NUTRITIONAL GENOMICS; NUTRIGENETICS & NUTRIGENOMICS

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Outline:

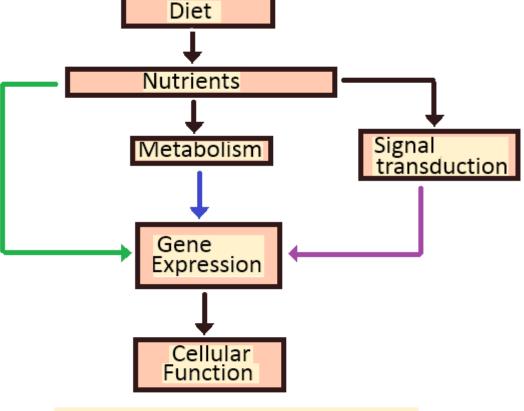
- Our idea about the topic
 - Categorizing Foods
 - Mechanisms of interactions
 - Definition of Nutrigenomics and Nutrigenetics
 - Future prospective
- Conclusion

Nutritional Genomics: Nutrigenetics & Nutrigenomics



Nutrient-gene interactions:

- Directly effects
- Indirectly effects



Disease progression or prevention

Bioactive Food Components

- Vitamins
 - Vitamin A
 - Vitamin D
 - Vitamin E
 - Vitamin C
 - Biotin
- Minerals
 - Calcium
 - Iron
 - Zinc
 - Selenium

Macronutients: Other food

- Fats
 - Fatty acids
 - Cholestrol
- Carbohydrates
 - Glucose
- Proteins
 - Amino acids

components:

- Flavonoids
- Polyphenols
- Xenobiotics

Minerals & Nutrigenomics

- Minerals
 - Iron
 - Calcium
 - Zinc
 - Iodine
 - Selenium
 - Magnesium
 - Potassium

 Participating in proteins and enzyme structures
 Cofactor functioning
 Interacting with transcription factors -> altering gene expression

As a signal in a cellular pathway

Iron

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Nutritional sources

Related Disease

Participating in biomolecules



Iron interacting with genome

IRE–IRP regulatory system: IRP1, IRP2

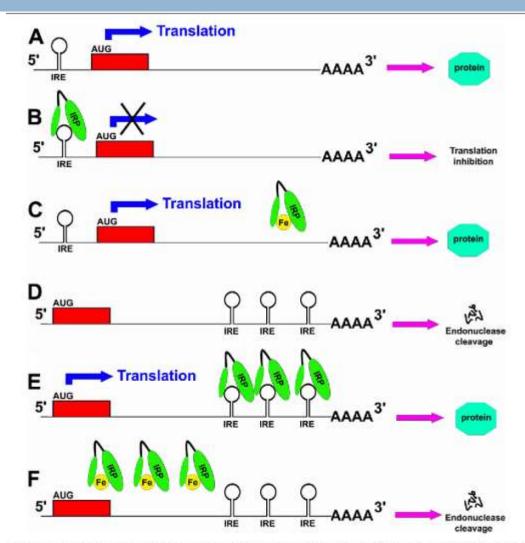
BMP/SMAD pathway signaling :
 SMAD1/5/8
 STAT3

IRE–IRP regulatory system ; An example of direct effect on gene expression

Iron Response
 Proteins:

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- Transcription factors
- controllers of vertebrate iron metabolism
- major iron homeostasis genes



station modulation by the IDD-IDE cionalise nathway a The translation of transcripts containing an IDE

Regulation of Iron Metabolism by Hepcidin An example of indirect effect through signal transduction

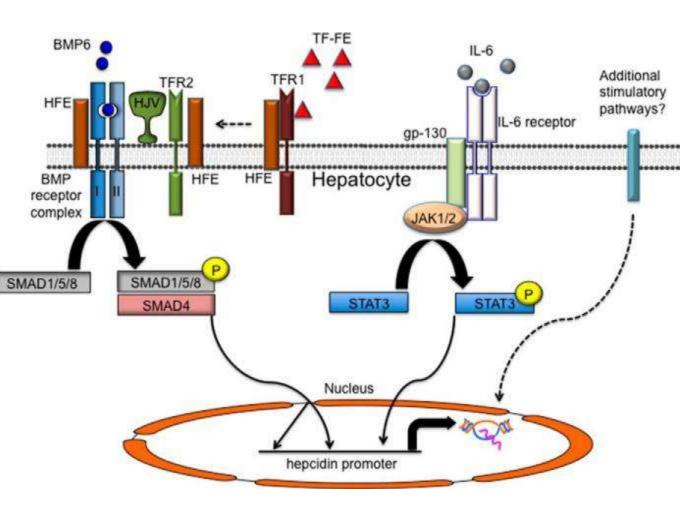
TF-FE & TFR1

BMP/SMAD pathway:

SMAD1/5/8

STAT3

regulating
 body iron
 homeostasis



Zinc

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Nutritional Related sources function

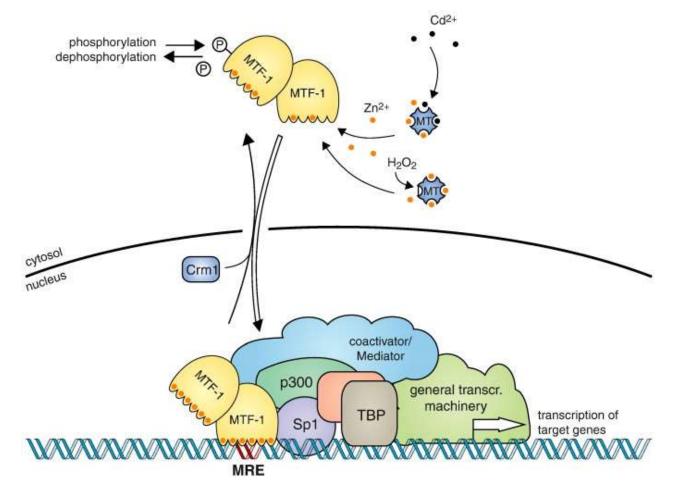
Related function in our body

cofactor function



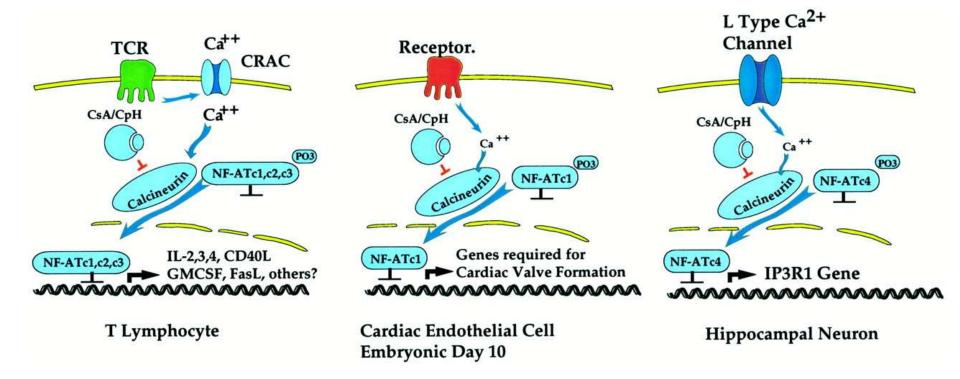
Zinc & gene expression

metal-responsive transcription factor 1 (MTF1)



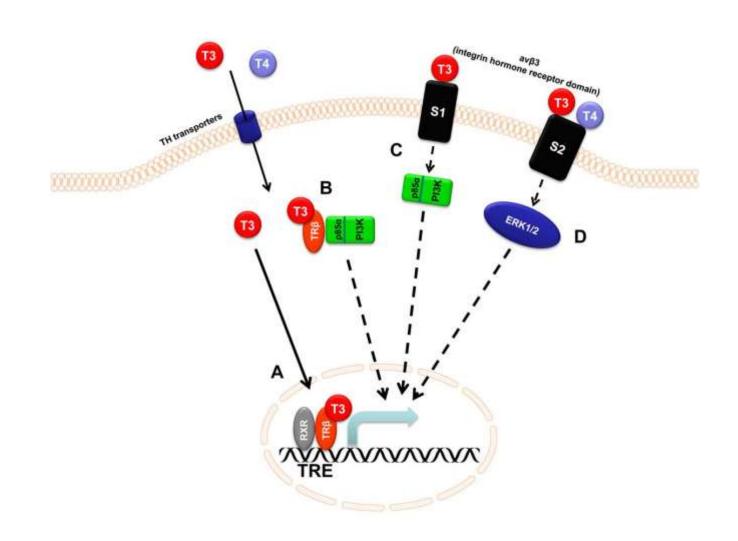
Calcium

- Transcription factors
 - calcineurin
 - NF-AT



lodine

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End of Part 1

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Any Question Please?!

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First Section Part 2:

Fatty Acids category Fat soluble vitamins category



Omega-3

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A polyunsaturated fatty acids (PUFA)

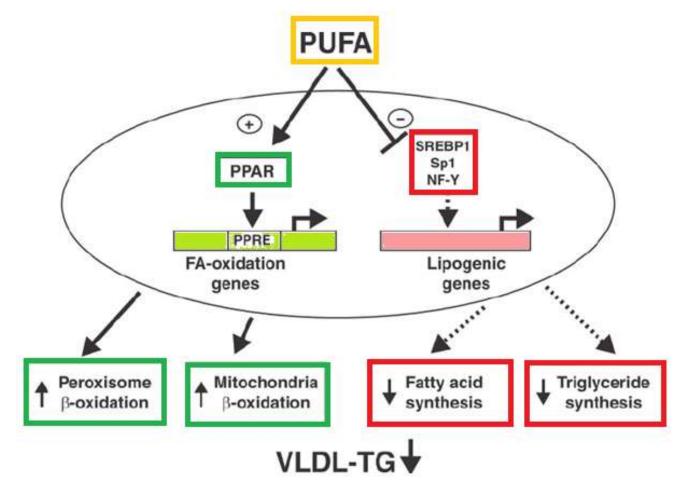
- Neuroprotective (Alzheimer's disease)
- cardiovascular heart disease
- immune function
- bone health
- muscle tonus
- Cancer
- general quality of life in aging



PUFA and modifying gene expression

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PPARs, SREBPs, LXR, HNF4, ChREBP



Carotenoids

Carotenoids (most notably beta-carotene)

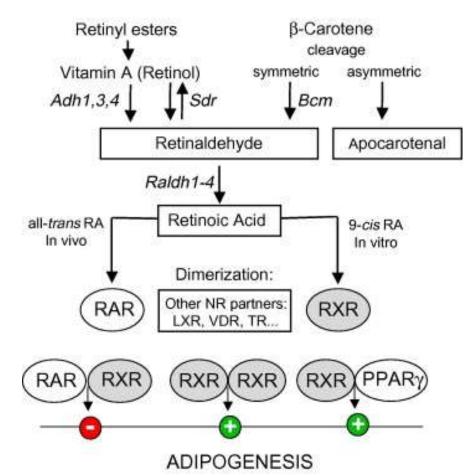
no role in the formation of vitamin A antioxidants and anti-inflammatory agents

- provitamin A , Vitamin A group (retinol, retinal, retinoic acid)
- promoting good vision (the retina of the eye), early atherosclerosis, cardiovascular disease ,skin aging and cancer development, immune system

Vitamin A and Gene Expression

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- a ligand for nuclear receptors (RAR, RXR)
- Esp. in the retinoic acid form



Vitamin D

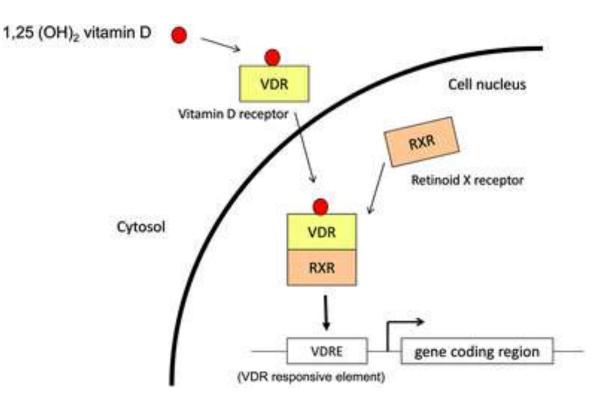
Regulating genes in:

- promoting intestinal calcium and phosphate absorption
- bone remodeling
- neuroprotective actions as Serotonin Production
- controlling cell growth and differentiation in a variety of tissues



1,25D/VDR signaling

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- 1,25-Dihydroxyvitamin D₃ (1,25D)
 - The endocrine metabolite of vitamin D
 - Vitamin D receptor (VDR)



Vitamin E

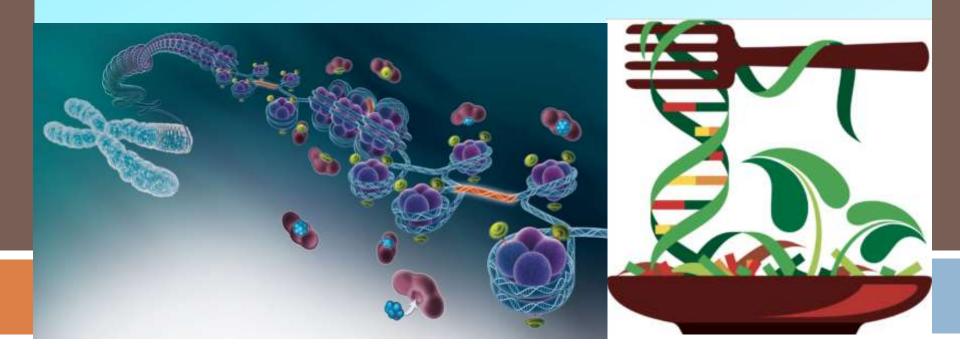
- vitamin E family (α, β, γ, δ) tocopherols and the corresponding tocotrienols
 Inflammatory/Immune Response
- pregnane X receptor (PXR), a nuclear receptor regulating a variety of drug metabolizing enzymes

End of Part 2

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Any Question Please?!

First Section Part 3: Affecting Epigenetic patterns as a way of nutrient gene interaction



Polyphenols

Flavonoids

- Resveratrol (3,5,4'-trihydroxy-trans-stilbene)
- Phytoalexins
- Tea catechins/epicatechins
 - epicatechin (EC),
 - epicatechin-3-gallate (ECG),
 - epigallocatechin (EGC),
 - epigallocatechin-3-gallate (EGC)

Genistein

- Phenolic Acids
- Lignans
- Stilbenes



Flavonoids & interaction with genome

acting via:

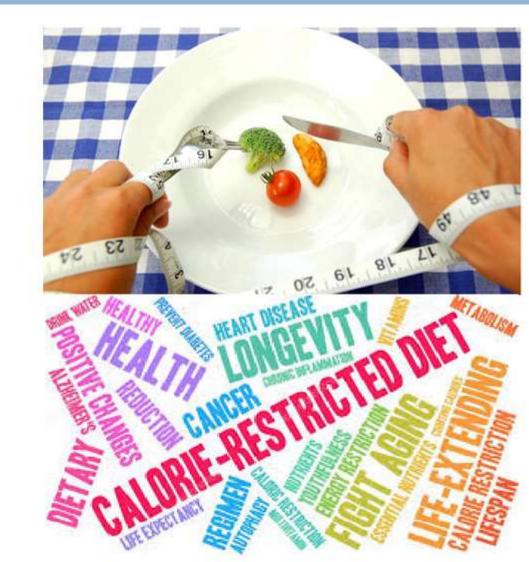
- Alter concentrations of reactive oxygen species
- The klotho gene, transcription factors
- Effect on intercellular signaling molecules including nitrous oxide and pro-inflammatory cytokines

Epigenetic mechanisms

Studied on:

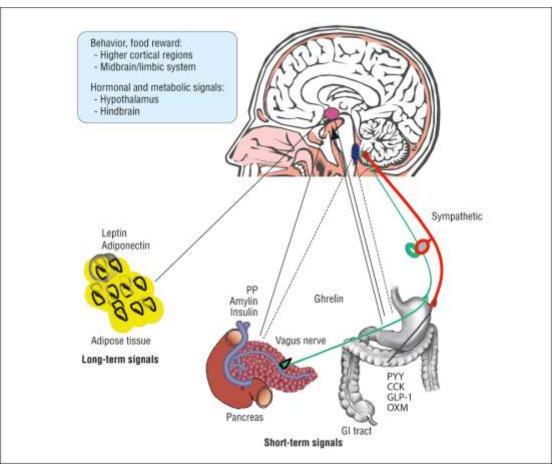
- Aging process
- Cancer
- Cardiovascular disease

Epigenetic Changes and Diet



Body Response to Nutritional State of Diet

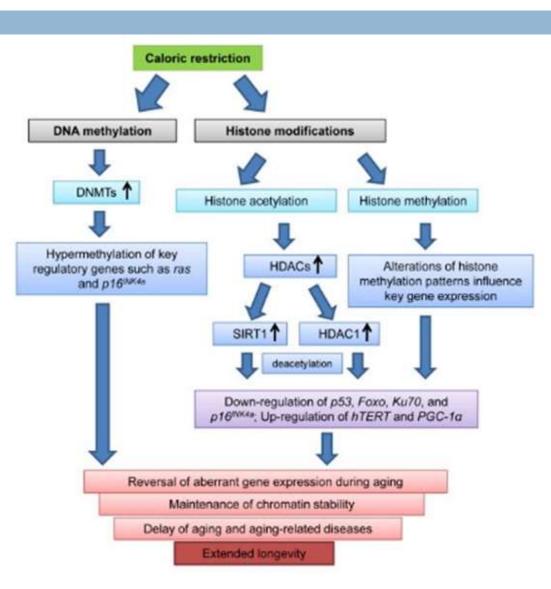
- □ Nutritional changes \rightarrow a complex signaling
 - Obesity \rightarrow resistance to the anorexigenic signals

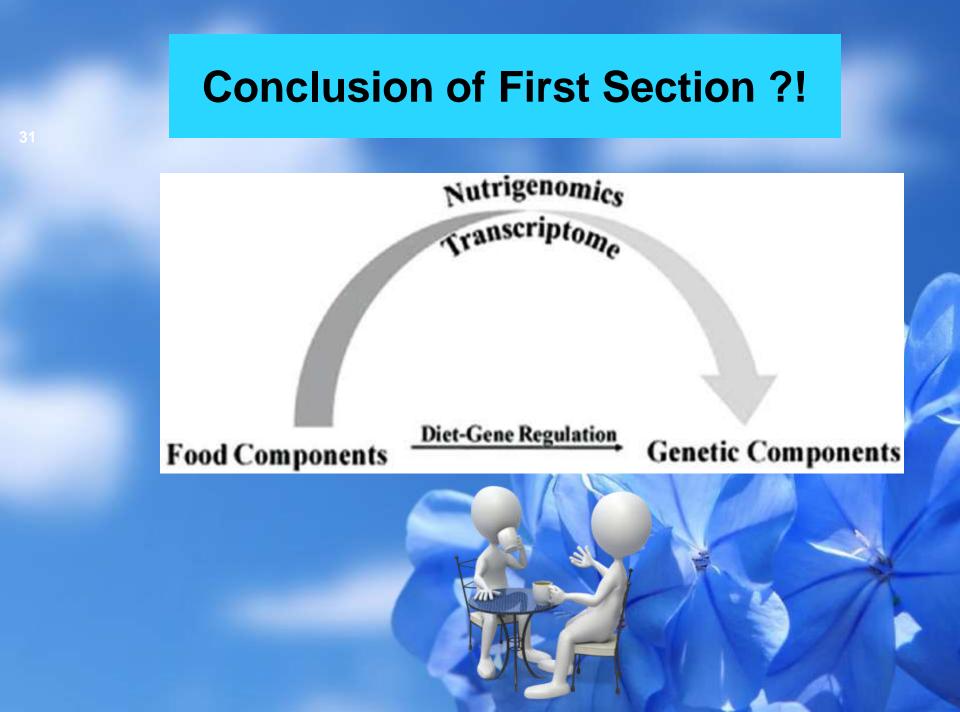


Calorie Restriction in Humans

Caloric restriction → Altering epigenetic processes via:

- DNA methylation
- Histone
 modification



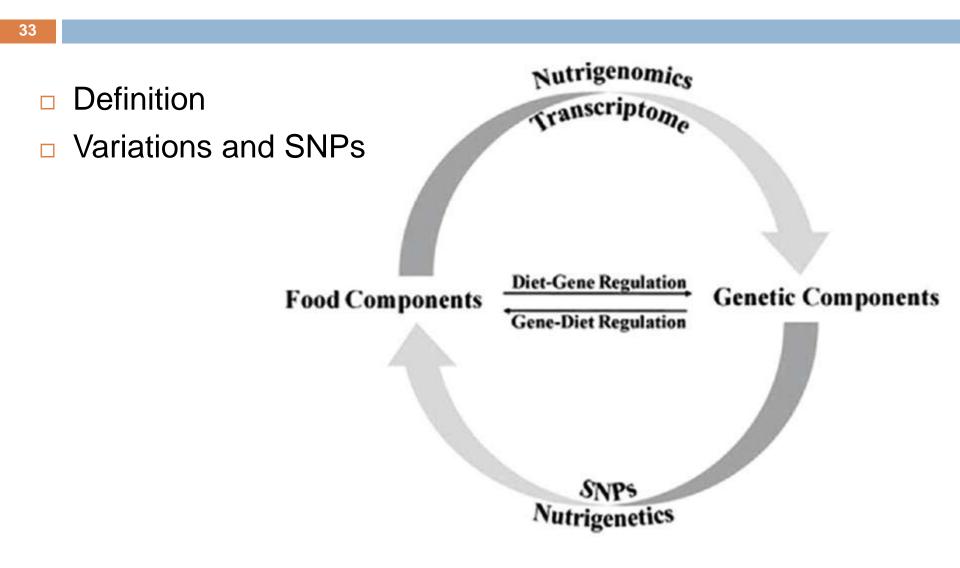


Section 2: Nutrigenetics

TERRET

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Nutrigenetics

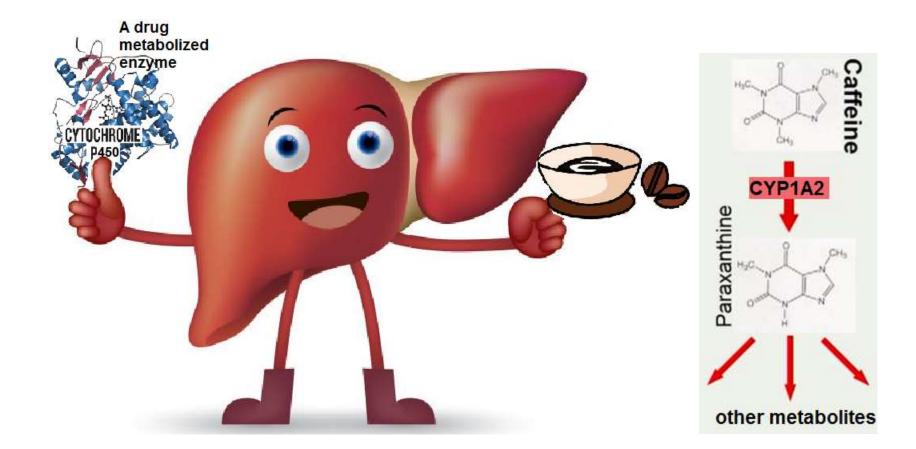


Coffee Story

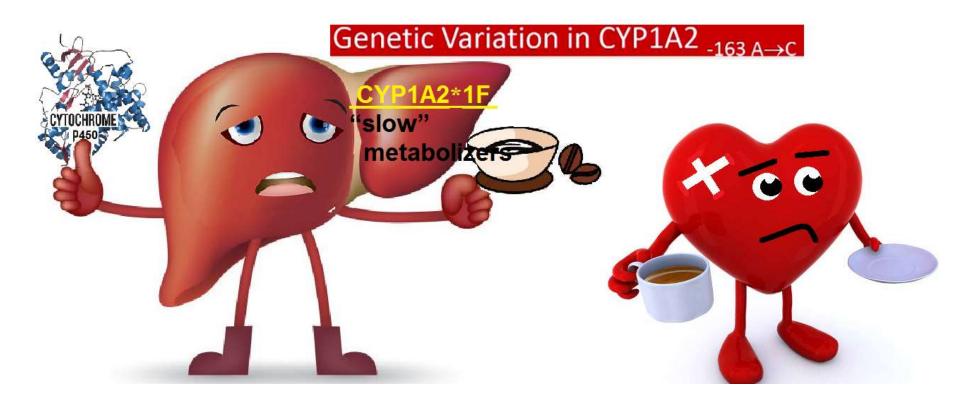


Metabolizing the Caffeine

polymorphic cytochrome P450 1A2 enzyme (CYP1A2)



Polymorphism of CYP1A2



Polymorphism of CYP1A2 Cont'd.

JAMA. 2006 Mar 8;295(10):1135-41.

Coffee, CYP1A2 genotype, and risk of myocardial infarction.

Cornelis MC¹, El-Sohemy A, Kabagambe EK, Campos H.

Author information

Cancer Epidemiol Biomarkers Prev. 2007 May;16(5):912-6.

The CYP1A2 genotype modifies the association between coffee consumption and breast cancer risk among BRCA1 mutation carriers.

Kotsopoulos J¹, Ghadirian P, El-Sohemy A, Lynch HT, Snyder C, Daly M, Domchek S, Randall S, Karlan B, Zhang P, Zhang S, Sun P, Narod SA.

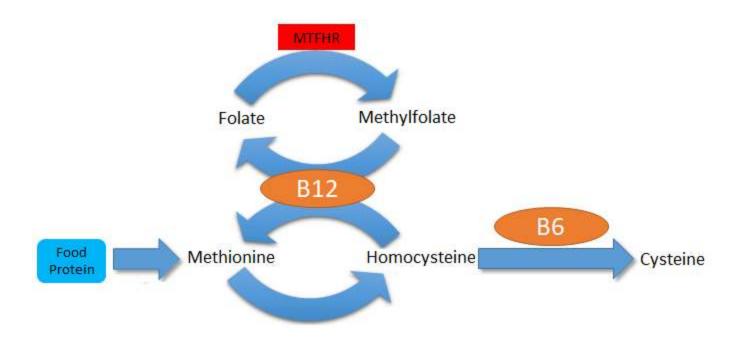
J Hypertens. 2009 Aug;27(8):1594-601. doi: 10.1097/HJH.0b013e32832ba850.

CYP1A2 genotype modifies the association between coffee intake and the risk of hypertension.

Palatini P¹, <u>Ceolotto G</u>, <u>Ragazzo F</u>, <u>Dorigatti F</u>, <u>Saladini F</u>, <u>Papparella I</u>, <u>Mos L</u>, <u>Zanata G</u>, <u>Santonastaso M</u>.

MTHFR (methyltetrahydrofolate reductase)

A common polymorphism: C677T SNP (Ala²²²Val)



plasma homocysteine



Genetic variation in *TAS1R2* (Ile191Val) is associated with consumption of sugars in overweight and obese individuals in 2 distinct populations^{1–3}

Karen M Eny, Thomas MS Wolever, Paul N Corey, and Ahmed El-Sohemy

Nutrition Journal

Research



Open Acces

Improved weight management using genetic information to personalize a calorie controlled diet

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Corresponding author

| Gene | Gene symbol | Polymorphism | %homozygote wild type | % heterozygote | % homozygote variant | HWE p < |
|--|-------------|--------------|--------------------------|----------------|-------------------------|---------|
| Angiotensin I converting enzyme | ACE | INS/DEL | 14.6% | 48.8% | 36.6% | 0.99 |
| Apolipoprotein C-III | APOC3 | 3175C>G | 73.3% | 20.0% | 6.7% | 0.17 |
| Cystathionine-beta-synthase | CBS | 699C>T | 53.5% | 41.9% | 4.7% | 0.81 |
| Cholesteryl ester transfer protein | CETP | 279G>A | 48.8% | 39.5% | 11.6% | 0.86 |
| Collagen, type I, alpha I | COLIAI | G Spl T | 58.1% | 34.9% | 7.0% | 0.94 |
| Glutathione S-transferase MI | GSTMI | Deletion (7) | 52.0% | 0.0% | 48.0% | N/A |
| Glutathione S-transferase pi | GSTPI | 313A>G | 57.8% | 33.3% | 8.9% | 0.68 |
| | | 341C>T | 56.8% | 34.1% | 9.1% | 1.00 |
| Glutathione S-transferase theta I | GSTTI | Deletion (*) | 86.0% | 0.0% | 14.0% | N/A |
| Interleukin 6 | IL6 | -174G>C | 66.7% | 33.3% | 0.0% | 0.37 |
| | | -634G>C | 86.0% | 14.0% | 0.0% | 0.89 |
| Lipoprotein lipase | LPL | 1595C>G | 69.8% | 27.9% | 2.3% | 1.00 |
| S-methyltetrahydrofolate- homocysteine methyltransferase reductase | MTRR | 66A>G | 19.0% | 45.2% | 35.7% | 0.90 |
| 5,10-methylenetetrahydrofolate reductase | MTHER | 1298A>C | 34.0% | 48.9% | 17.0% | 1.00 |
| | | 677 C>T | 48.0% | 44.0% | 8.0% | 0.95 |
| 5-methyltetrahydrofolate- homocysteine methyltransferase | MTR | 2756A>G | 59.5% | 33.3% | 7.1% | 0.86 |
| Nitric oxide synthase 3 (endothelial cell) | NOS3 | 894G>T | 44.2% | 44.2% | 11.6% | 1.00 |
| Peroxisome proliferator-activated receptor gamma | PPARG | Pro I 2Ala | 75.6% | 15.6% | 8.9% | 0.02 |
| Superoxide dismutase 2, mitochondrial | SOD2 | -28C>T | 10.0% | 54.0% | 36.0% | 0.57 |
| Superoxide dismutase 3, extracellular | SOD3 | 760C>G | 100.0% | 0.0% | 0.0% | 1.00 |
| Tumor necrosis factor | TNFa | -308G>A | 71.1% | 24.4% | 4.4% | 0.72 |
| Vitamin D receptor | VDR | C Tagl T | 23.3% | 46.5% | 30.2% | 0.91 |
| | | T BsmI C | 23.3% | 46.5% | 30.2% | 0.91 |
| | | T FokI C | 11.6% | 58.1% | 30.2% | 0.41 |

Table 2: Genes and polymorphisms tested in the nutrigenetic patient group.

Genotype frequencies in the study group and p-values for Hardy Weinberg Equilibrium (HWE) are shown. (*) the assay only measured presence or absence of the deletion so a HWE test is not applicable.

Association of polymorphic genes with response to nutrients

- TAS1R2 (Ile191Val)
- Glucose transporter type 2 (Glut-2)
- Tumor necrosis factor-alpha (TNF-α)
- Catechol-O-Methyltransferase enzyme (COMT)
- Apolipoprotein E (APO E)
- APOA1
- ...

End of the second section

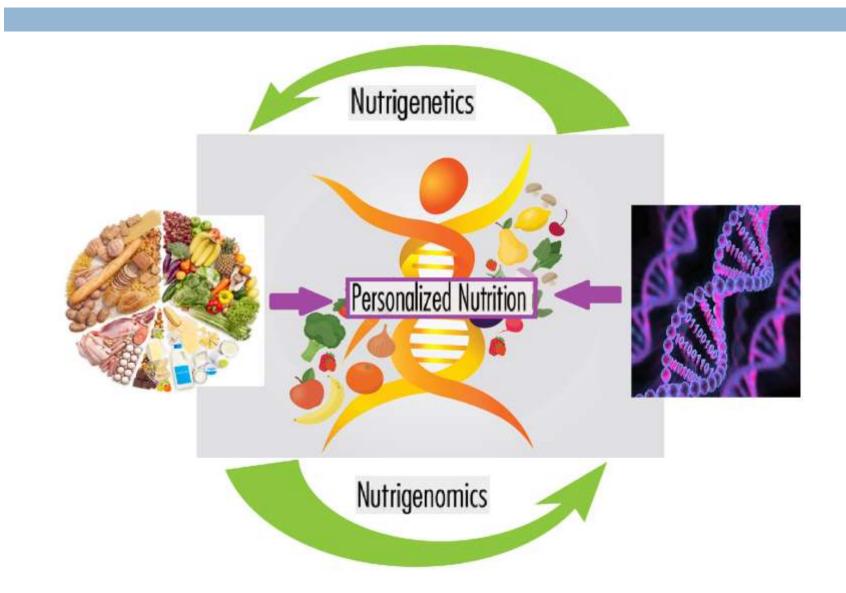
Third Section

FUTURE PROSPECTS

Disease Prevention & Health Promotion



Personalized Nutrition



Forth Section What I have learned?

The Forth Section Cont'd.

- Food sends informational signals to the genes.
- > Your genes are not your destiny.
- Food influence ingested behavior.

Food is more than calories, Food is information !

The Forth Section Cont'd.

- What's your definition of nutrigenomics and nutrigenetics?
- How Personalized Nutrition can be helpful in health promotion and disease prevention
- Nutrients and gene expression can lead to epigenetic altering?
- How nutrients can act as informational signals for our body?

Main References:

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- Fenech M, El-Sohemy A, Cahill L, Ferguson LR, French TA, Tai ES, Milner J, Koh WP, Xie L, Zucker M, Buckley M. Nutrigenetics and nutrigenomics: viewpoints on the current status and applications in nutrition research and practice. Lifestyle Genomics. 2011;4(2):69-89.
- Ramos-Lopez O, Milagro FI, Allayee H, Chmurzynska A, Choi MS, Curi R, De Caterina R, Ferguson LR, Goni L, Kang JX, Kohlmeier M. Guide for current nutrigenetic, nutrigenomic, and nutriepigenetic approaches for precision nutrition involving the prevention and management of chronic diseases associated with obesity. Lifestyle Genomics. 2017;10(1-2):43-62.
- Cornelis MC, EI-Sohemy A, Kabagambe EK, Campos H. Coffee, CYP1A2 genotype, and risk of myocardial infarction. Jama. 2006 Mar 8;295(10):1135-41.

Thanks For your attention