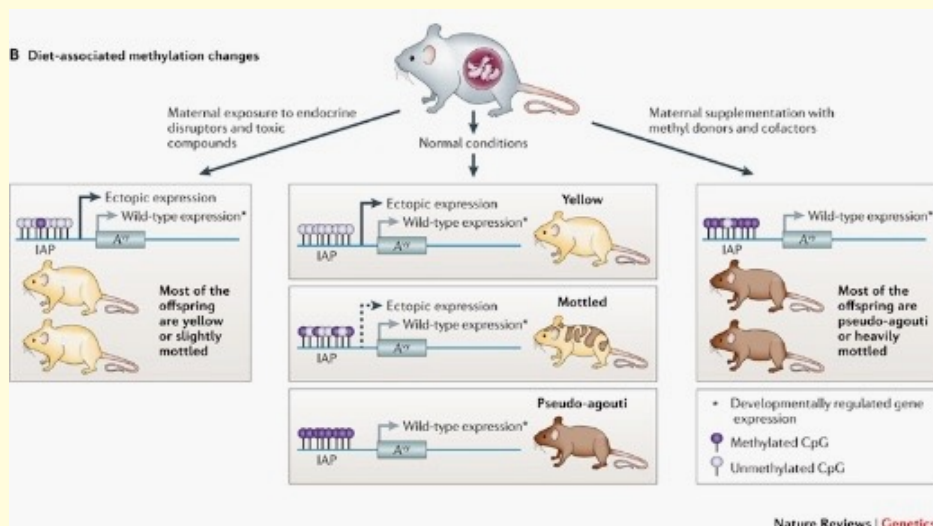
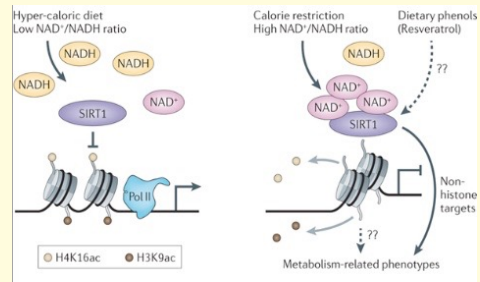
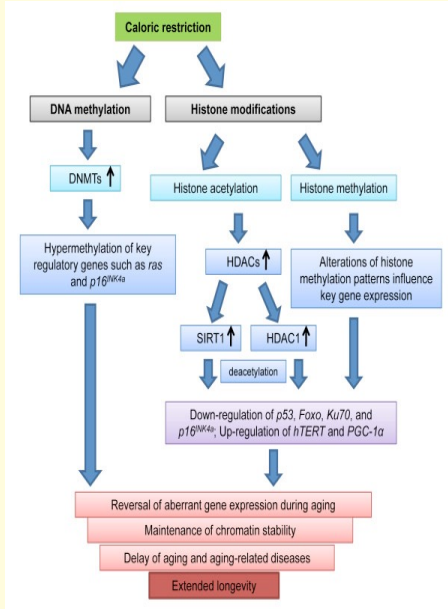


## ENVIRONMENT (NUTRITION) AND EPIGENETICS

### The “Agouti mouse” model

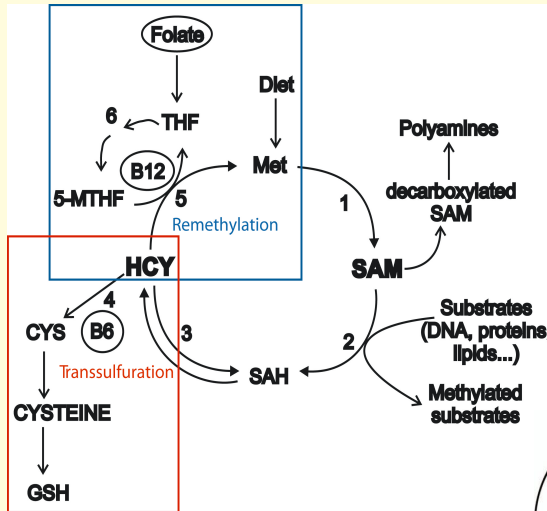


## Caloric restriction



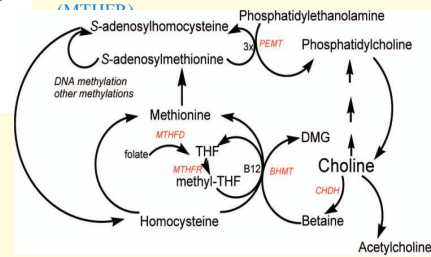
Yuanyuan et al., 2011

## One-carbon metabolism

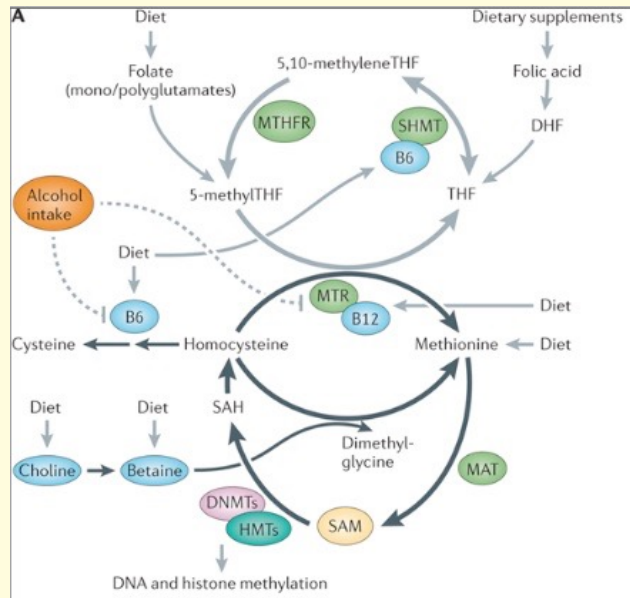


Met, Methionine  
 SAM, S-adenosylmethionine  
 SAH, S-adenosylhomocysteine  
 HCY, Homocysteine  
 CYS, Cystathionine  
 GSH, Glutathione  
 THF, Tetrahydrofolate  
 MTHF, methyltetrahydrofolate  
 B12, Vitamin B12  
 B6, Vitamin B6

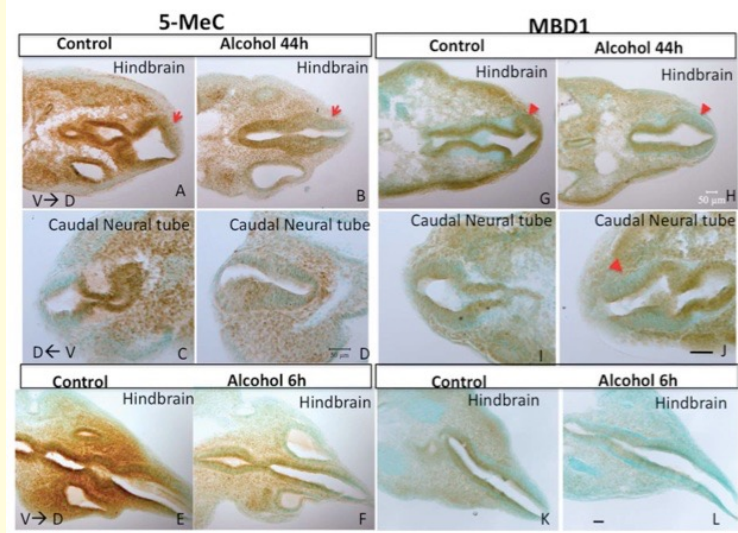
1, Methionine adenosyltransferase (MAT)  
 2, Methyltransferase(s)  
 3, SAH hydrolase  
 4, Cystathionine-β-synthase (CBS)  
 5, Methionine synthase  
 6, Methylene tetrahydrofolate reductase (MTHFR)



## Alcohol and B vitamins



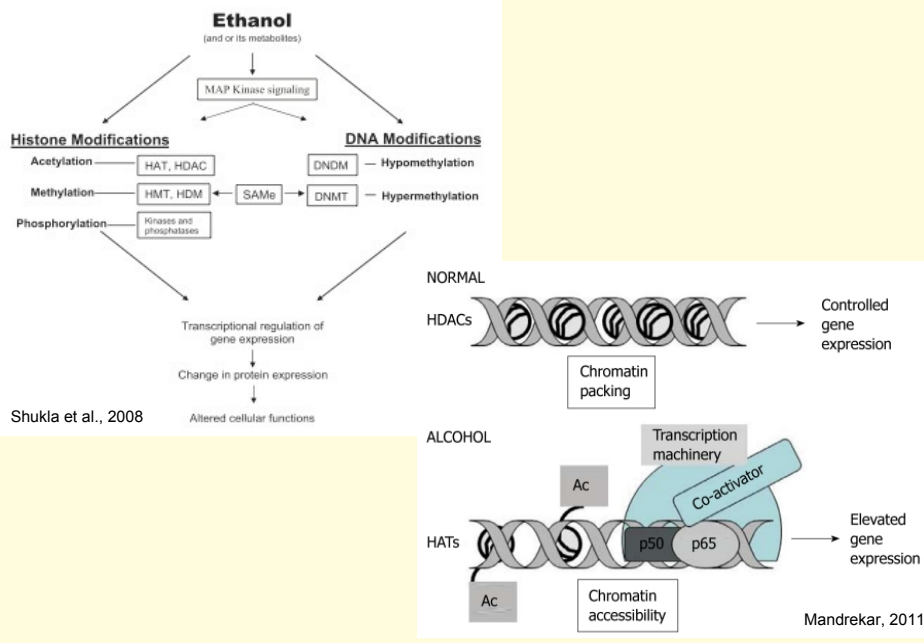
## Alcohol consumption and DNA methylation



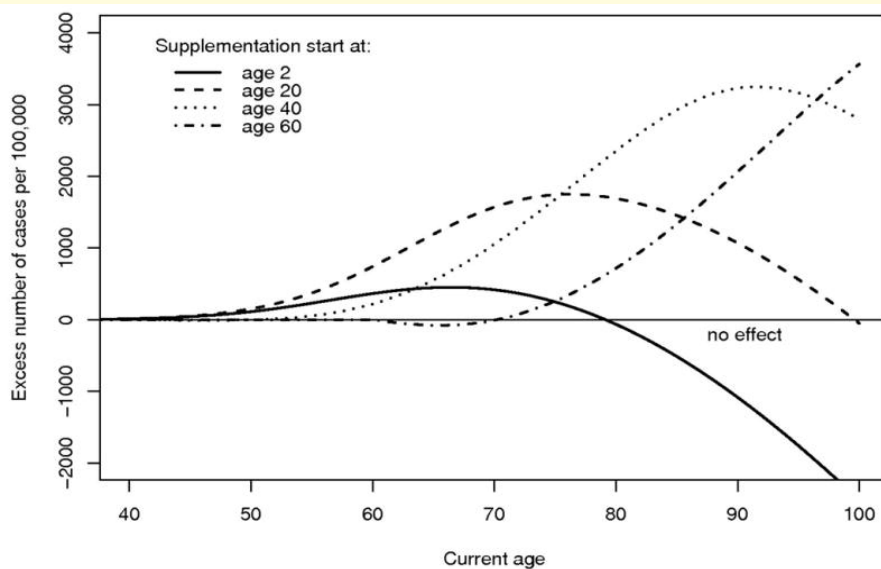
- Riduzione 5-metilcitosina.
- Ritardo nello sviluppo neurale.

Zhou et al., 2011

## Alcohol and histone acetylation



## Folate fortification and colorectal cancer



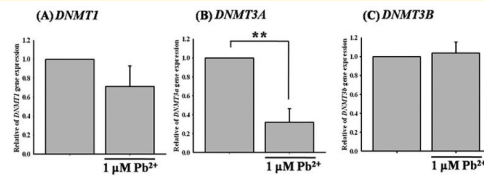
## Metalli

**Table 1.** An overview of the epigenetic effects of metal exposure

	DNA methylation	Histone modifications	ncRNA expression
Cadmium	✓	ND	✓
Arsenic	✓	✓	✓
Nickel	✓	✓	ND
Chromium	✓	✓[Cr (VI)]	ND
Methylmercury	✓	✓	ND
Lead	✓	ND	ND
Copper	ND	✓	ND
Organotin	✓	✓	ND

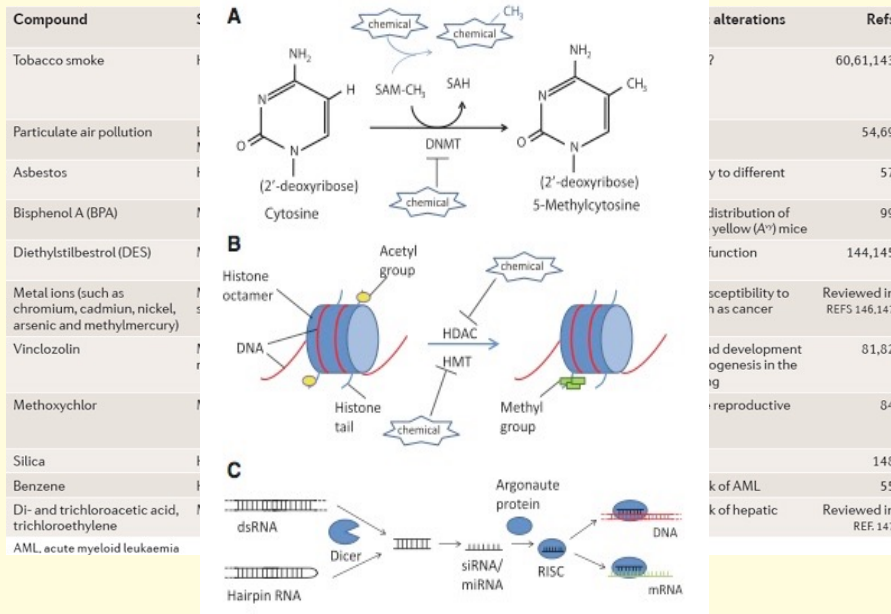
✓, Reported in the literature; ND, no data (no relevant data was available at the time of this publication).

Cheng et al., 2011

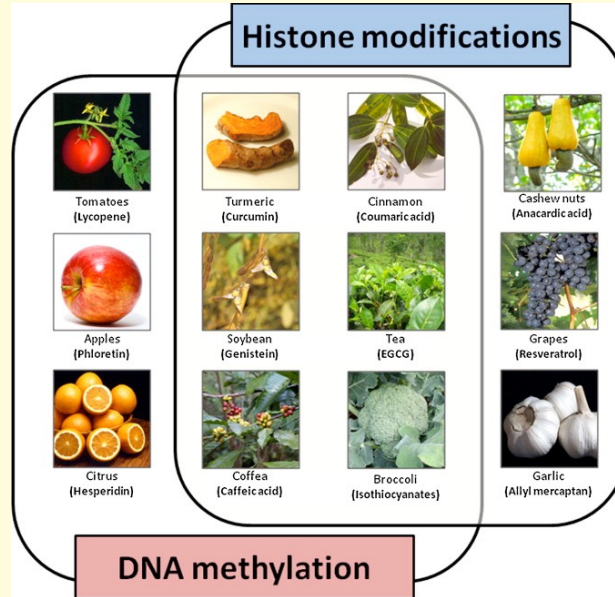


**Fig. 2.** DNMT gene expression in  $Pb^{2+}$ -treated cells. A431 cells were treated with  $1 \mu M Pb(NO_3)_2$  for 1 h, and total RNA was isolated. Gene expression of *DNMT1* (A), *DNMT3a* (B), and *DNMT3b* (C) were detected by real-time PCR. Values for mRNA levels are means  $\pm$  SEM. Statistical significance ( $*p < 0.05$  and  $**p < 0.01$ ) of the difference between control and  $Pb^{2+}$ -treated cells was determined by Student's t-test.

## Pollutants and pesticides



## “Epigenetic” vegetables



## EPIGENETICS (DNA METHYLATION) AND NEURODEGENERATION

## **“EPIGENETIC” DISEASES**

- **Fragile X Syndrome**
- **Rett Syndrome**
- **ICF Syndrome (Immunodeficiency, Centromeric instability and Facials abnormalities)**

• **Atherosclerosis**

## **Epigenetics in aging brain**

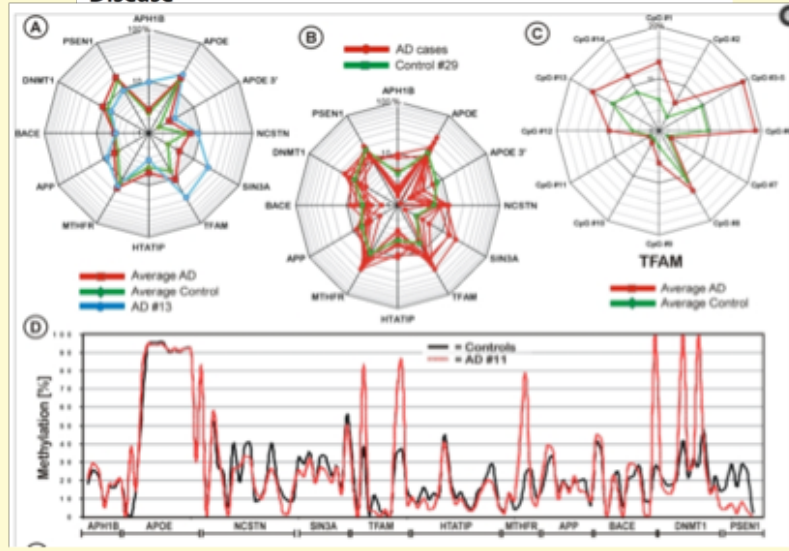
- > **Different cognitive disorders are associated with epigenetic dysregulation** (Graff and Mansuy, 2009)
- > **DNA methylation in individuals changes over time - hypomethylation** (Bjornsson et al., 2008)
- > **Early-life events may trigger biochemical pathways during aging through epigenetic modifications** (Lahiri et al., 2008)
- > **DNA methylation pattern in human brain seems to be area-specific** (Ladd-Acosta et al., 2007)
- > **DNA methylation pattern in human brain is dynamically changing during the lifespan - hypomethylation** (Siegmund et al., 2007)
- > **DNA methylation signatures in development and aging of the human prefrontal cortex** (Numata et al., 2012)
- > **Longitudinal changes in gene-specific DNA methylation** (Madrigano et al., 2012)
- > **Epigenetic regulation of BDNF affects aging** (Zeng et al., 2011)

## DNA methylation in AD subjects

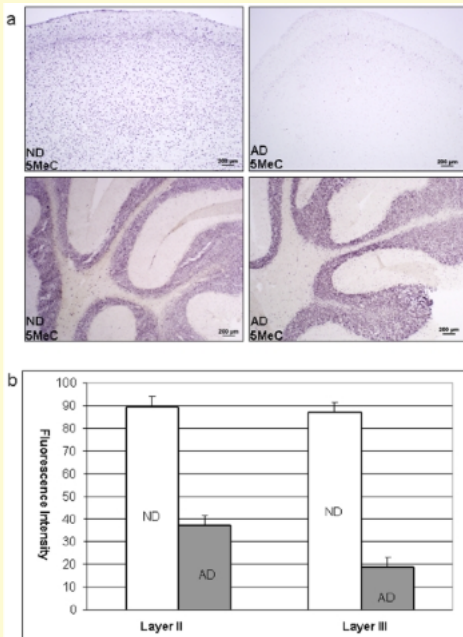
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PLoS one

### Age-Specific Epigenetic Drift in Late-Onset Alzheimer's Disease



## Differential DNA methylation in twins discordant for AD



DNMT1, MBD2 and other methylation “markers” are modulated in LOAD patients (Mastroeni et al., 2008)



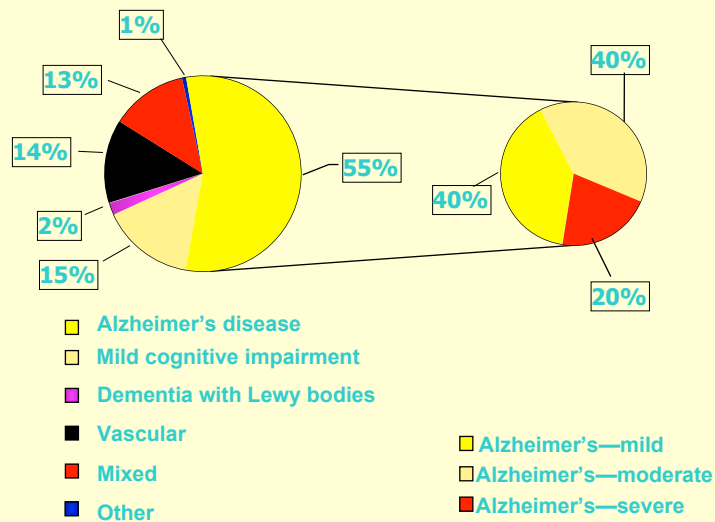
## Alzheimer's Disease (AD)



- Marta, I can't remember the name of that German guy I've lost my mind for...

- Alzheimer, grandma'

## Distribution of different forms of dementia



## Alzheimer Disease: AD

a progressive CNS disorder with a characteristic pathology

### Clinical features:

- memory deficits
- cognitive deficits
- alteration of motor and sensorial functions

### Pathological-anatomical features :

- extracellular deposits of  $\beta$ -amyloid protein (senile plaques)
- Intracellular fibrillar deposits of tau protein (neurofibrillary tangles)
- neuronal death (cortex, hippocampus, amygdala)

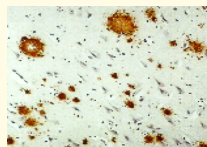
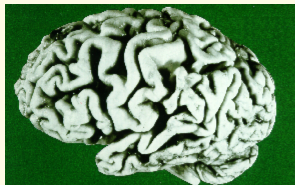
### Bio-molecular features ?? :

- increased APP expression
- increased  $\beta$ - e  $\gamma$ - secretase activity
- decreased  $\beta$ -amyloid clearance
- increased tau expression
- unbalance of tau phosphorylation/dephosphorylation

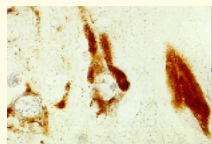
Many possible risk factors: the complex, non-Mendelian disease etiology suggests that an epigenetic component could be involved

## Alzheimer's Disease (AD)

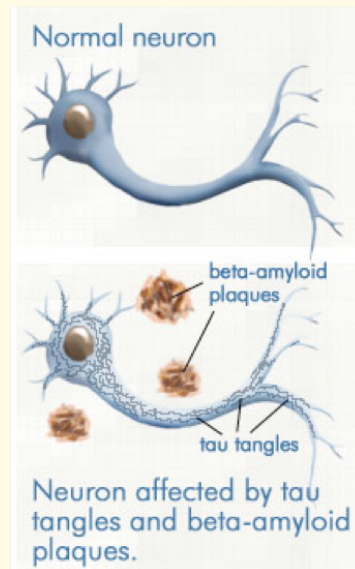
Brain atrophy



Senile Plaques  
(Amyloid  $\beta$ )

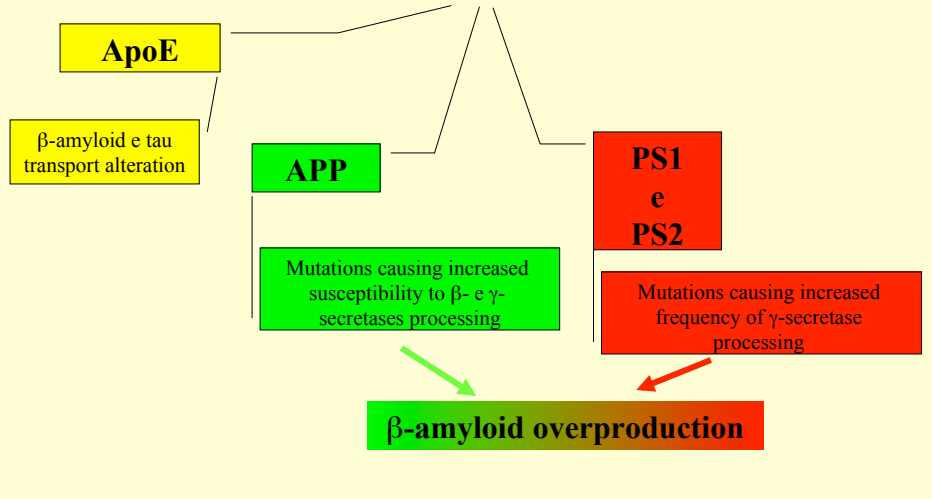


Neurofibrillary Tangles  
(Tau)



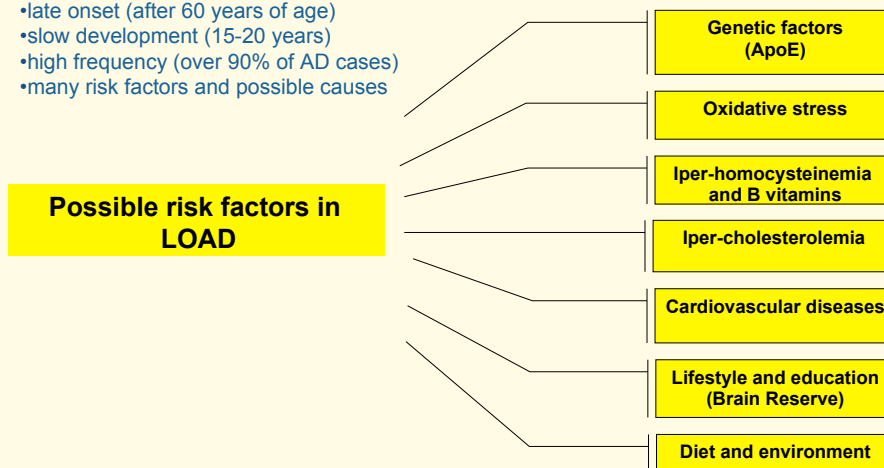
## AD: Genetic disease (FAD: familial AD)

- early onset (40-50 years of age)
- rapid development (2-3 years)
- autosomic dominant genetic pattern
- Due to mutations at specific genes



## AD: Sporadic disease (LOAD: Late Onset AD)

- late onset (after 60 years of age)
- slow development (15-20 years)
- high frequency (over 90% of AD cases)
- many risk factors and possible causes



The complex, non-Mendelian disease etiology suggests that an epigenetic component could be involved

## PSEN1: the amyloid hypothesis

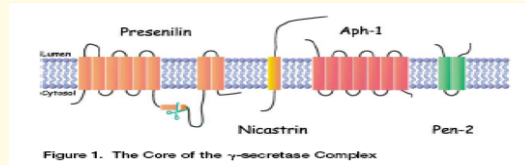
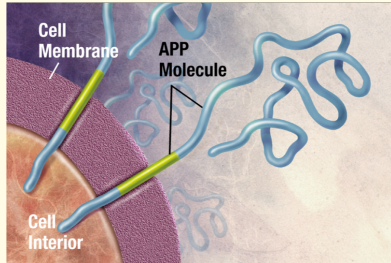
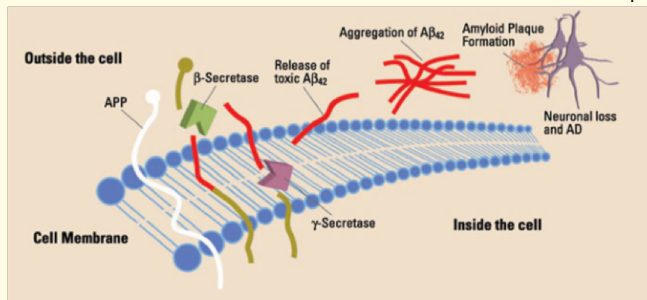


Figure 1. The Core of the  $\gamma$ -secretase Complex

PSEN1 is the main component of the gamma-secretase



## NUTRITION, EPIGENETICS AND

## NEURODEGENERATION

(a paradigmatic example)