



Genômica Nutricional e Doenças Cardiovasculares

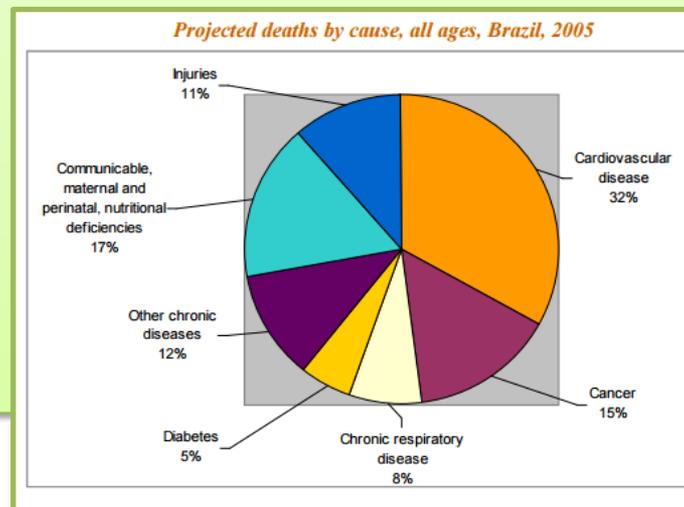
Ms. Mychelle Kytchia Rodrigues Nunes Duarte

A Doença Cardiovascular (DCV) - 30% das mortes globais (OMS).

No Brasil - 340.284 óbitos no ano de 2014,

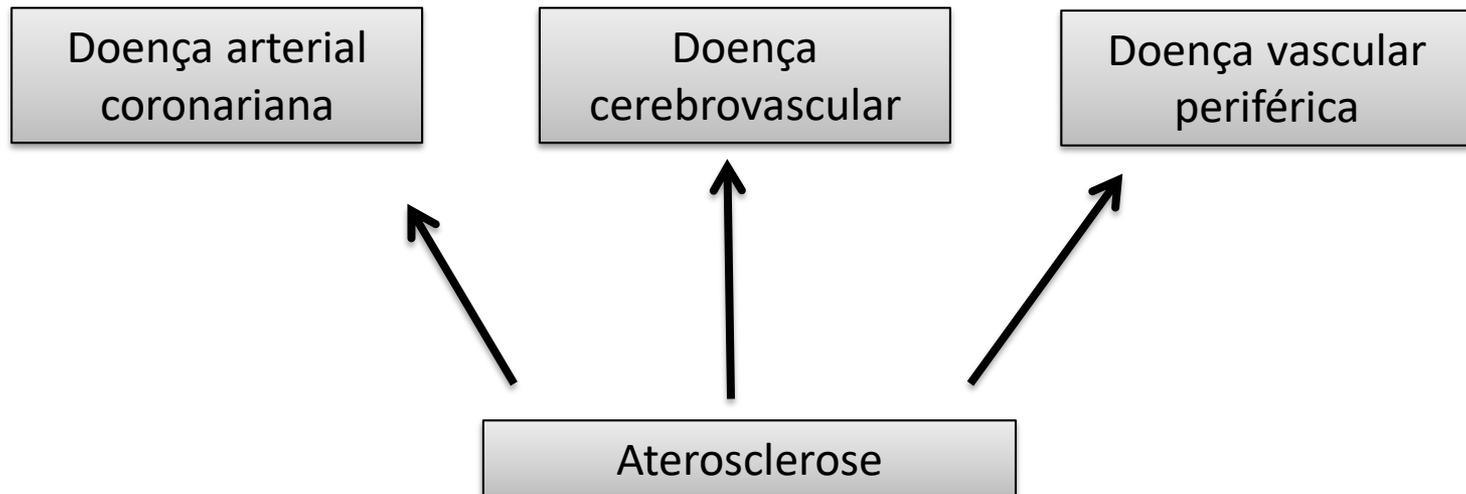
12.411 referente a doenças das artérias, das arteríolas e capilares segundo o o Código Internacional de Doenças (CID-10)

No Rio Grande do Norte no ano de 2014 - foram 5.164 óbitos devido as DCV.



As DCV alteram o funcionamento do sistema circulatório, formado por coração, vasos sanguíneos e linfáticos.

Podem ser classificadas em:



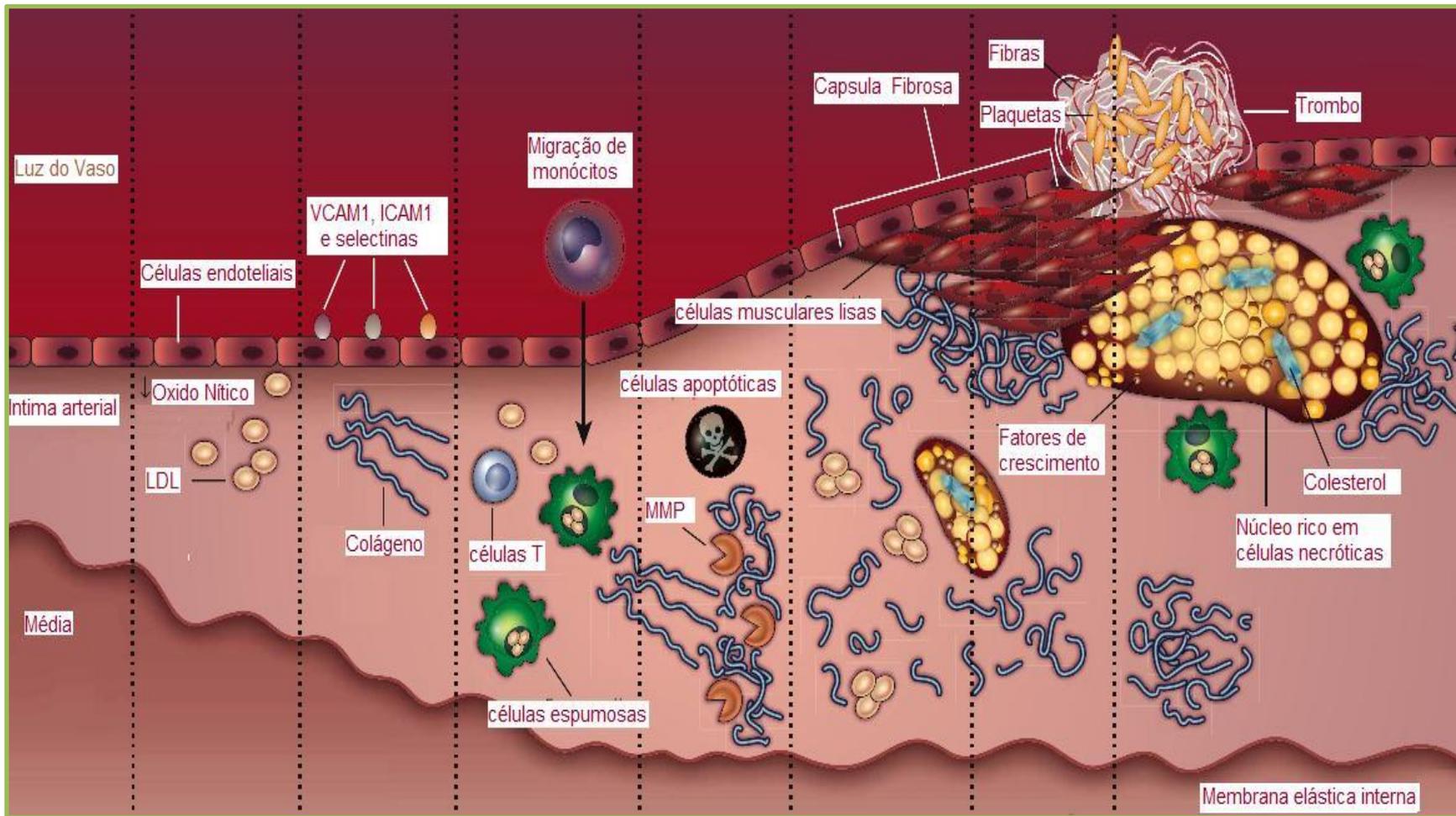


Figura 1. Representação da progressão do processo de aterosclerose. Adaptado de Sanz & Fayad, Nature, 2008.



BMC Med. 2014; 12: 158.
Published online 2014 Sep 25. doi: [10.1186/s12916-014-0158-6](https://doi.org/10.1186/s12916-014-0158-6)

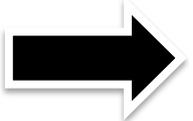
PMCID: PMC4199062

Dietary calcium intake and mortality risk from cardiovascular disease and all causes: a meta-analysis of prospective cohort studies

[Xia Wang](#), [Hongxia Chen](#), [Yingying Ouyang](#), [Jun Liu](#), [Gang Zhao](#), [Wei Bao](#)[✉] and [Maosheng Yan](#)[✉]

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Conclusions: This meta-analysis of prospective cohort studies suggests that dietary calcium intake is associated with cardiovascular mortality in a U-shaped manner and that high dietary calcium intake (>900 mg/day) is not associated with a decreased risk of all-cause mortality.



Eur J Nutr. 2012 December ; 51(8): 917–926. doi:10.1007/s00394-011-0268-2.

Nutrient and food intakes of middle-aged adults at low risk of cardiovascular disease: the international study of macro-/micronutrients and blood pressure (INTERMAP)



Food and food subgroup intake (g/1,000 kcal) by low cardiovascular disease (CVD) risk status^a, all participants (Japan, People's Republic of China, United Kingdom, United States), 1996–1999 the INTERMAP Study (*n* = 4,680)

Food subgroup ^b	Not low CVD risk (<i>n</i> = 3,923)	Low CVD risk (<i>n</i> = 757)	<i>P</i> value ^c
Total fruits	76.6 (28.0–145.5)	99.1 (48.9–168.6)	<0.001
Total vegetables	152.2 (104.4–212.1)	177.0 (124.0–244.0)	<0.001
Total grains	147.4 (91.2–283.9)	209.6 (120.4–332.7)	0.085
Pasta, rice, including recipes	54.7 (14.7–191.0)	128.8 (34.2–247.8)	<0.001
Nuts and legumes	3.9 (0.2–15.2)	3.0 (0.1–11.4)	0.024
Non-high-fat dairy (milk, yogurt, frozen yogurt, etc.)	40.5 (2.8–105.9)	49.5 (0.7–118.1)	0.020
High-fat dairy (cream, cheese, ice cream, milk, cheese, recipes, etc.)	5.4 (0.0–18.9)	3.3 (0.0–15.0)	<0.001
Fish, fish roe, shellfish	6.6 (0.0–30.6)	12.3 (0.0–35.4)	0.002
Poultry	9.0 (0.0–22.5)	6.9 (0.0–18.0)	<0.001
Beef, pork, veal, game meats	18.5 (7.4–34.6)	14.8 (4.9–27.9)	<0.001
Eggs	9.2 (2.3–19.0)	8.4 (2.0–17.5)	0.031
Processed meats	3.3 (0.0–10.9)	0.0 (0.0–7.4)	<0.001
Total visible fats (animal fats, margarines, table spreads, oils, shortenings, dressings)	13.1 (8.4–19.1)	13.3 (8.5–18.9)	0.874
Snacks, sweets	16.8 (5.6–30.7)	17.70 (6.2–33.4)	0.074
Alcoholic beverages	0.9 (0.0–86.5)	0.3 (0.0–18.4)	<0.001
Non-alcoholic beverages (excluding tea, coffee)	226.0 (0.0–576.0)	41.00 (0.0–485.7)	<0.001
Sugar-sweetened beverages ^d (UK and US samples only)	62.7 (0.0–188.7)	49.6 (0.0–142.8)	0.007
Total energy intake (kcal/day), mean (SD)	2,194.2 (633.2)	1,908.5 (514.7)	<0.001



CVD cardiovascular disease

Conclusions—Lower energy intake and differential intake of multiple specific nutrients and foods are characteristic of individuals at low risk for developing CVD. Identification of dietary habits associated with LR is important for further development of public health efforts aimed at reduction/prevention of CVD.

Lower Zinc Bioavailability May Be Related to Higher Risk of Subclinical Atherosclerosis in Korean Adults

Su Kyoung Jung^{1,2}, Mi-Kyung Kim^{1,2*}, Young-Hoon Lee³, Dong Hoon Shin⁴, Min-Ho Shin⁵, Byung-Yeol Chun⁶, Bo Youl Choi^{1,2}

J Am Soc Nephrol. 2009 Aug; 20(8): 1797–1804.

PMCID: PMC2723984

doi: [10.1681/ASN.2008060649](https://doi.org/10.1681/ASN.2008060649)

High Protein Intake Associates with Cardiovascular Events but not with Loss of Renal Function

Nynke Halbesma,* [Stephan J.L. Bakker](#),* [Desiree F. Jansen](#),† [Ronald P. Stolk](#),† [Dick De Zeeuw](#),‡ [Paul E. De Jong](#),* [Ronald T. Gansevoort](#),§* and for the PREVEND study group

Europe PMC Funders Group

Author Manuscript

***Cochrane Database Syst Rev.* Author manuscript; available in PMC 2014 September 15.**

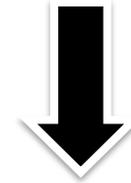
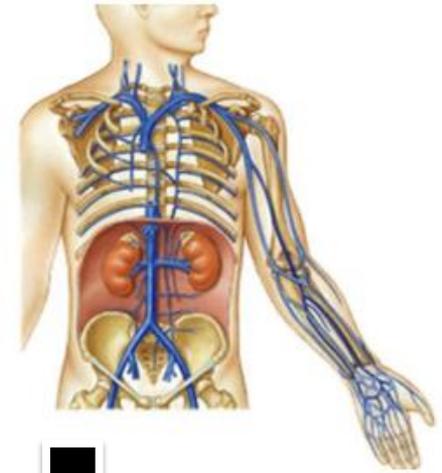
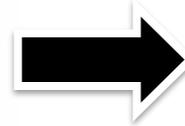
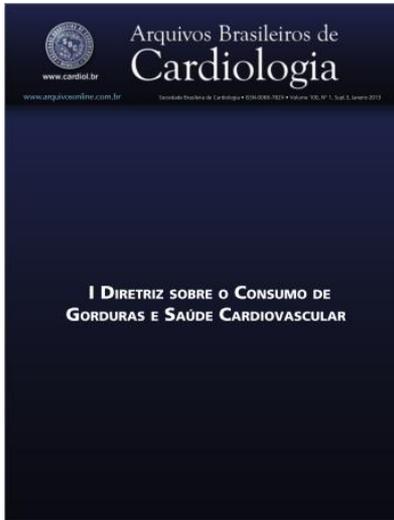
Published in final edited form as:

Cochrane Database Syst Rev. ; (7): CD002137. doi:10.1002/14651858.CD002137.pub2.

Reduced or modified dietary fat for preventing cardiovascular disease

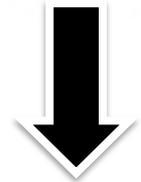
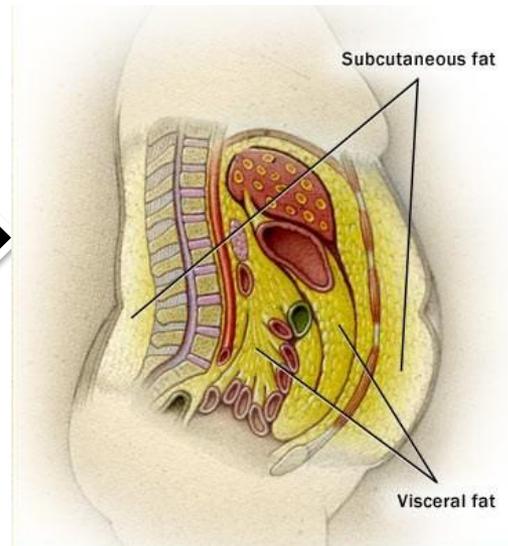
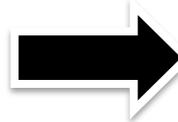
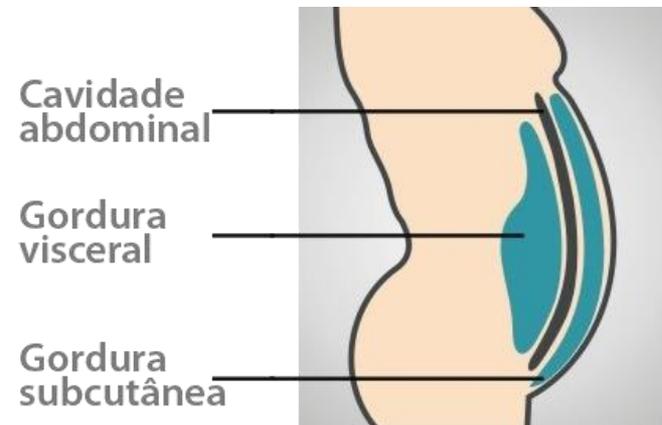
Lee Hooper¹, Carolyn D Summerbell², Rachel Thompson³, Deirdre Sills⁴, Felicia G Roberts⁵, Helen Moore², and George Davey Smith⁶





Saturated fatty acid	Monounsaturated fatty acid	Polyunsaturated fatty acid
 arachidic $C_{20}H_{40}O_2$	 erucic $C_{22}H_{42}O_2$	 arachidonic $C_{20}H_{32}O_2$
 stearic $C_{18}H_{36}O_2$	 oleic $C_{18}H_{34}O_2$	 linoleic $C_{18}H_{32}O_2$
 palmitic $C_{16}H_{32}O_2$		





OPEN ACCESS Freely available online

PLOS ONE

A Pilot Investigation of Visceral Fat Adiposity and Gene Expression Profile in Peripheral Blood Cells

Masaya Yamaoka¹, Norikazu Maeda^{1*}, Seiji Nakamura², Susumu Kashine¹, Yasuhiko Nakagawa¹, Aki Hiuge-Shimizu¹, Kohei Okita¹, Akihisa Imagawa¹, Yuji Matsuzawa³, Ken-ichi Matsubara², Tohru Funahashi¹, Iichiro Shimomura¹



Novel genes detected by transcriptional profiling from whole-blood cells in patients with early onset of acute coronary syndrome



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^a Faculty of Pharmaceutical Sciences, University of Sao Paulo, Brazil

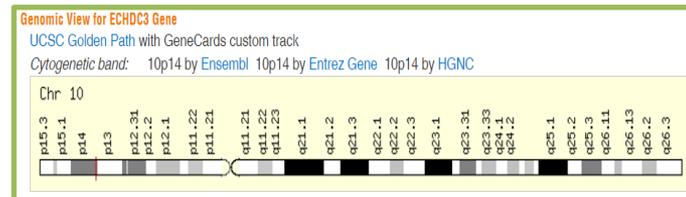
^b Department of Clinical and Toxicologic Analyses, Federal University of Rio Grande do Norte, Brazil

^c Group of Genomic Medicine, Galician Foundation of Genomic Medicine, CIBERER-University of Santiago, Spain

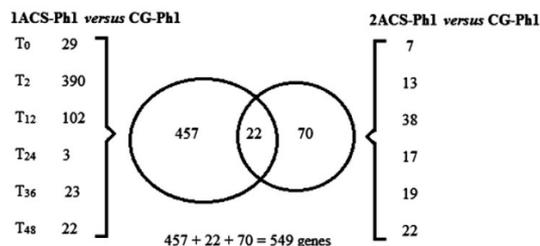
^d Genetics of Cardiovascular and Ophthalmologic Diseases, Hospital-University Complex of Santiago (CHUS), Spain

^e Functional Genomics Node National Institute for Bioinformatics (INB), Centro de Investigación Príncipe Felipe, Spain

^f Institute Dante Pazzanese of Cardiology, Brazil



Phase 1



Microarray gene expression

ADORA3, ALOX15, AREG, BCL2A1, BCL2L1, CA1, COX7B, CREM, CSF1, ECHDC3, FOLR3, GPX3, GSTM1, IL18R1, IL1R1, IL1RL1, IRS2, KCNE1, MMP9, MYL4, MYO5C, PER1, POLE2, RCN3, SAMSN1, SLC29A1, SLITRK4, SMPD3, THBS1, TLR4 and TREML4

Total = 31 genes

ALOX15, AREG, BCL2A1, BCL2L1, CA1, COX7B, ECHDC3, FOLR3, GSTM1, IL18R1, IL1RL1, IRS2, KCNE1, MMP9, MYL4, PER1, POLE2, SLC29A1, SMPD3, THBS1 and TREML4

Total = 21 genes

Phase 2

ACS-Ph2 versus CG-Ph2

ALOX15, AREG, BCL2A1, BCL2L1, CA1, COX7B, **ECHDC3**, IL18R1, IR2, KCNE1, MMP9, MYL4 and TREML4

Total = 13 genes

real-time PCR technical validation

real-time PCR biological validation

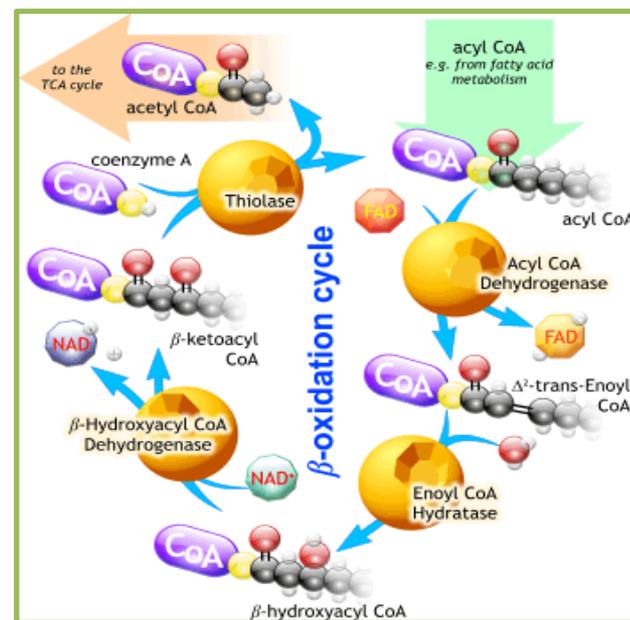
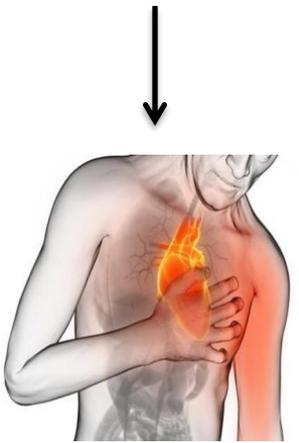
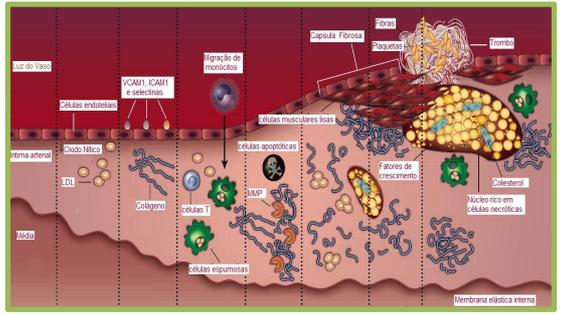
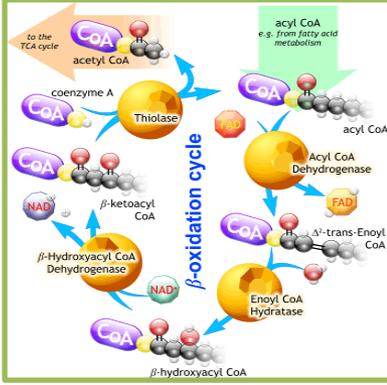
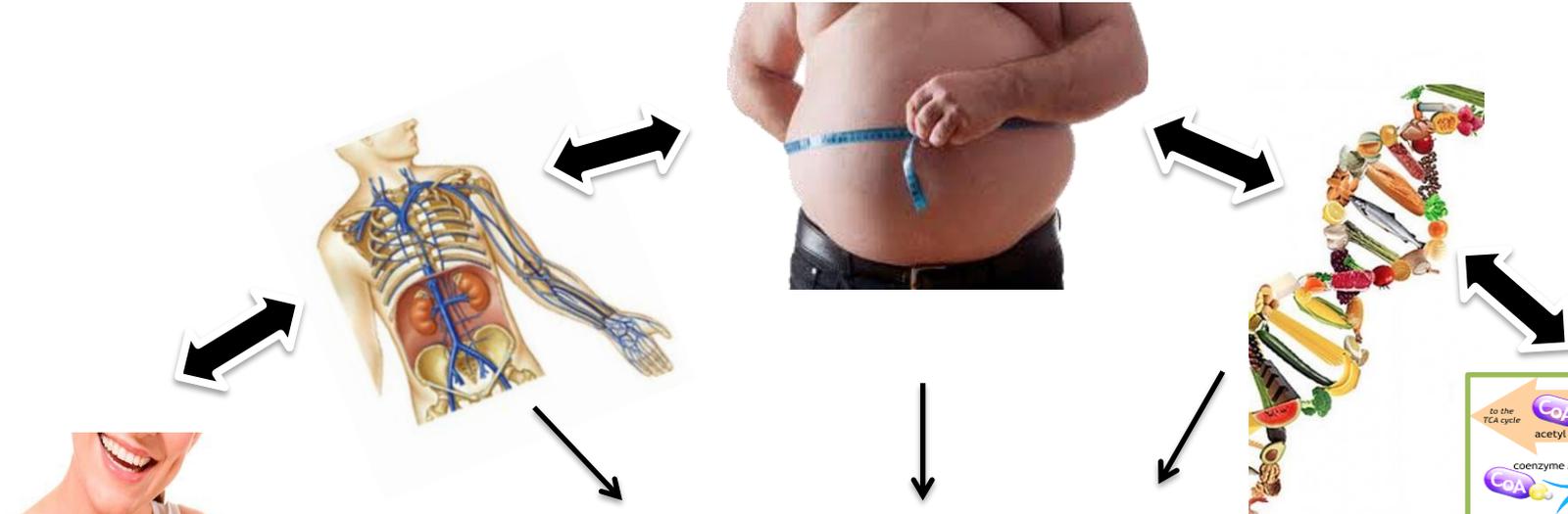


Fig. 1. Candidate gene expression biomarkers history of heart ischemia CG-Ph1: control group from phase 1; 1ACS-Ph1: patient with acute coronary syndrome without previously historic of heart ischemia from phase 1; 2ACS-Ph1: patient with acute coronary syndrome with previously history of heart ischemia from phase 1; CG-Ph2: control group from phase 2; ACS-Ph2: patient with acute coronary syndrome from phase 2; T₀: first stage; T₂: second stage; T₁₂: third stage; T₂₄: fourth stage; T₃₆: fifth stage; T₄₈: sixth stage.



O artigo intitulado “ ***The relationship of the oleic acid level and ECHDC3 mRNA expression with the extent of coronary lesion***” foi publicado no periódico “Lipids in Health and Disease”.

Duarte et al. *Lipids in Health and Disease* (2016) 15:144
DOI 10.1186/s12944-016-0312-6

Lipids in Health and Disease

RESEARCH

Open Access

The relationship of the oleic acid level and *ECHDC3* mRNA expression with the extent of coronary lesion



Mychelle Kytchia Rodrigues Nunes Duarte¹, Jéssica Nayara Góes de Araújo², Victor Hugo Rezende Duarte², Katiene Macêdo de Oliveira², Juliana Marinho de Oliveira³, Antonio Augusto Ferreira Carioca⁴, Raul Hernandez Bortolin², Adriana Augusto Rezende², Mario Hiroyuki Hirata⁵, Rosário Domingues Hirata⁵, Dan Linetzky Waitzberg⁶, Severina Carla Vieira Cunha Lima⁷, André Ducati Luchessi² and Vivian Nogueira Silbiger^{2*}

Identificar biomarcadores para o diagnóstico precoce das doenças cardiovasculares por meio da relação de biomoléculas e do estado nutricional com a extensão da lesão aterosclerótica.

Objetivos Específicos:

- Selecionar pacientes com risco cardiovascular, que tenham sido encaminhados para realização da cinecoronariografia pela primeira vez;
- Avaliar o perfil bioquímico, extensão da lesão aterosclerótica, concentração sérica de ácidos graxos, expressão do RNAm do *ECHDC3*, consumo alimentar e dietético em pacientes sem lesão e diferentes extensões de lesão aterosclerótica;
- Correlacionar a concentração sérica de ácidos graxos, o consumo alimentar e dietético, dados antropométricos, expressão do *ECHDC3* em pacientes sem lesão e diferentes extensões de lesão aterosclerótica;

Metodologia



HUOL
Hospital Universitário
Onofre Lopes

HEMODINÂMICA DO HUOL



Recrutamento

Inclusão

Pacientes
30-74 anos
Ambos os sexos

Realizar a
cinecoronariografia
pela primeira vez

Aprovado pelo Comitê de Ética
Protocolo 520/11
CAAE: 0001.0.051.294-11

Exclusão

Doença pré-estabelecida

Cardiomiopatia

Doença Hepática

Doença Renal

IAM

Distúrbios endócrinos,
com exceção de
diabetes

Metodologia



Primeira Casuística

Triagem de pacientes - 59

Aprovado pelo Comitê de Ética
Protocolo 520/11
CAAE: 0001.0.051.294-11

Mensuração da
pressão arterial

coleta de sangue
periférico

Cinecoronariografia

Antropometria

Friesinger

Peso e altura

Exames laboratoriais

Glicose

Creatinina

Colesterol

Uréia

Triglicérides

Ácido úrico

HDLc

AST

LDL

ALT

Concentração sérica
de ácidos graxos

Cromatografia
gasosa

Expressão gênica

Células mononucleares
Extração de RNA
cDNA
PCR real time

Metodologia



Segunda Casuística

Triagem de pacientes - 41

Aprovado pelo Comitê de Ética
Protocolo 520/11
CAAE: 0001.0.051.294-11

Coleta de sangue
periférico

Cinecoro-
nariografia

Consumo
Alimentar e
Dietético

Antropometria

Aferição de
pressão
arterial

Friesinger

2 -R24h

Virtual Nutri

Peso e altura

Perímetros

Composição
corpórea

Percentual de
gordura

Gordura Visceral

Exames laboratoriais

Expressão gênica

Glicose

Creatinina

Colesterol

Uréia

Triglicérides

Ácido úrico

HDLc

AST

LDL

ALT

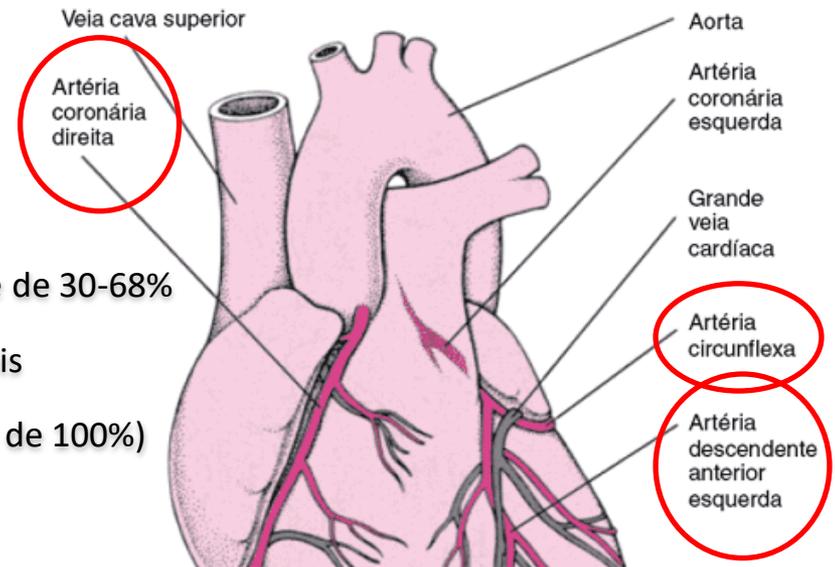
Células mononucleares
Extração de RNA
cDNA
PCR real time

Metodologia

Índice de Friesinger

Quantifica as lesões das artérias coronárias, utilizando uma variação de 0 a 15 pontos. Cada um das três artérias coronárias principais e seus ramos (descendente anterior, circunflexa e coronária direita) foi marcada separadamente, recebendo uma pontuação de zero a cinco:

- 0) nenhuma anormalidade arteriográfica
- 1) irregularidades da artéria ou estenose menor que 30%
- 2) estenose de 30-68%
- 3) múltiplas lesões ou pelo menos dois segmentos com estenose de 30-68%
- 4) estenose de 69-100%, exceto oclusão dos segmentos proximais
- 5) obstrução total (ou oclusão) de um segmento proximal (lesão de 100%)



Categorização	Sem lesão	Poucas lesões	Intermediárias lesões	Graves lesões
Pontuação	0	1-4	5-9	10-15

Metodologia

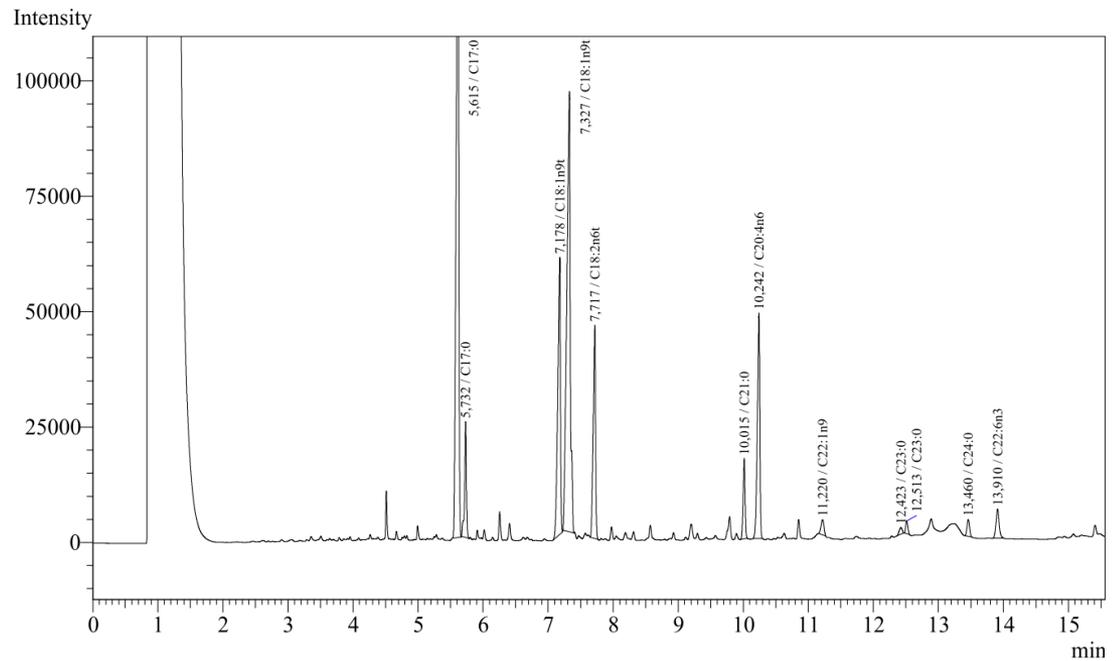
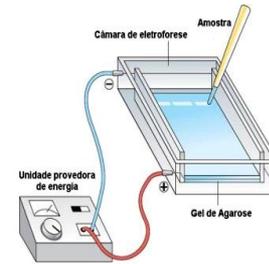


Figura 2. Exemplo do perfil sérico de ácidos graxos obtido através do CG-2010, Shimadzu (Paciente 37).

Metodologia

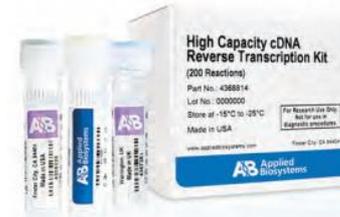
Extração do RNA TOTAL – células mononucleares



Agarose Gel Eletro. 1.0%
MOPS Buffer

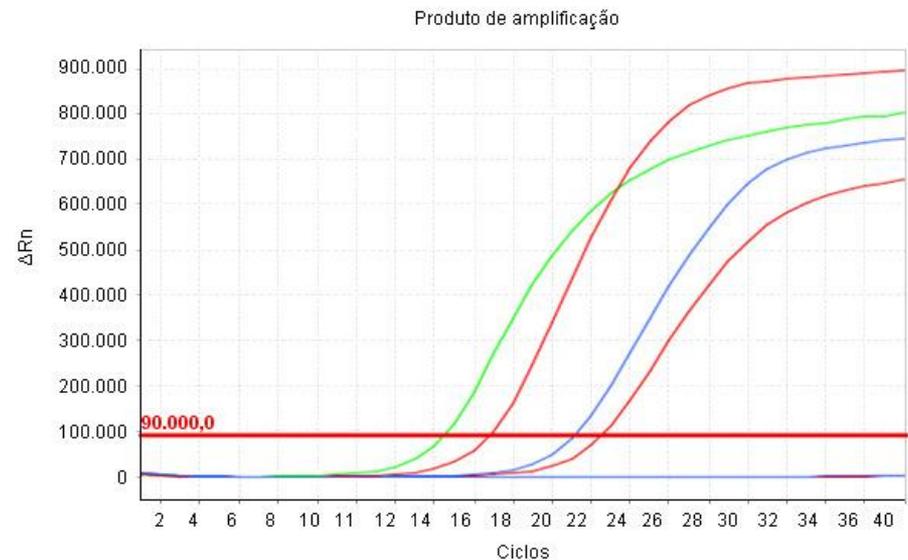
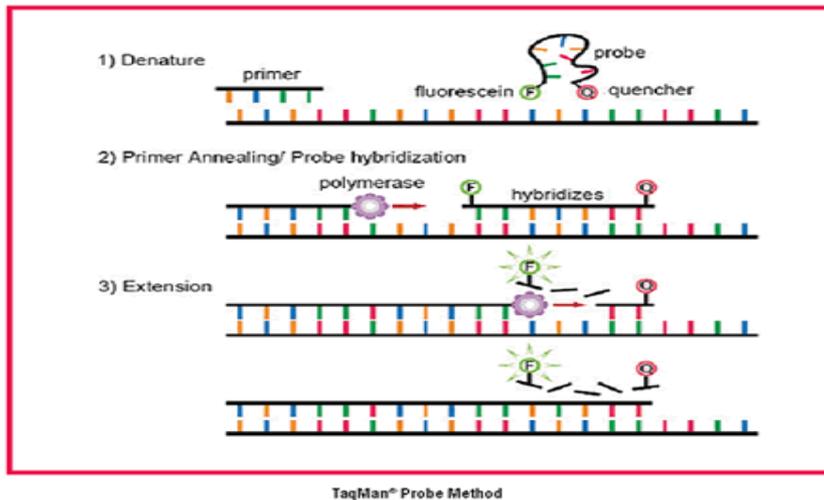


Qubit® 2.0
Fluorometer



Metodologia

Sistema TaqMan – Quantificação relativa



$$\Delta Ct = (Ct_{\text{gene alvo}} - Ct_{\text{controle endógeno}})$$

$2^{-\Delta Ct}$

ACTB

Metodologia

Consumo Alimentar e Dietético

The image shows a digital interface for a nutritional assessment system. On the left, there are two paper forms for 'CONSUMO ALIMENTAR - RECORDE DIÁRIO DE 24 HORAS'. The forms include fields for patient identification (name, sex, date of birth, address) and a grid for recording food intake over 24 hours. The grid is divided into sections for 'PERÍODO DA MANHÃ' (Morning) and 'PERÍODO DA TARDE' (Afternoon). The interface on the right is a web application with a header for 'virtual nutri Plus' and a navigation menu. A patient profile is displayed, showing a photo and personal information. Below the profile, there is a table for recording food intake data, similar to the paper forms on the left.

Os dados dietéticos da variabilidade, variabilidade intrapessoal foram ajustados pelo método de Nusser et al. 1996;

Em seguida, os resultados foram ajustados pela energia;

Foi utilizado a análise pela ANOVA para determinar a variabilidade intrapessoal e para estimar a variância intra e interpessoal foi utilizado a média quadrática;

Para controlar os fatores de confundimento inerentes a ingestão de energia total e remover as variáveis externas, os dados foram ajustados pela energia usando o método do resíduo.

Metodologia

Avaliação Antropométrica



Balança de Bioimpedância Elétrica (BIA)
tetrapolar da marca Omron HBF-514C
(Omron Healthcare, Japão)

Percentual de gordura

Percentual de gordura visceral

EUTRÓFICOS

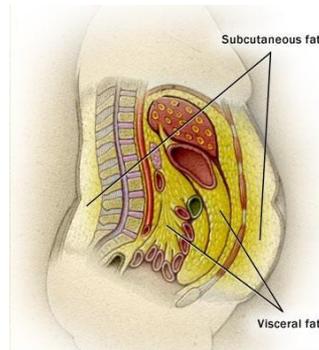
ALTO PG%

MUITO ALTO PG%

<9%

10-14%

>15%

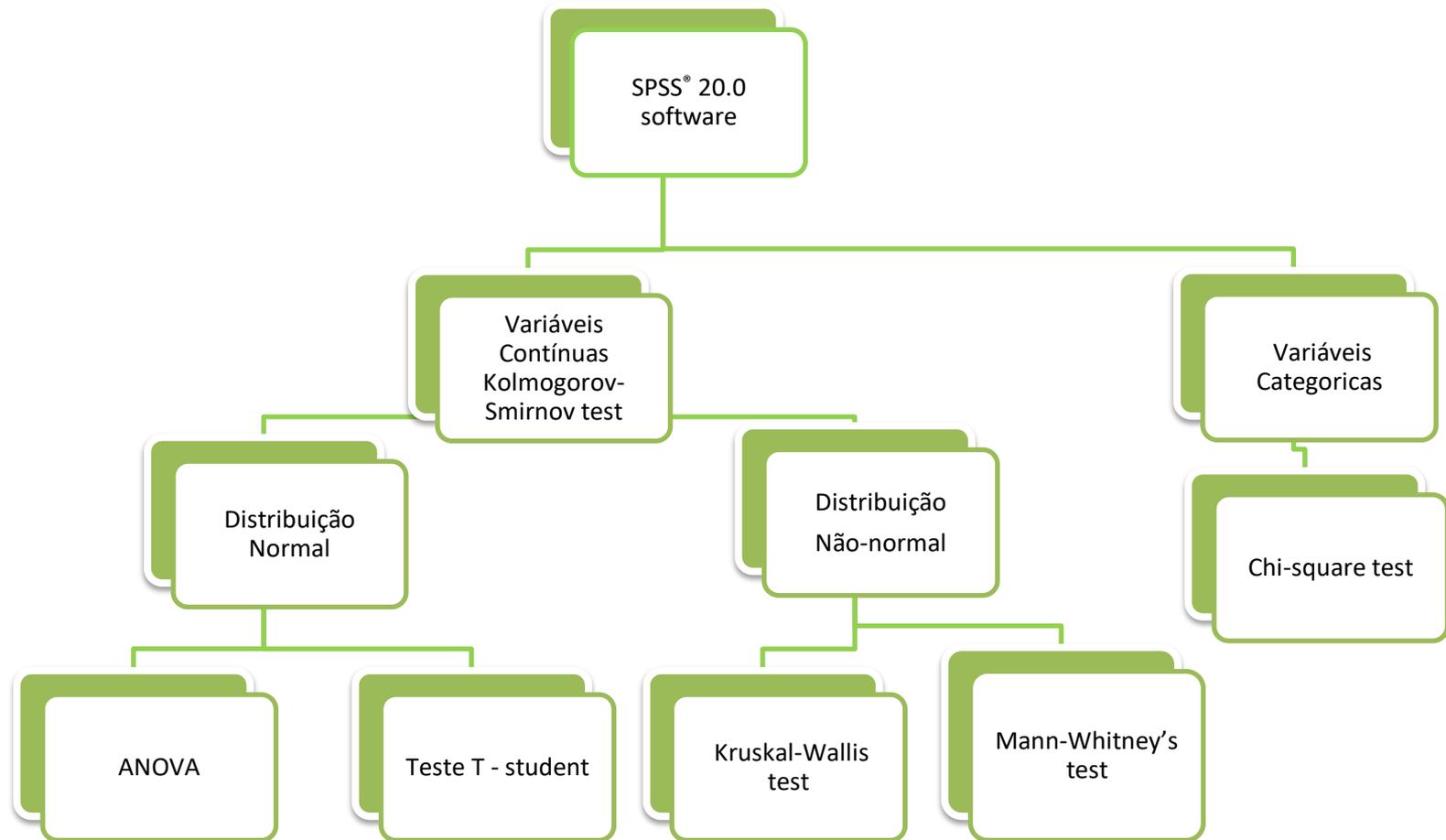


Metodologia

Avaliação Antropométrica



Metodologia



O nível de significância estatística aceitável foi $p < 0.05$.

Artigo – Resultados e Discussão

Table 1. Demographic, anthropometric and clinical data of patients classified according to the extent of coronary lesion

Variables	Total (n=59)	Without lesion (n=18)	Low lesion (n=17)	Intermediate lesion (n=17)	Major lesion (n=7)	<i>p-value</i>
Age, years	60.0 ± 9.0	55.0 ± 9.0	62.0 ± 9.0	63.0 ± 10	60.0 ± 7.0	0.080
Sex male, %	54.2	44.4	58.8	64.7	42.9	0.582
BMI, kg/m ²	26.82±5.65	27.86 ± 4.70	25.27 ± 8.52	26.92 ± 4.29	27.43 ± 1.13	0.630
Obesity, %	20.3	33.3	17.6	17.6	0	0.133
Dyslipidemia, %	91.5	77.8	100.0	94.1	100.0	0.080
Diabetes mellitus, %	30,5	33.3	23.5	35.3	28.6	0.884
Hypertension, %	79.7	77.8	76.5	82.4	85.7	0.944
Diastolic pressure, mmHg	84.0±20.0	81.0±8.0	80.0±24.0	90.0±18.0	90.0±32.0	0.355
Systolic pressure, mmHg	143.0±26.0	137.0±24.0	139.0±22.0	155.0±32.0	141.0±21.0	0.187
Alcoholism, %	18.6	27.8	17.6	5.9	28.6	0.350
Smoking, %	22.0	27.8	17.6	17.6	28.6	0.825
Physical activity, %	47.5	50.0	58.8	35.3	42.9	0.573

BMI, Body Mass Index; Data are presented as mean ± standard deviation for parametric samples and as median (interquartile range) for non-parametric samples. Categorical variables were compared by Chi-square test. Parametric analysis was performed by ANOVA way followed by test T. Non-parametric samples were performed by Kruskal-Wallis test followed by Mann Whitney test. P-values < 0.05 were considered statistically significant.

Artigo – Resultados e Discussão

Cont. Table 1. Demographic, anthropometric and clinical data of patients classified according to the extent of coronary lesion

Biochemical Analyses						
Glucose, mmol/l	93,0 mg/dL	4.75(3.89-14.26)	4.88(3.75-11.04)	5.55(3.72-21.87)	5.94(4.44-13.04)	0.443
Total cholesterol, mmol/l	178 mg/dL	4.36(3.08-6.50)	4.51(3.19-6.89)	5.05(2.95-7.85)	4.61(3.65-7.02)	0.555
HDL-cholesterol, mmol/l	35,5 mg/dL	0.95(0.57-1.45)	0.93(0.54-1.30)	0.91(0.41-1.32)	0.83(0.62-1.40)	0.669
LDL-cholesterol, mmol/l	103,3 mg/dL	2.38(1.49-4.57)	2.63(1.21-5.00)	3.26(1.21-5.74)	2.89(1.93-5.27)	0.419
Triglycerides, mmol/l	143 mg/dL	1.63(0.79-6.94)	2.25(0.86-6.96)	1.62(0.75-9.47)	1.45(0.85-3.24)	0.612
ALT, μ Kat/l	26 U/L	0.38 (0.17-1.57)	0.43 (0.25-1.04)	0.33 (0.17-2.19)	0.60 (0.25-1.45)	0.723
AST, μ Kat/l	31 U/L	0.52 (0.25-1.49)	0.43 (0.25-0.95)	0.52 (0.25-1.57)	0.60 (0.25 -1.57)	0.546
Ureia, mmol/l	35 mg/dL	5.93(3.34-9.52)	5.68(4.31-9.85)	5.93(3.37-10.35)	5.85(4.84-6.35)	0.693
Creatinine, μ mol/l	0,90 mg/dL	79.56(53.04-141.44)	79.56(53.04-123.76)	88.40(53.04-106.08)	70.72(17.68-106.08)	0.323
Uric acid, mmol/l	4,9 mg/dL	0.27(0.16-0.52)	0.28(0.11-0.43)	0.29(0.11-0.43)	0.33(0.14-0.37)	0.965

BMI, Body Mass Index; HDL-cholesterol, High density lipoprotein; LDL-cholesterol, Low density lipoprotein; AST, aspartate aminotransferase; ALT, Alanine transaminase. Data are presented as mean \pm standard deviation for parametric samples and as median (interquartile range) for non-parametric samples. Categorical variables were compared by Chi-square test. Parametric analysis was performed by ANOVA way followed by test T. Non-parametric samples were performed by Kruskal-Wallis test followed by Mann Whitney test. P-values < 0.05 were considered statistically significant.

Artigo – Resultados e Discussão

Table 2 –Serum fatty acid concentration according to the extent of coronary lesion

Fatty acids (%)	Total (n=59)	Without lesion (n=18)	Low lesion (n=17)	Intermediate lesion (n=17)	Major lesion (n=7)	<i>p-value</i>
SFA	44.12±3.85	44.70±4.18	43.79±4.46	43.93±2.74	43.93±4.40	0.904
Myristic (C14:0)	0.89±0.30	0.94±0.36	0.91±0.26	0.87±0.31	0.80±0.19	0.756
Palmitic (C16:0)	30.83±3.07	31.40±2.95	30.61±3.15	30.52±2.80	30.63±4.22	0.830
Stearic (C18:0)	12.40±1.89	12.36±2.15	12.27±2.10	12.54±1.70	12.50±1.30	0.979
MUFA	24.87±3.10	23.66±2.88	25.97±2.84a	25.77±2.96a	23.14±3.35	0.032
Palmitoleic (C16:1)	2.55±0.75	2.41±0.62	2.73±.63	2.56±0.89	2.47±0.98	0.658
Oleic (C18:1)	22.32±2.77	21.25±2.54	23.24±2.48a	23.22±2.73a	20.66±2.89	0.027
PUFA	31.01±4.78	31.65±4.46	30.24±5.37	30.30±3.99	32.94±6.01	0.529
n-6	26.77±4.29	27.26±3.66	25.96±5.13	26.46±3.54	28.26±5.53	0.628
Linoleic (C18:2)	18.06±3.84	18.64±3.29	17.23±4.35	18.10±3.30	18.47±5.39	0.742
Arachidonic (C20:4)	8.71±2.37	8.62±1.73	8.73±2.47	8.36±2.20	9.78±3.89	0.620
n-3	4.23±1.17	4.38±1.56	4.29±1.02	3.84±0.86	4.68±0.91	0.363
α-linolenic (C18:3)	0.54±0.20	0.56±0.20	0.53±0.21	0.52±0.16	0.57±0.31	0.916
EPA (C20:5)	0.46±0.18	0.53±0.24	0.42±0.15	0.43±0.16	0.52±0.11	0.223
DPA (C22:5)	0.83±0.23	0.83±0.21	0.86±0.17	0.80±0.28	0.85±0.33	0.891
DHA (C22:6)	2.39±0.93	2.46±1.26	2.48±0.82	2.10±0.67	2.74±0.73	0.410
n-6/n-3	6.77±2.12	6.80±1.93	6.61±2.96	7.14±1.63	6.16±1.29	0.765
SCD16	0.08±0.02	0.08±0.02	0.09±0.02	0.08±0.02	0.08±0.03	0.398
SCD18	1.85±0.40	1.77±0.38	1.95±0.38	1.90±0.45	1.68±0.36	0.376

EPA, Eicosapentaenoic; DHA, Docosahexaenoic; DPA, Docosapentaenoic SFA, saturated fatty acid; MUFA, monounsaturated fatty acid; PUFA, polyunsaturated fatty acid, n6, omega-6; n3, omega-3. Data are presented as mean ± standard deviation. Comparisons were performed by ANOVA way test followed by test t between pars. P-values < 0.05 were considered statistically significant. a, compared to Without lesion group.

Artigo – Resultados e Discussão

Int. J. Mol. Sci. **2013**, *14*, 18861-18880; doi:10.3390/ijms140918861

Oleic Acid Increases Synthesis and Secretion of VEGF in Rat Vascular Smooth Muscle Cells: Role of Oxidative Stress and Impairment in Obesity

Gabriella Doronzo, Michela Viretto, Cristina Barale, Isabella Russo, Luigi Mattiello, Giovanni Anfossi[†] and Mariella Trovati^{*}

Ma et al. *Lipids in Health and Disease* 2011, **10**:53
<http://www.lipidworld.com/content/10/1/53>

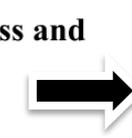
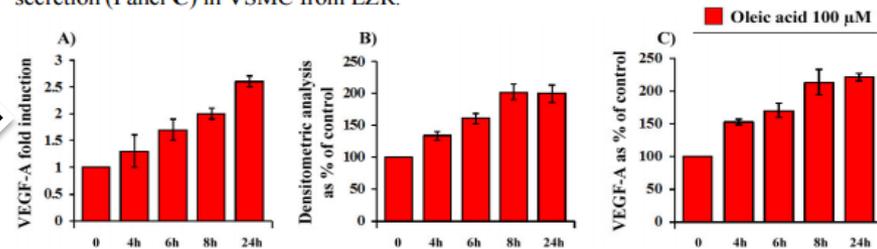


Figure 1. Time-dependent (4–24 h of incubation with 100 μ M oleic acid) influence of oleic acid on VEGF-A mRNA transcription (Panel A); protein synthesis (Panel B) and secretion (Panel C) in VSMC from LZR.



RESEARCH

Open Access

Oleic acid induces smooth muscle foam cell formation and enhances atherosclerotic lesion development via CD36

Shuangtao Ma[†], Dachun Yang[†], De Li, Bing Tang and Yongjian Yang^{*}

Camargo et al. *BMC Genomics* 2010, **11**:253
<http://www.biomedcentral.com/1471-2164/11/253>

RESEARCH ARTICLE

Open Access

Gene expression changes in mononuclear cells in patients with metabolic syndrome after acute intake of phenol-rich virgin olive oil

Antonio Camargo^{†1}, Juan Ruano^{†1}, Juan M Fernandez¹, Laurence D Parnell², Anabel Jimenez¹, Monica Santos-Gonzalez³, Carmen Marin¹, Pablo Perez-Martinez¹, Marino Uceda⁴, Jose Lopez-Miranda¹ and Francisco Perez-Jimenez^{*1}

Artigo – Resultados e Discussão

Pearson analysis showed a positive correlation between oleic acid with triglyceride levels ($r = 0.397$, $p = 0.002$), SCD16 ($r = 0.352$, $p = 0.006$) and SCD18 ($r = \mathbf{0.751}$, $p = 0.027$).

Nutrition 32 (2016) 88–94



ELSEVIER

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Applied nutritional investigation

Interaction of SNP in the CRP gene and plasma fatty acid profile in inflammatory pattern: A cross-sectional population-based study

Erica Oki M.Sc.^a, Marina M. Norde M.Sc.^a, Antônio A.F. Carioca M.Sc.^a,
Renata E. Ikeda B.Sc.^a, José M.P. Souza Ph.D.^b, Inar A. Castro Ph.D.^c,
Dirce M.L. Marchioni Ph.D.^a, Regina M. Fisberg Ph.D.^a, Marcelo M. Rogero Ph.D.^{a,*}

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Table 2

Plasma fatty acid (%) by inflammatory status

Plasma fatty acid (%)	Non-inflammatory cluster (n = 169)	Inflammatory Cluster (n = 93)	P-value*
SFA	39.44 (2.64)	39.89 (2.88)	0.209
Myristic (C14:0)	0.77 (0.56–1.06)	0.86 (0.54–1.21)	0.165
Palmitic (C16:0)	27.44 (2.55)	28.14 (2.68)	0.039
Stearic (C18:0)	11.15 (1.49)	10.81 (1.73)	0.098
MUFA	19.23 (16.37–21.72)	20.57 (17.48–23.76)	0.051
Palmitoleic (C16:1)	1.77 (1.35–2.24)	2.07 (1.05)	0.164
Oleic (C18:1)	17.38 (15.11–19.61)	18.68 (16.14–21.05)	0.058
PUFA	40.93 (4.81)	39.32 (4.85)	0.011
n-6	36.71 (4.58)	35.29 (4.63)	0.018
Linoleic (C18:2)	25.26 (3.89)	24.68 (4.20)	0.266
DGLA (C20:3)	2.45 (0.56)	2.43 (0.55)	0.761
Arachidonic (C20:4)	9.01 (2.19)	8.19 (1.75)	0.002
n-3	3.96 (3.61–4.64)	3.76 (3.45–4.43)	0.173
α -linolenic (C18:3)	0.61 (0.49–0.75)	0.65 (0.49–0.83)	0.247
Eicosapentaenoic (C20:5)	0.49 (0.36–0.63)	0.47 (0.35–0.60)	0.382
Docosapentaenoic (C22:5)	0.80 (0.65–0.99)	0.78 (0.70–0.87)	0.253
Docosahexaenoic (C22:6)	2.06 (1.70–2.46)	1.89 (1.54–2.42)	0.081
n3 HUFA	3.39 (2.92–4.01)	3.15 (2.68–3.77)	0.080
SCD-16 (C16:1/16:0)	0.06 (7.85–10.48)	0.07 (0.05–0.09)	0.299
SCD-18 (C18:1/C18:0)	1.55 (1.29–1.89)	1.74 (1.37–2.16)	0.045
D6D (C20:3/18:2 n-6)	0.07 (0.06–0.08)	0.07 (0.06–0.08)	0.343
D5D (C20:4/20:3)	3.76 (2.96–4.49)	3.32 (2.89–4.12)	0.047
n-6/n-3	9.22 (2.27)	9.20 (2.05)	0.953

DGLA, dihomo- γ -linoleic; HUFA, highly unsaturated fatty acid; MUFA, mono-unsaturated fatty acid; n3, omega-3; n6, omega-6; PUFA, polyunsaturated fatty acid; SCD, delta-9-desaturase; SFA, saturated fatty acid D6D, delta-6-desaturase; D5D, delta-5-desaturase

Values expressed as mean (standard deviation) or median (interquartile range)
* P-value for Student's *t* test. Values in bold are statistically significant ($P < 0.05$).

Artigo – Resultados e Discussão

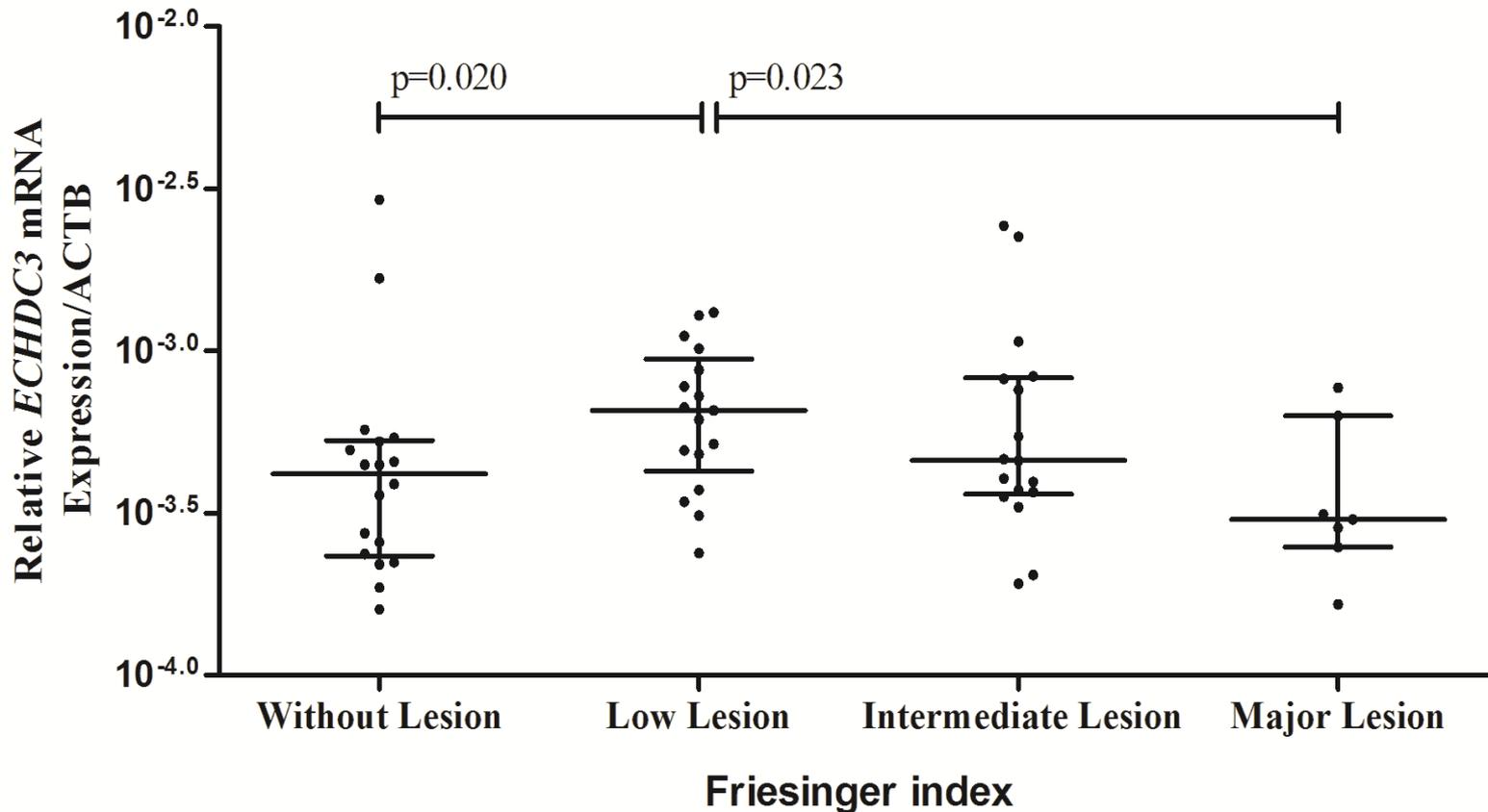


Figure 3. *ECHDC3* mRNA expression relative to *ACTB* expression according to the extent of coronary lesion. Data are presented as median and interquartile range. Relative expression of *ECHDC3* was calculated using the 2-deltaCT method. Statistically analysis was performed by Kruskal-Wallis test followed by Mann Whitney test.

Artigo – Resultados e Discussão

ECHDC3 gene expression positively correlated with SFA ($r = 0.259$, $p = 0.048$).

ORIGINAL RESEARCH

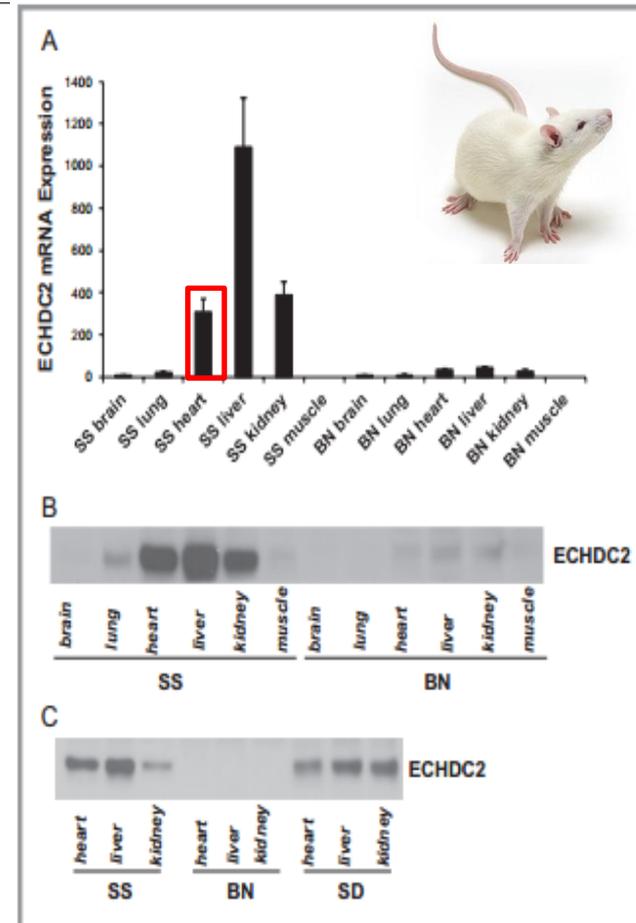


Enoyl Coenzyme A Hydratase Domain-Containing 2, a Potential Novel Regulator of Myocardial Ischemia Injury

Jianhai Du, PhD; Zhixin Li, PhD; Quan-Zhen Li, PhD; Tongju Guan, MSc; Qiuhui Yang, PhD; Hao Xu, PhD; Kirkwood A. Pritchard, Jr, PhD; Amadou K. S. Camara, PhD; Yang Shi, PhD

```

ECHDC2_RAT      MLRALPRALRLRRPWMSPGARGCAAHASTRTPEIQVQALTGPNQGITTEILMNRPHARNAL 60
ECHDC2_MOUSE    MLRVLPRALRLPCSWRFSGARDASHATRTPEIQVQALTGPNQGITTEILMNRPNARNAL 60
ECHDC2_HUMAN    MLRVL----CLLRPWRPLRARGCASDGAAGGSEIQVRALAGPDQGITTEILMNRPSARNAL 56
                ***.*      *.*      **.*:..:..:  .***:*.***:***:*****
ECHDC2_RAT      GNVFVSELLEALALQREDQQRVLLFRSAVKGVFCAGADLKERERMSAAEVGTFVQRLRG 120
ECHDC2_MOUSE    GNVFVSELLEALALQREDQQRVLLFRSAVKGVFCAGADLKEREQMSDVEVGTFFVQRLRG 120
ECHDC2_HUMAN    GNVFVSELLETLAQLREDQRVLLFRSGVKGVFCAGADLKEREQMSEAEVGVFVQRLRG 116
                *****:*****:*****:*****:*****:*****:*.**  .***.*****
ECHDC2_RAT      LMSEIAAFPAPTIAAMDGFALGGGLELALACDLRIAASSAVMGLIETTRGLLPAGGTTQR 180
ECHDC2_MOUSE    LMSEIAAFPVPTIAAMDGFALGGGLELALACDLRIAASSAVMGLIETTRGLLPAGGTTQR 180
ECHDC2_HUMAN    LMNDIAAFPAPTIAAMDGFALGGGLELALACDLRVAASSAVMGLIETTRGLLPAGGTTQR 176
                **.:***.*****:*****:*****:*****:*****:*****
ECHDC2_RAT      LPRCLGVALAKELIFTGRRLNGVQAHELGLVNHAVAQNEEGDAAHYHRALALAQEILPQAP 240
ECHDC2_MOUSE    LPRCLGVALAKELIFTGRRLNGAQARELGLVNHAVAQNEEGNAAYHRALALAQEILPQAP 240
ECHDC2_HUMAN    LPRCLGVALAKELIFTGRRLSGTEAHVGLVNHAVAQNEEGDAAHQARALALAQEILPQAP 236
                *****:*****:*.*:  *****:*****:***:*****
ECHDC2_RAT      IAVRLGKVAIDRGMVDIASGMAIEHMCYAQNIPTQDRLEGMAAFREKRPPKFFVGK 296
ECHDC2_MOUSE    IAVRLGKVAIDRGMVDIASGMAIEQMCYAQNIPTQDRLEGMAAFREKRAPKFFVGK 296
ECHDC2_HUMAN    IAVRLGKVAIDRGTQVDIASGMAIEGMCYAQNIPTDRLEGMAAFREKRTPKFFVGK 292
                *****  *****  *****:*****:*****:*****
    
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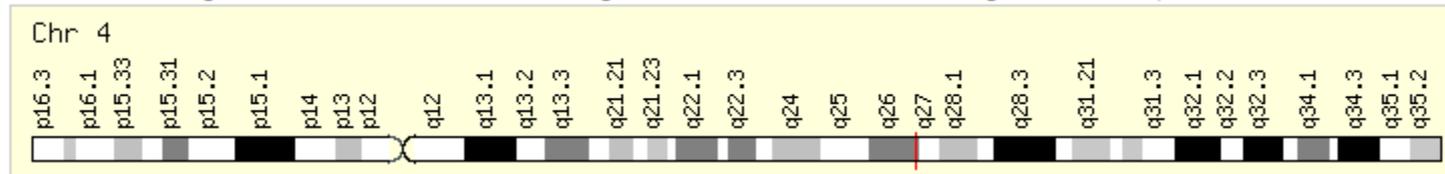


Doenças Cardiovasculares e Nutrigenética



Doenças Cardiovasculares e Nutrigenética

Gene
FABP2
4q28-q31



Polimorfismo na Ala54Thr (rs1799883)

Parâmetros antropométricos, perfil lipídico, resistência insulínica



PLoS One. 2016; 11(9): e0163421.

Published online 2016 Sep 29. doi: [10.1371/journal.pone.0163421](https://doi.org/10.1371/journal.pone.0163421)

Gene Polymorphisms of *FABP2*, *ADIPOQ* and *ANP* and Risk of Hypertriglyceridemia and Metabolic Syndrome in Afro-Caribbeans

[Laurent Larifla](#),^{1,2,*} [Christine Rambhojan](#),¹ [Marie-Odile Joannes](#),¹ [Suliya Maimaitiming-Madani](#),³ [Jean-Paul Donnet](#),¹ [Thérèse Marianne-Pépin](#),¹ [Roger Chout](#),¹ [Ronan Roussel](#),⁴ and [Lydia Foucan](#)¹

Larifla L, Rambhojan C, Joannes M-O, et al. Gene Polymorphisms of *FABP2*, *ADIPOQ* and *ANP* and Risk of Hypertriglyceridemia and Metabolic Syndrome in Afro-Caribbeans. Palmer ND, ed. *PLoS ONE*. 2016;11(9):e0163421. doi:10.1371/journal.pone.0163421.

Doenças Cardiovasculares e Nutrigenética



PLoS One. 2016; 11(9): e0163421.

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462 indivíduos afro-caribenhos

sem complicações cardiovasculares ou utilização de medicamentos

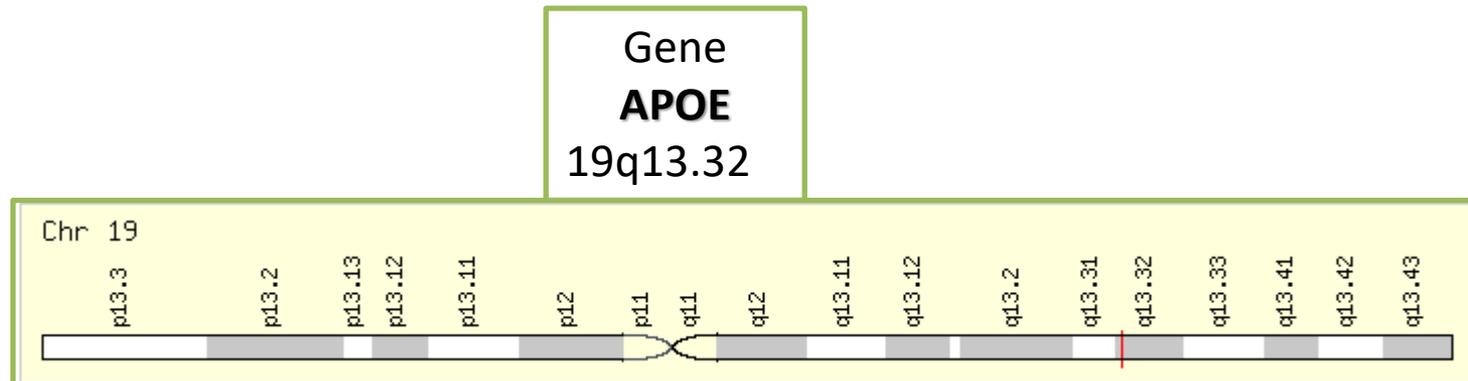


rs1799883- *FABP2* , rs1501299- *ADIPOQ* and rs5065- *ANP*
três genotipagem



Associação de variantes de genes *FABP2* , *ANP* e *ADIPOQ* com MetS
Efeito potencial sobre o metabolismo lipídico.

Doenças Cardiovasculares e Nutrigenética



Uma glicoproteína do soro que se encontra **associada aos quilomícrons circulantes, aos quilomícrons remanescentes e as VLDL**

A **contribuição** da APOE para a variabilidade nas **concentrações do colesterol** é importante.

rs429358 e rs7412 – mais estudados

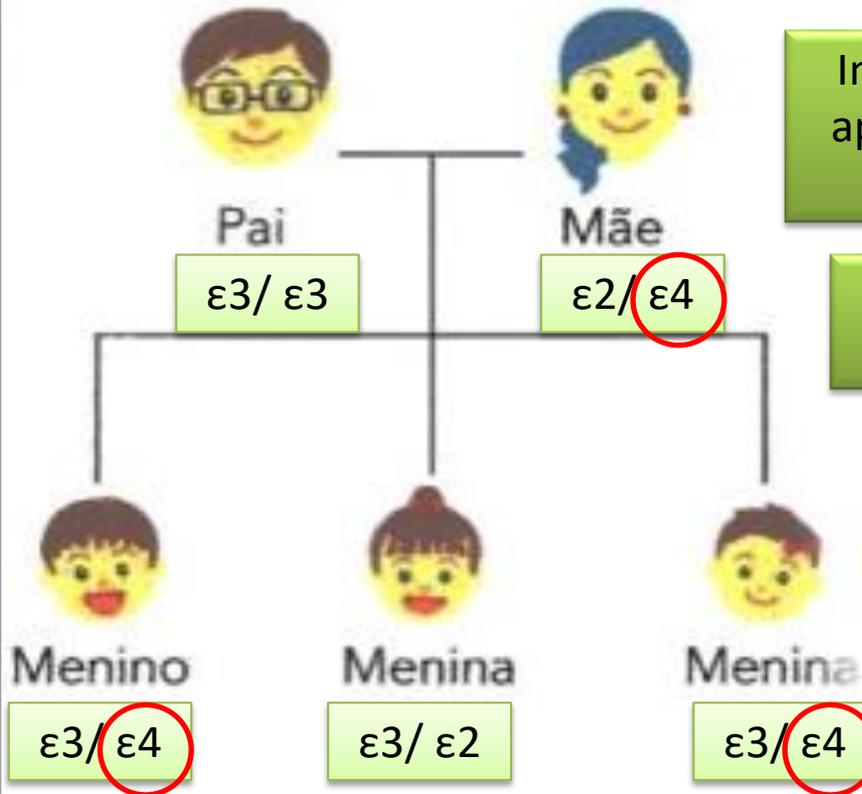
ε2 (10%) ε3 (60%) ε4 (30%)

-Minihane, AM; Jofre-Monseny, L.; Olano-Martin, E.; Rimbach, G. ApoE genotype, cardiovascular risk and responsiveness to dietary fat manipulation. Proceedings of the Nutrition Society. 2007; 66:183-97.

-Agrawal, S.; Mastana, S. Genetics of coronary heart disease with reference to APOAI-CII-AIV gene region. World Journal of Cardiology. 2014; 6(8): 755-763.

Doenças Cardiovasculares e Nutrigenética

Gene
APOE
19q13.32



Indivíduos que carregam o alelo ϵ_4 apresentam 40 a 50% de risco para DCV do que o ϵ_3/ϵ_3

Apresentam maiores quantidades de TG e LDL-c

Maior redução nas concentrações de TG ao consumirem altas doses de DHA (3,7g/dia)

-Minihane, AM; Jofre-Monseny, L.; Olano-Martin, E.; Rimbach, G. ApoE genotype, cardiovascular risk and responsiveness to dietary fat manipulation. Proceedings of the Nutrition Society. 2007; 66:183-97.

-Agrawl, S.; Mastana, S. Genetics of coronary heart disease with reference to APOAI-CII-AIV gene region. World Journal of Cardiology. 2014; 6(8): 755-763.

Doenças Cardiovasculares e Nutrigenética

Gene
APOE
19q13.32

Am J Clin Nutr. 2012 Dec;96(6):1447-53. doi: 10.3945/ajcn.112.043240. Epub 2012 Nov 7.

APOE genotype influences triglyceride and C-reactive protein responses to altered dietary fat intake in UK adults.

Carvalho-Wells AL¹, Jackson KG, Lockyer S, Lovegrove JA, Minihane AM.

Indivíduos adultos do Reino Unido
carreadores do alelo $\epsilon 4$
apresentam maior redução nas concentrações de TG
ao consumirem altas doses de **DHA (3,7g/dia)**
que o $\epsilon 3/ \epsilon 3$

Os autores sugerem que isso ocorra em razão da **maior atuação da isoforma APOE4 na remoção de remanescentes de VLDL.**

As **isoformas $\epsilon 2$ e $\epsilon 3$** - metabolismo **HDL**

-Carvalho-Wells AL, Jackson KG, Lockyer S, Lovegrove JA, Minihane AM. Am J Clin Nutr. 2012 Dec;96(6):1447-53. doi: 10.3945/ajcn.112.043240.

Doenças Cardiovasculares e Nutrigenética

Gene
APOE
19q13.32

Am J Clin Nutr. 2012 Dec;96(6):1447-53. doi: 10.3945/ajcn.112.043240. Epub 2012 Nov 7.

APOE genotype influences triglyceride and C-reactive protein responses to altered dietary fat intake in UK adults.

Carvalho-Wells AL¹, Jackson KG, Lockyer S, Lovegrove JA, Minihane AM.

indivíduos **ε4/ ε4**

Aumento das concentrações séricas **de PCR**
consumo de gorduras saturadas

indivíduos **ε4/ ε4**

alta sensibilidade no consumo de lipídios.

Doenças Cardiovasculares e Nutrigenética

Gene
APOE
19q13.32

[Biochem Genet.](#) 2010 Apr;48(3-4):342-55. doi: 10.1007/s10528-010-9331-6. Epub 2010 Jan 21.

The influence of nutrigenetics on the lipid profile: interaction between genes and dietary habits.

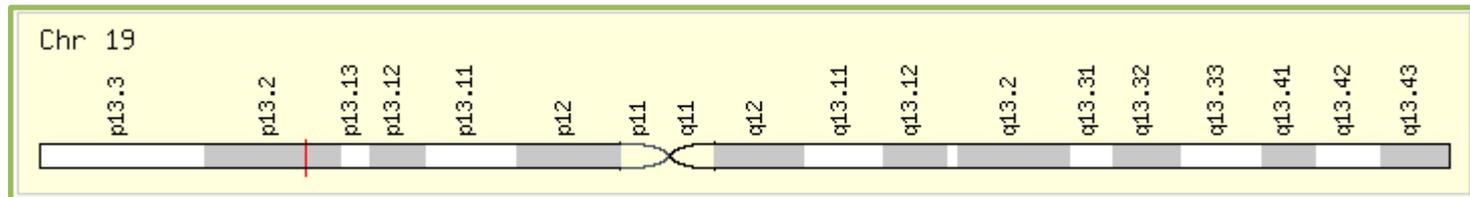
[de Andrade FM](#)¹, [Bulhões AC](#), [Maluf SW](#), [Schuch JB](#), [Voigt F](#), [Lucatelli JF](#), [Barros AC](#), [Hutz MH](#).

Carreadores do alelo $\epsilon 4$
São mais responsivos a mudanças no consumo de lipídios

Carreadores do alelo $\epsilon 2$
São mais responsivos a diminuição de carboidrato

Doenças Cardiovasculares e Nutrigenética

Gene
LDLr
19p13.2



Codifica a proteína LDLr – receptor de LDL

A expressão irá aumentar → Concentração intracelular de colesterol diminui



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[J Thromb Haemost.](#) Author manuscript; available in PMC 2009 Sep 1.

LDL Receptor Polymorphisms and the Risk of Coronary Heart Disease: the Atherosclerosis Risk in Communities Study

[N. Franceschini](#),¹ [H. Muallem](#),² [K.M. Rose](#),¹ [E. Boerwinkle](#),³ and [N. Maeda](#)²

-Franceschini N, Muallem H, Rose KM, Boerwinkle E, Maeda N. LDL Receptor Polymorphisms and the Risk of Coronary Heart Disease: the Atherosclerosis Risk in Communities Study. *Journal of thrombosis and haemostasis : JTH.* 2009;7(3):496-498. doi:10.1111/j.1538-7836.2008.03262.x.

Doenças Cardiovasculares e Nutrigenética

Gene
LDLr
19p13.2

SNPs avaliados – rs 1433099 (C>T) / rs 2738466 (A>G)

Frequência dos alelos T e G

27% (brancos), 54% (afro-americanos) – T

26% (brancos), 17% (afro-americanos) – G

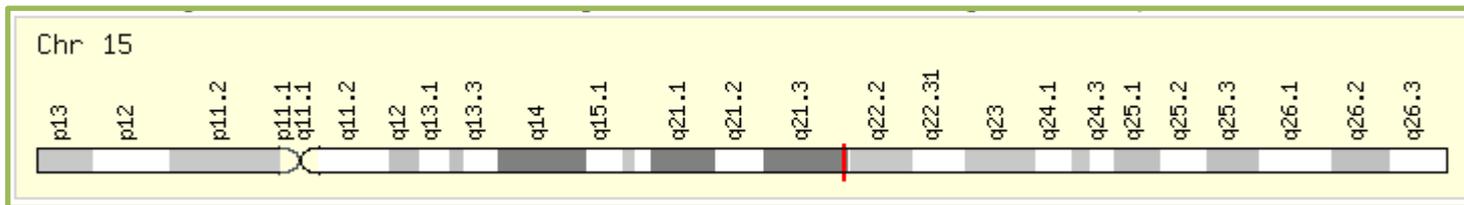
Afro-americanos – T (rs 1433099) – risco 23% DAC

TT (rs 1433099) – risco 47% DAC

SNP rs 2738466 – não foi associado a DAC

Doenças Cardiovasculares e Nutrigenética

Gene
LIPC
15q21-q23



É uma enzima sintetizada e secretada pelo fígado – metabolismo lipídico

A LIPC transforma a LDL em uma forma mais densa, menor → + aterogênica

Estudos contraditórios quanto ao papel aterogênico

- Nettleton JA, Steffen LM, Ballantyne CM, Boerwinkle E, Folsom AR. Associations between HDL-cholesterol and polymorphisms in hepatic lipase and lipoprotein lipase genes are modified by dietary fat intake in African American and White Adults . *Atherosclerosis*. 2007;194(2):e131-e140. doi:10.1016/j.atherosclerosis.2006.11.025.

Doenças Cardiovasculares e Nutrigenética

Gene
LIPC
15q21-q23



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Associations between HDL-cholesterol and polymorphisms in hepatic lipase and lipoprotein lipase genes are modified by dietary fat intake in African American and White Adults ^{1,2,3}

[Jennifer A. Nettleton](#), [Lyn M. Steffen](#), [Christie M. Ballantyne](#), [Eric Boerwinkle](#), and [Aaron R. Folsom](#)

- Nettleton JA, Steffen LM, Ballantyne CM, Boerwinkle E, Folsom AR. Associations between HDL-cholesterol and polymorphisms in hepatic lipase and lipoprotein lipase genes are modified by dietary fat intake in African American and White Adults . *Atherosclerosis*. 2007;194(2):e131-e140. doi:10.1016/j.atherosclerosis.2006.11.025.

Doenças Cardiovasculares e Nutrigenética

Gene
LIPC
15q21-q23



SNP -514 C>T (rs 1800588) – maiores concentrações HDL-c - TT

Foi avaliado no estudo a relação entre o SNP e o consumo alimentar

Indivíduos caucasianos – não houve interferência SNP sobre o perfil lipídico

Consumo de lipídios elevado:

HDL- c semelhante - Mulheres CC/Mulheres TT

HDL-c menor - Mulheres CT

- Nettleton JA, Steffen LM, Ballantyne CM, Boerwinkle E, Folsom AR. Associations between HDL-cholesterol and polymorphisms in hepatic lipase and lipoprotein lipase genes are modified by dietary fat intake in African American and White Adults . *Atherosclerosis*. 2007;194(2):e131-e140. doi:10.1016/j.atherosclerosis.2006.11.025.

Doenças Cardiovasculares e Nutrigenética

Gene	NCBI dbSNP	risco	resultado	análise
FABP2 (M)	rs1799883	A	GA	Alerta
FABP2 (F)	rs1799883	G	GA	Alerta
ApoE (SNPs)	rs429358/ rs7412	-	TT /CC	
ApoE genótipo		E4	E3/E3	Alerta
LDLR (C>T)	rs688	T	CT	Alerta
LDLR (G>T)	rs 6511720	T	GG	Ok
LDLR (C>T)	rs 5925	C	CT	Alerta
LIPC (A>G)	rs 2070895	A	GG	Ok
LIPC (C>T)	rs 1800588	C	CC	Cuidado

23 genes estudados

Doenças Cardiovasculares e Nutrigenética

- Dentre os SNPs avaliados a paciente **apresenta 6SNPs** que aumentam o risco para dislipidemia.
- Neste caso está presente **alto risco para hipercolesterolemia familiar (LDLR rs 688 e rs 5925)**.
- **Baixas concentrações de HDL** (LIPC rs 1800588)
- A paciente pode não apresentar melhora da HDL com suplementação de ácidos graxos (APOA1).

- **Sugestões:** Controle rigoroso dos lipídios, visto que vários genes podem favorecer o aumento do colesterol total, aumento de HDL e redução de LDL-c.
- Consumir nozes e castanha de forma moderada.
- Monitorar exames.

Doenças Cardiovasculares e Nutrigenética

“Os avanços em **genômica nutricional** têm favorecido a elaboração de **recomendações nutricionais mais específicas**, visando a redução do risco e o tratamento das DCVs.

Os resultados de **grandes estudos não podem ser extrapolados**, uma vez que esta relação depende de **fatores ambientais, como a própria alimentação, das características de cada população.**

Muitos esforços ainda são necessários para que o **cuidado individual personalizado seja alcançado**, uma vez que a gama de **genes e SNPs envolvidos nas DCNT é extensa.**”

Obrigada!



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