THAT NO PRODU

454 SIMILAR IN BOTH NON-STIMULATED AND STIMULATED HAEMOCYTES IN LEPIDOPTERAN L
455 PARTICULAR CASE, THE AUTHORS INDICATE THAT NO CAN BE CONTINUOUSLY SYNTHESIZED
456 INDUCED NOS FOR LONG PERIODS OF TIME (HOURS TO DAYS). THIS MEANS THAT THE EFFICIENCY
457 FOR KILLING PATHOGENS WITH NO MAY NOT ONLY RESIDE ON NO CONCENTRATION, BUT ALSO ON THE
458 DURATION OF NO SYNTHESIS (SEE LAURENT *et al*

459

460 NO IS PRODUCED DURING THE OXIDATION OF L-ARGININE (MÜLLER 1997). ARGININE IS AN AMINO
461 ACID, WHICH MUST BE OBTAINED FROM THE DIET (RIVERO 2006). ARGININE IS ALSO IMPORTANT FOR
462 SPERM MATURATION (OSANAI & CHEN 1993), EGG PRODUCTION (UCHIDA 1993), LONG TERM
463 MEMORY, CHEMOSENSORY (ANTENNAL LOBES, OLFACTION), AND VISUAL INFORMATION PROCESSING
464 (MÜLLER 1997). TRADE-OFFS ARE THUS EXPECTED AMONG THESE TRAITS AND THE IMMUNE

465

466 *Methodology*

467

468 BASAL LEVELS OF NO ARE NORMALLY USED, WHICH PROVIDES AN INCOMPLETE PICTURE OF
469 IMMUNE ABILITY. A BETTER ALTERNATIVE IS TO ASSESS NO PRODUCTION AFTER HOST INGESTION
470 OF PATHOGENS. THE STANDARD METHOD TO MEASURE NO IS TO USE THE GRINDING METHOD
471 THAT MEASURES NITRITE PRODUCTION – AN INDIRECT MEASURE OF NO (BREDT & SNYDER 1997).
472 USE OF CONTROLS SUCH AS AN L-NAME (A NOS-INHIBITORY ARGININE ANALOGUE; SEE RIVERO
473 2006) AND ITS INACTIVE ENANTIOMER D-NAME IS RECOMMENDED. CONTROLS ARE NEEDED
474 BECAUSE NITRITES CAN ALSO BE SUB-PRODUCTS OF OTHER REACTIONS NOT RELATED WITH NO
475 FOR THIS METHOD, HOWEVER, IT IS UNKNOWN WHETHER HAEMOLYMPH PROTEIN LOAD AFFECTS
476 STANDARDIZATION IS NEEDED BEFORE MEASUREMENT. THIS IS A TECHNICAL PROBLEM THAT
477 RESOLVED IN THE NEAR FUTURE. SIMILAR TO PO, MORE THAN A SINGLE INDIVIDUAL MAY BE
478 AN INDIVIDUAL HOST SAMPLE IS TOO SMALL.

479

480

481 *Lysozyme Activity and Antimicrobial Peptides*

482

483 LYSOZYME IS AN ENZYME WITH HYDROLYTIC ACTION MAINLY AGAINST THE PEPTIDOGL

484 GRAM CELL WALLS. IT CAN BE INDUCED, OR CONSTITUTIVELY EXPRESSED IN THE GUT TR

485 *al.* 1997), HAEMOCYTES AND FAT BODY (GILLISPIE AMPS SHOW GREAT STRUCTURAL486 DIVERSITY (MORE THAN 170 ISOFORMS HAVE BEEN FOUND IN INSECTS BARLET *et al*

487 PRODUCED SOON AFTER FOREIGN RECOGNITION (1-4 HOURS) WITH AN EFFICIENT PATHOC

488 KILLING ACTION (SCHMID-HEMPEL 2005B). SOME AMPS MAY REMAIN IN HAEMOLYMPH FO

489 TO THREE WEEKS (SCHMID-HEMPEL 2005B), WHICH IS CONVENIENT DURING SUBSEQUENT

490 ENCOUNTERS. AMPS ARE GENERALLY SHORT PEPTIDES, CONTAINING FEWER THAN 150–20

491 ACIDS (BULET. 2004), SO THEY ARE CONSIDERED ENERGY EFFICIENT, QUICK AND ECONOM

492 PRODUCE (OTVOS JR 2000). HOWEVER, THE OVERALL AMPS CONCENTRATION IN *Drosophila*493 HAEMOLYMPH CAN REACH 200 μ M (OTVOS JR 2000). AT THIS CONCENTRATION, ANTIMICRO

494 PEPTIDES DO SEEM COSTLY TO PRODUCE. IT IS POSSIBLE THAT RESOURCES (PROTEINS) GA

495 THROUGH ONTOGENY ARE INDISPENSABLE FOR ANTIMICROBIAL PEPTIDE GENERATION A

496 SYNTHESIS OF OTHER MOLECULES. THIS FACT COULD POTENTIALLY LEAD TO TRADE-OFF

497 IMMUNITY AND SYNTHESIS OF OTHER PRODUCTS (E.G. SPERMATOPHALYX PEPTIDES) AND

498 CONFORMATION.

499

500 *Methodology*

501

502 LYSOZYME ACTIVITY IS COMMONLY ASSAYED VIA THE CLEARANCE RATE OF BACTERIAL

503 USING HAEMOLYMPH. THE METHOD IS RELATIVELY SIMPLE: AFTER THE HAEMOLYMPH IS

504 AND MIXED WITH A BACTERIAL SOLUTION, A TURBIDITY ASSAY IS THEN PERFORMED (E.C
505 2004). SMALL ABSORBANCE VALUES INDICATE HIGH LYTIC ACTIVITY. ANOTHER WAY IS TO
506 DROPS OF HAEMOLYMPH TO BACTERIAL CULTURE, MEASURING THE AREA OF LYTIC ACTI
507 TIME (12 HOURS). THE LYTIC ACTIVITY APPEARS AS CLEAR CIRCULAR ZONES IN THE CULT
508 INDICATES THAT BACTERIA HAVE BEEN CLEARED; THE DIAMETER OF THE HOLE IS MEASU
509 AS THE INDICATOR OF LYTIC EFFECTIVENESS.

510

511 AMPS HAVE BEEN MEASURED BY INJECTION OR INGESTION OF COMPLETE BACTERIA, FUN
512 OR THEIR FRACTIONS (E.G. LPS, PEPTIDOGLYCANE). AMPS ACTIVITY IS ASSESSED IN THREE
513 MEASUREMENT OF THE CLEARANCE RATE OF A BACTERIAL SUSPENSION (VIA A TURBIDIT
514 TO THE LYSOZYME PROTOCOL DESCRIBED ABOVE); 2) MEASUREMENT OF THE RATE OF BA
515 GROWTH ON A PLATE, AFTER THE SAME BACTERIA HAVE BEEN INJECTED INTO THE HOST
516 HAEMOLYMPH HAS BEEN EXTRACTED AFTER SOME TIME; AND, 3) THE ANTIBACTERIAL ZONE
517 MEASUREMENT OF THE HOLE AREA, SIMILAR TO THE LYSOZYME PROTOCOL DESCRIBED A
518 PROBLEM WITH THESE DIFFERENT ASSAYS IS THAT ANTIMICROBIAL ACTIVITY CAN BE DU
519 MOLECULES AND NOT NECESSARILY AMPS. SINCE LYSOZYME NEEDS A PH 8 TO 6.5 FOR OP
520 ACTIVITY THIS CAN BE USED TO DISTINGUISH BETWEEN OTHER AMP ACTIVITY (RANTALA
521 2004, AHTIANEN ET AL 2005, BUT SEE DA SILVA ET AL 2000, ADAM & PARSOSN 2006).

522

523

524 **Some concluding remarks**

525

526 ONE INTENTION OF THIS REVIEW IS TO HIGHLIGHT THE ECOLOGICAL AND PHYSIOLOGICAL
527 INSECT IMMUNE MARKERS AND THEIR VALUE FOR ESTIMATING TRADE-OFFS BETWEEN IM
528 OTHER TRAITS. THUS, CURRENT (AND NEW) KNOWLEDGE OF IMMUNE PHYSIOLOGICAL M

529 MUST BE INCORPORATED INTO ECOLOGICAL DATA TO ALLOW FOR A DEEPER UNDERSTAN
530 REASONS FOR EVOLUTIONARY CORRELATIONS AND CONSTRAINTS. THE USE OF EACH ME
531 HERE MUST BE DETERMINED BY EACH RESEARCHER IN TERMS OF ITS OVERALL COST (IN T
532 CONSUMPTION OR PRICE), DIFFERENCES IN THE KIND OF PATHOGENS OR ELICITORS USED
533 IMMUNE RESPONSE, AND THE IMMUNE MARKER THAT CAN ADDRESS QUESTIONS VIA AN A
534 EXPERIMENTAL DESIGN. ANOTHER IMPORTANT CONSIDERATION FOR EACH RESEARCHER
535 TO MAKE REPEATED IMMUNE MEASUREMENTS ON THE SAME SAMPLE. UNFORTUNATELY
536 MEASUREMENT METHODS REFERRED HERE ARE NOT SUITABLE TO BE REPEATED MORE TH
537 EXAMPLE, PO AND NO CONTAINED IN HAEMOLYMPH EXTRACTION GRADUALLY DEGRADE
538 IN AN ULTRAFREEZER. ALSO, UNFREEZING AND RE-FREEZING SAMPLES ACCELERATE
539 DEGRADATION. OCCASIONALLY, HAEMOCYTE MEMBRANES CAN BECOME DISRUPTED EVI
540 SOLUTION OR CULTURE MEDIUM IF STORED FOR LONG PERIODS OF TIME (E.G. WEEKS TO M
541 RECOMMEND EXTRACTING ENOUGH SAMPLE TO DO ALL MEASUREMENTS REQUIRED FOR
542 EXPERIMENTAL DESIGN. IF A LARGE NUMBER OF INDIVIDUALS ARE REQUIRED, THEY (OR
543 CAN BE STORED AND MEASUREMENTS SHOULD BE ONLY DONE FOR THE EXACT NUMBER
544 CAN BE PROCESSED AND QUANTIFIED IN A SINGLE DAY.

545

546 FINALLY, THERE ARE TWO RELATED POINTS THAT WE WOULD LIKE TO HIGHLIGHT: (1) TH
547 IMMUNE MARKERS MENTIONED HERE ARE USUALLY SEEN AS RESPONSES TO ATTACK AN
548 PATHOGENS; AND (2) THE CORRELATIONS AMONG IMMUNE RESPONSES. AS MENTIONED B
549 NEGATIVE AND POSITIVE CORRELATIONS HAVE BEEN FOUND AMONG IMMUNE TRAITS, BU
550 SEEM TO BE HOST-SPECIFIC. COMMONLY, RESOURCES LIMITATION HAS BEEN USED TO EX
551 NEGATIVE CORRELATIONS, HOWEVER, THE NON-EXPRESSION OF AN IMMUNE EFFECTOR T
552 NECESSARILY MEANS THAT THERE IS NO RESPONSE. REACTIONS AND PRODUCTS OF IMM
553 ARE INTERCONNECTED AND THE KIND OF RESPONSE IS RELATED TO THE PATHOGENS VIR

554 MENTIONED BEFORE, NOT ALL PATHOGENS WILL INDUCE THE SAME REACTION, OR AT LE
555 MAGNITUDE OF RESPONSE CAN DIFFER DEPENDING ON THE HOST-PATHOGEN INTERACTIO
556 OCCASIONALLY TOLERANCE COULD BE THE BEST STRATEGY TO COPE WITH INFECTIONS.
557 OR TOLERANCE STRATEGIES ADOPTED BY AN INDIVIDUAL WILL DEPEND ON THE FORCE A
558 INFECTION. THERE COULD BE A CRITICAL HOST DAMAGE THRESHOLD, THUS WHEN THE IN
559 INFECTION IS BELOW THIS THRESHOLD (I.E. NO SIGNIFICANT DAMAGE TO THE HOST), THE
560 TOLERATE. ABOVE THE THRESHOLD, IT THEN PAYS TO ALLOCATE MORE INTO RESIST THE
561 EXAMPLE, PHAGOCYTIC ACTIVITY COULD BE TURNED ON, MEANWHILE PO ACTIVITY COU
562 DOWN REGULATED; IN THIS CASE PROBABLY THE HOST IS AVOIDING AUTOREACTIVITY M
563 (GENERATED THROUGH THE PO CASCADE). IN THIS EXAMPLE, THE IMMUNE STRATEGY IS
564 ORDER TO LIMIT PATHOGEN AND SELF DAMAGE, NEVERTHELESS THE PATHOGEN IS NOT E
565 SINCE IMMUNE SYSTEM IS HIGHLY COMPLEX, SEVERAL ASPECTS OF IMMUNE RESPONSE M
566 THEREFORE MEASURED TO COMPREHEND THE INSECT IMMUNITY STRATEGIES. ALSO, OCC
567 SOME IMMUNE RESPONSES ARE NOT SUITABLE TO TEST SOME HYPOTHESIS, FOR EXAMPL
568 CASCADE AND PO ACTIVITY OCCASIONALLY IS NOT AFFECTED BY THE NUTRITIONAL CON
569 RESPONSES, SUCH AS LYSOZYME, ARE AFFECTED (SEE JACOT *et al.* 2008, MORENO-GARCÍA
570 2010). THEREFORE, IT IS RECOMMENDED TO USE MORE THAN ONE IMMUNE MARKER WHEN
571 POSSIBLE. FINALLY, THE STARTING LINE IN INSECT IMMUNOECOLOGY IS THAT IT MUST IN
572 GREAT RANGE OF ENVIRONMENTAL FACTORS INVOLVED USING A TRADE-OFF FRAMEWO
573 ALLOCATION AND INVESTMENT STRATEGIES SHOULD BE STUDIED BY TAKING INTO ACCO
574 THE REAL COST IN NATURAL CONDITION AND THEIR EFFECTS IN THE EXPRESSION OF IMM
575 AND GENERATIONAL CHANGES.

576

577 IDEALLY, AN ECOLOGICAL STUDY OF INSECT IMMUNITY SHOULD START WITH THOUGHTF
578 REGARDING APPROPRIATE IMMUNE MARKERS. REGARDING METHODOLOGICAL ISSUES, IN

579 METHODOLOGICAL INFORMATION IS AVAILABLE BUT IT IS POSSIBLE THAT THE *status quo* C
580 EVOLUTIONARY ECOLOGY RESEARCHERS HAVE USED SUCH MARKERS LEADS OTHERS NO
581 ABOUT THE BEST WAY TO PROCEED. LASTLY, WE ENCOURAGE ECOLOGICAL AND EVOLUT
582 BIOLOGISTS ALIKE TO INCORPORATE USEFUL MOLECULAR TECHNIQUES (E.G. MICROASSA
583 QUANTITATIVE REVERSE TRANSCRIPTASE-PCR, ETC.) TO IDENTIFY THE POSSIBLE MECHAN
584 THE GENETIC BASIS UNDERLYING THE TRADE-OFFS BETWEEN IMMUNITY AND LIFE HISTO
585 HOPED THAT THIS INFORMATION PROVES CONSTRUCTIVE IN PAVING THE WAY TO A MOR
586 OF ECOLOGICAL IMMUNOLOGY.

587

588

589 **ACKNOWLEDGEMENTS**

590 WE THANK SALVADOR HERNÁNDEZ-MARTÍNEZ FOR PROVIDING ASSISTANCE WITH PROTO
591 HELPFUL SUGGESTIONS. WE THANK GLORIA TAVERA AND CHRIS ANDERSON FOR REVISIN
592 IN ENGLISH, AND PROVIDING THOUGHTFUL COMMENTS. MM-G WAS SUPPORTED BY THE F
593 EN CIENCIAS BIOMÉDICAS (CONACYT: GRANT NO. 172947), INSTITUTO DE ECOLOGÍA,
594 (UNIVERSIDAD NACIONAL AUTÓNOMA DE MÉXICO), AND THE CENTRO DE INVESTIGACIÓN
595 ENFERMEDADES INFECCIOSAS, INSTITUTO NACIONAL DE SALUD PÚBLICA, MÉXICO. AC-A V
596 SUPPORTED BY A GRANT PROVIDED BY PAPIIT-UNAM (PROJECT NO. 211506).

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For Review Only

TABLE 1. IMMUNE MARKERS USED IN INSECT ECOLOGICAL IMMUNITY STUDIES (SEE TEXT FOR FULL DISCUSSION)

IMMUNE MARKER	IMMUNE FUNCTION	METHODOLOGY	RECOMMENDATIONS	EXAMPLE REFERENCES
CUTICLE	ONE OF THE FIRST LINES OF DEFENSE AGAINST PATHOGENS; IT SENSES AND THE PRESENCE OF BACTERIAL FUNGI. USED FOR STRUCTURAL BARRIER QUANTIFICATION.	STRUCTURAL COLOR IS ASSESSED UNDER FLUORESCENT WHITE LIGHT. COLOURS ARE ANALYZED THROUGH IMAGES USING A GRAYSCALE VALUE.	CUTICULAR TRANSVERSE SECTIONS (PREPARATIONS (USING MICROTOME) FROM LEGS, THORAX AND ABDOMEN COULD DETERMINE CUTICLE RESISTANCE.	BARNES & SIVA-JOTHY 2000 THOMPSON <i>et al.</i> 2002 KORHONEN <i>et al.</i> 2005
HAEMOCYTE DENSITY	HAEMOCYTES MEDIATE PHAGOCYTOSIS, NODULATION, ENCAPSULATION/MELANIZATION AND RECOGNITION OF FOREIGN BODIES FOR CELLULAR IMMUNE RESPONSE QUANTIFICATION.	HAEMOCYTES EXTRACTION FOR COUNTING. HAEMOCYTES (SOME MORE COUNTED USING STAINED SUBSTRATE CHAMBER.	ADHESIVE PROPERTIES OF HAEMOCYTES ARE REDUCED BY ANTICOAGULANT BUFFERS. ACCURATE EXTRACTION OF HAEMOCYTES (AS MOST OF THEM REMAIN ATTACHED TO TISSUES), AN INJECTED PROTEASE INHIBITOR BLEND (E.G. PMSF, TLCK, LEUPEPTINE, EDTA) COULD BE USED.	RANTALA <i>et al.</i> 2000 USMANI & LOWE <i>et al.</i> 2001 FOECHT <i>et al.</i> 1994
PHAGOCYTOSIS	THE HAEMOCYTE PROCESSING AND KILL MICROBIAL PATHOGENS USED FOR CELLULAR IMMUNE RESPONSE QUANTIFICATION.	IN VITRO PHAGOCYTOSIS. FLUORESCENT BEADS, BACTERIA OR YEASTS ARE ATTACHED TO HAEMOCYTES <i>in vitro</i> . LABELED CELLS OR BEADS THEN QUENCHED WITH TRYPAN BLUE. PHAGOCYTE CELLS OR BEADS REMAIN FLUORESCENT. INGESTED CELLS ARE COUNTED MANUALLY OR USING IMAGE ANALYSIS SOFTWARE. PHAGOCYTTIC ACTIVITY IS GIVEN AS THE NUMBER/PROPORTION OF CELLS OR BEADS INGESTED PER μ L HAEMOLYMPH	USE OF SMALL PATHOGENS (BACTERIA, YEASTS) TO RECOMMENDED. PATHOGENS ARE ATTACHED TO HAEMOCYTES. INDIVIDUAL SIZE MAY INDUCE HIGHEST NODULE FORMATION AND IMMUNE RESPONSE. BACTERIAL AGGREGATIONS ALSO INDUCE SOME NODULE FORMATION. ASSESSMENT OF CELL STABILITY USING TRYPAN BLUE IS ALSO RECOMMENDED.	MAVROUDELIS <i>et al.</i> 1996 HINSWARD <i>et al.</i> 1998 MAVROUDELIS <i>et al.</i> 2005
NODULE/CAPSULE FORMATION	THIS RESULTS WHEN MULTIPLE HAEMOCYTES ATTACH TO AGGREGATIONS OF PATHOGENS. USED FOR CELLULAR IMMUNE RESPONSE QUANTIFICATION.	AN INJECTION OF PATHOGENS* (BACTERIA, PEPTIDOGLYCAN) IS NECESSARY FOR SOME TIME, HEMOLYMPH EXTRACTED OR HAEMOCYTES EXTRACTED AND THE NUMBER OF NODULES IS COUNTED.	CATEGORICAL MEASURES (SMALL, MEDIUM, LARGE), OR QUANTITATIVE MEASURES (WITH STAGE AND OCCLUSION MICROMETERS), IN ADDITION THE NUMBER OF NODULES IS COUNTABLE FOR NODULE QUANTIFICATION.	ALIBROOKMAN <i>et al.</i> 1989 KURTZ & SAUER 1999 KURTZ & SAUER 2008

- NYLON MONOFILAMENTS MIMIC PARASITE/PARASITOID INJECTIONS MANUALLY INSERTED THROUGH ABDOMINAL PLEURA OR SOFT BODY PART. QUALITATIVE MEASUREMENTS OF MELANIZATION QUANTIFICATION PRESENCE/ABSENCE OF VISIBLE MELANIZATION QUANTITATIVE MEASUREMENTS ARE NOT RECOMMENDED FOR SMALL INSECTS. MELANIN COVERAGE OR INTENSITY PERCENTAGES; MEAN GRAY SCALES USING IMAGE ANALYSIS SOFTWARE; DENSITOMETRY (USING RED+GREEN+BLUE FILTERS). (SIVA-JOTHY 2000; KOSKISAKI *et al.* 2004; SIMMONS *et al.* 2005)
- NYLON/SILICA BEADS MIMIC SMALL PATHOGEN INJECTION WITH MICROSYRINGE. THE USE OF DEAD, MEDIUM-SIZED PATHOGENS (E.G. YEAST) COULD BE USEFUL SINCE THEY ALSO INDUCE MELANIZATION AND ARE NATURALLY RECOGNIZED BY THE IMMUNE SYSTEM. (DADAMO 1999; SAMHOUCHE & KOELLA 2004)
- PHENOLOXIDASE (PO) A CONSTITUTIVE OXIDOREDUCTASE IN HAEMOLYMPH EXTRACTION. THE USE OF A PROTEASE INHIBITOR DURING THE EARLY STAGES OF MELANIN FORMATION; HIGHLY REACTIVE AND TOXIC INTERMEDIATE MOLECULES ARE PRODUCED. USE OF MICROPLATE WELL. SEVERAL REACTIVE SUPPRESSORS MAY INHIBIT HUMORAL RESPONSE QUANTIFICATION TAKEN IN A MICROPLATE READER FOR VARIOUS PERIODS (20-120MIN). HEMOLYMPH GRADATION/SPONTANEOUS ACTIVATION BEFORE MEASUREMENT (E.G. STANDARD PROTEIN ASSAY) TRYPSIN CAN BE USE TO ACTIVATE ALL PRO-PO PRESENT IN THE HAEMOLYMPH. (ROLDSWORTHY 2003; WILSON *et al.* 2003; WILSON 2005)

NITRIC OXIDE (NO) AN INDUCIBLE REACTIVE OXYGEN SPECIES PRODUCED BY THE IMMUNE SYSTEM DESTROYING INFECTIOUS AGENTS AND PROMOTING HOST DAMAGE. *Faraldo et al. 2005*

INDIRECT QUANTIFICATION USING ASSOCIATION WITH DEAD PATHOGENS. *Wernke & Fortiz 2004*

MOLECULE WITH AN IMPORTANT PROTECTION VIA THE GRIESS REACTION. *Faraldo et al. 2005*

DESTROYING INFECTIOUS AGENTS AND PROMOTING HOST DAMAGE. *Faraldo et al. 2005*

FOR HUMORAL RESPONSE NAPHTHYLETHYLENEDIAMINE ARE QUANTIFICATION. IN SOME CASES TRANSFERRED INTO A MICROPLATE WELL. GENERATION IS ACTIVATED BY ABSORBANCE IS RECORDED AT 540NM. INGESTION/INJECTION OF BACTERIA FROM COMMERCIAL KITS ARE AVAILABLE FOR CELL WALL COMPONENTS, E.G. LPS, RATE AND NITRITE MEASUREMENT. PEPTIDOGLYCAN), VIRUS, FUNGUS. THE USE OF L-NAME NOS INHIBITOR PARASITE INOCULATION. AS A CONTROL VERIFIES THAT QUANTIFIED NITRITES ARE A SUB PRODUCT OF ARGININE DEGRADATION BY NOS.

LYSOZYME ACTIVITY A CONSTITUTIVE AND INDUCIBLE ENZYME. THE CLEARANCE RATE OF BACTERIAL SUSPENSION USING SOLUTIONS IS AN IMPORTANT FACTOR IN THE ACTION AGAINST THE PEPTIDOGLYCAN. HAEMOLYMPH. EXTRACTED PH IS COMMONLY USED TO DISCRIMINATE AMONG LYTIC ACTIVITY. USED FOR HUMORAL RESPONSES. HAEMOLYMPH IS MIXED WITH AAMONG LYTIC ACTIVITY QUANTIFICATION. BACTERIAL SOLUTION AND CONTINUOUSLY ANTIMICROBIAL (PEPTIDE) MEASURED AT 492NM FOR A PERIOD OF TIME. SMALL ABSORBANCE VALUES INDICATE HIGH LYTIC ACTIVITY. IN THE LYTIC ZONE ASSAY, IN WHICH A BACTERIAL CULTURE IN SOLID AGAR MEDIA IS PINCHED, HOLES ARE FILLED WITH HAEMOLYMPH SAMPLE. DIAMETER OF CLEAR LYTIC ZONES IS THEN MEASURED.

Hurtz et al. 2000
Porta & Kortet 2003
Porta et al. 2004

ANTIMICROBIAL PEPTIDES INDUCIBLE SMALL PEPTIDES. THEY ASSAYED VIA THE CLEARANCE RATE OF AN INJECTION OF BACTERIA. A STANDARD BACTERIAL DOSE WAS INJECTED INTO THE HOST. EXTRACTED HAEMOLYMPH IS PLATED IN MEDIA AGAR AND CULTIVATED; PLATES ARE THEN SCORED FOR THE NUMBER OF BACTERIAL COLONIES FORMED BY USING A COLONY COUNTER. THIS MAY ALSO BE DONE VIA AN ANTIBACTERIAL ZONE ASSAY (AS USED IN LYZOZYME ACTIVITY PROTOCOL).

LYSOZYME GENERIC INHIBITOR (MORITZ & SCHMID-HEMPEL 2000) CAN BE USED TO DISTINGUISH AMONG ANTIMICROBIAL MOLECULES. (2) ASSAY METHODS.

DESTABILIZE BACTERIAL MEMBRANES. USED FOR HUMORAL RESPONSE QUANTIFICATION PROTOCOL). (2) ASSAY METHODS.

ACETYLGLUCOSAMINE) CAN BE USED TO DISTINGUISH AMONG ANTIMICROBIAL MOLECULES. (2) ASSAY METHODS.

* BEFORE INOCULATION, MINIMAL LETHAL AND/OR SUB-LETHAL DOSES MUST BE DETERMINED. OVERDOSES CAN PROMOTE SEPTIC SHOCK, RATHER THAN A MEASUREMENT OF ANTIBACTERIAL ACTIVITY.

** BECAUSE ANTIMICROBIAL ACTIVITY CAN BE DUE TO LYSOZYMES, REACTIVE OXYGEN SPECIES AND/OR ANTIMICROBIAL PEPTIDES, IMMUNE RESPONSE MEASUREMENTS AS HEMOLYTIC, LYZOZYME-LIKE OR ANTIBACTERIAL ACTIVITY.

970

971

972

DR. ROBERT FRECKLETON
EDITOR IN CHIEF
METHODS IN ECOLOGY AND EVOLUTION

DEAR DR. FRECKLETON:

THANKS VERY MUCH FOR YOUR COMMUNICATION REGARDING THE REVIEWERS' COMMENTS TO OUR
*Current immunity markers in insect ecological immunology: assumed trade-offs and
methodological issues*. WE ARE HAPPY TO KNOW THAT SUCH COMMENTS WERE FAIRLY POSITIVE. WE
ARE GRATEFUL WITH THE REVIEWERS AS THEIR COMMENTS HAVE BEEN VERY ENRICHING. WE HAVE PREPARED
A REVISION ACCORDING TO THESE COMMENTS.

YOU WILL FIND IN DETAIL HOW WE HAVE ADDRESSED SUCH CHANGES. SUCH CHANGES CAN BE TRACED
DIRECTLY TO THE LINE NUMBERS THAT ARE ALSO INDICATED BELOW.

Reviewer 1

1- ONE FIRST GENERAL COMMENT IS RELATED TO IMMUNITY EVOLUTION. WE AGREE THAT IMMUNITY
IS EVOLVING. HOWEVER, THE INSECT IMMUNE SYSTEM IS A SET OF CELLS, MOLECULES AND REACTION
FEATURES NOT ONLY RESIST PATHOGEN INVASION, BUT ALSO LIMITS THE DAMAGE CONSEQUENCE OF
INFECTION (THIS IDEA IS NOW MENTIONED IN LINE 85-88). ALL THESE FEATURES CHARACTERIZE THE
SYSTEM OF INSECTS AND CAN CHANGE AMONG GENERATIONS. IN CONSEQUENCE, WE CAN STATE THAT THE
IMMUNE SYSTEM IS CONTINUOUSLY EVOLVING. REVIEWER IS ALSO CONCERNED ABOUT THE HOW THESE
MARKERS ARE RELATED WITH RESISTANCE AGAINST PATHOGENS. AT THE INSECT IMMUNE MECHANISMS
WE MENTIONED THE EFFECT OVER THE PATHOGEN OF EVERY IMMUNE RESPONSE (WHICH ARE REACTION
IMMUNE MARKERS) MENTIONED HERE. ALSO, ALONG THE TEXT WE MENTIONED THE EFFECTS OF THESE
ON THE PATHOGEN (E.G. SEE LINES 104-105, 109-110, 136-146) OR ITS RELATION WITH HOST DEFENCE
(PLEASE SEE LINES 175-176, 203-206).

2- A SECOND GENERAL COMMENT IS RELATED TO THE EMPHASIS OF THE ECOLOGY AND THE RATIO OF
OFFS. THE OCCURRENCE OF THESE TRADE-OFFS HAVE BEEN PREVIOUSLY AND ACCURATELY PROVEN BY
RESEARCHES. OUR INTENTION IS NOT TO GIVE A RESUME OF THESE PAPERS, WHICH IS THE REASON WE
GIVE DETAILS OF EACH ONE OF THESE STUDIES. OUR MAIN INTENTION IN THIS REVIEW IS TO PROPOSE
BE TESTED) ABOUT THE PHYSIOLOGICAL RELATIONSHIP BETWEEN IMMUNITY AND OTHER LIFE HISTORY
INTERMEDIATE CHARACTERS, THAT ARISE TRADE-OFFS. HOWEVER, WE COMPLETELY AGREE WITH THE REVIEWER
THAT IS NECESSARILY TO DISTINGUISH THE LEVEL OF THE TRADE-OFFS DETECTED. FOR THIS REASON
PARAGRAPH ENTITLED *Trade-offs among immune responses* HAS BEEN INCLUDED IN THE INSECT IMMUNE
MECHANISMS SECTION (SEE LINES 148-162).

3- WE DELETE THESE LINES IN ORDER TO AVOID CONFUSION.

4- THE REVIEWER IS CONCERNED ABOUT THE METHODS REPEATABILITY. FOR THIS REASON A NEW
PARAGRAPH WAS INCLUDED IN THE CONCLUSION (SEE LINES 534-544). WE APPRECIATE THIS COMMENT.

5- THE REVIEWER NOTICE THAT SOME METHODS (E.G. GENETICS, PROTEOMICS) ARE NOT DISCUSSED.
EXPLANATION FOR THIS HAS BEEN INCLUDED IN THE INTRODUCTION (SEE LINES 65-72).

DETAILS

- 6- SEXUAL SELECTION IS NOW DEFINED AS THE “DIFFERENTIAL MATING AND FERTILIZATION SUCCESS”
- 7- WE REFER AS SHORT DEVELOPMENTAL TIMES NOW (LINE 59).
- 8- PARENTHESIS WAS CHECKED
- 9- WE DIFFER FROM THE REVIEWER’S OPINION RESOLVING THAT MELANIN IS A POLYMER NON-TOXIC MOLECULES PRODUCED THROUGH THE PO CASCADE ARE (LINES 108-110).

Reviewer 2

- 1- WE HAVE CHANGED THE LINES OF THE ABSTRACT, IN ORDER TO AVOID CONFUSION WHEN WE REFER TO THE USE OF IMMUNE MARKERS. SEE NOW LINES 27-30.
- 2- WE AGREE THAT THE IMMUNE MARKER DOES NOT HAVE A COST, BUT THE IMMUNE FUNCTION REQUIRES THE MARKER HAS. THIS PARAGRAPH WAS RE-WRITTEN TO AVOID CONFUSION (PLEASE SEE LINES 60-63).
- 3- WE NOW GIVE EXAMPLES AND REASONS OF THE IMMUNE ASSAYS THAT CAN NOT BE SUITABLE FOR ALL SPECIES. NOW SEE LINES 65-72.
- 4- THE REVIEWER IS COMPLETELY CORRECT. WE REWRITE THIS LINE (SEE LINES 80-81).
- 5- INSECT PO PATHWAY WAS CHECKED. WE AGREE THAT CATECHOLAMINES ARE BETTER PO SUBSTRATES. HOWEVER PO ALSO HYDROXYLATE TYROSINE, WHICH IS A PRECURSOR OF THE PO CASCADE IN INVERTEBRATES. SOME MORE DETAILS OF THIS CASCADE ARE GIVEN IN LINES 125-130.
- 6- WE REFER AS CUTICLE DARKNESS NOW. SEE LINE 167.
- 7- ROBB ET AL (2003) REFERENCE HAS BEEN INCLUDED TO EXPLAIN THAT IN SOME SPECIES CUTICLE DARKNESS IS NOT RELATED TO DISEASE RESISTANCE (SEE LINE 193-194)
- 8- THE REVIEWER IS ENTIRELY CORRECT. WOUND REPAIR CONSIDERED AS AN IMMUNE RESPONSE. SEE LINE 111-112.
- 9- WE NOW GIVE EXAMPLES AND REFERENCES ABOUT NON-IMMUNE FUNCTIONS OF HAEMOCYTES (LINES 206-208). THIS SECTION WAS ALSO RE-WRITTEN TO AVOID CONFUSION (SEE LINES 203-213).
- 10- WE EXPLAIN METHODS TO OBTAIN HAEMOCYTES. HOWEVER, WE STATE THAT THE METHOD USED AND THE USE OF ANTICOAGULANTS MUST BE CONSIDERED PROPERLY BY EACH RESEARCHER (LINES 226-229). THE REVIEWER IS CONCERNED ABOUT THE PROBLEMS GENERATED BY THE USE OF ANTICOAGULANTS AND THE DILUTION FACTOR. BEFORE EXTRACTION EACH RESEARCHER COMPLETES

ESTABLISHES THE AMOUNT OF MEDIUM IN WHICH SAMPLE (WITH OR WITHOUT ANTICOAGULANT) OBTAINED AND DILUTION FACTOR IS DETERMINED BEFORE EXTRACTION. AS WE CONSIDER THE PRACTICE WE DID NOT INCLUDE THIS ISSUE IN THE TEXT.

- 11- AS PERFECTLY SUGGESTED BY THE REVIEWER, WE REFER NOW THE EXISTENCE OF HAEMATOP IN SOME INSECTS (PLEASE SEE LINES 289-290).
- 12- WE DETAILED THE ENHANCING INTERACTION BETWEEN HAEMOLYMPH AND PO ACTIVITY (SEE LINES 342).
- 13- WE HAVE RE-WRITTEN THIS LINE IN ORDER TO MAKE IT CONSISTENT WITH THE WHOLE SENTENCE (SEE LINES 347).
- 14- WE STATE THAT THE USE OF IMPLANTS ARE A SUITABLE METHOD TO TEST FOR TRADE-OFFS (PLEASE SEE LINES 345-347). AS ACCURATELY SUGGESTED BY THE REVIEWER, WE DISCUSS NOW THE PROBABLE DIFFERENCES WHEN USING DEAD OR LIVE PATHOGENS. ALSO WE GIVE AN EXAMPLE TO ILLUSTRATE THIS DIFFERENCE (PLEASE SEE LINES 352-359).
- 15- THE REVIEWER WAS COMPLETELY RIGHT. WE WERE WRONG WHEN WE STATE THAT THERE CAN BE A DEFICIENCY WITHOUT A MELANIN DEFICIENCY. THIS LINE IS CORRECTED NOW. SEE LINES 385-386.
- 16- WE AGREE WITH THE REVIEWER THAT PO CAN BE PRODUCED IN OTHER TISSUES, AND THIS IDEA IS NOW WRITTEN IN THE TEXT (SEE LINES 378-382). NEVERTHELESS, WE ATTEMPTED TO MAKE A HYPOTHESIS RELATED TO THE PO TRANSPORTATION TO OTHER TISSUES AND THE POSSIBLE BASE OF A TRADE-OFF. WE DO NOT HAVE ANY EVIDENCE IF THIS IDEA IS SPECIES-SPECIFIC. HOWEVER, WE ARE AWARE THAT THIS HYPOTHESIS NEEDS FURTHER EXPLORATION; THIS HAS BEEN STATED IN THE TEXT (SEE LINES 383-384).
- 17- THE REVIEWER IS CORRECT; TYROSINE (TYR) CAN BE STORED AS MULTIPLE INTERMEDIATE MOLECULES. THIS IS NOW STATED IN THE TEXT (SEE LINES 403-405). HOWEVER, TRADE-OFFS CAN BE DETECTED IF THERE IS ENOUGH TYR (AND ITS SUBPRODUCTS) IS GATHERED THROUGH ONTOGENY (PLEASE SEE LINES 406-407). AS SUGGESTED BY THE REVIEWER, NOW WE STATE THAT PO MEASURES NEED TO BE INTERPRETED IN THE CONTEXT OF THE SPECIE'S TYROSINE METABOLISM, PLEASE SEE LINES 406-407.
- 18- THE PEROXIDE SUPPRESSOR HAS BEEN STATED IN THE TEXT (SEE LINE 422).
- 19- DETAILS OF WHY CONTROLLING TOTAL PROTEIN LOAD IS NOW EXPLAINED. PLEASE SEE LINES 426-428.
- 20- REVIEWER IS COMPLETELY CORRECT. WE NOW STATE THAT NO CAN BE CONTINUOUSLY SYNTHESIZED AND INDUCED NOS FOR LONG PERIODS OF TIME (SEE LINES 454-456).
- 21- THE REVIEWER IS CONCERNED ABOUT THE USE OF MODERN METHODS (E.G. GENE EXPRESSION ANALYSIS). AN EXPLANATION FOR THIS HAS BEEN INCLUDED IN THE INTRODUCTION (SEE LINES 65-72). ALSO, A RECOMMENDATION WAS ALSO INCLUDED IN THE CONCLUSION SECTION (SEE LINES 581-586).
- 22- WE APPRECIATE THE REVIEWER ADVICE. HOWEVER, IN THE INTRODUCTION WE POINT OUT THAT THE METHODS MENTIONED IN THE MANUSCRIPT ARE SUMMARIZED IN TABLE 1. WE CONSIDER REDUCING AND REFER TABLE 1 IN EVERY IMMUNE MARKER MENTIONED IN THE TEXT.

- 23- WE AGREE THAT THE JH CAN BE A COMPLICATE MODEL TO EXEMPLIFY THE HANDICAP HYPOTHESIS. WE WANTED TO DEMONSTRATE THE CORRECT USE OF AN IMMUNE ASSAY. AS PROPERLY SUGGESTED BY THE REVIEWER, THE WHOLE SECTION WAS REMOVED. REVIEWER ASKED FOR AN EXAMPLE WHERE AN IMMUNE MARKER IS NOT USEFUL TO ANSWER A RESEARCH QUESTION. WE DID NOT REFER TO ANY PAPER SINCE OUR AIM WAS NOT UNDERESTIMATE THE CURRENT RESEARCH IN INSECT ECOLOGICAL IMMUNITY. THE SCOPE OF THE PAPER IS TO HIGHLIGHT IMMUNE MARKERS THAT HAVE BEEN CAREFULLY AND PROPERLY USED IN PREVIOUS WORKS. FOR THESE REASONS WE DELETED OR RE-WROTE LINES WHERE THIS CONFUSING OR NEGATIVE TONE WAS USED (SEE LINES 566-572). NEEDLESS TO SAY, WE ARE VERY GRATEFUL WITH THESE THOUGHTFUL COMMENTS.
- 24- THE SCHNEIDER'S WORK HAS NOW BEEN USED TO EXPLAIN NOT ONLY THE USE OF IMMUNE MARKERS TO RESIST PATHOGENS, BUT ALSO TO TOLERATE AND AVOID DAMAGE (PLEASE SEE LINES: 354-359). IN ADDITION TO THIS, WE ALSO MENTION THE NEED OF MULTIPLE MEASURES OF IMMUNE RESPONSE NOT ONLY TO BALANCE TRADE-OFFS WITH OTHER TRAITS, BUT TRADE-OFFS AMONG IMMUNE RESPONSES (SEE LINES: 500-505). THANK THE REVIEWER FOR THIS EXCELLENT SUGGESTION.

Reviewer 3

- 1- THE REVIEWER IS CONCERNED ABOUT THE NEGATIVE TONE OF THE ARTICLE. OUR AIM IS TO NOT UNDERESTIMATE THE CURRENT METHODOLOGIES IN INSECT ECOLOGICAL IMMUNITY. THE SCOPE OF THE PAPER IS TO HIGHLIGHT IMMUNE MARKERS THAT HAVE BEEN CAREFULLY AND PROPERLY USED IN PREVIOUS WORKS. FOR THESE REASONS WE DELETED OR RE-WROTE LINES WHERE THIS NEGATIVE TONE WAS USED (FOR EXAMPLE, SEE LINES: 310-314, 344-347, 526-530). WE AGREE WITH THE REVIEWER THAT THE IMMUNE MARKER IS NOT THE OBJECT OF STUDY ITSELF. COMMONLY THE ECOLOGICAL AND EVOLUTIONARY CONSEQUENCE OF THE TRADE-OFFS ARE DISCUSSED, BUT THE POSSIBLE PHYSIOLOGICAL LINK AMONG IMMUNITY AND OTHER TRAITS ARE NOT CONSIDERED. WE BELIEVE THAT THIS LINK CAN BE USEFUL FOR DISCUSSIONS IN ECOLOGICAL IMMUNITY RESEARCHES. FOR THESE REASONS, WE MAKE (WITH THE AVAILABLE INFORMATION BUT IMMUNOLOGICAL AND ECOLOGICAL) HYPOTHESES RELATING TO THE POSSIBLE BASE OF A TRADE-OFF AMONG IMMUNITY AND OTHER TRAITS. HOWEVER, WE ARE AWARE THAT THESE HYPOTHESES NEED FURTHER EXPLORATION; THIS HAS BEEN STATED IN THE TEXT (SEE LINES 500-505).
- 2- AS RECOMMENDED BY THE REVIEWER, WE CAREFULLY REVISED THE IMMUNOLOGY LITERATURE CITED HERE. WE WERE WRONG WHEN WE STATED THAT USING THE PHRODO ASSAY IS BETTER THAN THE PHRODO ASSAY, THIS LINE WAS RE-WRITTEN TO AVOID CONFUSIONS (SEE LINES 263-265). WE THANK THE REVIEWER FOR POINTED OUT THIS MISTAKE. HOWEVER, IT SEEMS THAT THE REVIEWER FOUND OTHER PLACES WHERE FURTHER IMPROVEMENT CAN BE MADE. WE ARE OPEN TO SOLVE THESE ISSUES IF THEY ARE OUR REVIEWER'S CONCERN.
- 3- WE AGREE THAT IT IS NOT RELEVANT TO TEST AMP SPECIFICITY, FOR THIS REASONS WE DELETED THE TEXT AS WELL IN TABLE 1. WE HAVE DONE SOME RECOMMENDATION BASED IN OUR OWN EXPERIMENTAL DATA. FOR EXAMPLE, IN ONE OF OUR INSECT MODELS, CUTICULAR THICKNESS, BUT NOT COLOUR, WAS RELEVANT.

AN IMMUNE CHALLENGE (UNPUBLISHED DATA). NONETHELESS, MEASUREMENTS COULD BE NOT A TASK, IT IS POSSIBLE TO GATHER A GREAT NUMBER OF INDIVIDUAL MEASUREMENTS, AND ITS COST IS ELEVATED. OF COURSE, THE USE OF EVERY METHOD MENTIONED HERE MUST BE DETERMINED BY THE RESEARCHER NOT ONLY IN TERMS OF ITS OVERALL COST, BUT ALSO IN THE QUESTION TO BE ANSWERED. THIS IDEA HAS BEEN INCLUDED IN TEXT, PLEASE SEE LINES 530-534).

- 4- ONE OF OUR AIMS IN THIS REVIEW IS TO GIVE AN OVERVIEW OF THE MOST FREQUENTLY-USED IMMUNOLOGICAL MARKERS APPLIED IN THE FIELD OF EVOLUTIONARY ECOLOGY OF INSECTS. AS MENTIONED BEFORE, WE HAVE DONE SOME RECOMMENDATIONS NOT ONLY TRYING TO IMPROVE TECHNIQUES BUT ALSO GAVE SOME POSSIBLE SOLUTIONS; ALL BASED IN THE EXISTENT LITERATURE, AND OUR OWN EXPERIENCE. WE CONSIDER THAT WE HAVE PROPOSED THE USE OF NEW AND FEASIBLE TECHNIQUES, FOR EXAMPLE, MEASUREMENT OF THICKNESS, PHRODO ASSAYS, NO MEASUREMENTS. WE AGREE WITH THE REVIEWER IN THE SENSATION THAT THE USE OF PATHOGEN MUST BE CONSIDERED, THIS IDEA HAS BEEN INCLUDED IN THE TEXT (PLEASE SEE LINES 213, 364-366, 553-555). ALSO, THE REVIEWER IS COMPLETELY CORRECT, MAXIMAL IMMUNE RESPONSE IS NOT NECESSARILY RELATED WITH MAXIMUM FITNESS, EVEN SOMETIMES, THE ABSENCE OF IMMUNE RESPONSE CAN BE ADVANTAGEOUS. THIS IDEA IS MENTIONED IN THE CONCLUSION (LINES 560-565).
- 5- THE REVIEWER IS CONCERNED ABOUT THE USE OF NON PATHOGENS TO INDUCE AN IMMUNE RESPONSE. OUR INTENTION WAS TO MAKE SOME RECOMMENDATIONS NOT TO CONSIDER THEM AS INAPPROPRIATE FOR STUDIES IN THE FIELD OF ECOLOGICAL IMMUNOLOGY. WE DELETED WORDS OR RE-WROTE LINES TO AVOID A CONFUSING CONNOTATION, FOR EXAMPLE PLEASE SEE NOW LINES 253-260, 310-314, 344-347. WE COMPLETELY AGREE WITH THE REVIEWER THAT THE USE OR NOT USE OF NON LIVING PARTICLES IS AN EXCELLENT ISSUE TO DISCUSS. HOWEVER, OUR PRINCIPAL AIM IN THIS PAPER WAS TO DISCUSS MEASUREMENTS AND PROPOSED THEORIES (SOME OF THEM TO BE TESTED) ABOUT THE PHYSIOLOGICAL LINK WITH IMMUNITY (MEASURED WITH THE MARKERS MENTIONED IN THE TEXT) AND OTHER LIFE HISTORY TRAITS. WE DO NOT INCLUDE THIS SUBJECT IN THE MS. WE DO NOT WANT TO MAKE FEEL THE REVIEWER DISAPPOINTED TO SEE THAT THE NEW MANUSCRIPT DOES NOT INCLUDE THIS SUGGESTION, BUT WE BELIEVE THAT THIS ISSUE COULD BE PROPERLY FOR AN EXTEND DISCUSSION (EVEN A REVIEW FOR PUBLICATION).
- 6- AN ADDITIONAL COLUMN WITH REFERENCES HAS BEEN INCLUDED IN TABLE 1. SOME SHORTENING OF THE TEXT WAS DONE WHEN POSSIBLE. WE APPRECIATE THIS USEFUL SUGGESTION.

THE PAPER HAS BEEN REVISED BY TWO ENGLISH SPEAKING PERSONS

FINALLY, SOME NEW REFERENCES HAVE BEEN ADDED.

PLEASE, DO NOT HESITATE TO CONTACT ME IF THERE ARE FURTHER ISSUES TO DISCUSS.

SINCERELY,

MIGUEL MORENO-GARCÍA