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“LIPOPROTEÍNA Lp(a) Y RIESGO
CARDIOVASCULAR: DETECCIÓN DE
ISOFORMAS PATÓGENAS DE Lp(a) POR UN
NUEVO MÉTODO DE UNIÓN A LA FIBRINA”

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DIRECTOR DE TESIS:
DR. EDMUNDO CHÁVEZ COSSIO

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AVENIDA DE
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DOCTORADO EN CIENCIAS BIOMEDICAS

FACULTAD DE MEDICINA

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DIRECTOR GENERAL DE ADMINISTRACION
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PRESENTE.

Por medio del presente me permito informar a usted que en la reunión del Subcomité Académico de Doctorado en Ciencias Biomédicas que se llevo a cabo el día 9 de octubre del presente año, se acordó designar el siguiente jurado para examen de Doctorado en Ciencias Biomédicas de la M. en C. PEÑA DIAZ AURORA DE LA con número de expediente 91542 y número de cuenta 7152037-1 con la tesis titulada: "LIPOPROTEÍNA Lp(a) Y RIESGO CARDIOVASCULAR: DETECCIÓN DE ISOFORMAS PATÓGENAS DE Lp(a) POR UN NUEVO MÉTODO DE UNIÓN A LA FIBRINA", dirigida por el Dr. Edmundo Chávez Cossío.

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**LIPOPROTEÍNA Lp(a) Y RIESGO
CARDIOVASCULAR: DETECCIÓN DE
ISOFORMAS PATÓGENAS DE Lp(a)
POR UN NUEVO MÉTODO DE UNIÓN
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LIPOPROTEINA (a), Lp(a) Y ENFERMEDAD ATEROTROMBÓTICA.

Este trabajo de tesis se realizó con la colaboración de:

- El Instituto Nacional de la Salud y de la Investigación Médica (INSERM), Unidad 143 Trombosis y Biología Vascular, Hospital de Bicêtre y la Unidad 146 , Facultad de Medicina Xavier Bichat EPI 9907, Paris, Francia.
- El Instituto Nacional de Cardiología "Ignacio Chávez", (INC), México, D.F.
- El Instituto Nacional de Antropología e Historia, (INAH), México, D.F.
- El Centro de Ingeniería Genética y Biotecnología. División de Inmunotecnología y Diagnósticos. El Instituto Nacional de Angiología, La Habana, Cuba.

En el departamento de Hematología del INC, se identificó la inhibición competitiva del plasminógeno por la Lp(a), con esta finalidad se empleó un modelo "in vitro". El modelo permite ponderar la participación de la Lp(a) como inhibidor del plasminógeno y obtener un perfil de la inhibición fibrinolítica de un individuo por la presencia de Lp(a). La concentración de Lp(a) y las isoformas de apo(a) se identificaron en el departamento de Endocrinología del INC.

En el Centro de Ingeniería Genética y Biotecnología en la Habana, Cuba, se diseñó y obtuvo un anticuerpo monoclonal anti apo-(a).

El Instituto Nacional de Antropología e Historia identificó y facilitó el acceso para obtener las muestras de sangre de las comunidades mazahuas y mayas. Recibimos también el apoyo de la Secretaría de Salud del Estado de Sinaloa para

obtener las muestras de la comunidad de mayos, poblados de Capomos y Tehuecos.

El Instituto Nacional de Neurología remitió, para su estudio, a un grupo de individuos con enfermedad vascular cerebral, que se incluyó dentro del grupo de estudio.

Durante el primer año de trabajo de tesis, del 15 de enero de 1997 al 15 de agosto de 1997, recibí una beca del sistema ECOS-ANUIES, para realizar una estancia de investigación en la U 143 del INSERM, Hospital de Bicetre, Francia. Durante el periodo mencionado se pusieron a punto las técnicas que permiten identificar la unión de apo(a) a la fibrina, que actualmente se aplican con éxito en el INC.

Del 15 de octubre al 15 de diciembre de 1999 recibí una beca de la Dirección General de Postgrado de la UNAM para realizar una segunda estancia de investigación en la U 143 del INSERM, Hospital de Bicetre, Francia.

RESUMEN

Este trabajo de tesis propone una nueva estrategia para explorar la capacidad trombogénica de la Lp(a), a través del estudio funcional de la unión de la apo(a) a la fibrina como indicador de la asociación de la Lp(a), al proceso aterotrombótico.

Con esta finalidad se puso a punto una técnica que permite evaluar la unión de apo(a) a la fibrina. Se estudiaron también 248 individuos, 105 con cardiopatía isquémica (CI) y 52 con enfermedad cerebrovascular (EVC) de origen trombótico y 91 controles.

El grupo de EVC tuvo una alta unión de apo(a) a la fibrina (0.268 ± 0.15 nmol/L) en comparación con el grupo de CI (0.155 ± 0.12 nmol/L), y con el grupo control (0.158 ± 0.09 nmol/L) $p < 0.0001$; lo que sugiere que existen diferencias particulares en los mecanismos de regulación anti-trombótica en los lechos arteriales cerebrales y coronarios.

Es también importante resaltar que existen diferencias en la concentración y en la presencia de distintas isoformas en poblaciones genéticamente diferentes.

SUMMARY

This thesis proposes a well-characterized model of a fibrin surface to develop a functional approach for the detection of pathogenic Lp(a). The assay is based on the competitive binding of Lp(a) and plasminogen for fibrin, and quantifies fibrin-bound Lp(a). High Lp(a) binding to fibrin is correlated with decreased plasmin formation. In a transversal case-control study we studied 248 individuals, 105 had a history of ischemic cardiopathy (IC), 52 had cerebro-vascular disease (CVD) of thrombotic origin, and 91 were controls.

The remarkably high apo(a) fibrin-binding in CVD (0.268 ± 0.15 nmol/L) as compared to IC (0.155 ± 0.12 nmol/L) suggests the existence of peculiar and poorly understood differences in pro- or anti-thrombotic mechanisms in either cerebral and/or coronary arteries.

Our results demonstrated that Lp(a) fibrin-binding and are associated with athero-thrombotic disease

INTRODUCCIÓN

Las enfermedades aterotrombóticas, cardiovasculares o cerebrovasculares, son una de las principales causas de morbilidad y mortalidad en los países occidentales. En nuestro país, la Secretaría de Salud publicó en su página electrónica (<http://www.salud.gob.mx>) que en el año 2000 la enfermedad isquémica del corazón fue la segunda causa de muerte y que la enfermedad cerebrovascular la sexta causa, en la población general.

Tanto factores genéticos como ambientales dan origen a la aterosclerosis; algunos de ellos como la lipoproteína de baja densidad (LDL), la hipertensión arterial, la diabetes y el tabaquismo se han estudiado ampliamente y se puede, actualmente, conocer el riesgo que conlleva presentar alguno o varios de estos factores con el de padecer enfermedades aterotrombóticas.

Sin embargo, existen otros factores que actualmente se están conociendo y evaluando. A este grupo pertenece la lipoproteína (a) o Lp(a). La Lp(a) tiene una estructura muy parecida a la LDL, ambas poseen un núcleo de lípidos de composición similar; rodeado por la apolipoproteína B-100, la Lp(a) presenta además otra apolipoproteína, apo(a), que le confiere características antigénicas y fisiopatológicas diferentes, favorece el desarrollo de la placa aterosclerosa y favorece el desarrollo y permanencia de un trombo, por lo que se considera el punto común entre la aterosclerosis y la trombosis.

La presencia de una LDL en la estructura de Lp(a) se asocia al desarrollo acelerado de la aterosclerosis. La similitud entre la estructura de la apolipoproteína (a) y el plasminógeno, precursor de la plasmina, se asocia con una permanencia mayor de depósitos de fibrina en sitios de lesión endotelial y menor generación de plasmina, lo que acentúa las propiedades aterogénicas y trombogénicas de la Lp(a).

Bajo esta perspectiva, se realizaron numerosos estudios epidemiológicos que describen una asociación positiva entre la elevada concentración plasmática de Lp(a) y un incremento en las enfermedades cerebrovasculares, cardiovasculares, la reestenosis de puentes coronarios, la reoclusión de angioplastías y el desarrollo prematuro de aterosclerosis, en ocasiones asociada a altas concentraciones de LDL y/o a bajas concentraciones de lipoproteínas de alta densidad (HDL). La mayoría de estudios prospectivos confirman estos resultados, aún cuando no existe un acuerdo en el límite de corte para considerar una concentración plasmática normal; algunos autores consideran que es de 20 mg/dL y otros hasta 30 mg/dL. También se han empleado técnicas y diferentes anticuerpos en su detección, monoclonales o policlonales, lo que ha generado divergencias en los resultados.

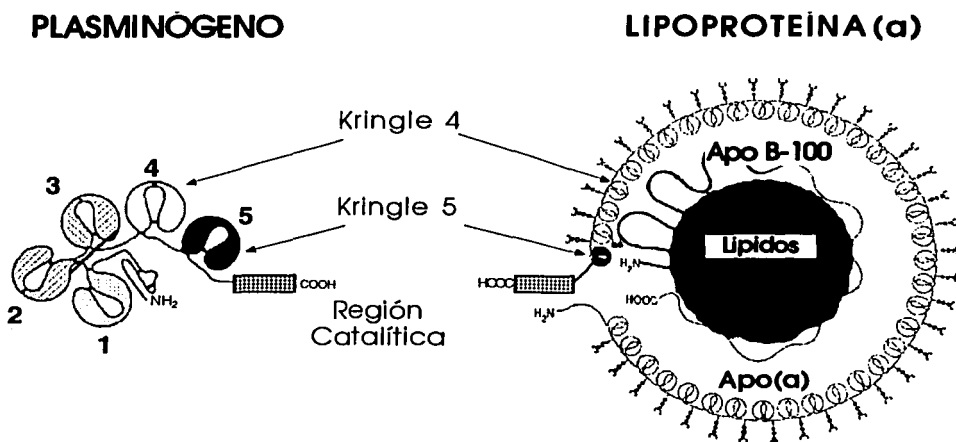
Sin embargo, otros estudios no encuentran asociación entre la Lp(a) y la enfermedad arterial coronaria.

Esta discrepancia en los resultados podría ser entre varios motivos, el reflejo de la gran heterogeneidad estructural de la molécula de apo(a), que se traduce en una

heterogeneidad funcional como inhibidor competitivo del plasminógeno, disminuyendo la formación de plasmina.

Desde 1963, Kåre Berg, identificó la presencia de Lp(a) en el plasma¹; sin embargo, hasta 1987 Eaton y col² identificaron parcialmente la secuencia de la glicoproteína apo(a) que posteriormente clonaron³, demostrando el gran parecido en su estructura terciaria entre uno de los componentes de la Lp(a), la glicoproteína apo(a), con el plasminógeno, el precursor de la plasmina, Figura 1.

FIGURA 1.



Relación entre la estructura química de Lp(a) y del plasminógeno. En Lp(a) hay un número variable de kringles tipo 4, un kringle tipo 5, una región catalítica serina-proteasa que en Lp(a) no puede activarse.

Este hallazgo despertó el interés de numerosos grupos de investigación que encontraron el punto común entre la aterosclerosis y la trombosis^{4, 5, 6, 7, 8, 9, 10}.

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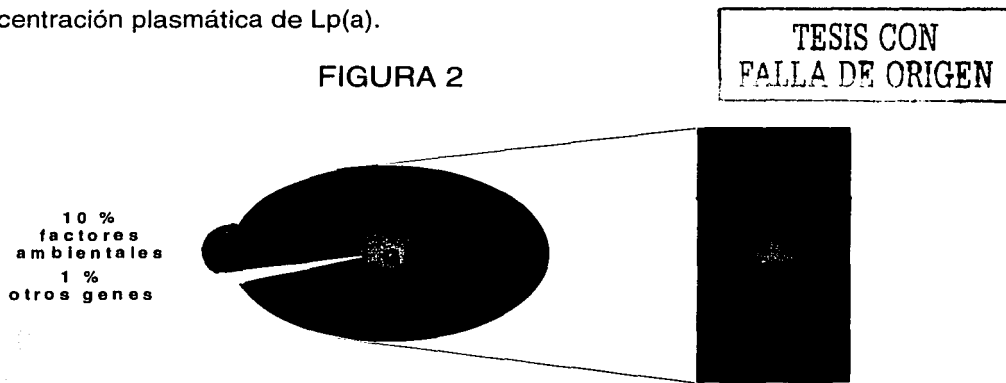
No todos los organismos sintetizan apo(a). Su presencia se identifica en los humanos, en los primates del viejo mundo, en algunos primates del nuevo mundo ¹¹, y en el puerco espín ¹².

FUNCIÓN

No se conoce con certeza cuál puede ser su función en el organismo. Aporta colesterol desde el hígado a los órganos que sintetizan hormonas esteroidales ¹³ y a las células de los tejidos en proceso de reparación ¹⁴. Además, tampoco se conoce la función de la apo(a) que se encuentra en los testículos y en el cerebro de manera independiente, es decir, sin formar parte de la molécula de Lp(a) ¹⁵.

CONCENTRACIÓN PLASMÁTICA

La concentración plasmática de Lp(a), que depende de su síntesis hepática ^{16, 17}, varía de un individuo a otro en un límite aproximado de <10 mg/dL a > 100 mg/dL; Es independiente de otros factores como la dieta, el colesterol, la obesidad, el tabaquismo, y se mantiene con variaciones discretas a lo largo de la vida ^{18, 19}, la Figura 2 muestra que el gene de apo(a) participa en casi el 90% en la regulación de la concentración plasmática de Lp(a).



Generalmente, un individuo hereda de una manera autosómica codominante dos isoformas de apo(a) que pueden identificarse en el laboratorio por la técnica de SDS-PAGE y posteriormente por inmunotransferencia, empleando anticuerpos monoclonales o policlonales contra apo(a). Por este método se identifican cerca de 37 isoformas, que según la clasificación de Uterman se denominan F (faster), B (=apoB-100), S1(slow), S2, S3, S4, dependiendo de su velocidad de migración en comparación con apoB-100²⁰.

Las isoformas de apo(a) muestran una relación inversa entre el peso molecular y la concentración plasmática de Lp(a)²¹, probablemente porque a medida que se incrementa el tamaño de apo(a) se incrementa la dificultad para secretarse de la célula, como ocurre en la línea celular HepG2 de hepatocarcinoma humano²².

El método no permite distinguir diferencias entre las isoformas de pesos moleculares cercanos, aunque recientemente, se describe la posibilidad de identificar con esta técnica, isoformas en un rango muy amplio de pesos moleculares con la ayuda de una referencia que contiene recombinantes de apo(a) de diferentes pesos moleculares y que correlaciona con la técnica de electroforesis de campo pulsado ($r= 0.97$)²³ que se emplea para identificar los diferentes genotipos que codifican para la apo(a)²⁴. Con la técnica de campo pulsado se describieron, en un estudio de población caucásica americana, 19 alelos diferentes²⁵.

Algunas hormonas pueden modificar su concentración plasmática; la hormona tiroidea²⁶, los estrógenos^{27, 28, 29, 30} y esteroides anabólicos la reducen³¹, y la

hormona de crecimiento la incrementa³²; pero no puede modificarse por medicamentos hipolipemiantes o por la dieta^{18, 33}.

La concentración plasmática de la Lp(a) se incrementa en padecimientos como el síndrome nefrótico, la artritis reumatoide y transitoriamente después de un infarto al miocardio o de intervenciones quirúrgicas^{34, 35, 36}.

Existen variaciones en la concentración promedio y en la presencia de isoformas en poblaciones genéticamente diferentes^{37, 38, 39, 40}. Los individuos negros tienen al menos, el triple de concentración plasmática de Lp(a) que los blancos^{41, 42, 43, 44}.

CONCENTRACIÓN PLASMÁTICA DE LP(a) EN DIVERSAS POBLACIONES

Con la finalidad de identificar a las poblaciones en las que la Lp(a) represente un factor de morbilidad y mortalidad importante y resaltar su impacto en la salud pública, se han llevado a cabo numerosos estudios en poblaciones sanas. En la Tabla 1 se muestra algunos de estos estudios^{45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66}.

En el curso del trabajo de tesis analizamos los trabajos de poblaciones publicados en los últimos diez años, que aparecen en medline. Como criterio de búsqueda se empleó las siguientes palabras clave: Lp(a), ethnicity, population.

Encontramos que a pesar del gran interés que el tema ha cobrado en los últimos años no existe un criterio para el manejo pre-analítico de la muestra, ni estándar ni técnica de referencia, ni un consenso en el empleo de anticuerpos monoclonales o policlonales para determinar la concentración plasmática de Lp(a). Entre los

puntos que señalan la dificultad que existe para comparar los diferentes estudios, resalta también la ausencia de estudios en la población latinoamericana y china. En el mapa que se muestra en la Figura 3 se señalan los estudios resultantes de la búsqueda y que emplearon la técnica de Elisa, la más empleada para determinar la concentración plasmática de Lp(a). El diámetro de los círculos representa el número de sujetos estudiados. El círculo que representa a la población mexicana corresponde a nuestro estudio realizado en el grupo de mestizos, mazahuas, mayas y mayos.

TABLA 1

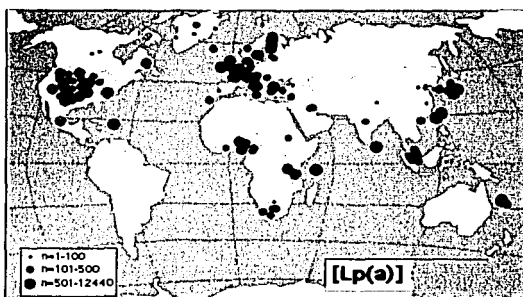
Ref	Población	Concentración de Lp(a)
46	USA Blancos y Negros	tienen en promedio, el doble de la concentración que los blancos.
47	Húngaros	No existe diferencia de concentración promedio entre hombres y mujeres. Las mujeres tienen una distribución bimodal. Sólo 9.4% de la población en general supera 30 mg/dL.
48	Italianos	75% de la población tiene una concentración promedio menor a 10 mg/dL.
49	Alemanes, Gahananos, Chinos	La concentración promedio de los gahaneses es el doble que la de los alemanes y 1.65 veces mayor que la de chinos. Las diferencias interétnicas se asocian a las isoformas de apo(a) de alto PM.

- 50 Tibetanos, Coreanos, Los nigerianos tienen una concentración 2.7 veces Chinos, Nigerianos, más alta que los tibetanos. Sin correlación con Belgas. otras variables estudiadas.
- 51 Tiroleeses, Alemanes, Los enfermos tienen mayor concentración que los Galeses, Israelitas, controles, excepto en los israelitas. En los Chinos de Singapur, alemanes no se midió concentración. Indués de Singapur. El gen de apo(a) es un factor determinante en la Grupo enfermos concentración de Lp(a) y confirma su papel como coronarios factor de riesgo en CI.
- 52 Japoneses y Chinos Correlación inversa entre la concentración y el tamaño de la isoforma de apo(a) en ambas poblaciones.
- 53 Españoles 38% de la población estudiada con concentración mayor a 30 mg/dL
- 54 Negros de La concentración promedio de Lp(a) es el doble en Sychelles, Blancos los negros que en los blancos. Correlación positiva Suizos con colesterol total, LDL colesterol y Apo B. Sin correlación con alcohol, tabaco o IMC.
- 55 Caucásicas y afro Las mujeres afro caribeñas tienen mayor caribeñas en el 3er concentración de Lp(a) que las mujeres caucásicas trimestre embarazo
- 56 Mexicanos Correlación inversa entre la concentración y el

- residentes en EUA tamaño de la isoformas de apo(a).
- 57 Caucásicos y japoneses y Diferencias significativas de los subtipos de alelos entre las poblaciones estudiadas y en pacientes con IAM.
- 58 Esquimales y Daneses caucásicos y Correlación inversa entre la concentración y el tamaño de las isoformas de apo(a) en las 3 poblaciones.
- 59 Chinos e hindúes residentes en Singapur. Recién nacidos e hindúes en Singapur. Hindúes tienen mayor concentración que los chinos, dato que concuerda con la tasa de mortalidad de sus respectivas poblaciones adultas en Singapur.
- 60 Niños españoles 15.1 % de la población estudiada tiene concentración arriba de 30 mg/dL.
- 61 Griegos El percentil 77 de los hombres y 66 de las mujeres, tiene menos de 30 mg/dL de Lp(a).
- 62 Pigmeos y Bantús 41% de los hombres pigmeos y 52% de las mujeres pigmeas tiene una concentración mayor a 30 mg/dL. En los bantús fue de 47% y 55% en hombres y mujeres, respectivamente. No hay diferencias significativas entre las 2 poblaciones. Sin correlación con otras variables estudiadas.
- 63 Mexicanos Análisis segregacional sugiere la existencia de otro

- residentes en EUA gen que participa con otro 41% en el control de la concentración.
- 64 Negros Sudafricanos Sin correlación entre el tamaño de la isoforma de apo(a) y la concentración de Lp(a).
- 65 Sicilianos Resultados similares a los de poblaciones caucásicas.
- 66 Estonianos y Rusos Estonianos mayor concentración de Lp(a) que los rusos.
- 67 Taiwaneses Correlación positiva con la edad, LDL, fibrinógeno. Correlación negativa con TG, bajo HDL, IMC, resistencia insulina en hombres. Menopausia no cambia la concentración de Lp(a)?, anticonceptivos la disminuyen.
-

FIGURA 3



Distribución geográfica de los estudios de la concentración plasmática de Lp(a)

ESTRUCTURA QUÍMICA

La composición de la Lp(a) es parecida a la de lipoproteínas de baja densidad o LDL. Ambas lipoproteínas contienen colesterol, triacilglicerol y fosfolípidos, que pueden disolverse y transportarse en el plasma, gracias a la presencia de una proteína, apoB-100, que rodea al grupo de lípidos y colesterol⁶⁷.

TABLA 2

Propiedades	Lp(a)	LDL
Diámetro A°	269 ± 12	206 ± 29
Movilidad electroforética	pre beta	Beta
Punto isoeléctrico	4.9	5.6
Peso molecular (x10 ⁶)	3.08	2.93
Densidad hidratada	1.055 - 1.12	1.02 - 1.063
Apolipoproteínas	Apo(a), Apo B-100	Apo B-100
Proteínas (%)	27- 30.9	22.4
Colesterol (%)	7.9	8.5
Éster de colesterol (%)	37.1	40.7
Triacilglicerol (%)	19	21.3
Fosfolípidos (%)	5	7.1

La diferencia principal entre la molécula de LDL y de Lp(a) radica en la presencia de apo(a) que se une a través de un puente disulfuro con apoB-100 entre Cys en posición 69 del KIV-9 de apo(a) y Cys 3734 de apoB-100; la unión se estabiliza

por puentes de hidrógeno e interacciones van der Waals en otras regiones de ambas proteínas^{68, 69, 70}. La apoB-100⁷¹ tiene la misma estructura y conformación en la molécula de Lp(a) y en LDL. La proporción de apo(a):apoB-100 es de 1:1⁷², pero considerando sus parámetros fisicoquímicos podrían existir entidades en proporción 2:1⁶⁹.

Como es frecuente en las lipoproteínas plasmáticas, Lp(a) puede tener diferentes tamaños, con pesos entre 800 y 1300 kDa y por lo tanto diferente densidad; estas diferencias reflejan, en menor medida, la composición del núcleo de lípidos y, principalmente, del polimorfismo estructural de apo(a)⁷³.

Existen dificultades técnicas para obtener la forma nativa de apo(a) y las predicciones de su estructura secundaria sugieren ausencia de α -hélice⁷⁴. Sin embargo, después de reducir los enlaces sulfhidrilo y por la técnica de dicroísmo circular, se observó 8 % de α -hélice, 21% de hoja β y 71% de arreglo al azar⁷⁵.

La apo(a) pertenece a la familia de serina proteasas, al igual que el plasminógeno, la protrombina, el activador tisular del plasminógeno, el activador del plasminógeno tipo urocinasa y el factor XII. Estas proteínas derivan de un gen ancestral común a todas ellas.

La apo(a) es muy parecida al plasminógeno (Figura 1). Los genes que codifican para ambas proteínas se encuentran muy cercanos, en el cromosoma 6, banda q26-27²⁰. Ambos genes se orientan cabeza con cabeza a 50 kb de distancia⁷⁶, 56 en la región terminal 5', que en el caso de apo(a) presenta un polimorfismo que puede expresarse modificando la eficacia en la transcripción y da lugar a

diferencias en la concentración plasmática de Lp(a) no sólo entre individuos sino, como ya se ha mencionado, entre diferentes grupos étnicos.

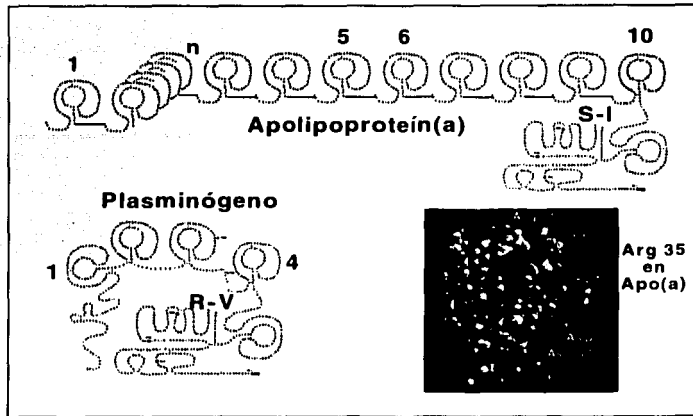
El gen de la apo(a) puede tener diferentes tamaños, cada uno corresponde al número de veces que se presenta una secuencia de 5.5 kb cuyo número puede variar entre 12 a 51. La región promotora del gen de apo(a) tiene, río arriba, un fragmento de 1 kb de tamaño cuya secuencia presenta varios sitios potenciales de interacciones. Con la interleucina 6 (IL6) presenta siete sitios, hecho que permite explicar el incremento pasajero en la concentración plasmática de Lp(a) durante estados de inflamación aguda y tres sitios con elementos específicos de transcripción hepática (HNF-1, CEBP, y LF-A1).

La apo(a) y el plasminógeno tienen una región catalítica con una similitud de 94%, pero que en el caso de apo(a) carece de la capacidad de activarse y de tener una función enzimática a causa de la presencia de arginina en lugar de serina en el sitio de activación.

El plasminógeno y la apo(a), tienen también, diferente número de módulos llamados kringles, Figura 4. A través de los kringles se reconocen y se unen a otras macromoléculas y/o a sitios específicos de la membrana celular ⁷⁷.

Los kringles se unen entre ellos por regiones interkringles, que son segmentos de 26 a 36 aminoácidos ricos en serina, prolina y/o treonina; cada región interkringle tiene 6 sitios potenciales para O-glicosilaciones .

FIGURA 4

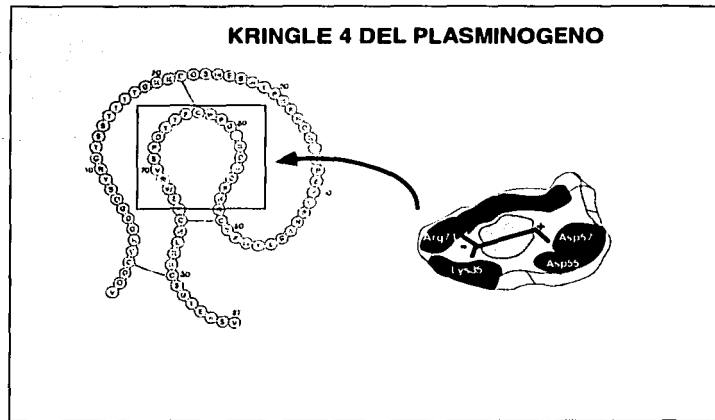


Estructura de plasminógeno y de Apo(a). R-V indica el sitio de hidrólisis del plasminógeno en Apo(a) el sitio S-I no se hidroliza; n indica el número de KIV tipo 2.

En la función de reconocimiento participa una estructura globular que se genera en el interior del kringle y que rodea a una región hidrófoba formada por varios aminoácidos aromáticos que se estabilizan a través de puentes de hidrógeno y separan a un grupo catiónico de un grupo aniónico ⁷⁸. Esta región se conoce como el sitio de unión a la lisina, o LBS Figura 5, por sus siglas del término en inglés lysine-binding-site. Sus características estructurales generan una geometría relativamente rígida a la que selectivamente tienen acceso y pueden unirse ligandos alifáticos o aromáticos de 6.8 Å, del tipo de ácidos ω -amino-carboxílicos como es el ácido ϵ -aminohexanóico o compuestos análogos .

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FIGURA 5



Representación esquemática del sitio LBS del kringle 4 del plasminógeno.

El plasminógeno tiene 5 tipos de kringles muy similares entre sí, pero con pequeñas diferencias en el sitio de unión a la lisina, que modifican el grado de afinidad por diferentes ligandos. El K1 del plasminógeno tiene un LBS de gran afinidad; el polo catiónico se forma con Arg 35 y Arg-71, el polo aniónico se forma con Asp-55 y Asp-57⁷⁹.

El LBS del K4 del plasminógeno tiene una afinidad intermedia; es una región hidrófoba en forma de V que genera una topografía en la que se alinean los anillos aromáticos de Phe-64, Trp-62 y Trp-72 que separan al grupo aniónico, formado por Asp-55 y Asp-57 del catiónico, formado por Lys-35 y Arg-71⁸⁰.

POLIMORFISMO DE APOLIPOPROTEINA (a)

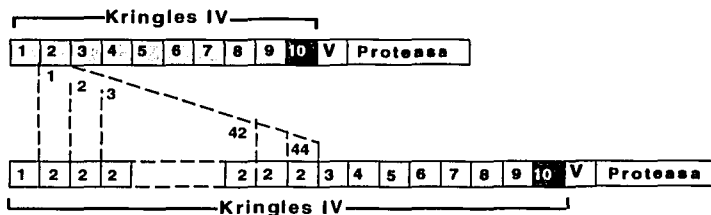
La apo(a) comparte con el plasminógeno el kringle V y un número variable de kringles IV, Figura 6.

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No todos los kringles IV de apo(a) son iguales entre sí; se clasifican en 10 subtipos diferentes⁸¹;

FIGURA 6

Polimorfismo Estructural de la Apo(a)



Posibles variaciones en el tamaño de Apo(a) dependiendo del número de kringles IV tipo 2

En la molécula de apo(a) cada uno está presente una sola vez y únicamente el kringles IV-2 se presenta un número diferente de veces, lo que da origen a una heterogeneidad estructural y a isoformas de diferente tamaño con pesos moleculares entre 280 y 800 kDa. De todos ellos, el kringles IV-10 es el más parecido al kringles 4 del plasminógeno; el sitio de unión de alta afinidad a la lisina se forma por un polo aniónico Asp-55 y Asp-57 y por un polo catiónico Arg-71 y Arg-35; entre ambos polos hay un microambiente hidrófobo que se forma por la presencia de tres aminoácidos aromáticos Trp-62, Phe-64 y Trp-72. El KIV-10 de la apo(a) difiere del K-4 del plasminógeno, por la presencia de Arg en lugar de Lys en la posición 35. Este kringles tiene un papel preponderante en la unión de Lp(a) a la lisina presente en la fibrina⁸², de esta forma impide que el plasminógeno tenga

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acceso a las redes de fibrina y se active por el activador tisular del plasminógeno^{83, 84, 85, 86}, generándose una insuficiencia fibrinolítica que favorece la aterosclerosis y la trombosis .

Otra fuente de polimorfismo de apo(a) son las glicosilaciones; cada kIV de apo(a) tiene un sitio potencial para una N-glicosilación y, considerando las glicosilaciones interkringles, 30% de cada mol de apo(a) corresponde a carbohidratos: manosa, galactosa, galactosamina, glucosamina y ácido siálico, en relaciones aproximadas de 3:7:5:4:7, respectivamente⁷⁵ .

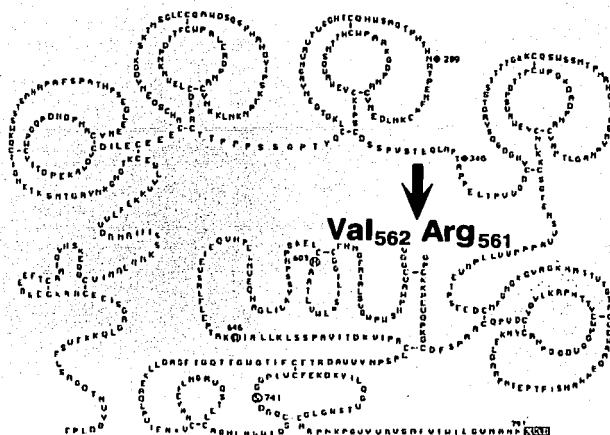
FIBRINOLISIS

En el interior de los vasos sanguíneos la fibrina se deposita para consolidar las reacciones hemostáticas. Es un proceso benéfico que permite reparar y devolver sus funciones al endotelio vascular después de una lesión; sin embargo, cuando la permanencia de la fibrina no es autolimitada, favorece el desarrollo de la aterosclerosis.

El sistema fibrinolítico destruye los depósitos de fibrina en el interior de los vasos sanguíneos, tanto los remanentes de la actividad hemostática como los que se van formando y acumulando durante la evolución de la placa aterosclerosa. Gracias a un balance entre los activadores del plasminógeno y los diferentes tipos de inhibidores de estos activadores, su respuesta se inclina entre el reposo y la actividad fibrinolítica.

Como resultado de la activación fibrinolítica se genera plasmina, Figura 7.

FIGURA 7



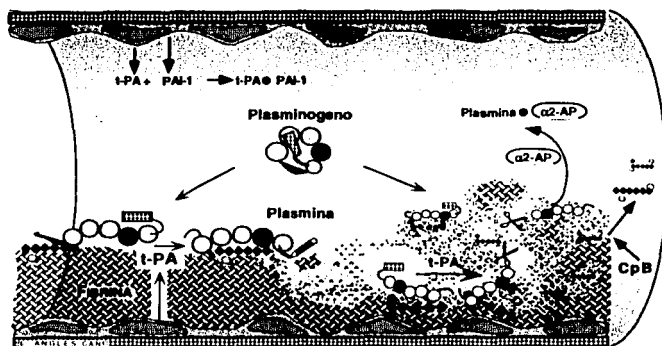
La flecha indica el sitio de escisión del plasminógeno para generar plasmina

La plasmina es una enzima proteolítica que puede ejercer su efecto sobre numerosos substratos plasmáticos. Esto no ocurre así, gracias a que es la propia fibrina la que localiza y señala el sitio de la activación fibrinolítica. Entre sus redes se une el plasminógeno, precisamente en el sitio de alta afinidad o LBS y permite y orienta adecuadamente la unión de su activador tisular (t-PA). Así se genera plasmina en el interior del depósito de fibrina, que actúa escindiéndola y a la vez poniendo al descubierto otros sitios de afinidad, residuos de lisina carboxi-terminales, para amplificar su respuesta. Una vez que se desintegra la fibrina, el mecanismo se detiene, el endotelio vascular no libera más activador. En este punto los remanentes de plasmina y de activador tisular, que alcanzan la

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circulación, se encuentran con sus inhibidores respectivos, la alfa2-antiplasmina y el inhibidor del activador tisular del plasminógeno tipo 1 (PAI-1). Figura 8.

FIGURA 8



Representación esquemática de la fibrinólisis

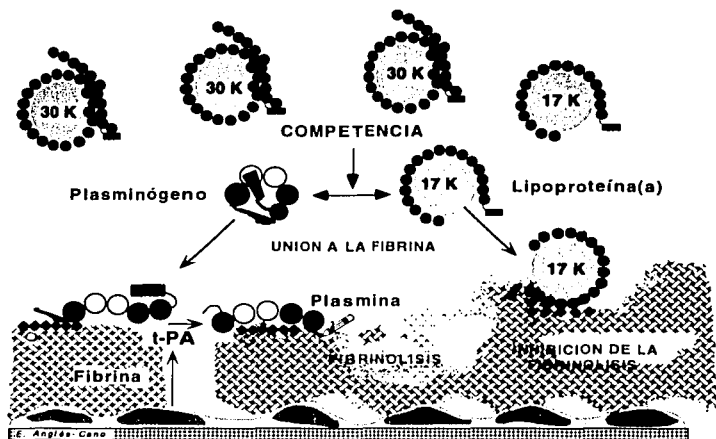
Lp(a) INHIBIDOR COMPETITIVO DEL PLASMINÓGENO

La semejanza entre el plasminógeno y la apolipoproteína (a) permite que las diferentes isoformas de apo(a) compitan por los sitios de afinidad (LBS) de la fibrina con el plasminógeno.

Sin embargo, no todas las isoformas de apo(a) muestran la misma afinidad. Esta depende del tamaño de la isoforma y de su concentración plasmática^{87, 88} Figura 9.

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FIGURA 9

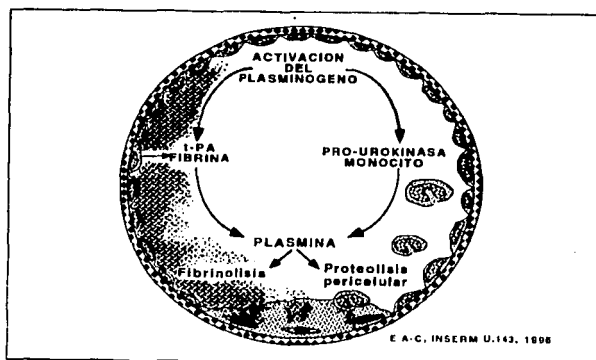


Mecanismo de acción de la lipoproteína Lp(a). El polimorfismo estructural de la apolipoproteína apo(a) influye en el efecto antifibrinolítico de la Lp(a). Las isoformas de talla más pequeña poseen mayor afinidad por la fibrina y producen un efecto más pronunciado.

El plasminógeno y las diferentes isoformas de Lp(a)⁸⁹, compiten también por los residuos de lisina en los receptores de la superficie de las células endoteliales⁹⁰, en los monocitos U937⁹¹, en las plaquetas⁹², en las células mononucleares y en modelos *in vitro* de matrices que simulan la membrana extracelular⁹³. La Figura 10 muestra esquemáticamente la formación de plasmina, tanto sobre la superficie de fibrina como sobre los monocitos.

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FIGURA 10



El plasminógeno se activa tanto sobre la superficie de fibrina como sobre la superficie celulares.

Los dos activadores del plasminógeno: activador tipo tisular activa la fibrinólisis y el activador tipo urocinasa activa la proteólisis pericelular.

Otro mecanismo más que altera el balance del sistema fibrinolítico, es la disminución de la síntesis del activador tisular del plasminógeno (t-PA) y el incremento de la síntesis del inhibidor del activador tisular del plasminógeno (PAI-1) ⁹⁴ que se observa en cultivos de células endoteliales expuestas a Lp(a).

Lp(a) y ATEROGÉNESIS.

La aterosclerosis es un proceso multifactorial y crónico ⁹⁵. Las arterias más afectadas son las de mediano o gran calibre, particularmente en los sitios que están expuestos a las fuerzas de cizalla. Las alteraciones en el metabolismo de los lípidos, las lesiones o alteraciones en el funcionamiento de las células endoteliales y la pérdida de equilibrio entre el depósito de fibrina y la fibrinólisis,

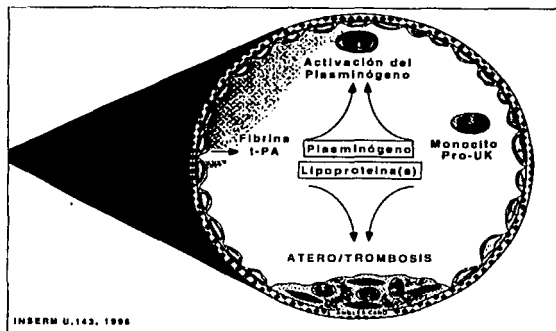
son los principales factores que se identifican como precursores de la placa aterosclerótica. Los dos primeros favorecen el inicio ⁹⁶ mientras que la insuficiencia fibrinolítica interviene principalmente en el crecimiento de la placa aterosclerótica⁹⁷. Las lesiones de la íntima favorecen la migración y la proliferación de células, la acumulación de lípidos, la aceleración en la síntesis de la matriz extracelular y el desequilibrio en la síntesis de factores quimiotácticos y de crecimiento. Todas estas acciones originan las lesiones ateroscleróticas que se clasifican en tres categorías:⁹⁸

La lesión precoz, que se caracteriza por un depósito en la íntima de lípidos (principalmente de baja densidad o LDL), presencia de macrófagos y de células espumosas. En la lesión intermedia existe ya proliferación celular de macrófagos, de células espumosas, de células musculares lisas que provienen de la media y que éstas últimas adquieren la capacidad de sintetizar ciertas proteínas de la matriz extracelular. En las lesiones avanzadas se pueden identificar deformaciones de la pared arterial en las que hay depósitos de lípidos que se rodean de una capa fibrosa formada por proteoglicanos, células espumosas y células musculares lisas. Una fisura en este sitio permite el contacto del factor tisular con el factor VII de la coagulación y se inician los mecanismos procoagulantes que culminan con la formación de un trombo que puede o no obstruir completamente el flujo sanguíneo, y dar un curso acelerado y en ocasiones fatales a estos padecimientos.

La Lp(a) favorece la aterogénesis a través de varios mecanismos; los macrófagos fagocitan a la Lp(a) ⁹⁹ y se depositan en el subendotelio, se transforman en células

espumosas y dan lugar a deformaciones que disminuyen la luz de los vasos sanguíneos.

FIGURA 11

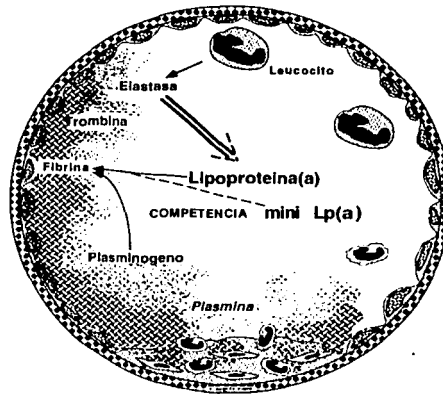


La Lp(a) favorece la aterotrombosis y disminuye la fibrinólisis.

En las células endoteliales de arteria coronaria en cultivo, se observa que la Lp(a) estimula la expresión de moléculas de adhesión vascular 1 (VCAM-1) y de selectina-E, proceso que desencadena la atracción de los macrófagos¹⁰⁰. Otro mecanismo propuesto se relaciona con la disminución de la generación de plasmina que tiene las siguientes consecuencias: a) permanencia prolongada de depósitos de fibrina, con el consecuente incremento en el depósito de colesterol y formación de la placa aterosclerosa y b) disminución de la activación, por hidrólisis parcial con plasmina, del factor de crecimiento transformante β (Transforming Growth Factor o TGF- β), factor que limita el crecimiento de las células del músculo liso vascular.

Se ha propuesto¹⁰¹ que durante los episodios inflamatorios los leucocitos, macrófagos y células T, secretan enzimas proteolíticas. Los fragmentos de Lp(a)/Apo(a) que se forman bajo estas circunstancias, pueden tener una mayor capacidad aterosclerosa como lo representa la figura 12

Figura 12



Por la acción de enzimas proteolíticas se forman partículas de Lp(a) con mayor actividad aterosclerótica.

Lp(a) INHIBE LA FIBRINOLISIS

La elevada concentración plasmática de Lp(a) no siempre interfiere con la fibrinólisis normal. Las isoformas de apo(a) muestran diferente actividad antifibrinolítica, por lo que se debe subrayar la importancia de considerar la actividad antifibrinolítica de las isoformas en la predicción de la enfermedad cardiovascular¹⁰².

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Se han llevado a cabo diversas estrategias de estudio *in vitro*. Empleando un modelo de fibrina en fase sólida ^{103, 104, 105, 106, 107}, ha sido posible identificar de manera sensible y específica la inhibición de la fibrinólisis por las isoformas de apo(a).

Estos estudios han permitido concluir el mecanismo y las diferentes variables que intervienen en la inhibición competitiva del plasminógeno, por diferentes isoformas nativas de Lp(a) o por formas recombinantes de apo(a). Estos trabajos muestran que:¹⁰⁸

- 1) La afinidad de Lp(a) por la fibrina muestra una relación inversa con el tamaño de las isoformas de apo(a), que pueden tener constantes de disociación (Kd) entre 50 y 500 nmol/L ⁸⁷
- 2) Tanto el plasminógeno como la Lp(a) compiten por los mismos sitios de unión, lo que corresponde a un mecanismo de inhibición competitiva saturable.
- 3) El potencial antifibrinolítico de Lp(a) depende de la afinidad y de la concentración de cada una de las 2 isoformas de apo(a) presentes en el plasma¹⁰⁸.

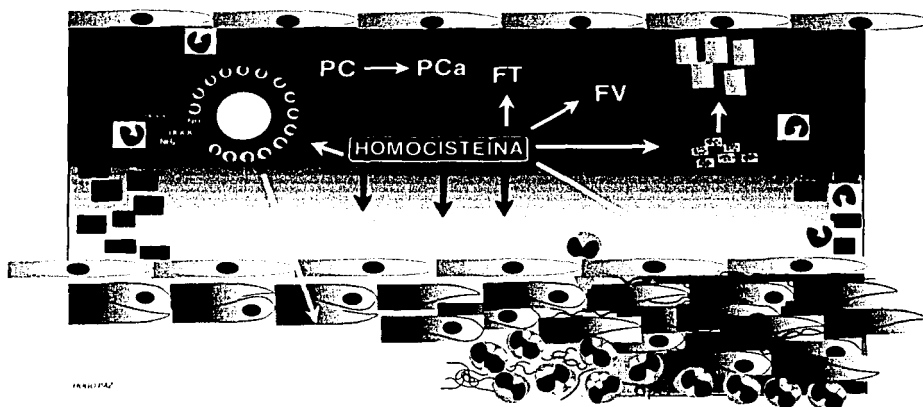
Lp(a) Y HOMOCISTEINEMIA

La hiperhomocisteinemia se asocia con un incremento de las enfermedades trombóticas y aterosclerosas¹⁰⁹. La trombosis se favorece porque altera varios mecanismos de regulación antitrombótica de la hemostasia.

Los mecanismos fisiológicos que regulan la hemostasia permiten, junto con el sistema fibrinolítico, que la actividad pro- y anti-coagulante mantenga un equilibrio.

Entre los mecanismos anticoagulantes naturales se encuentra: la antitrombina III, que inhibe directamente a los factores enzimáticos de la coagulación activados (IIa, IXa, Xa) ; el sistema de la Proteína C y su cofactor la proteína S, que inhibe a los cofactores activados de la coagulación (Va, VIIIa), y el inhibidor del factor tisular.

Figura 13



PC, proteína C; PCa, proteína C activada; FT, factor tisular; FV, cofactor V de la coagulación. PGI₂, prostaciclina o prostaglandina I₂, NO, óxido nítrico; Tm, trombomodulina.

La homocisteína incrementa la actividad del factor tisular, disminuye la expresión y la actividad de la trombomodulina que es indispensable para la activación de la proteína C, disminuye la actividad anticoagulante de la antitrombina III así como la unión del activador tisular del plasminógeno a su receptor en la superficie de las células (anexina II)¹¹⁰ y también el incremento de la concentración de homocisteína plasmática, se refleja como un incremento en la unión de Lp(a) a la fibrina ¹¹¹., Figura 13

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Lp(a) e INHIBIDOR DEL FACTOR TISULAR

Otro mecanismo recientemente propuesto¹¹² por el que Lp(a) favorece la trombosis intravascular, contempla la unión directa, dependiente de la concentración, al inhibidor del factor tisular. El inhibidor del factor tisular (IFT) modula la actividad catalítica del factor VII de la coagulación y es reconocido como un mecanismo fisiológico anticoagulante. El inhibidor del factor tisular se localiza en las células endoteliales y la interacción Lp(a)/IFT sugiere que existe un mecanismo procoagulante que se localiza en la placa aterosclerosa.

ESTUDIOS CLÍNICOS

Bajo esta perspectiva, se realizaron numerosos estudios epidemiológicos, que describen una asociación positiva entre la elevada concentración plasmática de Lp(a) y un incremento en las enfermedades cerebrovasculares¹¹³, cardiovasculares, la reestenosis de puentes coronarios, la reoclusión de angioplastías y el desarrollo prematuro de aterosclerosis asociada a altas concentraciones de LDL y/o a bajas concentraciones de HDL^{114, 115}. La mayoría de estudios prospectivos confirman estos resultados^{116, 117, 118, 119, 120, 121, 122}, aún cuando no existe un acuerdo en el límite de corte para considerar una concentración plasmática normal; algunos autores consideran que es de 20 mg/dL¹²¹, otros hasta 30 mg/dL¹¹⁶ y que el empleo de anticuerpos diferentes, monoclonales o policlonales, genera divergencias en los resultados⁵.

Sin embargo, otros estudios no encuentran asociación entre la Lp(a) y la enfermedad arterial coronaria^{123, 124}. Esta discrepancia en los resultados podría

ser el reflejo de la gran heterogeneidad estructural de la molécula de apo(a), que se traduce en una heterogeneidad funcional como inhibidor competitivo del plasminógeno por la fibrina, disminuyendo la formación de plasmina.^{7, 125.}

Craig y col ¹²⁶ hicieron un meta-análisis de los diferentes estudios prospectivos que estudian a la Lp(a) como factor de riesgo de la enfermedad arterial coronaria en población caucásica, poniendo especial atención en los criterios de selección de los trabajos a fin de que los resultados de los diferentes estudios fueran comparables. La tabla 3 muestra que, los individuos que desarrollaron enfermedad arterial coronaria tenían una mayor concentración de Lp(a) que los controles^{116, 117, 118, 119, 124, 127, 128, 129, 130, 131, 132, 133}. Sugieren que la variabilidad de los resultados puede ser efecto del manejo pre-analítico de la muestra, en especial a la temperatura que se almacenó la muestra. Concluyen que la Lp(a) es un factor prospectivo de riesgo independiente en la enfermedad isquémica coronaria.

TABLA 3

Ref	País	N Casos	N Controles	Cociente [Lp(a)] mg/L Casos/Controles	ES
128	Finlandia				
	Hombres	97	148	0.68	0.050
	Mujeres	97	121	1.24	0.049
118	Alemania	107	5124	2	0.038
129	Alemania	33	828	1.8	0.084
130	Reino Unido	49	192	1.4	0.045
131	Finlandia	138	130	1.18	0.044
132	Dinamarca	74	190	1.32	0.078
120	EUA	233	390	1.22	0.021
125	EUA	296	296	1	0.018
134	EUA				
	Hombres	90	90	1.97	0.132
	Mujeres	44	44	1.34	0.179
117	Suiza	26	109	1.61	0.070
133	Islandia	104	1228	1.35	0.034
119	Reino Unido	229	1145	1.54	0.033

Craig et al. Clinical Chemistry 1998; 44: 2301-2306

JUSTIFICACION DEL ESTUDIO

Por la inconsistencia en los resultados entre los diferentes estudios prospectivos para señalar a la Lp(a) como un factor de riesgo en la enfermedad arterial vascular, es importante evaluar el impacto de la Lp(a) como factor de riesgo a través de su capacidad para inhibir la fibrinolisis. Por esta razón es necesario el empleo de un método funcional que identifique la unión de Lp(a) a la fibrina en los diferentes sujetos de estudio.

HIPÓTESIS

La cantidad de apo(a) unida a la fibrina, las isoformas de Lp(a) de bajo peso molecular, se encuentran aumentadas en las muestras plasmáticas de sujetos con enfermedad arterial vascular en comparación con sujetos sin ésta.

OBJETIVO DEL ESTUDIO

- 1.-Determinar cuál es la cantidad de apo(a) unida *in vitro* a la fibrina, en las muestras plasmáticas de sujetos con enfermedad arterial vascular, en comparación con sujetos sin ella.
- 2.- Analizar qué tipo de isoformas de Lp(a) se unen en mayor proporción a la fibrina en las muestras plasmáticas de sujetos con enfermedad arterial vascular, en comparación con sujetos sin ella .
- 3.- Determinar si existe una relación entre la concentración plasmática de Lp(a) y la cantidad de apo(a) unida *in vitro* a la fibrina, en las muestras plasmáticas de sujetos con enfermedad arterial vascular, en comparación con sujetos sin ésta.

DISEÑO DEL ESTUDIO

TRANSVERSAL ANALITICO

Por el control de la maniobra: observación.

Por la captación de la información: prolectivo.

Por la medición de la maniobra en el tiempo: transversal.

Por la presencia de un grupo control: comparativo.

Por la asignación de sujetos al grupo de estudio y control : no aleatorio.

Por la capacidad en la aplicación y evaluación de las maniobras: abierto y ciego simple.

POBLACIÓN DEL ESTUDIO

Individuos con enfermedad isquémica coronaria que acudieron a la consulta externa del Instituto Nacional de Cardiología, que cumplieron con los criterios de selección y consintieron, por escrito, en participar en el estudio.

Individuos con enfermedad vascular cerebral que acudieron a la consulta externa del Instituto Nacional de Neurología y fueron remitidos al Instituto Nacional de Cardiología, cumplieron con los criterios de selección y consintieron, por escrito, en participar en el estudio.

Población sin antecedentes de enfermedad isquémica que acudieron a donar sangre al Instituto Nacional de Cardiología.

CASOS

Criterios de inclusión:

- Antecedente de enfermedad vascular coronaria, infarto agudo del miocardio, angina estable o inestable.
- Antecedente de infarto cerebral de origen trombótico.
- En caso de ser necesario, los enfermos estaban bajo tratamiento con fármacos hipolipemiantes, antihipertensivos, hipoglucemiantes o antiagregantes plaquetarios.

Criterios de no inclusión:

- ▶ Evento trombótico en el mes anterior al estudio.
- ▶ Hipertensión sin control.
- ▶ Déficit neurológico con demencia
- ▶ Embolia cerebral de origen cardíaco o consecuente con una angiografía coronaria.
- ▶ Enfermedad hepática aguda o crónica.
- ▶ Enfermedad renal aguda o crónica.
- ▶ Enfermedad hemorrágica.
- ▶ Procedimientos quirúrgicos, 3 meses previos al estudio.
- ▶ Procesos infecciosos graves, 2 meses previos al estudio.

CONTROLES

Donadores de sangre sin antecedentes de enfermedad isquémica cerebral o coronaria.

Definición operativa de los criterios de exclusión

1) Enfermedad hepática

Incremento en: bilirrubina directa > 1.5 mg/dL
 bilirrubina indirecta > 1.0 mg/dL
 fosfatasa alcalina > 70 U/l
 transaminasa oxalacética > 25 U/l
 transaminasa pirúvica > 20 U/l

Tiempo de protrombina prolongado >20 %

Cuando esté consignado en el expediente por médico del Instituto Nacional de Cardiología.

2) Enfermedad renal

Creatinina > 1.5 mg/dL

Cuando esté consignado en el expediente por un médico del Instituto Nacional de Cardiología.

3) Enfermedad hemorrágica

Tiempo de protrombina prolongado > 20 %

Tiempo de tromboplastina parcial prolongado > 20 %

Fibrinógeno < 1.5 g/L

Cuando esté consignado en el expediente por un médico del Instituto Nacional de Cardiología.

4) Trombosis

Dímeros D-D positivos. Cuando esté consignado en el expediente por un médico del Instituto Nacional de Cardiología.

DEFINICIÓN OPERATIVA DE VARIABLES

Variables independientes

Enfermedad aterotrombótica, variable nominal.

Categorías:

1 Enfermedad isquémica coronaria (CI). Diagnóstico clínico, electrocardiográfico o por angiografía coronaria. Se considera positivo cuando esté consignado en el expediente por un cardiólogo del Instituto Nacional de Cardiología.

2 Enfermedad vascular cerebral (EVC). Diagnóstico clínico, o por estudio doppler o tomografía computarizada. Se considera positivo cuando esté consignado en el expediente por un neurólogo del Instituto Nacional de Neurología y se remita para inclusión al estudio en el Instituto Nacional de Cardiología.

Variables dependientes

1 Concentración plasmática de Lp(a). Se estimó en mg/dL. Es una variable continua con escala de medición de proporción.

2 Lp(a) unida a la fibrina. Se estimó la cantidad de apo(a) unida a las placas de fibrina extrapolando cada resultado a una curva de calibración que se construye con un estándar de recombinantes apo(a) desde 10 a 34 kringles. Variable continua con escala de medición de proporción.

3 Isoformas de Lp(a). Se reconocieron comparando su migración contra un estándar de recombinantes apo(a) desde 10 a 34 kringles. Variable discreta en escala de medición nominal.

Variables de confusión

1 Colesterol total, triacilgliceroles, lípidos de baja densidad, lípidos de alta densidad, variables continuas en escala de medición de proporción, (mg/dL).

2 Antecedentes heredo-familiares, variable nominal.

3 El fibrinógeno, el factor VII de la coagulación, el volumen de sedimentación globular, los leucocitos, son variables continuas con escala de medición de proporción.

Variables modificadoras, variables continuas en escala de medición de proporción.

1 Presión arterial. Se registrará en mm Hg con el individuo en posición sedente, después de 20 min. de reposo.

2 Circunferencia de la cintura. Se medirá con cinta métrica , se registrará en cm.

3 Circunferencia de la cadera. Se medirá con cinta métrica , se registrará en cm.

4 Índice de masa corporal. Se calculará dividiendo el peso entre el cuadrado de la talla (Kg/m^2).

Variables universales, variables continuas, escala de medición de proporción.

1) La edad se registrará en años.

2) El peso se registrará en Kg

3) La talla se medirá de pie y sin calzado con una cinta métrica colocada en la pared. Se registrará en cm .

DESCRIPCION OPERATIVA DEL ESTUDIO

Se revisarán los expedientes un día previo a su consulta y a los candidatos se les informará del estudio y se les invitará a participar previa carta de consentimiento informado.

Los enfermos que aceptaron ingresar al estudio se sometieron a exámenes de escrutinio (si no se han practicado en el transcurso del último año), para descartar enfermedades hepática, renal o trastornos hematológicos.

A los enfermos que cumplieron con los criterios de selección se les aplicó un cuestionario que incluyó una ficha de identificación, antecedentes familiares, antecedentes personales.

ASPECTOS ÉTICOS

Ya que los participantes en el estudio se sometieron a una punción venosa adicional a su manejo habitual, implicando un riesgo mayor al mínimo, se les informó sobre las características y objetivos del estudio.

MATERIAL, MÉTODOS Y PROCEDIMIENTOS

GRUPO DE ESTUDIO

Estudiamos 248 individuos, 157 enfermos y 91 controles. 105 enfermos con cardiopatía isquémica, infarto del miocardio, o con angina estable/inestable. 52 enfermos con enfermedad vascular cerebral de origen trombótico, Tabla 4

TABLA 4

Categorías	Número Individuos	Porcentaje
EVC	52	21
CI	105	42.3
Infarto del miocardio	70	28.2
Angina estable/inestable	35	14.1
Controles	91	36.6
Total	248	100

MUESTRA DE SANGRE

Se obtuvo de cada individuo con 12 horas de ayuno, después de permanecer sentado 20 minutos, una muestra de 15 mL de sangre por punción venosa con una jeringa estéril desechable. La muestra de sangre se dividió en 3 alícuotas iguales. La primera se colocó en un tubo A de plástico que contenía citrato de sodio (3.8%) en una relación 1:9 como anticoagulante, para determinar los

tiempos de coagulación, fibrinógeno y dímeros D-D. La segunda se colocó en un tubo B de vidrio, sin anticoagulante para la determinación de química sanguínea. La tercera se colocó en un tubo de plástico C que contenía EDTA 3 mM como anticoagulante para determinar concentración de Lp(a), isoformas, grado de unión a la fibrina. Los tubos se centrifugan a 2000 x g durante 20 min. a 4°C.

El plasma del tubo C se separó en varias alícuotas. A las alícuotas destinadas al estudio de Lp(a) se adicionaron inhibidores de proteasas (Trasylol®, Bayer Farma, Puteaux, Francia). Las muestras se congelaron a -70°C hasta efectuar las pruebas, máximo 6 meses. Se analizaron en bloques con el fin de disminuir la variabilidad ínter ensayo.

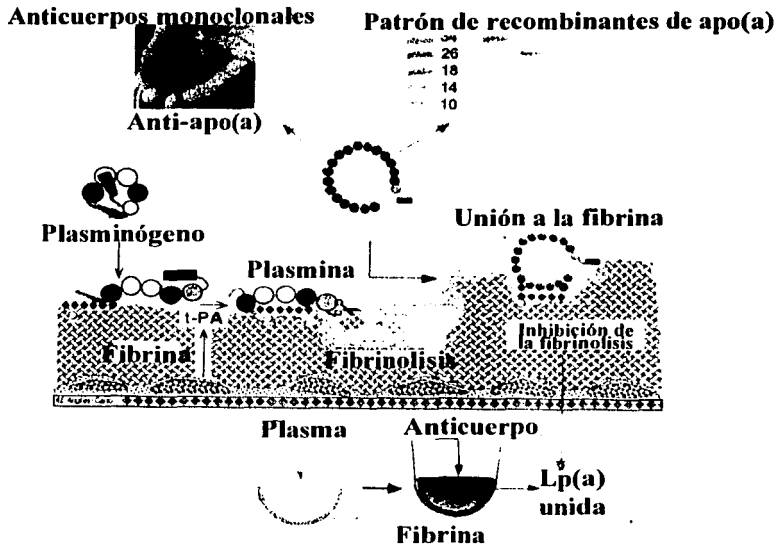
UNIÓN DE APO(a) A LA FIBRINA

Se pusieron a punto las técnicas para obtener un modelo experimental que permite identificar *in vitro* la unión de la Lp(a) a la fibrina.

El modelo distingue la inhibición competitiva entre la Lp(a) y el plasminógeno como se muestra en la Figura 14, que se traduce en una incompetencia fibrinolítica.

La malla de fibrina que se forma en los pozos de la placa presenta sitios expuestos de lisina de 6.8 Å, a los que selectivamente tienen acceso y pueden unirse moléculas que presenten una estructura globular que se genera en el interior de los kringles.

FIGURA 14



Representación esquemática del método para identificar la unión de apo(a) a la fibrina

Este modelo tiene las siguientes características:

Superficie de fibrina $121 \text{ mm}^2 = 410 \text{ fmol de fibrina/cm}^2$. La superficie que se cubre con un volumen de $50 \text{ }\mu\text{l}$ corresponde a 88 mm^2 .

El fibrinógeno no se desprende de la placa de PVC gracias a la presencia de polímeros de glutaraldehído que lo fijan y le proporcionan una flexibilidad adecuada para permitir el acceso de las proteínas afines.

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La presencia de fibrina en las placas se puso en evidencia con un anticuerpo monoclonal, Y18, cuyo epítipo se sitúa en el fragmento 1-51 de la cadena A α del fibrinógeno; la reactividad del anticuerpo disminuye a medida que se incrementa el tiempo de reacción entre el fibrinógeno y la trombina. Con la ayuda del anticuerpo FDP-14, (colaboración con el Dr Nieuwenhuizen, Gaubius Institute, Leyden, Holanda), se identificó el fragmento E, producto de fragmentación del fibrinógeno¹³⁴ y el anticuerpo DD3B6/22 identificó los dímeros D-D; producto de fragmentación de la fibrina, (proporcionado por el Dr Rylatt, Agen Biomedical LTD, Brisbane, Australia).

Para hacer las determinaciones de unión de apo (a) a la fibrina, se lavaron las placas de fibrina tratadas con plasmina con amortiguador A (0.05 mmol/L de fosfato de sodio, pH 7.4, 0.08 nmol/L cloruro de sodio, 0.01 % thymerosal, 0.01% Tween 20, 2mg/mL albúmina bovina).

Se depositó por duplicado en cada pozo 50 μ l de plasma (dilución 1:16 en amortiguador A + 40 mg/mL albúmina bovina)

Se incubó durante 12 h a 4°C con la finalidad de permitir el equilibrio entre la unión de plasminógeno/apo(a) a la fibrina.

Se retiraron las proteínas que no se unieron lavando la placa con amortiguador A.

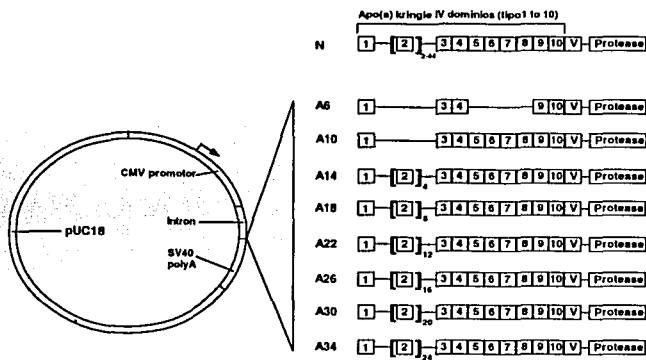
Se adicionó 50 μ l de un anticuerpo monoclonal anti-apo(a)¹³⁵ conjugado con peroxidasa.

Se incubó la placa 2 h a 37°C

Se lavó la placa con amortiguador A y se adicionan 50 µl de una solución de 1 mg/mL de ácido 2,2'-azino-bis(3-etilbenzotiazolina)-6 sulfónico (ABTS, Boehringer Mannheim) para revelar la coloración.

El cambio en absorbencia (▲ A 405nm/min) se midió en un lector de placas (MR 500, Dynatech). La cantidad de apo(a) unida a la fibrina se obtuvo extrapolando los valores obtenidos a una curva patrón elaborada con un calibrador de referencia que contenía cantidades equimoleculares de isoformas recombinantes de apo(a) de 10 a 34 kringle23, Figura 15.

FIGURA 15



Representación esquemática de las isoformas de apo(a)

CONCENTRACIÓN PLASMÁTICA DE Lp(a)

Método inmunonefelométrico, reactivos y equipo de Beckman Co, Palo Alto Ca.

EUA.

ISOFORMAS DE Apo(a)

Método de inmunoelectroforesis con transferencia en papel de nitrocelulosa que comprende los siguientes puntos:

- ▶ Reducción de los enlaces sulfhidrilo que unen a la apolipoproteína (a) con el resto del núcleo lipoproteico que conforma a la lipoproteína (a).
- ▶ Depósito y migración de las muestras en gel de acrilamida.
- ▶ Transferencia de las proteínas del gel a papel de nitrocelulosa.
- ▶ Las isoformas de apo(a) se localizan con un anticuerpo monoclonal dirigido contra apo(a) marcado a la peroxidasa cuya presencia se revela con 4 cloronaftol.

Las proteínas presentes en el plasma son identificadas tomando como referencia un estándar que contiene diferentes isoformas recombinantes de apo(a) de masa molecular conocida²³, Figura 13.

Cada imagen de inmunotransferencia se digitalizó (Molecular Analyst TM/PC) y se analizó la densidad de la banda (Densitometric scanning Bio Rad GS-670).

Cada uno de los datos densitométricos de las isoformas (una o dos) se sumó y consideró como el 100%.

ANÁLISIS ESTADÍSTICO

Los promedios, medias, desviaciones estándar y rangos intercuartiles se calcularon para todas las variables.

Los límites de los cuartiles se calcularon tomando en consideración a toda la población (CI, EVC y controles). Las diferencias en la proporción de enfermos y

controles en cada cuartil se analizó con la prueba de chi-cuadrada para variables discretas.

Para las variables continuas, las comparaciones entre los grupos fueron evaluadas con análisis paramétrico o no paramétrico de acuerdo con la distribución de la variable.

Para las variables no paramétricas se empleó la prueba de LSD (least significant difference test) o la prueba de U de Mann-Whitney.

Para eliminar la posible interferencia de las diferentes edades entre los grupos se empleó el análisis de covarianza.

Los resultados de la distribución de la concentración de Lp(a) y de la unión de Lp(a) a la fibrina, se compararon con la prueba de Kolmogorov-Smirnov.

Para determinar la asociación entre la concentración de Lp(a), la unión de Lp(a) a la fibrina y los fenotipos de las isoformas de apo(a) con la enfermedad isquémica aterotrombótica, se comparó el número de individuos entre el primero y el último cuartil. Las diferencias se estimaron por tablas de contingencia, razones de momios e intervalos de confianza 95%.

Todas las pruebas estadísticas se hicieron empleando el programa SPSS-10.1 (EUA).

RESULTADOS

El grupo de pacientes con evidencia clínica de enfermedad isquémica, no exhibe diferencias significativas con el grupo control respecto al género y al índice de masa corporal Tabla 4 . Sin embargo la edad difiere significativamente en ambos grupos de enfermos con respecto al control, motivo por el cual, aún cuando la concentración plasmática de Lp(a) no se modifica con la edad, los resultados que se muestran en la Tabla 5 se ajustaron a esta variable (análisis ANCOVA).

TABLA 5

Características demográficas de la población estudiada

	CONTROL	EVC	CI
EDAD (años)	40.2±13.1	53.4±10.5*	62.5±10.6*
IMC Kg/m ²	26.3±5.6	27.0±3.5	27.2±3.4
Cintura/Cadera	0.88±0.069	0.93±0.056	0.94±0.44

*ANOVA $p < 0.001$, IMC=índice de masa corporal

TABLA 6

Concentración plasmática de Lp(a) y Unión de apo(a) a la fibrina

Variable	Control	EVC	CI
Concentración de Lp(a) mg/dL	16.3±18	20.3±22.8	20.5±25.29
Unión de apo(a) a la fibrina nM	0.158±0.09	0.268±0.15*	0.155±0.12

*ANCOVA $p < 0.0001$ contra grupo control y CI

La unión de apo(a) a la fibrina en el grupo de EVC se incrementó de manera muy significativa ($p < 0.001$) comparada tanto con el resultado del grupo control como con el del grupo de CI.

La Tabla 7 ilustra la distribución expresada en porcentaje tanto en enfermos como en los controles en cada cuartil. Las variables que se muestran son: la concentración plasmática de Lp(a), la unión de apo(a) a la fibrina y los fenotipos de apo(a).

En el grupo control tanto la concentración de Lp(a) como la unión de apo(a) a la fibrina, tienen una distribución no paramétrica, el mayor porcentaje de individuos se encuentra en los primeros cuartiles.

El grupo de CI muestra una distribución bimodal, el mayor porcentaje de individuos se encuentra en los dos primeros cuartiles y no disminuye gradualmente como en el caso del grupo control.

El grupo de EVC difiere notablemente tanto del grupo de CI como del grupo control, el mayor porcentaje de individuos se encuentra en los últimos cuartiles tanto para la concentración de Lp(a) como para la unión de apo(a) a la fibrina.

TABLA 7

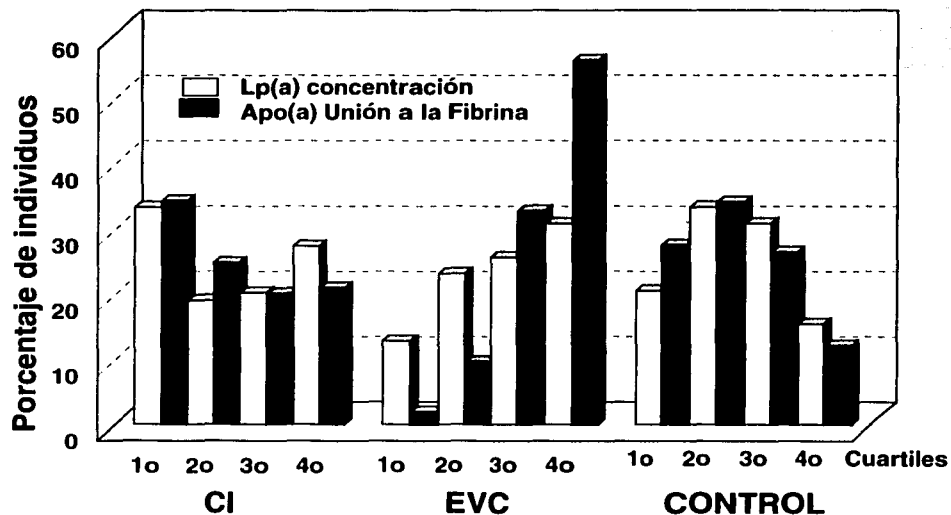
Porcentaje de individuos en cada cuartil para las variables de concentración plasmática de Lp(a), Unión de apo(a) a la fibrina y fenotipos de apo(a). El fenotipo 1 es la isoforma con mayor concentración en los sujetos heterocigotos.

Cuartil	1	2	3	4	P*
Concentración Lp(a) mg/dL	<6.1	6.1-13.8	13.9-22.45	>22.45	
CI (%)	33.3	19	20.2	27.4	0.074
EVC (%)	12.8	23.1	25.6	38.5	<0.001
CONTROL (%)	20.5	33.3	30.8	15.4	0.009
Unión de Apo (a) (nM)	<0.098	0.098-0.162	0.163-0.231	>0.231	
CI (%)	34.3	24.8	20	21	0.089
EVC (%)	1.9	9.6	32.7	55.8	<0.001
CONTROL (%)	27.5	34.1	26.4	12.1	0.003
Apo(a) fenotipo 1 (K4)	<20	20-23	23.1-26	>26	
CI (%)	20.8	35.1	28.6	15.6	0.006
EVC (%)	28.9	18.4	21.1	31.6	0.102
CONTROL (%)	23.1	18.5	32.3	26.2	0.187
Apo(a) fenotipo2 (K4)	<16.5	16.6-19	19.1-21	>21	
CI (%)	28.3	28.2	17.4	26.1	0.217
EVC (%)	24.1	17.2	31	27.6	0.118
CONTROL (%)	20.6	20.6	29.4	29.4	0.332

P <0.05 prueba de chi-cuadrada*

La Figura 16 muestra gráficamente los datos de la Tabla 7

FIGURA 16



La Tabla 8 muestra los resultados de la prueba de razón de momios y los intervalos de confianza 95. Las variables de concentración plasmática de Lp(a), unión de apo(a) a la fibrina y fenotipos de apo(a) en los tres grupos de estudio CI, EVC y control.

TABLA 8

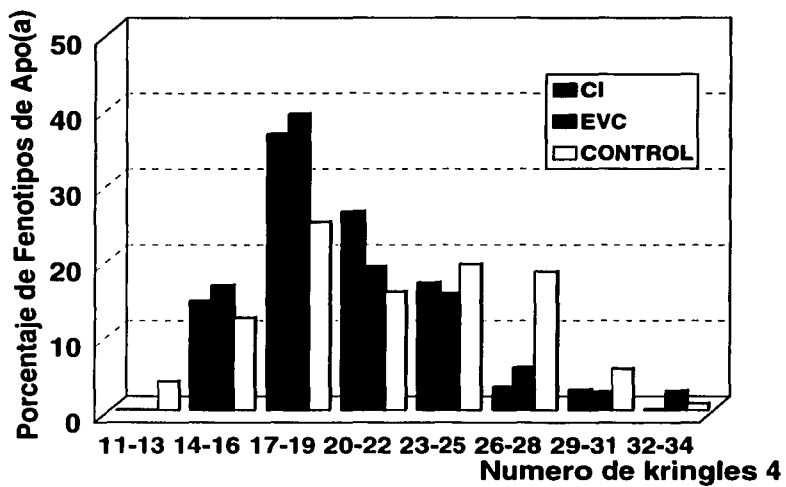
Razón de momios e intervalos de confianza 95% para las variables de concentración plasmática de Lp(a), Unión de apo(a) a la fibrina y Fenotipos de apo(a)

	Cuarto Cuartil	Primer Cuartil	RM (95 % IC)
Lp(a) (mg/dL)			
CI (%)	27.4	33.3	1.10 (0.43-2.78)
EVC (%)	38.5	12.8	4 (1.13-14.09)
CONTROL (%)	15.4	20.5	
Unión de Apo(a) (nM)			
CI (%)	21.0	34.3	1.39 (0.57-3.37)
EVC (%)	55.8	1.9	65.9 (7.95-546.8)
CONTROL (%)	12.1	27.5	
Apo(a) Fenotipo 1			
CI (%)	15.6	20.8	0.66 (0.24-1.84)
CONTROL (%)	26.2	23.1	
EVC (%)	31.6	28.9	0.96 (0.33-2.84)
Apo(a) Fenotipo 2			
CI (%)	26.1	28.3	0.64 (0.19-2.24)
EVC (%)	27.6	24.1	0.80 (0.20-3.25)
CONTROL (%)	29.4	20.6	

En el grupo de enfermos estudiados no se observa una asociación significativa entre la cardiopatía isquémica y la concentración de Lp(a) o de la unión de apo(a) a la fibrina. En contraste, el grupo de EVC muestra diferencias muy significativas tanto de la concentración de Lp(a) como de la unión de apo(a) a la fibrina. También se distingue una tendencia de asociación entre las enfermedades isquémicas con las isoformas de bajo peso molecular (aunque por el número de individuos, la observación no es significativa).

Empleando un método de análisis que relaciona la contribución de cada uno de los fenotipos a la concentración total de Lp(a), se aprecia que las isoformas, (Figura 17), menores a K22 son más frecuentes en la enfermedad isquémica coronaria y en la enfermedad vascular cerebral, que en los controles (76.86 %, 74.32 % y 56.3 %, respectivamente). El análisis nos permitió distinguir que los fenotipos K17-K19 son los más frecuentes tanto en la enfermedad isquémica como en la población control.

FIGURA 17



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DISCUSIÓN Y CONCLUSIONES

La afinidad de Lp(a) por la fibrina muestra una relación inversa con el tamaño de las isoformas de apo(a), que pueden tener constantes de disociación (K_d) entre 50 y 500 nmol/L⁸⁷; y con la concentración de Lp(a).

La relación entre la estructura y la actividad antifibrinolítica de la Lp(a) ha permitido identificar que cada isoforma tiene una afinidad diferente por la fibrina, muestra una relación inversa con el tamaño y por lo tanto, concentraciones totales similares pueden exhibir diferente potencial anti-fibrinolítico y atero-trombótico. En acuerdo con estos hallazgos, recientemente se publicó¹³⁶ que la presencia de isoformas de bajo peso molecular es un indicador importante para la evolución de la aterosclerosis estenótica.

Hasta la fecha ha sido difícil establecer el riesgo asociado entre la Lp(a) y la enfermedad isquémica coronaria y la enfermedad cerebral arterial. El motivo principal ha sido la ausencia de un método funcional que refleje la actividad antifibrinolítica y atero-trombótica, consecuencia de la presencia de los diversos fenotipos y de la concentración de cada uno de ellos.

Aunado a este inconveniente, los diferentes estudios epidemiológicos se han hecho midiendo la concentración plasmática de Lp(a) en ausencia de un método y de una referencia internacional, lo que no ha permitido obtener resultados comparables y las diferencias que provienen del polimorfismo genético han sido difíciles de identificar. Dos estudios prospectivos han sido particularmente

importantes en los que no fue posible distinguir diferencias significativas entre la población isquémica coronaria y la población control.

Sin embargo, a pesar de estos antecedentes, un meta-análisis en donde se analizaron 27 estudios prospectivos, contrastando los resultados de Lp(a) plasmática en el tercio superior contra los del tercio inferior, concluyó que el riesgo relativo de padecer una enfermedad coronaria si se tiene una concentración elevada de Lp(a) es de 1.6 (IC 95% 1.4-1.8, $p < 0.0001$)¹³⁷

Esta nueva estrategia para estudiar a la Lp(a) nos ha permitido identificar que no en todos los individuos representa el mismo riesgo patogénico. La originalidad del método reside en su capacidad para señalar a los sujetos en los cuáles la presencia de una determinada isoforma de apo(a) puede disminuir su capacidad fibrinolítica. Muestra un perfil fibrinolítico que depende de la presencia de Lp(a).

Bajo las condiciones del método el mecanismo antagonista competitivo de la Lp(a) sólo puede modificarse por la presencia de análogos de la lisina⁸⁷

Nuestro sistema experimental simula el comportamiento del plasminógeno y de la Lp(a), cuando, en condiciones fisiológicas, se enfrentan a la fibrina.

Este trabajo de tesis aporta evidencia de: a) que el efecto antifibrinolítico de la Lp(a) tiende a asociarse (en nuestro trabajo sin significado estadístico) a las isoformas de bajo peso molecular, b) que los enfermos con enfermedad vascular cerebral tienen una alta asociación con la unión de apo(a) a la fibrina, 65.9 (7.95-546.8).

Este inesperado resultado, concuerda con la hipótesis propuesta por Jürgens y col¹³⁸ de que la Lp(a) es el factor de riesgo más importante para la enfermedad cerebrovascular.

Aún más, nuestros resultados sugieren que existen mecanismos antitrombóticos diferentes entre la arteria coronaria y la cerebral en los enfermos estudiados.

Recientemente, se propuso un modelo que aporta evidencias de las diferencias pro- y anti-coagulantes, entre las células endoteliales de un lecho vascular a otro¹³⁹. Confirma la existencia de diferencias regionales en el balance hemostático.

La trombomodulina es una proteína que está más involucrada en mantener el balance hemostático en los vasos de los pulmones y del corazón que en los vasos del hígado. Mientras que los activadores del plasminógeno, tipo tisular y tipo urocinasa son importantes para mantener el balance hemostático en los tres órganos.

En el cerebro, no se conocen cuáles son los mecanismos fisiológicos que tienen un mayor peso para mantener el balance hemostático. Los resultados de este trabajo de tesis, que mostraron una asociación muy grande entre la unión de la Lp(a) a la fibrina y la enfermedad cerebrovascular, podrían ser un factor que permita orientar estudios futuros para esclarecer los mecanismos fisiopatológicos de la aterotrombosis en este lecho vascular.

PERSPECTIVAS

La mayoría de los diferentes estudios epidemiológicos resaltan la asociación entre la concentración plasmática de la lipoproteína (a) y el riesgo de padecer enfermedad cardiovascular y cerebrovascular, sobre todo cuando se asocian otros factores de riesgo como son el colesterol de lipoproteínas de baja densidad y el tabaquismo. Pero es importante resaltar que existen diferencias en la concentración y en la presencia de isoformas en poblaciones genéticamente diferentes. Identificar el comportamiento de inhibición de la fibrinólisis por las diferentes isoformas de Lp(a) representa una estrategia de estudio que podrá permitir identificar a los individuos y a los grupos étnicos con incompetencia fibrinolítica por la presencia de Lp(a).

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TESIS CON
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FUNCTIONAL APPROACH TO INVESTIGATE Lp(a) IN ISCHAEMIC HEART AND CEREBRAL DISEASES

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Running head: Assay to detect Lp(a) fibrin binding capacity

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Abstract

Background. Lp(a), a major cardiovascular risk factor, contains a specific apolipoprotein, apo(a), which by virtue of structural homology with plasminogen inhibits the formation of plasmin, the fibrinolytic enzyme. A number of clinical reports support the role of Lp(a) as a cardiovascular or cerebral risk factor, and experimental data suggest that it may contribute to atherothrombosis by inhibiting fibrinolysis.

Design. A well characterized model of a fibrin surface and an apo(a) specific monoclonal antibody were used to develop a functional approach to detect pathogenic Lp(a). The assay is based on the competitive binding of Lp(a) and plasminogen for fibrin, and quantifies fibrin-bound Lp(a). High Lp(a) binding to fibrin is correlated with decreased plasmin formation. In a transversal case-control study we studied 248 individuals, 105 had a history of ischemic cardiopathy (IC), 52 had cerebro-vascular disease (CVD) of thrombotic origin, and 91 were controls.

Results. The remarkably high apo(a) fibrin-binding in CVD (0.268 ± 0.15 nmol/L) as compared to IC (0.155 ± 0.12 nmol/L) suggests the existence of peculiar and poorly understood differences in pro- or anti-thrombotic mechanisms in either cerebral and/or coronary arteries.

Conclusions Our results demonstrated that Lp(a) fibrin-binding and small Apo(a) isoforms are associated with athero-thrombotic disease.

Key words: fibrinolysis, lipoprotein (a), plasminogen, ischemic cardiopathy, cerebrovascular disease.

Introduction

Atherothrombosis is a leading cause of serious morbidity and death due to myocardial infarction and cerebrovascular accidents. Within this context, the Lp(a) has emerged as a major risk factor and a clear association between high plasma Lp(a) and ischemic cardiopathy (IC) or cerebrovascular disease (CVD) has been demonstrated in several case-control and prospective studies [1]. Lp(a) plasma concentration depends on its hepatic synthesis [2, 3] and varies from one individual to another within an approximate range of <10 mg/dL to > 100 mg/dL; it is independent of factors such as diet, cholesterol, obesity, and smoking, and is maintained within small variations during the life span [4]. Lp(a) is an LDL-like particle containing apo(a), a distinctive plasminogen-like glycoprotein. Apo(a) consists of a serine-proteinase region and several kringle domains, derived from those of plasminogen but presenting some distinctive features [5]. Only plasminogen's kringles 5 (one copy) and 4 (multiple copies) are represented in apo(a). The copies of apo(a) kringle 4 are not alike, ten different types have been recognized [6]. Each type is present only once, except for kringle 4 type 2, which appears in multiple copies originating a structural heterogeneity and different size isoforms (molecular weight between 280 and 800 kDa) that are in inverse correlation with Lp(a) plasma concentration [7]. The serine-proteinase region of apo(a) contains the same residues that constitute the potential active site in plasminogen, however a Ser-Ile substitution at the Arg-Val activation cleavage site renders apo(a) insensitive to plasminogen [5,8].

On the basis of these homologies and activation differences, several studies have pointed out that Lp(a) has an antifibrinolytic capacity. The mechanism by which Lp(a) inhibits fibrinolysis remains controversial, *in vitro* studies [9, 10] have demonstrated that Lp(a) competes with plasminogen for similar fibrin binding sites. The formation of a ternary complex between plasminogen, tissue-type plasminogen activator (t-PA) and fibrin is required for efficient plasminogen activation [11], this activation mechanism is reduced when apo(a)/Lp(a) is bound to fibrin [12,13].

It has also been shown that solution phase interactions between apo(a) and plasminogen inhibit the binding of plasminogen to plasmin-modified fibrin surfaces [14]. However, a mechanism that explains the direct role of Lp(a) in promoting intravascular thrombosis has not been identified as yet. Recently [15], it has been described that Lp(a) binds to tissue factor pathway inhibitor (TFPI), with consequent TFPI inhibition. TFPI main role is to modulate factor VIIa/tissue factor catalytic activity and it is now recognized as a physiological anticoagulant. TFPI is present in the vascular smooth muscle cells [15] and the Lp(a)-TFPI interaction suggests a procoagulant mechanism in the atherosclerotic plaque.

These features suggest that Lp(a) has both, antifibrinolytic and prothrombotic capacities.

In spite of these molecular interaction data, and epidemiological evidence, the general approach for the study of Lp(a) as a vascular risk factor has until now, been limited to the determination of Lp(a) plasma concentration. A normal value for plasma concentration has not been agreed upon, some authors consider this to be up to 20 mg/dL [16] and others up to 30 mg/dL [17], pointing out that the use of different

monoclonal or polyclonal antibodies, generates differences in results [18]. In order to assess concordance in results among different methods, a calibrator and a reference ELISA method has been evaluated and demonstrated to be insensitive to apo(a) isoform size heterogeneity [19].

In the present work, we propose a novel approach that allows identification of individuals with a high Lp(a) binding capacity to plasmin degraded fibrin surfaces. This functional method that allows unambiguous estimation of the effect of Lp(a) on fibrinolysis is based on the competitive binding of Lp(a) and plasminogen to fibrin surfaces [10].

We hypothesize that the direct assessment of the ability of Lp(a) to bind to fibrin and decrease thereby plasminogen binding may provide a better and straightforward estimation of its potential atherothrombotic risk.

We performed a transversal case-control study which pointed out Lp(a) fibrin binding association with CVD.

Methods

Study group

Patients were selected among subjects attending the out-patient clinics of the National Institutes of Cardiology and Neurology (Tlalpan, D.F., Mexico). A total of 248 individuals were studied, 157 patients and 91 controls. The patients had a history of ischaemic cardiopathy n=105 (66.9 %), including myocardial infarction (67.7%), and stable/unstable angina (32.3%) or thrombotic cerebro-vascular disease n=52 (33.1%).

The diagnosis of ischemic cardiopathy was made by the clinical and

electrocardiographic studies as well as by coronary angiography. The diagnosis of cerebral vascular disease was based on the clinical signs of transient cerebral ischemia or cerebral infarction, as well as doppler and computerized tomography. The patients selected had none of the following exclusion pathologies criteria: uncontrolled arterial hypertension, hepatic or renal failure, neurological deficit with dementia, cerebrovascular emboli of cardiac origin or after carotid angiography, and recent (less than two months before the blood sample collection) myocardial infarct or surgical interventions. Patients with clinical evidence of ischemic disease were, if necessary, under pharmacological treatment in order to control hyperlipidemia, hypertension, diabetes mellitus or platelet aggregation.

Controls were healthy volunteers with no history of IC or CVD. Written consent was obtained from all participants, the study was approved by the Ethical Committee and the procedures were in accordance with the guidelines established by our institutions.

Blood samples

Fasting venous blood (up to 10 mL) was drawn from the forearm into sterile polypropylene tubes containing 4 mmol/L EDTA as anticoagulant. The samples were immediately centrifuged at 2000 x g for 15 min at 4°C. Plasma was removed by aspiration, supplemented with 100 KIU/mL aprotinin (Trasylol®, Bayer Pharma, Puteaux, France) and stored in aliquots at -70°C for a maximum of six months.

Lp(a) fibrin-binding assay

The assay is based on the ability of Lp(a) to compete with plasminogen for fibrin binding either in purified systems [10] or at the plasma/fibrin interface [20]. Conditions for this competition in plasma have been established [20] and advantageously used to develop

a functional assay to detect pathogenic Lp(a). Fibrin surfaces used as a template for the competitive interaction between plasminogen and Lp(a) present in plasma, were prepared and characterized as previously described [21,22]. Buffer A (0.05 mmol/L sodium phosphate, pH 7.4, 0.08 mmol/L NaCl, 0.01% (w/v) Thymerosal, 0.01 % (w/v) Tween 20) supplemented with BSA where indicated, was used in all experiments. To perform the assay, a volume of 50 μ L of plasma (diluted 1:16 in buffer A supplemented with 40 mg/mL BSA) was incubated in duplicate with fibrin surfaces for 12 h at 4°C, to allow plasminogen/Lp(a) equilibrium competition for fibrin binding. Unbound proteins were removed by washing three times with buffer A supplemented with 2 mg/mL of bovine serum albumin and a volume of 50 μ L of a peroxidase-conjugated monoclonal mouse antibody, directed against apo(a) [23], was added. This antibody does not cross-react with plasminogen. After 2 h at 37°C the plate was washed and a solution of 1 mg/mL 2,2'-azino-bis(3-ethylbenzthiazoline)-6-sulphonic acid (ABTS, Boehringer Mannheim) was used for colour development. The change in absorbance ($\Delta A_{405/min}$) was measured with a microtitration plate counter (MR 5000, Dynatech). The amount of Lp(a) bound is expressed in nmol/L of apo(a) by referring to a calibration curve. The standard curve was obtained from a reference standard composed of equimolar amounts of five recombinant apo(a) isoforms of different lengths (10 to 34 kringles) [24].

Lp(a) plasma concentration

Lp(a) plasma concentration was measured employing an immunonephelometric method (reagents and equipment from Beckman Co, Palo Alto CA, USA).

Apo(a) phenotyping

Apo(a) isoforms were identified as described [24] using an immunoblot procedure and a standard composed of recombinant apo(a)s containing 10, 14, 18, 26 and 34 kringles. Each apo(a) band image of the immunoblots was digitalized (Molecular Analyst TM/PC) and analyzed by densitometric scanning (Bio-Rad GS-670). The densitometric data from each band was expressed as a percentage relative to the sum of isoform (one or two) considered as 100%. The molar concentration of each apo(a) isoform was then calculated by relating their molecular mass and relative proportion in plasma with the concentration of Lp(a). For statistical purposes (see below and Tables 2 and 3) isoforms present at the highest concentration were considered as phenotype 1 whereas phenotype 2 represented the low concentration isoform in heterozygous subjects.

Statistical Analysis

For all variables, the mean and SD or the median and range inter-quartile were calculated. Quartiles cut-off values for Lp(a) concentration, apo(a) fibrin-binding capacity, and apo(a) phenotypes were calculated from the entire study population (IC, CVD and controls) and the proportion of patients with IC, CVD and control interquartil was analyzed by χ^2 test. Differences between groups were determined by χ^2 for discrete variables. For continuous variables comparisons between groups were evaluated with parametric or non-parametric analysis of variance according to the variable distribution. Least significant difference test (LSD) for parametric variables or U test of Mann-Whitney for non-parametric variables were used to determine significant differences between groups. Covariance analyses were also used to assess the

potentially confusing effect of age. Lp(a) concentrations and Lp(a) binding distribution were compared by the Kolmogorov-Smirnov test.

The association of Lp(a) plasma concentration, apo(a) binding and apo(a) isoforms with IC and CVD in the fourth quartile versus the first quartile was evaluated by contingency tables, odd ratios and 95% confidence intervals were calculated.

All statistical analyses were performed with the SPSS-10.1 Package (SPSS, USA).

Results

The group of patients with clinical evidence of ischaemic disease did not significantly differ from the group of control subjects, regarding gender and body mass index (Table 1). The age was significantly higher in both ischaemic disease groups compared to the control group ($p < 0.001$). Even if the Lp(a) plasma concentration is not age dependent, the results for Lp(a) concentrations and apo(a) binding were adjusted by age (ANCOVA analyse). By ANCOVA the results showed that apo(a) binding was significantly higher in the CVD patients as compared to IC patients and the control group. No differences were found in the Lp(a) concentrations among the studied groups of patients. All haemostatic variables tested (PT, PTT, TT) were normal in all subjects, (data not shown).

Table 2 illustrates the percentage of patients and controls in each quartile of Lp (a) concentration, apo(a) fibrin binding and apo(a) phenotype. Lp(a) concentration and apo(a) fibrin binding were skewed to lower values in controls, IC Lp(a) concentration showed an increment in the fourth quartile and an inverse relation was observed with large apo(a) phenotype.

CVD subjects differ substantially from controls and IC, increasing values for Lp(a) concentration and apo(a) fibrin binding were observed among the higher quartiles.

Table 3 shows the possible association between ischaemic cardiopathy or cerebrovascular disease and either Lp(a) concentration, apo(a) binding, or apo(a) phenotypes. The ischaemic cardiopathy was not associated with Lp(a) levels, apo(a) fibrin binding or apo(a) phenotypes in this particular group of Mexican patients. The cerebrovascular disease was significantly associated with high Lp(a) plasma concentration and high apo(a) fibrin binding. The presence of IC or CVD was associated, though not statistical significant to small size apo (a) isoforms.

Figure 1 shows the more important contribution, though not statistical significant, of low molecular weight phenotypes, less than 22 kringles, to Lp(a) plasma concentration in IC and CVD than in controls, (76.86%, 74.32% and 56.3%, respectively). This analysis allowed us to notice that K17-K19 phenotypes are more frequent in IC and CVD than in controls.

Furthermore, analyzing (in the same manner) only the subjects with high apo(a) fibrin-binding, the contribution of low molecular weight phenotypes, less than 19 kringles, to plasmatic Lp(a) concentration became more evident in IC (80.15 %), CVD (61.97%) and in controls, (56.65%), providing further evidence of the greater antifibrinolytic potential associated with small size isoforms and high Lp(a) concentration.

Discussion

Lp(a) affinity for fibrin shows an inverse relationship with the size of the apo(a) isoform within a dissociation constant (Kd) range from 50 to 500 nmol/L [10]. The importance of this structural and functional relationship is the fact that the Lp(a) antifibrinolytic potential depends, firstly, on the affinity of each of the two apo(a) isoforms found in plasma and, secondly, on Lp(a) concentration [10,25,26]. Therefore, similar Lp(a)

concentrations may display differences in atherothrombotic risk, depending on apo(a) size and affinity for fibrin. In agreement with these findings, it has been recently reported that small apo(a) isoforms are strong indicators of advanced stenotic atherosclerosis [27]. However, the lack of a method to evaluate the functional heterogeneity of Lp(a) is indeed a main obstacle for the correct assessment of coronary artery disease and CVD risk associated with Lp(a). This fact and the diversity of assay methods to measure Lp(a) plasma concentration may explain the discrepancy of some prospective studies that do not find a correlation between Lp(a) plasma level and coronary artery disease or CVD [28,29,30,31, 32].

Despite these drawbacks, a recent meta-analysis of 27 prospective coronary heart disease studies [1] concluded that individual concentrations of plasma Lp(a) in the highest third of baseline measurements versus those in the lowest third, result in a combined risk ratio of 1.6 (95% CI 1.4 to 1.8, $p < 0.00001$). The recognition that Lp(a) pathogenicity may not be alike in all individuals encouraged us to develop a new strategy for the evaluation of Lp(a). The original approach of this method to the pathogenesis of Lp(a)-related vascular disease, resides in its capacity to unveil the functional behaviour of apo(a) with regard to fibrinolysis impairment. This procedure is based on the antifibrinolytic mechanism of Lp(a) and takes into account both the affinity of apo(a) isoforms for fibrin and the plasma concentration of Lp(a).

A pathophysiological patient profile due to Lp(a) fibrin-binding is thus generated.

Under the conditions of the assay, the competitive binding of Lp(a) and plasminogen can only be affected by lysine-analogues [10]. Our experimental system mimics the possible behavior of plasminogen and Lp(a) when challenged by fibrin formation and

fibrinolysis activation. Its relevance for in vivo biology and disease has been previously evaluated in the pathophysiological model represented by the nephrotic syndrome [20,33]. The variations in plasminogen and Lp(a) plasma concentrations observed in this pathology result in significant differences in plasmin formation. We provided direct evidence that the more Lp(a) binds to fibrin, the less plasminogen is bound and plasmin is generated. Plasminogen binding to fibrin was shown to be decreased as a result of Lp(a) plasma concentration and isoform size. This mechanism results in low plasmin formation and a potential antifibrinolytic effect.

In the present work our findings provide experimental evidence for: 1) the potential antifibrinolytic effect associated, though not statistically significant, with small size isoforms of apo(a) in IC and CVD, and 2) the strong association between Lp(a) fibrin-binding and our CVD group. The high apo(a) fibrin binding observed in CVD patients as compared to controls is indicative of an efficient competition between plasminogen and Lp(a) for fibrin binding sites. This finding suggests that under conditions of fibrin challenging in the vasculature or in tissues, Lp(a) may inhibit plasmin formation, an important condition for the pathophysiology of atherothrombosis. Because of the small CVD case number it is not reasonable to draw final conclusions from our study regarding the unexpectedly high final Odds ratios results as there is only one case in the first quartile. However, there is a significantly increased relation between Lp(a) fibrin-binding and CVD. Our results agree with the hypothesis that elevated plasma Lp(a) is the primary factor associated with the presence of ischemic cerebrovascular disease as reported by Jürgens et al. [34].

Furthermore, our data suggest the existence of peculiar and poorly understood differences in antithrombotic mechanisms in cerebral and coronary arteries in these patients. Further studies will be necessary to determine whether these differences are of ethnical and/or environmental origin. In fact, a well documented model that provides evidence for the differences in the endothelial-cell-derived anticoagulant and procoagulant activities from one vascular bed to another, has been recently proposed [35]. The model supports the possibility of the existence of regionally distinct haemostatic balances. As a matter of fact, thrombomodulin is more important in maintaining the haemostatic balance in the lungs and heart than in the liver, whereas tissue-type plasminogen activator (t-PA) and urokinase-type plasminogen activator (u-PA) are important antithrombotic mechanisms in all three vascular beds. In the brain the factors maintaining haemostatic balance have not been identified as yet and the significant association between CVD and the antifibrinolytic effect of Lp(a) in this vascular bed could be an important clue.

Figure legends

Figure 1. Distribution and frequency of apo(a) phenotypes by kringle number in plasma Lp(a) from patients with ischaemic cardiopathy (IC), cerebrovascular disease (CVD) and controls.

Table 1

Demographic data, Lp(a) plasma concentration and apo(a) fibrin binding

Variable	Control	CVD	IC
Gender ⁺			
Male (%)	78.6	76.1	82.5
Female (%)	21.4	23.9	17.5
Age (years) [§]	40.2±13.1	53.4±10.5*	62.5±10.6*
Body Mass Index (Kg/m ²) [§]	26.3±5.6	27.0±3.5	27.2±3.4
Lp(a) (mg/dL) [†]	16.3±18	20.3±22.8	20.5±25.29
Apo(a)binding (nmol/L) [†]	0.158±0.09*	0.268±0.15	0.155±0.12**

⁺ The results are mean ± SD. CVD: cerebrovascular disease, IC: ischaemic cardiopathy,

[§]ANOVA, *<0.001 vs control. [†]ANCOVA **p<0.0001 Vs CVD

Table 2.

Distribution of patients with ischaemic cardiopathy (IC), cerebrovascular disease (CVD), and control subjects according to Lp(a) concentration, Apo(a) binding and Apo(a) phenotypes .

Quartile	1	2	3	4	P*
Lp(a) concentration	<6.1	6.1-13.8	13.9-22.45	>22.45	
IC (%)	33.3	19	20.2	27.4	0.074
CVD (%)	12.8	23.1	25.6	38.5	<0.001
CONTROL (%)	20.5	33.3	30.8	15.4	0.009
Apo (a)binding (nM)	<0.098	0.098-0.162	0.163-0.231	>0.231	
IC (%)	34.3	24.8	20	21	0.089
CVD (%)	1.9	9.6	32.7	55.8	<0.001
CONTROL (%)	27.5	34.1	26.4	12.1	0.003
Apo(a) phenotype 1 (No. of K4)	<20	20-23	23.1-26	>26	
IC (%)	20.8	35.1	28.6	15.6	0.006
CVD (%)	28.9	18.4	21.1	31.6	0.102
CONTROL (%)	23.1	18.5	32.3	26.2	0.187
Apo(a) phenotype 2 (No. of K4)	<16.5	16.6-19	19.1-21	>21	
IC (%)	28.3	28.2	17.4	26.1	0.217
CVD (%)	24.1	17.2	31	27.6	0.118
CONTROL (%)	20.6	20.6	29.4	29.4	0.332

IC= ischaemic cardiopathy, CVD=cerebrovascular disease * χ^2 test.

Table 3
 Association between ischaemic cardiopathy or cerebrovascular disease and Lp(a) parameters

	Highest Quartile	Lowest Quartile	OR (95 % CI)
Lp(a) (mg/dL)			
IC (%)	27.4	33.3	1.10 (0.43-2.78)
CONTROL (%)	15.4	20.5	
CVD (%)	38.5	12.8	4 (1.13-14.09)
CONTROL (%)	15.4	20.5	
Apo(a) Binding (nM)			
IC (%)	21.0	34.3	1.39 (0.57-3.37)
CONTROL (%)	12.1	27.5	
CVD (%)	55.8	1.9	65.9 (7.95-546.8)
CONTROL (%)	12.1	27.5	
Apo(a) Phenotype 1			
IC (%)	15.6	20.8	0.66 (0.24-1.84)
CONTROL (%)	26.2	23.1	
CVD (%)	31.6	28.9	0.96 (0.33-2.84)
CONTROL (%)	26.2	23.1	
Apo(a) Phenotype 2			
IC (%)	26.1	28.3	0.64 (0.19-2.24)
CONTROL (%)	29.4	20.6	
CVD (%)	27.6	24.1	0.80 (0.20-3.25)
CONTROL (%)	29.4	20.6	

haemic cardiopathy (n=105), CVD=cerebrovascular disease (n=52), OR= Odds ratio,

CI=confidence intervals, Apo(a) phenotypes by kringle number

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LIPOPROTEIN(a) POLYMORPHISM IN MEXICAN AMERINDIAN POPULATIONS

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Abstract. Lp(a) is a complex particle of unknown function mainly present in primates. It consists of a low-density lipoprotein and apo(a), a polymorphic glycoprotein homologous to plasminogen, the precursor of plasmin the fibrinolytic enzyme. Apo(a) is, however, an inactive zymogen that inhibits fibrinolysis through binding to fibrin. By virtue of its composite structure, Lp(a) is currently considered as a link between atherosclerosis and thrombosis. Marked inter-ethnic differences in Lp(a) concentration probably related to the genetic polymorphism of apo(a) has been reported. In this work we have evaluated the Lp(a) pattern in three Amerindian groups (Mayos n=68, Mazahuas n=62 and Mayas n=36) and 41 Mestizo subjects (admixture of Amerindian and Spaniards). Lp(a) plasma concentrations, apo(a) phenotypes and fibrin binding were analyzed. The Mestizo group presented the less skewed distribution and the highest median Lp(a) concentration (13.25 mg/dL) relative to Mazahuas (8.2 mg/dL), Mayas (8.25 mg/dL) and Mayos (6.5 mg/dL). There was not significant difference between groups in the prevalence of Lp(a) concentrations greater than the conventional cut-off for cardiovascular risk (> 30 mg/dL). Phenotype distribution was different in Mayas and Mazahuas as compared to the Mestizo group. The higher Lp(a) fibrin binding was found in the Maya population. There was an inverse relationship between the size of apo(a) polymorphs and both Lp(a) levels and Lp(a) binding to fibrin in all groups. In conclusion, the Lp(a) levels and distribution in Amerindians is highly skewed toward lower levels and differs significantly from the Mestizo ethnic group.

INTRODUCCION

The lipoprotein Lp(a) is a low density lipoprotein (LDL)-like particle mainly found in primate species and currently considered as a major risk factor for cerebrum and cardiovascular atherosclerotic disease (Djurovic and Berg, 1997). Lp(a) composition is indeed similar to that of the atherogenic LDL particle in terms of lipid core composition (cholesterol, triglycerides, phospholipids) and the apolipoprotein Apo B-100. The distinctive feature of Lp(a) is the glycoprotein apo(a), coded by a gene that arose from a duplicate copy of the plasminogen gene during primate evolution (Lawn et al., 1995). Apo(a) is therefore homologous to plasminogen, the precursor of plasmin the fibrinolytic enzyme (McLean et al., 1987). Plasminogen contains five different types of a structure termed kringle consisting of 80 to 90 amino acid residues arranged in a disulfide-bridge triple loop structure. Kringles 1 and 4 of plasminogen allow binding to fibrin, thus ensuring its *in situ* transformation into plasmin by plasminogen activators and thereby the lysis of a fibrin clot. Apo(a) contains 10 different types of plasminogen-like kringle 4 (K-IV) and a copy of each, kringle 5 and the serine-protease region. Although apo(a) shares with plasminogen its fibrin-binding function, it cannot be transformed into a plasmin-like serine-protease. Lp(a) may, therefore, compete with plasminogen for binding to fibrin and inhibit fibrinolysis, linking thereby the pathophysiological process of atherosclerosis and thrombosis (de la Peña-Díaz et al., 2000).

More than 30 apo(a) isoforms have been identified (Marcovina et al., 1993) that differ by the number of copies of K-IV type 2 which, in any given individual, is determined by the apo(a) gene (Boerwinkle, 1992). This polymorphism could be involved in determining Lp(a) levels as epidemiological evidence indicates an inverse relationship between Lp(a) concentration and the size of apo(a) (Gaubatz et al., 1990). The apo(a) gene is indeed a

major determinant of Lp(a) concentrations in plasma, it accounting for greater than 90% of its variations (Boerwinkle et al., 1992). A similar inverse relationship has also been found between apo(a) size and the ability of Lp(a) to compete with plasminogen for binding to fibrin (Hervio et al. 1996).

Differences in Lp(a) concentration and apo(a) phenotype distribution have been described in Caucasian, African and Asian groups (Sandolzer et al., 1991; Wong et al., 1999; Sorensen et al., 1994). Data in Amerindian populations have not been reported until now. The existence of different ethnic groups in Mexico and the prevalence of an important Mestizo population prompted us to investigate the distribution of Lp(a) concentration, apo(a) kringle IV phenotypes and the binding of Lp(a) to fibrin in these populations. The Mestizo can be distinguished from Amerindians by higher Lp(a) concentration from a less skewed distribution.

Subjects. A total of 240 non-related healthy subjects, women and men from four different ethnic groups, were studied. All subjects were selected at random from healthy individuals living in four states from Mexico. Anthropometric data are shown in Table 1. The historical, cultural and linguistic characteristics were different for each group as determined by the Instituto Nacional de Antropología e Historia de México. Plasma samples were obtained from Mayos (north-west of the country at Capomito, Estado de Sinaloa), Mazahuas (Santa Rosa and Pueblo Nuevo, Estado de Mexico) and Mestizos (Mexico City). The Maya samples were obtained from Tahdzibichen, near Merida, Yucatan. The present study was approved by the ethical committee of our institutions. Informed consent was obtained from all individuals. After an overnight fast, blood (10 ml) was drawn from the antecubital vein from each apparently healthy subject. Blood was collected into sodium

EDTA tubes (1mg/ml) and kept on ice until separation of plasma within the next 2 hours by centrifugation at 2000 g for 20 minutes at 4°C. Samples of plasma were supplemented with a proteinase inhibitor (Aprotinin 100 IU/ml, Trasylol[®] Bayer) and stored at -70°C until analysis (less than 6 months from sampling).

METHODS

Plasma Lp(a) concentrations. Lp(a) in plasma was measured by kinetic immunonephelometry (Gillery et al., 1993) using a commercial antibody in an automated Array protein system (Beckman Co, Palo Alto, CA, USA). The intra-assay and inter-assay coefficients of variation were less than 8%.

Apo(a) Kringle IV isoforms determination.

Sample preparation. Plasmas were adjusted to 15 mg/dl of Lp(a) or directly diluted 1:2 in sample buffer made of 0.34 M N-ethylmorpholine in 1.5 M Tris pH 8.8, 0.83 M 2-mercaptoethanol and 17.8% bromophenol blue in 10% glycerol. A mix of five recombinant apo(a) isoforms containing 10, 14, 18, 26 and 34 kringles were used as standard (Anglés-Cano et al., 1999). Samples and the standard were supplemented with 0.1% SDS and boiled for 5 minutes in a water bath.

Electrophoresis on a 3.75% and 6% discontinuous polyacrylamide gel and immunoblotting were performed as previously described (Anglés-Cano et al., 1999). A volume of 12 µl of plasma or recombinant apo(a) was applied to the gel and electrophoresis was run at room temperature using 35V for the first half an hour, followed by 70 V for another 6 hrs.

Apo(a) size determination. Blots were scanned in a GS-670 Bio-Rad densitometer using the Molecular Analyst^{TM/PC} software. The procedure sensitivity was 50 ng. The apo(a) isoform

size number was determined by reference to a standard curve made by plotting the relative migration of five recombinant apo(a) isoforms against the log of the number of kringle units. The inter and intra-assay coefficients of variation were less than 2%.

Lp(a) fibrin-binding assay. The experiments were performed with fibrin surfaces prepared and characterized as previously described (Fleury and Anglés-Cano, 1991). The procedure to assess binding of plasma Lp(a) to fibrin in competition with plasminogen was essentially as described (Soulat et al., 2000) with minor modifications. The amount of Lp(a) bound was expressed in nanomoles of apo(a) by referring to a calibration curve built with a reference standard composed of equimolar amounts of 5 recombinant apo(a) isoforms of different length (10 to 34 kringles) (Anglés-Cano et al., 1999)

Statistical methods. The overall shape of the distribution of plasma Lp(a) and apo(a) kringle IV phenotypes were compared between populations using the Kolmogorov-Smirnov test. Because of the highly skewed distribution of Lp(a) levels in the four ethnic groups, non-parametric Kruskal-Wallis tests were used to compare the median Lp(a) concentrations between groups. To estimate the influence of the apo(a) size polymorphism on Lp(a) plasma levels, Spearman rank correlation coefficients (Rs) were calculated. The prevalence of higher Lp(a) levels (> 30mg/dL) were compared with the χ^2 test. Multiple group means were compared by analysis of variance (ANOVA). The Statistical Package for Social Science (SPSS) V. 10 was used.

RESULTS

General characteristics of the four ethnic groups are shown in Table 1. There were no associations between Lp(a) and either age, weight, height, waist, BMI or gender; no adjustment was therefore necessary for the analysis.

Plasma Lp(a) concentrations. The distribution of plasma Lp(a) levels in each of the four ethnic groups is shown in figure 1. In all four groups the distribution is skewed toward low concentrations. The plasma levels of Lp(a) were less skewed in the Mestizo group than in the other groups. The median Lp(a) concentration was significantly higher in Mestizos than in the three indigenous groups (Table 2). The prevalence of Lp(a) greater than 30 mg/dL, the conventional cut-off for atherosclerotic risk, was slightly higher in Mazahuas than in the other groups (Table 2).

Apo(a) phenotypes. **Twenty six different apo(a) phenotypes were identified between all ethnic groups.** Apo(a) phenotype distribution in each of the four ethnic groups is shown in Figure 2. Phenotype distribution was significantly different in Mayas and in Mazahuas as compared to the Mestizo group. Differences in distribution were also observed between Mazahuas and Mayos. There was an inverse relationship between Lp(a) levels and apo(a) kringle number in Mestizos, Mazahuas and Mayos (Table 3); surprisingly, the Maya population studied showed a direct relationship between apo(a) kringle IV number and Lp(a) concentration.

Lp(a) binding to fibrin. Mean and median for Lp(a) binding to fibrin surfaces was significantly higher in Mayas when compared to each of the other ethnic groups (Table 2). Lp(a) median binding in Mazahuas was significantly lower than binding in Mestizos. An inverse relationship was found between Lp(a) binding and apo(a) kringle number in all populations (Table 3).

DISCUSSION

Several reports have suggested that plasma Lp(a) concentration is determined by the apo(a) gene locus (Kraft et al., 1992, Boerwinkle, 1992). However, variations among populations have been observed. As a matter of fact, in Caucasians the apo(a) gene size explains 40-60% of plasma Lp(a) concentration, whereas in other ethnic groups it varies from as low as 28% in Sudanese to as high as 70% in Chinese (Scholz et al., 1999). Indeed, the number of kringle IV type 2 that define the apo(a) polymorphism is inversely related not only to the plasma concentration of Lp(a) (Utermann et al., 1987) but also to its ability to bind to fibrin (Hervio et al., 1996). Therefore, genetic susceptibility to the effects of Lp(a) in a given population may be better assessed by estimation of the relationship between these parameters. The availability of a reliable fibrin-binding assay and a well defined standard composed of recombinant apo(a) isoforms of known kringle number, allowed us to perform the first population study integrating Lp(a) concentration determination, apo(a) kringle IV number analysis and the functional fibrin-binding assay. The Lp(a) concentration frequency distribution in the Mexican indigenous group was skewed to levels (<10 mg/dl) lower than those observed in Mestizo, Caucasian, Chinese, Black African, African-American and Mexican-American populations (Haffner et al., 1992; Gaw et al., 1994; Kraft et al., 1996). Another major finding in this study is the significant difference in Lp(a) plasma concentration between Mestizo and indigenous populations. The Mestizo population currently living in the Mexican Republic arose after the Spanish colonization started in 1502. Note that the mean concentration of Lp(a) found in the Mexican Mestizo group (16.6 ± 18 mg/dl) is similar to that found in Spanish populations (18.2 ± 14.8 mg/dl) (GomezGerique et al., 1996; Muros et al., 1996). However, the prevalence of Lp(a) greater

than the conventional cut-off value usually used to define cardiovascular risk (>30 mg/dl) is higher in apparently healthy European/Caucasian populations (15-20 %) than in the Mexican groups (Table 2). This difference is probably related to a less skewed distribution of Lp(a) concentrations in the Spanish population relative to the distribution observed in our study and reflects in a certain way cross genetic influences resulting from the admixture of Amerindians with Spaniards. The distribution of apo(a) phenotypes was bimodal in Mestizos and multimodal in the indigenous Amerindian groups. In agreement with this finding is the observation by Haffner et al. (1992) that Lp(a) concentrations in Mexican-Americans with greater Native Amerindian genetic admixture, are close to the values found in our Amerindian populations.

An interesting finding was the direct correlation (Table 3) between apo(a) phenotype and Lp(a) concentration in the Maya population we have studied. This finding suggests that the number of repetitive sequences in the apo(a) gene coding for kringle IV type 2, and therefore the size of apo(a), is not a major determinant of Lp(a) concentration in this population. Mayas also showed the greatest fibrin-binding but no correlation was found with apo(a) size. Because these observations may be of great value to our understanding of the influence of apo(a) size on both Lp(a) concentration and binding to fibrin, we are currently performing another study in a Maya population from Guatemala.

In conclusion, Lp(a) distribution in Mexican Amerindians is highly skewed toward lower levels relative to the Mestizo population. Apo(a) size distribution in Amerindians also differs significantly from the apo(a) size distribution in Mestizos. Our data provide further support to a genetic hypothesis for the difference in Lp(a) parameters among Amerindians and Mestizos.

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Figure Legends

Fig. 1. Distribution of lipoprotein Lp(a) concentrations in the populations studied.

Difference between Mestizos and Amerindians is significant ($p < 0.001$)

Fig. 2. Distribution of apolipoprotein(a) isoforms by kringle number. The X axis indicates the total number of plasminogen-like kringles including the single copy of kringle V.

Kolmogorow)Smirnov Z test: Mayas $p = 0.05$, Masahuas $p < 0.001$, vs Mestizos.

Table 1. Anthropometric characteristics of populations studied.

	Mayos	Mayas	Mazahuas	Mestizos
n =	68	36	60	76
Age (years)	55.8 ± 14.5	39.3 ± 21.3	19.35 ± 10.9	34.0 ± 9.9
Weight, Kg	65.6 ± 12.7	NA	43.2 ± 12.7	71.0 ± 13.4
Height, cm	167 ± 12.2	NA	147.3 ± 9.7	162.0 ± 26.0
Waist, cm	95.5 ± 9.1	NA	NA	86.8 ± 14.5
BMI, kg/m²	23.6 ± 4.5	NA	19.0 ± 4.0	26.1 ± 3.8

n = number of subjects studied.

Values are Mean ± SD

BMI, body mass index. NA, not available.

Table 2. Lp(a) plasma concentrations and fibrin-binding by ethnic group.

	n	Lp(a) Concentration			Fibrin-Binding	Fibrin-binding
		Mean \pm SD	Median	>30 mg/dl [¶]	Mean \pm SD	Median
Mayos	68	10.5 \pm 1.3	6.55	7.4	0.153 \pm 0.12	0.10
Mayas	36	10.6 \pm 8.5	8.25	5.6	0.257 \pm 0.14**	0.28**
Mazahuas	60	11.6 \pm 12	8.20	11.7	0.146 \pm 0.14	0.09
Mestizos	76	16.6 \pm 18*	13.25**	7.5	0.155 \pm 0.09	0.14

n = number of subjects studied.

Lp(a) concentration is expressed in mg/dl. Lp(a) binding is expressed in nanomol bound.

[¶]Proportion of subjects with Lp(a) concentration >30mg/dl, the conventional cut-off value for cardiovascular risk. Values in %.

Mean * = p < 0,05 vs. Mestizos. ** p < 0,01 vs. Mayas

ANOVA LSD post hoc Median: p < 0,001

Table 3. Correlation between apo (a) size and Lp(a) fibrin-binding and concentration.

Ethnic Group	Apo (a) phenotype/Lp (a) level	Apo(a) phenotype/fibrin-binding
Mayos	-0.361*	-0.07
Mayas	0.541*	-0.261
Mazahuas	-0.007	-0.248
Mestizo	-0.422**	-0.003

• p < 0.05

**p < 0.005

REVIEW ARTICLE

Lipoprotein Lp(a) and Atherothrombotic Disease

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High plasma concentrations of lipoprotein (a) [Lp(a)] are now considered a major risk factor for atherosclerosis and cardiovascular disease. This effect of Lp(a) may be related to its composite structure, a plasminogen-like inactive serine-proteinase, apoprotein (a) [apo(a)], which is disulfide-linked to the apoprotein B100 of an atherogenic low-density lipoprotein (LDL) particle. Apo(a) contains, in addition to the protease region and a copy of kringle 5 of plasminogen, a variable number of copies of plasminogen-like kringle 4, giving rise to a series of isoforms. This structural homology endows Lp(a) with the capacity to bind to fibrin and to membrane proteins of endothelial cells and monocytes, and thereby inhibits binding of plasminogen and plasmin formation. This mechanism favors fibrin and cholesterol deposition at sites of vascular injury and impairs activation of transforming growth factor-beta (TGF- β) that may result in migration and proliferation of smooth muscle cells into the vascular intima. It is currently accepted that this effect of Lp(a) is linked to its concentration in plasma, and an inverse relationship between apo(a) isoform size and Lp(a) concentrations that is under genetic control has been documented. Recently, it has been shown that inhibition of plasminogen binding to fibrin by apo(a) from homozygous subjects is also inversely associated with isoform size. These findings suggest that the structural polymorphism of apo(a) is not only inversely related to the plasma concentration of Lp(a), but also to a functional heterogeneity of apo(a) isoforms. Based on these pathophysiological findings, it can be proposed that the predictive value of Lp(a) as a risk factor for vascular occlusive disease in heterozygous subjects would depend on the relative concentration of the isoform with the highest affinity for fibrin. © 2000 IMSS. Published by Elsevier Science Inc.

Key Words: Lipoprotein(a), Atherosclerosis, Thrombosis, Fibrinolysis.

Introduction

In 1963, Kåre Berg identified the presence of lipoprotein(a) [Lp(a)] in plasma (1); nevertheless, it was not until 1987 when Eaton et al. (2) partially identified the apoprotein(a) [apo(a)] glycoprotein sequence. The protein was later cloned (3) and showed a strong structural similarity between one of the Lp(a) components, apo(a) glycoprotein, and the plasmin precursor, plasminogen. This finding stimulated the interest of different research groups who found a common point between atherosclerosis and thrombosis (4–11). Different epidemiological studies were performed from

this perspective and identified a positive correlation between high Lp(a) plasma concentration and an increase in cerebrovascular (12) and cardiovascular diseases, as well as coronary restenosis, postangioplasty reocclusion, and premature development of atherosclerosis related to high low-density lipoprotein (LDL) concentrations and/or low high-density lipoprotein (HDL) concentrations (13,14). Most prospective studies have confirmed these results (15–21). Even when a normal value for plasma concentration has not been agreed upon, some investigators considered this to be 20 mg/dL (20) and others suggested 30 mg/dL (15), pointing out that the use of different antibodies, monoclonal or polyclonal, generates differences in results (5).

Nevertheless, other studies do not find a relationship between Lp(a) and coronary arterial disease (22,23). This discrepancy may be a reflection of the large structural hetero-

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entity of the apo(a) molecule, which is conducive to a functional heterogeneity as a plasminogen competitive inhibitor of fibrin because it lowers the formation of plasmin (24).

Not all organisms synthesize apo(a). Its presence has been identified in humans, in some Asian, European, and American primates (25), and in hedgehogs (26).

Function

The function of apo(a) in the organism is unknown. It provides cholesterol from the liver to organs that synthesize steroid hormones (28) and to tissues for cell repair (29). Furthermore, the function of apo(a) found in testicles and brain is unknown and is independent from Lp(a); that is, it functions without being part of the Lp(a) molecule (30).

Concentration

Lp(a) plasma concentration, which depends on its hepatic synthesis (31,32), varies from one individual to another within an approximate range of <10 mg/dL to >100 mg/dL; it is independent on other factors such as diet, cholesterol, obesity, and smoking, and is maintained within small variations throughout the lifespan (33,34). In general, an individual inherits, in a codominant autosomic fashion, two apo(a) isoforms that may be identified by sodium dodecylsulfate polyacrylamide gel electrophoresis (SDS-PAGE) and, subsequently, by immunotransference, making use of monoclonal or polyclonal antibodies against apo(a). By means of this method, almost 37 isoforms have been identified, which, according to the Utermann classification are: F (faster); B (i.e., apoB-100); S1 (slow), and S2, S3, or S4, depending on their migration velocity, in comparison with that of apoB-100 (35). The method does not allow for distinction of differences between isoforms of close molecular weight. Nevertheless, it has recently become possible to visualize the isoforms with the help of a reference, accounting for apo(a) recombinant isoforms with different molecular weights, correlating ($r = 0.97$) with the technique of pulsed-field electrophoresis (36). With the use of this reference it has been possible to obtain a linear relationship between the log r -apo(a) kringle number and the relative migration using SDS-PAGE, successfully identifying apo(a) isoforms over a wide range of molecular sizes. Pulsed-field electrophoresis has been employed to identify different genotypes codified for apo(a) (37). With the aforementioned technique, 19 different alleles have been described in a U.S. study of a population of whites (38).

Apo(a) isoform size accounts for an inverse correlation with Lp(a) plasma concentration (39), probably because, as apo(a) size increases, less protein is secreted from the cell, as occurs in the case of the human hepatocarcinoma cell line HepG2 (40).

Some hormones can modify Lp(a) plasma concentration, such as estrogens (41-44); anabolic steroids might reduce it (45) and growth hormone can increase it (46), but it cannot be modified by lipid-lowering medication or by diet (33,47). The plasma concentration of Lp(a) increases in diseases such as diabetes mellitus, nephrotic syndrome, rheumatoid arthritis, and in a transitory fashion after myocardial infarction or surgical intervention (5,33,34,48-50).

There are variations both in its average concentration and in the abundance of isoforms in genetically different populations (51-54). Lp(a) plasma concentration in blacks is at least three times greater than in whites (55-58).

Chemical Structure

Lp(a) composition is similar to that of LDLs. Both of these lipoproteins contain cholesterol, triglycerides, and phospholipids that may be dissolved and transported by plasma, due to the presence of a protein, apoB-100, which surrounds the lipid group and cholesterol (Figure 1).

The difference between them is that Lp(a) contains another glycoprotein, apo(a), which is bound to apoB-100 by a disulfide bridge between Cys in the 69 position for KIV-9 of apo(a) and Cys 3734 for apoB-100; the union is stabilized by hydrogen bonds and van der Waals interactions in other areas of both proteins (59-61). ApoB-100 (62) has the same structure and conformation in Lp(a) and the LDL molecule. The ratio apo(a):apoB-100 is 1:1 (63), and in considering their physicochemical variables there may be entities in a 2:1 ratio (64).

As usually occurs with some plasma proteins, Lp(a) may have different sizes with weights, ranging between 800 and 1300 kDa, and thus different densities. These differences are a reflection, although to a lesser extent, of lipid core composition, and especially of apo(a) structural polymorphism (65).

There are technical difficulties involved in obtaining apo(a) in its native form, and predictions of its secondary structure suggest an absence of an α -helix (66). Nevertheless, after reducing sulphydryl binding, and by means of the circular dichroism technique, 8% α -helix, 21% β -sheet, and 71% random arrangements have been observed (67). Apo(a) belongs to the serine protease family, together with plasminogen, prothrombin, tissue plasminogen activator, urokinase-type plasminogen activator, and factor XII. These proteins derive from an ancestral gene common to all of them. Apo(a) is very similar to plasminogen. Genes codifying both proteins are very close, in chromosome 6, band q26-27 (35). The genes are a 50-kb distance apart in a head-to-head position (68,69) in the terminal region 5'. In the case of apo(a), the gene presents a polymorphism, which can be expressed by modifying the efficiency of the transcription. This, in turn, originates differences in Lp(a) plasma concentration, not only between individuals but between different ethnic groups (68). The apo(a) gene may have different sizes, each

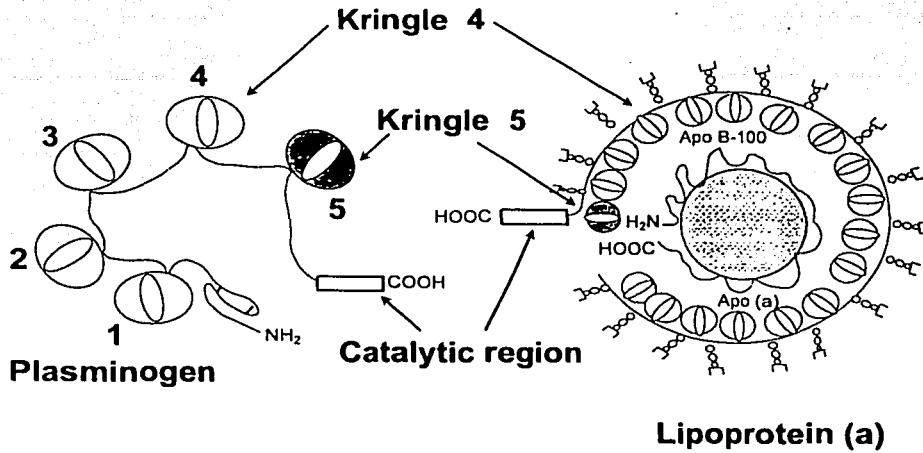


Figure 1. Chemical structure of lipoprotein(a) and plasminogen.

corresponding to the number of times in which a 5.5-kb sequence is present, with its number varying between 12 and 51 (3). The gene-promoting region of apo(a) possesses, in a distal position, a 1-kb fragment with a sequence showing sites with a different interaction potential. For interleukin-6 (IL-6), it shows seven sites; this conformation could explain the increase in Lp(a) plasma concentration during acute inflammation states and the three sites with specific elements (HNF-1, CEBP, and LF-A1) for hepatic transcription (69).

Apo(a) and plasminogen contain a protease region with a 94% similarity, but in the case of apo(a) it lacks the ability to become active and perform its enzymatic function due to the presence of arginine instead of serine at the activation site (3).

Plasminogen and apo(a) also include a different number of modules, called kringles. Through kringles they bind and recognize other macromolecules and/or specific sites in the cell membrane (73,74).

Kringles connect with each other by interkringle regions that are segments of 26-36 serine-, proline-, and/or threonine-rich amino acids (3); each interkringle region contains six potential O-glycosylation sites.

The kringle recognition function includes the participation of a structure generated in the inner loop surrounding a hydrophobic region formed by different aromatic amino acids that are stabilized by means of hydrogen bonds and separate a cationic from an anionic group (73). This region is known as the lysine-binding site (LBS). Its structural characteristics generate a relatively rigid geometry that allows selective access and binding of 6.8-Å aliphatic or aromatic ligands of ω -amino-carboxylic acid type, such as ω -amino-hexanoic acid or similar compounds (74). Plasminogen comprises five very similar kringle types, but with small differences at the lysine-binding site, which modify the degree of

affinity for different ligands. Plasminogen K I has a high-affinity LBS; the cationic pole has Arg-35 and Arg-71, and the anionic pole has Asp-55 and Asp-57 (74,75). The LBS in plasminogen K IV has an intermediate affinity; its V-shaped hydrophobic region generates a topography in which the aromatic rings of Phe-64, Trp-62, and Trp-72 separate the anionic group, which is formed by Asp-55 and Asp-57, and from the cationic group by Lys-35 and Arg-71 (76).

Apolipoprotein (a) Polymorphism

Apo(a) shares kringle V and a variable number of kringle IVs with plasminogen (35). Not all kringles IV of apo(a) are alike; they are classified into 10 different subtypes (77). In the apo(a) molecule, each is present only once, except for kringle IV-2, which appears on multiple copies, originating structural heterogeneity and different size isoforms that account for molecular weights of between 280 and 800 kDa.

Kringle IV-10 is most similar to the kringle IV of plasminogen, with the high-affinity, lysine-binding site being formed by Asp-55 and Asp-57 in the anionic pole, and by Arg-71 and Arg-35 in the cationic pole. Between the two poles there is a hydrophobic microenvironment that is formed by three aromatic amino acids: Trp-62, Phe-64, and Trp-72. KIV-10 of apo(a) differs from K-IV of plasminogen due to the presence of Arg instead of Lys in position 35. This kringle has a very important role in the Lp(a) union with lysine (78), preventing plasminogen access to the fibrin clot, thus blocking the action of the tissue plasminogen activator (79-82). This generates fibrinolytic insufficiency that, in turn, promotes atherosclerosis and thrombosis.

Another apo(a) polymorphism source is glycosylation:

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(apo(a) KIV has a potential site for N-glycosylation. If N-linked glycosylation is considered, 30% of each molecule (apo(a)) corresponds to carbohydrates as follows: mannose; galactose; galactosamine; glucosamine, and sialic acid ratios of close to 3:7:5:4:7, respectively (83).

a) Plasminogen Competitive Inhibitor

The fibrinolytic system destroys fibrin deposits in blood vessels, either those remaining from hemostatic activity or those formed and accumulated during the development of atherosclerotic plaque. Because there is a balance between plasminogen activators and different types of inhibitors of these activators, its response varies either toward a pro-thrombotic condition or a fibrinolytic activity.

As a result of fibrinolytic activation, plasmin is generated, a proteolytic enzyme that may have effects on different plasma substrates. In the present case, however, this does not occur, because fibrin itself locates and signals the fibrinolytic activation sites. Within the fibrin mesh, plasminogen is bound precisely at the lysine-binding site and properly directs the binding of its tissue activator (i.e., tissue plasminogen activator). Therefore, plasmin is generated in the inner section of the fibrin deposit, breaking and exposing other affinity sites such as terminal carboxylate residues, to amplify its response. Once fibrin is disintegrated, the mechanism stops, and the vascular endothelium does not release additional activator; the remaining circulating plasmin and tissue activator find their corresponding inhibitors, α 2-antiplasmin and type 1 tissue plasminogen activator inhibitor (PAI-1).

The similarity between plasminogen and apolipoprotein(a) allows different apo(a) isoforms to compete with plasminogen for fibrin affinity sites. The affinity of each isoform depends on its size and its plasma concentration (7,84,85); in addition, the plasmin formed at the surface of fibrin may vary with modifications of the concentration of Lp(a) *in vivo* (94).

Plasminogen and the different Lp(a) isoforms (86) also compete for lysine residues on the surface receptors of endothelial cells (81,87), U937 monocytes (87,88), platelets (89), mononuclear cells (86), and on matrices for *in vitro* models simulating the extracellular membrane (90).

Another mechanism that alters the fibrinolytic system balance is either the decrease or increase in synthesis of the tissue plasminogen activator (t-PA) or of the tissue plasminogen activator inhibitor (PAI-1), respectively (33,91), which is observed in cultures of endothelial cells exposed to Lp(a).

Lp(a) and Atherogenesis

Lp(a) favors atherogenesis through different mechanisms: macrophages phagocytize Lp(a) (92), migrate, and settle in the subendothelium, becoming transformed into foam cells, generating deformities that decrease the lumen of the blood

vessels. It has been observed that, in cultures of endothelial cells from the coronary artery, Lp(a) stimulates the expression of vascular adhesion molecule-1 (VCAM-1) and selectin E, a process that triggers attraction to macrophages (93). Another proposed mechanism is related to a decrease in plasmin generation that accounts for: (a) prolonged permanence of fibrin deposits (74), with the consequent increase of cholesterol deposits and formation of atherosclerotic plaque, and (b) decrease of activation, by partial hydrolysis with plasmin, of TGF- β , which prevents the growth of vascular smooth muscle cells.

Lp(a) Inhibits Fibrinolysis Depending on Apo(a) Isoforms

A high plasma concentration of Lp(a) does not always interfere with normal fibrinolysis. Apo(a) isoforms show different antifibrinolytic activity (84,85), hence the importance of taking into account the antifibrinolytic activity of the isoforms in the prediction of cardiovascular diseases (95).

Many different strategies have been applied in *in vitro* studies. While making use of a solid-phase fibrin model (96-100), it has been possible to identify, with high specificity and sensitivity, fibrinolysis inhibition by apo(a) isoforms. These studies have permitted identification of the mechanisms and different variables involved in plasminogen competitive inhibition, due either to the different Lp(a) native isoforms (84) or through apo(a) recombinant forms (24).

These studies demonstrate the following: (1) Lp(a) affinity for fibrin shows an inverse relationship with the size of the apo(a) isoform within a K_d range of 50-500 nM (84); (2) both plasminogen and Lp(a) compete for the same binding sites, which corresponds to a saturable competitive inhibition mechanism, and (3) Lp(a) antifibrinolytic potential depends on the affinity and concentration of each of the two apo(a) isoforms found in plasma (101).

Lp(a) and Homocysteinemia

Hyperhomocysteinemia is related to an increase in the incidence of thrombotic and atherosclerotic diseases (102). Thrombosis is favored because of alterations in different antithrombosis-regulation mechanisms; in addition, it increases tissue factor activity, lowers the expression and activity of thrombomodulin necessary for protein C activation, and lowers the anticoagulant activity of antithrombin III as well as the binding capacity of the tissue plasminogen activator to its receptor on cell surfaces (anexin II) (103). In addition, an increased plasma homocysteine concentration manifests itself as an increase in Lp(a) binding to fibrin (104,105).

Conclusions

Most epidemiological studies point out the relationship between lipoprotein (a) plasma concentration and the risk of suffering cardiovascular and cerebrovascular diseases, espe-

cially when other risk factors are present, such as high levels of low-density lipoproteins and smoking. It is also important to point out that there are differences in the concentration and in the presence/nonpresence of some isoforms within genetically different populations. Therefore, identification of fibrinolysis inhibition behavior of different Lp(a) isoforms necessitates applying a strategy to the study that will allow for the identification of individuals and ethnic groups suffering from fibrinolytic deficiency due to the presence of high concentrations of Lp(a). This line of investigation represents a challenge to the various disciplines in studying the role of Lp(a) in the pathogenesis and progression of atherosclerosis.

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Lipoproteína (a), ¿es un factor de riesgo en la enfermedad aterotrombótica coronaria?

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Palabras clave: Lipoproteína (a). Aterotrombosis. Enfermedad arterial coronaria. Fibrinólisis.
Key words: Lipoprotein (a). Atherothrombosis. Coronary artery disease. Fibrinolysis.
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A pesar de que numerosos estudios *in vitro* e *in vivo* confirman la participación de la Lp(a) en los procesos fisiopatológicos de la aterosclerosis y de que los estudios retrospectivos consistentemente confirman la asociación entre Lp(a) y la enfermedad isquémica coronaria y cerebral, algunos estudios prospectivos han reportado resultados inconsistentes.

Varios motivos contribuyen en esta controversia, entre ellos destacan la falta de un estándar universal de referencia, el método analítico de estudio, las condiciones en el manejo de la muestra y sobre todo las características estructurales distintivas de la partícula de la lipoproteína (a) o Lp(a). La Lp(a) está formada por una lipoproteína de baja densidad o LDL¹ que contiene un núcleo de colesterol, triglicéridos y fosfolípidos, rodeado de una proteína apoB-100 a la cual se une la glicoproteína apo(a). Como es frecuente en las lipoproteínas plasmáticas, la Lp(a) puede tener diferentes tamaños; estas diferencias se deben, en menor medida, a la composición del núcleo de lípidos y, principalmente, al polimorfismo estructural de apo(a).² La apo(a) pertenece a la familia de serino proteasas, al igual que el plasminógeno, la protrombina, el activador tisular del plasminógeno, el activador del plasminógeno tipo urocinasa y el factor XII. Estas

proteínas derivan de un gene ancestral común a todas ellas.

La apo(a) es muy parecida al plasminógeno (Fig. 1). Los genes que codifican para ambas proteínas se encuentran muy cercanos, pero el gene de la apo(a) tiene la peculiaridad de tener diferentes tamaños que dependen del número de veces que se repite una secuencia de 5.5 kb. El tamaño de las isoformas de la apo(a) tienen una relación inversa con la concentración plasmática de la Lp(a)³ y a medida que se incrementa el tamaño de apo(a), se incrementa la dificultad para secretarse de las células hepáticas, lo que da lugar a diferencias en la concentración plasmática de Lp(a), no sólo entre individuos, sino entre diferentes grupos étnicos. Algunas hormonas pueden modificar su concentración plasmática: la hormona tiroidea,⁴ los estrógenos,³⁻⁶ y los esteroides anabólicos la reducen⁷ en tanto la hormona de crecimiento la incrementa,¹⁰ pero no puede modificarse por medicamentos hipolipemiantes o por la dieta.¹¹

La apo(a) y el plasminógeno, el precursor de la plasmina, tienen una región proteasa muy similar. La presencia de arginina en lugar de serina en el sitio de activación impide que la apo(a) pueda ser activada y por ende que esta región exprese una actividad enzimática.

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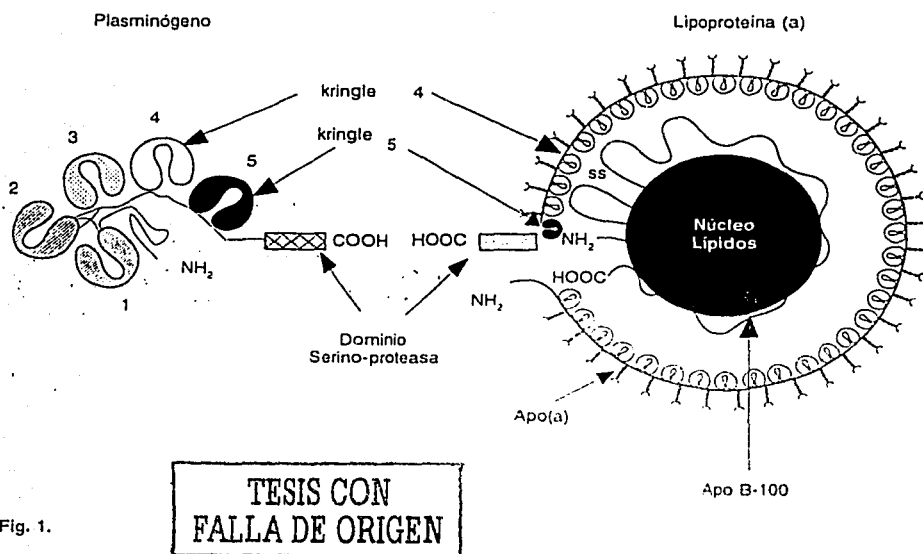
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Esta similitud estructural y diferencia enzimática despertó el interés de numerosos grupos de investigación que encontraron en la Lp(a) el punto común entre la aterosclerosis y la trombosis.¹²⁻¹⁴ Bajo esta perspectiva, se realizaron numerosos estudios epidemiológicos, que describen una asociación positiva entre la elevada concentración plasmática de Lp(a) y un incremento en las enfermedades cerebrovasculares,¹⁹ cardiovasculares, la reestenosis de puentes coronarios, la reoclusión de angioplastias y el desarrollo prematuro de aterosclerosis asociada a altas concentraciones de LDL y/o a bajas concentraciones de HDL.^{20,21} La mayoría de estudios prospectivos confirman estos resultados.²²⁻²⁴ Aun cuando no existe un acuerdo en el límite de corte para considerar una concentración plasmática normal, algunos autores consideran que es de 20 mg/dL, otros hasta 30 mg/dL y que el empleo de anticuerpos diferentes, monoclonales o policlonales, generan divergencias en los resultados. Sin embargo, otros estudios no encuentran asociación entre la Lp(a) y la enfermedad arterial coronaria.^{29,30} Esta discrepancia en los resultados podría ser el reflejo de la gran heterogeneidad estructural de la molécula de apo(a), que se traduce en una heterogeneidad funcional como inhibidor competitivo del plasminógeno por la fibrina, disminuyendo así la formación de plasmina.^{15,31}

Craig y col.³² hacen un meta-análisis de diferentes estudios prospectivos en población caucásica, evalúan a la Lp(a) como factor de riesgo en la enfermedad arterial coronaria, ponen especial atención en los criterios de selección de los trabajos a fin de que los resultados fueran comparables. Concluyen que en los individuos que desarrollaron enfermedad arterial coronaria la concentración de Lp(a) es mayor que en el grupo control (excepto en los pacientes masculinos de un estudio)³³ y sugieren que la variabilidad puede ser efecto del manejo pre-analítico de la muestra, especialmente de la temperatura en que ésta haya sido almacenada.

Recientemente³⁴ un meta-análisis de 27 estudios prospectivos de Lp(a) y enfermedad isquémica coronaria, concluye que independientemente del lugar geográfico del estudio, edad promedio de los individuos, tiempo de duración del estudio, métodos de ensayo y grado de ajuste de las variables de confusión, el riesgo relativo es de 1.7 (intervalo de confianza 95% de 1.4-1.9, $2P < 0.00001$) para 18 estudios en población general que incluyeron un total de 4,044 casos, y un riesgo relativo de 1.3 (intervalo de confianza 95 de 1.1-1.6, $2P < 0.001$) para 9 estudios que incluyeron un total de 1,392 individuos con antecedentes de enfermedad coronaria. Este meta-análisis incluyó algunos de los trabajos analizados por



Craig y col. pero con un criterio diferente de evaluación; comparó la concentración plasmática de Lp(a) en el tercil superior contra el tercil más bajo de distribución.

En ambos meta-análisis resalta sin lugar a dudas, la asociación que existe entre la Lp(a) y la enfermedad isquémica coronaria, frecuentemente con isoformas de bajo peso molecular de apo(a);³⁵ sin embargo hasta la fecha no es posible afirmar si este hecho es un factor directo de riesgo aterotrombótico o si el riesgo se favorece a través de permitir una mayor concentración plasmática de Lp(a).

El hecho de que la Lp(a) inhibe la fibrinólisis, ofrece un mecanismo fisiopatológico congruente con ambas situaciones;³⁶ explica porqué una alta concentración de Lp(a) se refleja en diferentes grados de riesgo aterotrombótico que dependen de las isoformas de apo(a)³⁷ y pone de manifiesto la importancia cualitativa y cuantitativa de las interacciones entre la apo(a), el plasminógeno y la fibrina.

Empleando un modelo de fibrina en fase sólida,³⁸⁻⁴² ha sido posible identificar de manera sensible y específica la inhibición de la fibrinólisis por las isoformas de apo(a). Los estudios concluyen que el potencial anti-fibrinolítico de Lp(a)

depende de la concentración y afinidad de cada una de las dos isoformas de apo(a)³³ y muestran una relación inversa con el tamaño de la isoforma en un rango de 50 a 500 nM. Este dato se comprueba al observar que las isoformas de apo(a) de bajo peso molecular son indicadores importantes en la aterosclerosis obstructiva.³⁵

El estudio de la Lp(a) en la enfermedad aterotrombótica sigue representando un reto multidisciplinario, en donde una estrategia objetiva de estudio subraye la importancia de considerar la actividad antifibrinolítica de las isoformas en la predicción de la enfermedad cardiovascular.⁴⁴ Deberán corregirse también, para determinar la concentración plasmática de Lp(a), la falta de acuerdo que existe para el manejo pre-analítico de la muestra, el empleo de un estándar y técnica de referencia y el empleo de anticuerpos monoclonales o policlonales.

Es también importante llevar a cabo estudios en poblaciones sanas, con la finalidad de identificar a las poblaciones en las que la Lp(a) represente un factor de morbilidad y mortalidad importante y conocer su impacto en la salud pública. Resalta la ausencia de estudios en la población latinoamericana a pesar del gran interés que el tema ha cobrado en los últimos años.

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Inhibition of Fibrinolysis by Lipoprotein(a)

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ABSTRACT: A high plasma concentration of lipoprotein Lp(a) is now considered to be a major and independent risk factor for cerebro- and cardiovascular atherothrombosis. The mechanism by which Lp(a) may favour this pathological state may be related to its particular structure, a plasminogen-like glycoprotein, apo(a), that is disulfide linked to the apo B100 of an atherogenic LDL-like particle. Apo(a) exists in several isoforms defined by a variable number of copies of plasminogen-like kringle 4 and single copies of kringle 5 and the catalytic region. At least one of the plasminogen-like kringle 4 copies present in apo(a) (kringle IV type 10) contains a lysine binding site (LBS) that is similar to that of plasminogen. This structure allows binding of these proteins to fibrin and cell membranes. Plasminogen thus bound is cleaved at ARG₅₆₁-VAL₅₆₂ by plasminogen activators and transformed into plasmin. This mechanism ensures fibrinolysis and pericellular proteolysis. In apo(a) a Ser-Ile substitution at the Arg-Val plasminogen activation cleavage site prevents its transformation into a plasmin-like enzyme. Because of this structural/functional homology and enzymatic difference, Lp(a) may compete with plasminogen for binding to lysine residues and impair, thereby, fibrinolysis and pericellular proteolysis. High concentrations of Lp(a) in plasma may, therefore, represent a potential source of antifibrinolytic activity. Indeed, we have recently shown that during the course of the nephrotic syndrome the amount of plasminogen bound and plasmin formed at the surface of fibrin are directly related to *in vivo* variations in the circulating concentration of Lp(a) (*Arterioscler. Thromb. Vasc. Biol.*, 2000, 20: 575-584; *Thromb. Haemost.*, 1999, 82: 121-127). This antifibrinolytic effect is primarily defined by the size of the apo(a) polymorphs, which show heterogeneity in their fibrin-binding activity—only small size isoforms display high affinity binding to fibrin (*Biochemistry*, 1995, 34: 13353-13358). Thus, in heterozygous subjects the amount of Lp(a) or plasminogen bound to fibrin is a function of the affinity of each of the apo(a) isoforms and of their concentration relative to each other and to plasminogen. The real risk factor is, therefore, the Lp(a) subpopulation with high affinity for fibrin. According to this concept, some Lp(a) phenotypes may not be related to atherothrombosis and, therefore, high Lp(a) in some individuals might not represent a risk factor for cardiovascular disease. In agreement with these data, it has been recently reported that Lp(a) particles containing low molecular mass apo(a) emerged as one of the leading risk conditions in advanced stenotic atherosclerosis (*Circulation*, 1999, 100: 1154-1160). The predictive value of high Lp(a) as a risk factor, therefore,

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depends on the relative concentration of Lp(a) particles containing small apo(a) isoforms with the highest affinity for fibrin. Within this context, the development of agents able to selectively neutralise the antifibrinolytic activity of Lp(a), offers new perspectives in the prevention and treatment of the cardiovascular risk associated with high concentrations of thrombogenic Lp(a).

KEYWORDS: Fibrin surface; Fibrinogen; Plasminogen; Plasmin; Kringle domains; Lysine binding site; Tissue-plasminogen activator; Urokinase; Atherosclerosis; Thrombosis; Atherothrombosis; Monocytes; Macrophages; Fibrinolysis; Plasminogen activation; Plasminogen activator inhibitor.

INTRODUCTION

The presence of the lipoprotein Lp(a) in plasma was first identified by Berg in 1963 as an LDL-like particle.¹ Lp(a) composition is indeed similar to that of LDL in terms of cholesterol, triglycerides, phospholipids, and apo B100. It was not until 1987 when Eaton *et al.*² and Mc Lean *et al.*³ identified apo(a), the unique and distinctive component of Lp(a), as a member of the plasminogen gene family,⁴ both by amino acid sequence analysis and cDNA cloning. The apo(a) glycoprotein indeed showed a strong structural homology with plasminogen, the plasmin precursor. Apo(a) is disulfide linked to the apo B100 of the LDL-like particle. These findings stimulated the interest of different research groups that found in Lp(a) a common point between atherosclerosis and thrombosis.^{5,6} Different epidemiological studies were performed from this perspective and identified high Lp(a) plasma concentration as a major and independent cardiovascular risk factor.⁷ Although some studies did not find a relationship between Lp(a) and coronary heart disease,^{8,9} most prospective studies (reviewed in Ref. 10) confirmed the initial results and extended its application to coronary restenosis, post-angioplastic occlusion, as well as cerebrovascular accidents, and premature development of atherosclerosis. The mechanism by which Lp(a) favors the atherogenic and/or thrombogenic processes is not clearly understood. However, Lp(a) has been found associated with fibrin deposits in atherosclerotic plaques.¹¹ Moreover, recent data indicate that the large structural heterogeneity of the apo(a) molecule is conducive to a functional heterogeneity as a plasminogen competitive inhibitor for fibrin binding that lowers the formation of plasmin and thereby decreases fibrinolysis.^{12,13} In order to understand the molecular basis of this mechanism we analyze the structural aspects of plasminogen and apo(a) interactions with fibrin and consider recent evidence indicating that Lp(a) behaves as an inhibitor of fibrinolysis.

STRUCTURAL BASIS FOR THE ANTIFIBRINOLYTIC ACTIVITY OF LP(A)

Plasminogen and Apolipoprotein(a): General Structure

Apo(a) and plasminogen are constituted by structural domains called kringles and a serine-proteinase region (see FIGURE 1). The kringle structure was first described in prothrombin and is found in several copies in proteins that evolved from a common ancestral gene: plasminogen, apo(a) and hepatocyte growth factors.⁴ Kringles

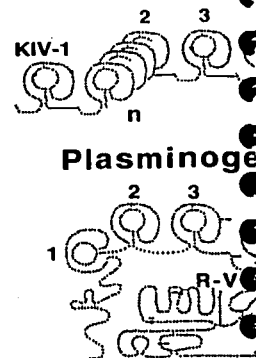


FIGURE 1. Structural homology (adapted from Petersen *et al.*, *J. Biol. Chem.* 261: 11111-11114, 1986) of five kringle domains and a serine-proteinase region of plasminogen kringle 5 (KV) and plasminogen kringle 4 (KIV). These kringle domains are present in ten different types (KIV-1 to KIV-10) of apo(a) isoforms. The presence of several apo(a) isoforms facilitates the Arg561-Val562 plasminogen cleavage by Ser-11e, S-1 (see FIG. 2).

are sequences of 80 to 90 amino acids, stabilized by three disulfide bonds. Kringles 1 to 5, differ from each other in that they show high affinity for fibrin, while kringles 6 to 10 show low affinity. The interaction between lysine residues in fibrin and the lysine binding sites of plasminogen allows plasminogen to bind to fibrin.

Apo(a) contains a variable number of kringle domains. The homology with kringle 4 of plasminogen is followed by a single copy of plasminogen kringle 4 (KIV-10) with 94% homology with the corresponding region of plasminogen, which is connected by regions of 26-31 amino acids. These regions are alike; they are classified as type I. Each of these regions appears in a variable number of copies in different apo(a) isoforms, that account for 30% of the total weight, including 30% weight due to a specific lysine residue that ensures the covalent

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taining small apo(a) context, the develop- rino-lytic activity of ment of the cardio- bogenic Lp(a).

Plasmin; Kringle or; Urokinase; Ath- tes; Macrophages; or inhibitor.

st identified by Berg in imilar to that of LDL in B100. It was not until (a), the unique and dis- n gene family,⁴ both by (a) glycoprotein indeed the plasmin precursor. particle. These findings ind in Lp(a) a common epidemiological studies p(a) plasma concentra- Although some studies rt disease,^{8,9} most pro- results and extended its usion, as well as cere- rosclerosis. The mecha- ogenic processes is not ted with fibrin deposits that the large structural onal heterogeneity as a owers the formation of nderstand the molecular plasminogen and apo(a) iting that Lp(a) behaves

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nains called kringles and ture was first described at evolved from a com- rowth factors.⁴ Kringles

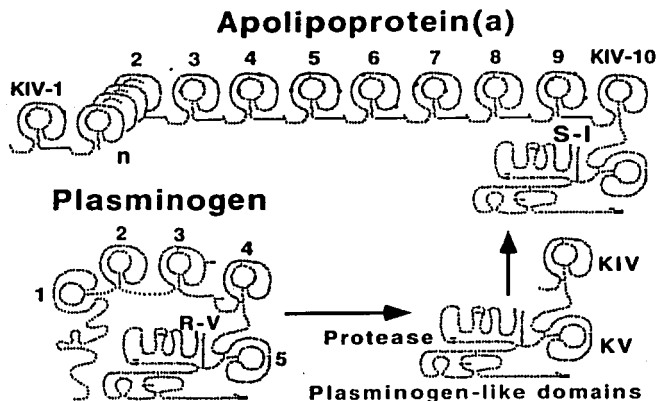


FIGURE 1. Structural homologies between apolipoprotein(a) and plasminogen (adapted from Petersen *et al.*, *J. Biol. Chem.*, 1990, **265**: 6104-6111). Plasminogen consist of five kringle domains and a serine-protease region. Apo(a) is constituted by single copies of plasminogen kringle 5 (KV) and of the protease region and by multiple copies of plasminogen kringle 4 (KIV). These KIV copies are not identical and have been classified¹⁸ in ten different types (KIV-1 to KIV-10). A variable number of KIV-2 determines the existence of several apo(a) isoforms. All other KIV types are present in single copies. R-V indicates the Arg561-Val562 plasminogen cleavage site, which in apo(a) has been substituted by Ser-Ile, S-I (see Fig. 2).

are sequences of 80 to 90 amino acid residues organized in a triple-loop structure, stabilized by three disulfide bridge.¹⁴ The kringle domains of plasminogen, designated 1 to 5, differ from each other. Kringles 1 and 4 contains lysine binding subsites that show high affinity for lysine residues in fibrin.^{15,16} The specific interactions between lysine residues in fibrin or cell membrane proteins and the lysine binding subsites of plasminogen allow plasminogen binding and activation.¹⁷

Apo(a) contains a variable number of kringle domains that share 61-75% homology with kringle 4 of plasminogen.^{2,3} The kringle 4-like repeats of apo(a) are followed by a single copy of plasminogen kringle 5 and a protease domain that shares 94% homology with the corresponding domain of plasminogen. Kringles are interconnected by regions of 26-36 serine-, proline-, and/or threonine-rich amino acids with six potential O-glycosilation sites. Not all plasminogen-like kringle 4 of apo(a) are alike; they are classified in 10 different types,¹⁸ hereafter referred as KIV-1 to KIV-10 (see FIG. 1). Each of these kringles is present only once, except KIV-2, which appears in a variable number, originating structural heterogeneity and different size isoforms, that account for molecular weights between about 280 and 800 kDa including 30% weight due to glycosilation.¹⁹ KIV-9 possesses an additional cysteine residue that ensures the covalent binding between apo(a) and Cys3734 of apo B-100

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and, thereby, the formation of the Lp(a) particle.²⁰ By sequence comparison and molecular modeling²¹ it has been shown that KIV-10 contains a lysine binding site (LBS) similar to that of plasminogen kringle 4, and that slightly modified LBS are present in KIV-5 to KIV-8. These kringle copies may confer to apo(a) binding capabilities similar to those of plasminogen.

Structure of the Lysine Binding Site

The structure of the lysine binding site, an elongated hydrophobic depression lined with aromatic residues that separate by 6.8 Å a cationic group from an anionic group positioned at opposite ends of the trough, has been well defined.²² Its structural characteristics generate a relatively rigid geometry that allows selective access and binding of C-terminal lysine residues in fibrin, as well as aliphatic or aromatic ligands of ω-amino carboxylic acid type. Based on information derived from biochemical,²³ mutagenesis,²⁴ crystallographic,^{25,26} and spectroscopic²⁷ studies, an appropriate set of amino acids that are key components for lysine binding has been identified (see TABLE 1). Thus, the LBS of plasminogen K1 consists of three aromatic residues, Tyr64/Trp62/Tyr72, lining the hydrophobic trough of the LBS and the double charged anionic, Asp55/Asp57, and cationic, Arg35/Arg71 centres located at opposite ends of the hydrophobic trough. In apo(a) only KIV-10 contains an LBS with similar characteristics that is identical to that of plasminogen kringle 4 except for a conservative Arg→Lys35 substitution also present in plasminogen kringle 1.²⁸

TABLE 1. Key components of the lysine-binding site

Kringle	Anionic pole		Hydrophobic aromatic residues			Cationic pole	
Pg K1	Asp 55	Asp 57	Tyr 64	Trp 62	Tyr 72	Arg 35	Arg 71
Pg K4	Asp 55	Asp 57	Phe 64	Trp 62	Trp 72	Lys 35	Arg 71
Pg K5	Asp 55	Asp 57	Tyr 64	Trp 62	Trp 72	Ile 35	Leu 71
KV	Asp 55	Asp 57	Tyr 64	Trp 62	Trp 72	Ser 35	Leu 71
KIV-10	Asp 55	Asp 57	Phe 64	Trp 62	Trp 72	Arg 35	Arg 71
KIV-9	Asp 55	Gly 57	Tyr 64	Trp 62	Trp 72	Arg 35	Arg 71
KIV-8	Asp 55	Glu 57	Tyr 64	Trp 62	Trp 72	Arg 35	Arg 71
KIV-7	Asp 55	Glu 57	Tyr 64	Trp 62	Trp 72	Arg 35	Arg 71
KIV-6	Asp 55	Glu 57	Tyr 64	Trp 62	Trp 72	Arg 35	Arg 71
KIV-5	Asp 55	Glu 57	Tyr 64	Trp 62	Trp 72	Arg 35	Arg 71
KIV-4	Asp 55	Val 57	Tyr 64	Trp 62	Trp 72	Arg 35	Arg 71
KIV-3	Asp 55	Val 57	Tyr 64	Tyr 62	Trp 72	Arg 35	Arg 71
KIV-2	Asp 55	Val 57	Tyr 64	Tyr 62	Trp 72	Arg 35	Arg 71
KIV-1	Asp 55	Val 57	Tyr 64	Tyr 62	Trp 72	Arg 35	Arg 71

NOTE: Pg. Plasminogen kringles are designated K1, K4, and K5. KV and KIV-1 to KIV-10 correspond to apo(a) kringles.

Both plasminogen K4 and apo(a) K1 by two important substitutions are, therefore, specific structure apo(a) KIV-10. The critical argumented; its hydrophobic indol and serves as an obvious mark

The Serine-Protein

Serine-proteinases are synth activated by proteolytic pro enzymes.²⁹ These include bind enzyme (S₁-S_n and S'₁-S'_n) cleavage site on the substrate Arg₅₆₁-Val₅₆₂, peptide scissile active plasmin.³¹ This cleavage must restrict the conformation disulfide bond is of important sequence corresponding to this of plasminogen that is strictly bond in apo(a) consist of Ser-1 recognition by activators (F) plasminogen prevent its cleava

We have recently consider residues at, respectively, P-P of specific fibrinolytic tests age site by activators and gene sis, r-apo(a) mutant plasmid HepG2 cells; culture superna and the corresponding r-Lp were treated with u-PA or r

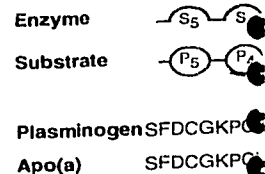


FIGURE 2. Scheme of the region in apo(a). Amino acid 585, according to Petersen *et al.* responding sequence in apo(a) activators cleave plasminogen (the arrow). In apo(a), amino plasminogen, however the clea

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sequence comparison and gains a lysine binding site. Slightly modified LBS are reported to apo(a) binding capacity.

hydrophobic depression of a group from an anionic site well defined.²² Its structure allows selective access as aliphatic or aromatic nature derived from biophysical studies,²⁷ an lysine binding has been shown to consist of three aromatic residues of the LBS and the Arg71 centres located at KIV-10 contains an LBS similar to plasminogen kringle 4 except for plasminogen kringle 1.²⁸

Res	Cationic pole	
2	Arg 35	Arg 71
2	Lys 35	Arg 71
2	Ile 35	Leu 71
2	Ser 35	Leu 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71
2	Arg 35	Arg 71

KV and KIV-1 to KIV-1C

Both plasminogen K4 and apo(a) KIV-10 however, differ from plasminogen kringle 1 by two important substitutions, Phe64 and Trp72, instead of Tyr64 and Tyr72, that are, therefore, specific structure requirements for the LBS of plasminogen K4 and apo(a) KIV-10. The critical role of this residue in lysine binding has been well documented; its hydrophobic indole is unusually exposed in the LBS of K4 and KIV-10 and serves as an obvious marker of the binding site.²⁵

The Serine-Proteinase Domain in Plasminogen and Apo(a)

Serine-proteinases are synthesized and secreted as single-chain zymogens that are activated by proteolytic processing through specific interactions with active enzymes.²⁹ These include binding forces between subsites of the active site of the enzyme (S_1-S_n and $S'_1-S'_n$) and amino acid residues (P_1-P_n and $P'_1-P'_n$) of the cleavage site on the substrate³⁰ (see FIGURE 2). For instance, cleavage of the $P_1-P'_1$, Arg₅₆₁-Val₅₆₂ peptide scissile bond in plasminogen by its activators yields two-chain active plasmin.³¹ This cleavage site is located in a small disulfide-bounded loop that must restrict the conformation around this bond. It has been shown, indeed, that this disulfide bond is of importance for the specificity of plasminogen activation.³² The sequence corresponding to this loop is comprised in the P_2-P_{17} / $P'_2-P'_{23}$ sequence of plasminogen that is strictly preserved in apo(a); in contrast, the $P_1-P'_1$ peptide bond in apo(a) consist of Ser-Ile instead of Arg-Val,² a substitution that may impair recognition by activators (Fig. 2). Indeed, the introduction of these substitutions in plasminogen prevent its cleavage by activators.³³

We have recently considered the possibility that introduction of both Arg and Val residues at, respectively, $P-P'_1$ by site directed mutagenesis together with the use of specific fibrinolytic tests may ensure potential recognition of this activation cleavage site by activators and generation of plasmin-like activity.³⁴ To test this hypothesis, r-apo(a) mutant plasmids encoding this sequence were expressed in 293 cells and HepG2 cells; culture supernatants containing wildtype r-apo(a), Arg-Val r-apo(a), and the corresponding r-Lp(a)-like particles were thus obtained. These products were treated with u-PA or t-PA in the presence of cofactors known to stimulate

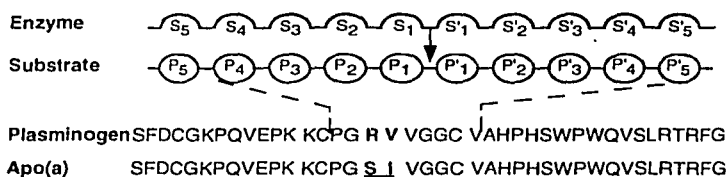


FIGURE 2. Scheme of the plasminogen activation cleavage site and the corresponding region in apo(a). Amino acid sequences of the cleavage site in plasminogen (residues 544-585, according to Petersen *et al. J. Biol. Chem.* 1990, 265, 6104-6111) compared to the corresponding sequence in apo(a) (residues 4291-4332, according to Ref. 3). Plasminogen activators cleave plasminogen at the $P_1-P'_1$ (Arg-Val) scissile peptide bond (indicated by the arrow). In apo(a), amino acid residues P_2-P_{17} and $P'_2-P'_{23}$ are identical to those of plasminogen, however the cleavage site is substituted by Ser-Ile.

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plasminogen activation by t-PA (fibrin and CNBr-fragments of fibrinogen) or u-PA (6-Ahx, a lysine analogue). However, neither cleavage of apo(a) nor plasmin-like activity could be demonstrated on treatment of Arg-Val r-apo(a)/r-Lp(a) mutants with plasminogen activators, either in the presence or in the absence of cofactors known to induce changes in plasminogen conformation or to stimulate plasminogen activation.

These results strongly suggest that to promote efficient cleavage of plasminogen, t-PA recognises complex structural elements on the surface of native plasminogen that are distant from those at P4-P2'.³⁵ The exact nature of these secondary interactions remains unknown. Because r-apo(a) contains multiple copies of kringle 4 and lacks copies of kringles 1 to 3 of plasminogen, it may not possess the potential for such enzyme-substrate interactions, that are independent of the binding cleft around P1; r-apo(a) Arg-Val may fail, thereby, to induce a better fit of the cycle peptide loop containing the Arg-Val bond with the activators. The presence of multiple kringle repeats in apo(a) does not substitute the specific conformation present in plasminogen that is required for the selective and specific recognition by activators. It is, therefore, apparent that besides the absence of the Arg-Val site, apo(a) has undergone further evolutionary changes which prevent its activation towards a plasmin-like protease. Because of this important enzymatic difference, the ability of selected kringle-4 copies of apo(a) to compete with plasminogen for binding to fibrin and cell surfaces results in decreased fibrinolysis and impaired pericellular proteolysis.^{36,37}

INHIBITION OF PLASMINOGEN BINDING TO FIBRIN, THE MAJOR MECHANISM OF ACTION OF LP(A)

Physiological fibrinolysis involves heterogeneously catalyzed reactions that proceed at the fibrin/plasma interface, where fibrin provides a surface to which tissue plasminogen activator (t-PA) and plasminogen bind specifically.^{15,17} Molecular assembly of these proteins results in a ternary complex that efficiently generates plasmin on the surface of fibrin and thereby triggers the dissolution of a clot.³⁸

Initial limited degradation of the surface of fibrin by plasmin unveils carboxy-terminal lysine residues and increases the local concentration of plasminogen, a process that amplifies and accelerates the degradation of fibrin.³⁹ This mechanism is mediated by specific interactions of lysine residues in fibrin with the kringle domains of plasminogen.⁴⁰ In a plasma milieu, the progression of this process is markedly influenced by α_2 -antiplasmin, the specific plasmin inhibitor, and by active carboxy-peptidase B, an exopeptidase that cleaves Arg and Lys residues at C-terminal position.⁴¹ The number of carboxy-terminal lysine residues is thus limited and, thereby, the amount of bound plasminogen.¹⁷ On the other hand, the blockade of such residues by isolated plasminogen kringle 4 has been shown to interfere competitively with clot lysis by a mechanism involving binding to lysine-fibrin residues.⁴² The presence in apo(a) of kringle modules structurally related to those of plasminogen endows Lp(a) with the ability to compete with plasminogen for binding to fibrin. The effect of Lp(a) on plasminogen binding to fibrin and cell surfaces has been studied by several groups.^{37,43-45} A number of experimental *in vitro* studies resulted in convincing evidence that Lp(a) binds to the fibrin surface and cell membranes and, thereby, competes with plasminogen so as to inhibit its activation.^{46,47} We have shown that, indeed, both free recombinant apo(a) (see FIGURE 3) and apo(a) in Lp(a) particles (see

FIGURE 3. Determination of plasminogen to fibrin (as indicated in Ref. 12) formed in the absence and in the presence of Lp(a) (18 kringles). For each concentration of Lp(a), the amount of plasminogen bound, and K_d , the dissociation constant. The graph depicts the reciprocal of the bound plasminogen versus the reciprocal of the free plasminogen. The inhibition constant ($K_i = 4$ nmol/L) is indicated by the abscissa.

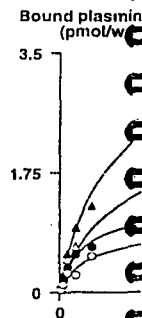


FIGURE 4. Inhibition of plasminogen binding to fibrin by Lp(a) (as indicated in Ref. 12) containing a trace amount of fibrin in the presence of 0 (Δ), the Langmuir equation allowed to be fitted to the data (bound) and K_d (the dissociation constant) was determined. By plotting the reciprocal of the bound plasminogen versus the reciprocal of the free plasminogen, the inhibition constant $K_i = 32$ nmol/L was determined (not shown).

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f fibrinogen) or u-PA (a) nor plasmin-like (a)/r-Lp(a) mutants absence of cofactors imulate plasminogen

age of plasminogen, native plasminogen secondary interactions of kringle 4 and less the potential for binding cleft around the cycle peptide loop of multiple kringle present in plasminogen by activators. It is, e, apo(a) has under- towards a plasmin- the ability of selected ling to fibrin and cell ular proteolysis.^{36,37}

FIBRIN, LP(A)

d reactions that pro- face to which tissue ally.^{15,17} Molecular efficiently generates tion of a clot.³⁸ in unveils carboxy- plasminogen, a pro-³⁹ This mechanism in with the kringle n of this process is ibitor, and by active esidues at C-terminal as limited and, there- blockade of such res- erere competitively brin residues.⁴² The ose of plasminogen nding to fibrin. The has been studied by resulted in convinc- branes and, thereby, We have shown that, 1 Lp(a) particles (see

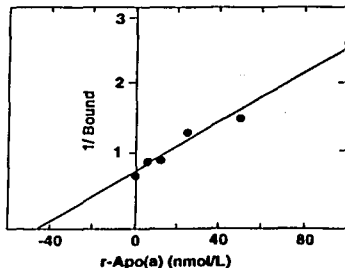


FIGURE 3. Determination of the inhibition constant of apo(a) for the binding of plasminogen to fibrin (as indicated in Ref. 48). Plasminogen binding experiments were performed in the absence and in the presence of various concentrations of recombinant apo(a) (18 kringles). For each concentration of apo(a) the binding parameters (B_{max} , the maximum bound, and K_d , the dissociation constant) were determined according to the Langmuir equation. The graph depicts the reciprocal of B_{max} as a function of the concentration of apo(a). The inhibition constant ($K_i = 44$ nmol/L) was calculated from the *intercept* of the straight line with the abscissa.

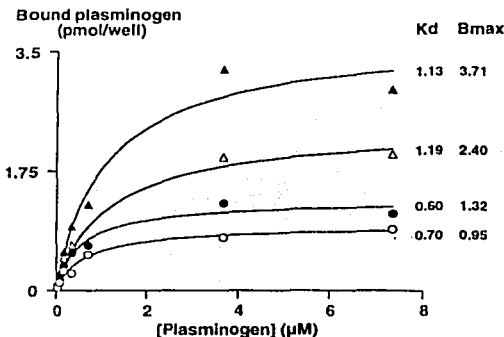


FIGURE 4. Inhibition of the binding of plasminogen to fibrin by purified 18 kringles Lp(a) (as indicated in Ref. 12). Increasing concentrations of plasminogen (0 to 8 μ mol/L) containing a trace amount of radiolabeled plasminogen (5 nmol/L) were incubated with fibrin in the presence of 0 (▲), 50 (△), 100 (●) and 200 (○) nmol/L Lp(a). Data fitted to the Langmuir equation allowed calculation of B_{max} (maximum amount of plasminogen bound) and K_d (the dissociation constant of the interaction) for each Lp(a) concentration. By plotting the reciprocal of the B_{max} versus the concentration of added Lp(a) an inhibition constant $K_i = 32$ nmol/L was determined at the *intercept* of the linear regression curve with the abscissa (not shown).

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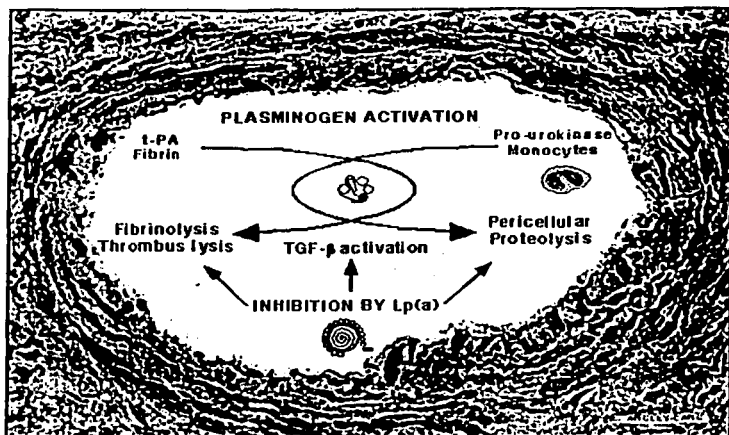


FIGURE 5. Competitive inhibition of the binding and activation of plasminogen by Lp(a) on fibrin and cell surfaces. This competitive mechanism results in inhibition of fibrinolysis, pericellular proteolysis and TGF- β activation. The different components of the plasminogen activation system and Lp(a) are represented in the intravascular space of a vessel section.

FIGURE 4) efficiently inhibit ($K_i = 44$ nmol/L and 30 nmol/L, respectively) the binding of plasminogen to fibrin^{12,48} (see FIGURE 5). Such unique behaviour was attributed to the fibrin-binding properties conferred by the kringle 4 repeats of apo(a).⁴⁸ Plasminogen and the different Lp(a) isoforms also compete for lysine residues on the surface receptors of endothelial cells,³⁷ platelets,⁴⁹ and mononuclear cells.⁴⁵ Thus, Lp(a) interferes with the evolution of fibrinolysis at the surface of fibrin, endothelial cells, monocytes, and platelets through binding of apo(a), an eternal zymogen that decreases the local concentration of plasminogen and cannot be transformed into an active enzyme. Most of the effects of Lp(a)—persistence of fibrin deposits, accumulation of cholesterol and proliferation of smooth muscle cells in the intima, are related to a decrease in plasmin activity (FIG. 5). Hypofibrinolysis and cholesterol accumulation are a direct consequence of the presence of Lp(a) at the surface of fibrin and cell membranes: apo(a) inhibits plasmin formation and the LDL components favors cholesterol accumulation.

OTHER ANTIFIBRINOLYTIC AND ANTIPROTEOLYTIC EFFECTS OF LP(A)

Lp(a) may stimulate the expression of PAI-1 and inhibit the synthesis of t-PA by endothelial cells in culture.^{50,51} Thus, inhibition of t-PA by PAI-1 and low t-PA antigen levels may enhance Lp(a)-dependent hypofibrinolysis by decreasing the amount of

t-PA available for the activation of apo(a), the generation of plasmin. Insufficient activation of plasminogen converts muscle cells into the intima.

ANTIFIBRINOLYTIC EFFECTS

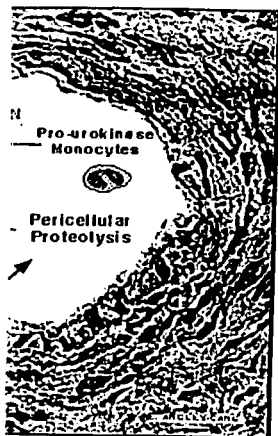
To date there is no direct evidence of plasmin binding and activation in experimental studies. Occlusive disease of large arterial segments with low plasmin levels.⁵⁴ Biomedical studies on endogenous and t-PA inhibition by apo(a) in the presence of apo(a) have been shown to exhibit a dose-dependent inhibition of plasminogen activation by injection of human apo(a) in a transgenic mouse model.⁵⁷

In humans Lp(a) plasma levels vary from one individual to another, from 10 mg/dl to more than 100 mg/dl, and are associated with cholesterol, obesity, and hypertension. However, there are large variations in Lp(a) plasma levels obtained in this human population. Plasminogen binding and plasminogen binding and measured the effect of Lp(a) on plasminogen activation at the surface of fibrin (see FIGURE 6). A direct inverse relationship between Lp(a) concentration and plasmin formation has thus been observed, which may decrease fibrinolysis.

LP(A) INHIBITS FIBRINOLYSIS

In general, an individual has a specific Lp(a) isoform; only a reduced number of isoforms exist. The size of the Apo(a) isoform is related to the Lp(a) concentration.⁶³ In general, the size of the Lp(a) isoform and, therefore, the size of the Lp(a) molecule. A difference in

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activation of plasminogen by Lp(a) results in inhibition of fibrinolysis, at components of the plasminogenolytic space of a vessel section.

101/L, respectively) the binding behaviour was attributed to 4 repeats of apo(a).⁴⁸ Plasminogen lysine residues on the surface of mononuclear cells.⁴⁵ Thus, Lp(a) coats the surface of endothelial cells, forming a protective layer that decreases plasminogen activation. Plasminogen is transformed into an active plasmin, which degrades fibrin deposits, accumulation of which in the intima, are related to atherosclerosis and cholesterol accumulation. The surface of fibrin and cell membrane components favors cholesterol

t-PA available for the activation of plasminogen. It has also been shown that Lp(a) inhibits the activation of TGF- β ⁵² and that, in transgenic mice expressing human apo(a), the generation of plasmin and thereby the activation of TGF- β are decreased.⁵³ Insufficient activation of TGF- β may result in migration and proliferation of smooth muscle cells into the intima, an important mechanism in atheroma plaque formation.

ANTIFIBRINOLYTIC ACTIVITY OF APOLIPOPROTEIN(A) *IN VIVO*

To date there is no definitive evidence that Lp(a) interferes with plasminogen binding and activation *in vivo*. Some evidence has been obtained from a few experimental studies. Occlusive arterial thrombosis with incorporation of Lp(a) into damaged arterial segments was observed in cynomolgus monkeys with high Lp(a) plasma levels.⁵⁴ Biemond *et al.*⁵⁵ studied the effect of a recombinant form of apo(a) on endogenous and t-PA mediated lysis in an *in vivo* model of experimental thrombosis; endogenous thrombolysis but not t-PA induced lysis was significantly reduced in the presence of apo(a). In contrast, transgenic mice that express human apo(a) have been shown to exhibit reduced t-PA induced lysis of pulmonary emboli produced by injection of human platelet-rich plasma clots.⁵⁶ However, the most conclusive report to this respect concerns the elimination of the pathogenic activity of apo(a) by disruption of the KIV-10 lysine binding site using genetic engineering in transgenic mice.⁵⁷

In humans Lp(a) plasma concentration, which depends on its hepatic synthesis,⁵⁸ varies from one individual to another within an approximate range of less than 10 mg/dl to more than 100 mg/dl; it is independent from other factors, such as diet, cholesterol, obesity, and smoking and is maintained within small variations during the life span. However, during the course of the nephrotic syndrome, large individual variations in Lp(a) plasma concentration have been observed.⁵⁹ Data have been obtained in this human model system,^{60,61} indicating that Lp(a) interferes with plasminogen binding and activation under conditions fashioned *in vivo*. We have measured the effect of individual variations in the concentration of Lp(a) on plasminogen activation at the surface of fibrin and have determined the existence of an inverse relationship between plasmin formation and quantity of Lp(a) that is bound (see FIGURE 6). A direct relationship between binding of Lp(a) to fibrin and the decrease in plasmin formation due to modifications *in vivo* of the Lp(a) plasma concentration has thus been established.^{60,61} These studies strongly suggest that apo(a) may decrease fibrinolysis *in vivo*.

LP(A) INHIBITS FIBRINOLYSIS DEPENDING ON APO(A) ISOFORMS

In general, an individual inherits, in a codominant autosomic fashion, two apo(a) isoforms; only a reduced number of individuals are homozygous for the apo(a) trait.⁶² The size of the Apo(a) isoforms accounts for an inverse correlation with Lp(a) plasma concentration.⁶³ In general, the smaller the hypervariable region of the apo(a) allele and, therefore, the size of the apo(a) isoform, the higher is the plasma concentration of Lp(a). A difference in the distribution of apo(a) isoforms between patients with

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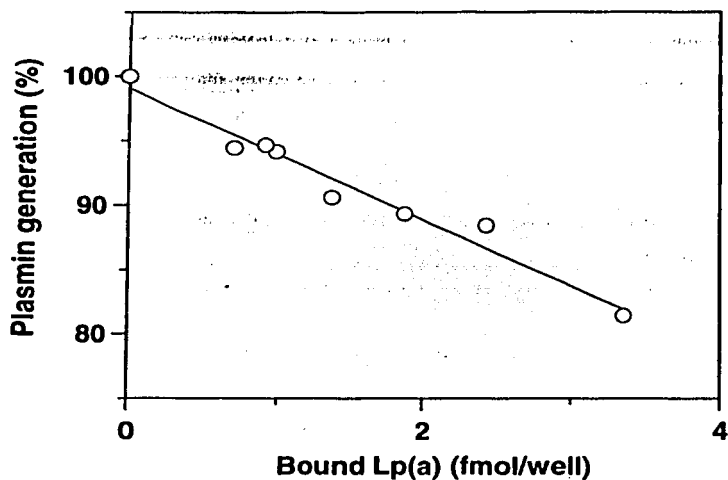


FIGURE 6. Inhibition of plasmin generation by Lp(a). Varying concentrations of Lp(a) (18 kringles) were incubated with 200 nmol/L of Glu-plasminogen (final concentration) on fibrin bound-t-PA. Plasminogen activation was followed at 37°C by measuring the amidolytic activity of plasmin with a synthetic substrate (CBS 00-65, Stago). At the end of the activation, the plate was washed and bound Lp(a) was detected with a radiolabeled antibody directed against apo(a); bound radioactivity was transformed in fmol of Lp(a). Plasminogen activation expressed as a percentage relative to the maximum of plasmin generated in the absence of Lp(a), is plotted against the amount of Lp(a) bound. Data represent the mean of three determinations. A linear correlation ($r = 0.985$) was found between the inhibition of plasmin generation and the increase in the amount of Lp(a) bound.

atherosclerosis and a control population has been recently reported.⁶⁴⁻⁶⁶ Low molecular mass isoforms were found more frequently in subjects with high Lp(a) concentration and a history of myocardial infarction or intermittent claudication. Thus, short apo(a) alleles may favor atherogenesis by increasing the concentration of Lp(a). However, the difference in allele distribution between patients at risk and controls is not always observed, and the inverse relationship between apo(a) size and Lp(a) concentration is not linear, thus suggesting the existing of a functional diversity among apo(a) isoforms. Indeed, some plasmas with a high Lp(a) concentration may fail to induce a decrease in fibrinolysis.⁴⁴ The question remains as to whether short apo(a) isoforms are *per se* an atherothrombotic risk factor, especially when associated with high Lp(a) plasma concentration. Recent experimental and clinical evidence provide arguments that favor this hypothesis.

Since the atherogenic potential conferred by the kringle Lp(a) phenotypes may have an affinity for fibrin. We demonstrated in subjects containing a single, different apo(a) allele, a significant effect and that this heterogeneity in low molecular mass showed a significant effect suggesting that the variation in the ability to bind to fibrin.⁶⁷ In heterozygosity, differences in affinity for fibrin were demonstrated.¹³ Under these conditions, the function of the affinity of apo(a) isoforms relative to each other and to the prevalence of low molecular mass isoforms in advanced stenotic atherosclerosis.

Our findings of an inverse relationship support the hypothesis that the affinity for fibrin. According to the data related to atherogenesis and

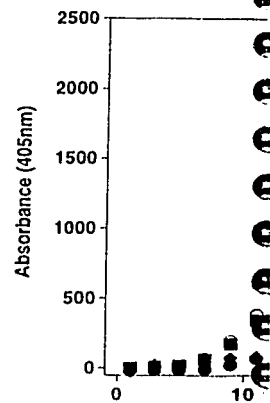
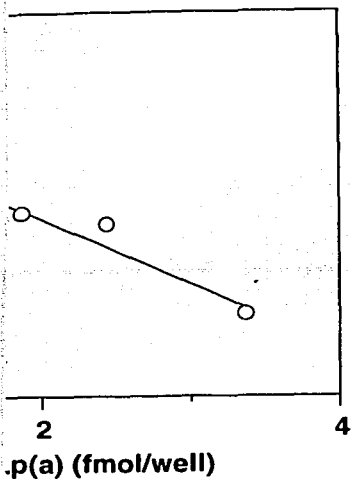


FIGURE 7. Inhibition of plasminogen activation by distinct Lp(a) particles. Plasminogen was activated by distinct Lp(a) particles and the detection of the amidolytic activity at 405 nm as a function of time. Data represent the mean of three determinations of an apo(a) isoform containing 18 kringles ($\Delta A_{405}/\text{min}$).

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tion by Lp(a). Varying concentrations of 100/L of Glu-plasminogen (final concentration was followed at 37°C by measuring the substrate (CBS 00-65, Stago). At the end of Lp(a) was detected with a radiolabeled antibody was transformed in fmol of Lp(a). Plasmin to the maximum of plasmin generated amount of Lp(a) bound. Data represent the inhibition ($r = 0.985$) was found between the inhibition the amount of Lp(a) bound.

been recently reported.⁶⁴⁻⁶⁶ Low molecular weight in subjects with high Lp(a) concentration or intermittent claudication. Thus, short Lp(a) increasing the concentration of Lp(a). The relationship between patients at risk and controls is not clear. The relationship between apo(a) size and Lp(a) concentration is not clear. The existence of a functional diversity among Lp(a) isoforms with a high Lp(a) concentration may fail to explain the relationship. The question remains as to whether short apo(a) Lp(a) is a risk factor, especially when associated with high Lp(a). Experimental and clinical evidence provide

Since the atherogenic potential of Lp(a) is related to the lysine binding properties conferred by the kringle 4 repeats of apo(a), we have explored the possibility that Lp(a) phenotypes may have different functional properties with regard to their affinity for fibrin. We demonstrated that Lp(a) particles purified from homozygous subjects containing a single, distinct isoform of apo(a) display different antifibrinolytic effect and that this heterogeneity is related to apo(a) size polymorphism. Isoforms of low molecular mass showed the highest affinity for fibrin (see FIGURE 7). This finding suggests that the variable number of kringles in apo(a) influence its ability to bind to fibrin.⁶⁷ In heterozygous subjects, representing more than 94% of the population,⁶² differences in affinity for fibrin of each of the Lp(a) particles have been demonstrated.¹³ Under these conditions, the net antifibrinolytic effect of Lp(a) is a function of the affinity of each of the Lp(a) isoforms and of their concentration relative to each other and to plasminogen.¹² This mechanism is in agreement with the emergence of low molecular weight apo(a) phenotypes as a leading risk condition of advanced stenotic atherosclerosis.⁶⁸

Our findings of an inverse relationship between isoform size and affinity for fibrin support the hypothesis that the real risk factor is the Lp(a) population with high affinity for fibrin. According to this concept some Lp(a) phenotypes might fail to be related to atherogenesis and, therefore, some individuals with high Lp(a) would not

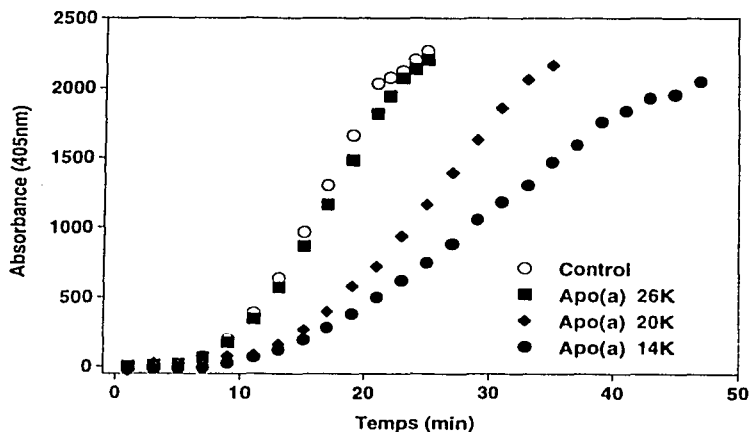


FIGURE 7. Inhibition of the formation of plasmin on fibrin by different Lp(a) isoforms. Plasminogen was activated by fibrin-bound t-PA in the presence of 0 or 300 nmol/L of distinct Lp(a) particles containing either 26, 20, or 14 kringles. Chromogenic substrate detection of the amidolytic activity of plasmin is expressed by the change in absorbance at 405 nm as a function of time. The amount of plasmin formed was lower in the presence of apo(a) isoforms containing 14 and 20 kringles as indicated by lower initial velocities ($\Delta A_{405}/\text{min}$).

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be at risk of coronary heart disease. Taken altogether, these results suggest that, besides the quantitative factor, an important qualitative effect must be considered in the atherothrombotic role of Lp(a). This new concept about the functional heterogeneity of Lp(a) adds a new dimension to the evaluation of the predictive value of Lp(a) as a risk factor for cardiovascular disease.

CONCLUSION

The similarity between plasminogen and apolipoprotein (a) allows different apo(a) isoforms to compete with plasminogen for fibrin affinity sites. The affinity of each isoform depends on its size and its plasma concentration.^{19,47,67} Indeed, we have clearly shown that the influence of Lp(a) on fibrinolysis depends on the high affinity of small apo(a) isoforms for fibrin and on their concentration relative to plasminogen. Lp(a) particles containing low molecular weight apo(a) isoforms have the most profound influence on fibrinolysis by acting as a prominent competitive antagonist of plasminogen. As a result, the plasmin formed at the surface of fibrin may also vary with modifications of the concentration of Lp(a) *in vivo*.⁶⁰ An antifibrinolytic mechanism may therefore explain the pathophysiological effects of Lp(a) by relating both high concentrations of Lp(a) and small apo(a) isoforms. This mechanism is in agreement with the emergence of low molecular weight apo(a) phenotypes as a leading risk condition of advanced stenotic atherosclerosis.⁶⁸ These data indicate that, to evaluate the potential risk associated with Lp(a) levels in heterozygous subjects, it is the relative concentration of the Lp(a) subpopulation having an apo(a) isoform with the highest affinity for fibrin and not the absolute Lp(a) concentration that should be considered.

ACKNOWLEDGMENTS

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