

Epigenetics, Evolution and Development

**Behavioral, Metabolic and Immune
Implications**

**George P. Chrousos,
University of Athens**

Evolution vs. Development =

The Evo-Devo Field

Phylogeny vs. Ontogeny

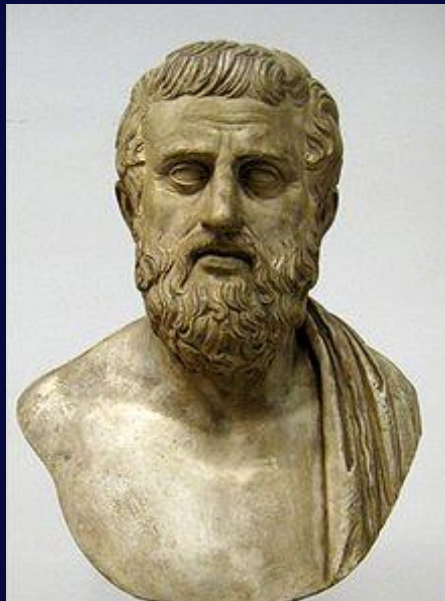
Genetics vs. Epigenetics ,

Genetics vs. Epigenetics

- **Complex Systems**
 - **Evolutionary and Developmental Stressors**
 - **Stress and Evo-Devo**
-

***“Πολλά τα δεινά κ’ουδέν ανθρώπου
δεινότερον πέλλει....”***

***‘There are many wonderful things and
nothing is more wonderful than the
human ...’***



Σοφοκλής

Sophocles

496-406 BCE

HUMAN COMPLEXITY: **POST(EPI)GENOMIC ERA**

Human genome:

About 3 billion bases

About 20 thousand protein-coding genes

About 18 thousand ncRNA-coding genes

**About 100-140 thousand transcripts
(mRNA, ncRNA)**

About 200-260 thousand proteins

Single nucleotide polymorphisms (snp' s or snv' s),
microsatellites or copy number variants :

About >25 million snp' s (snv' s), 1.5 million indels

About 20 million microsatellites

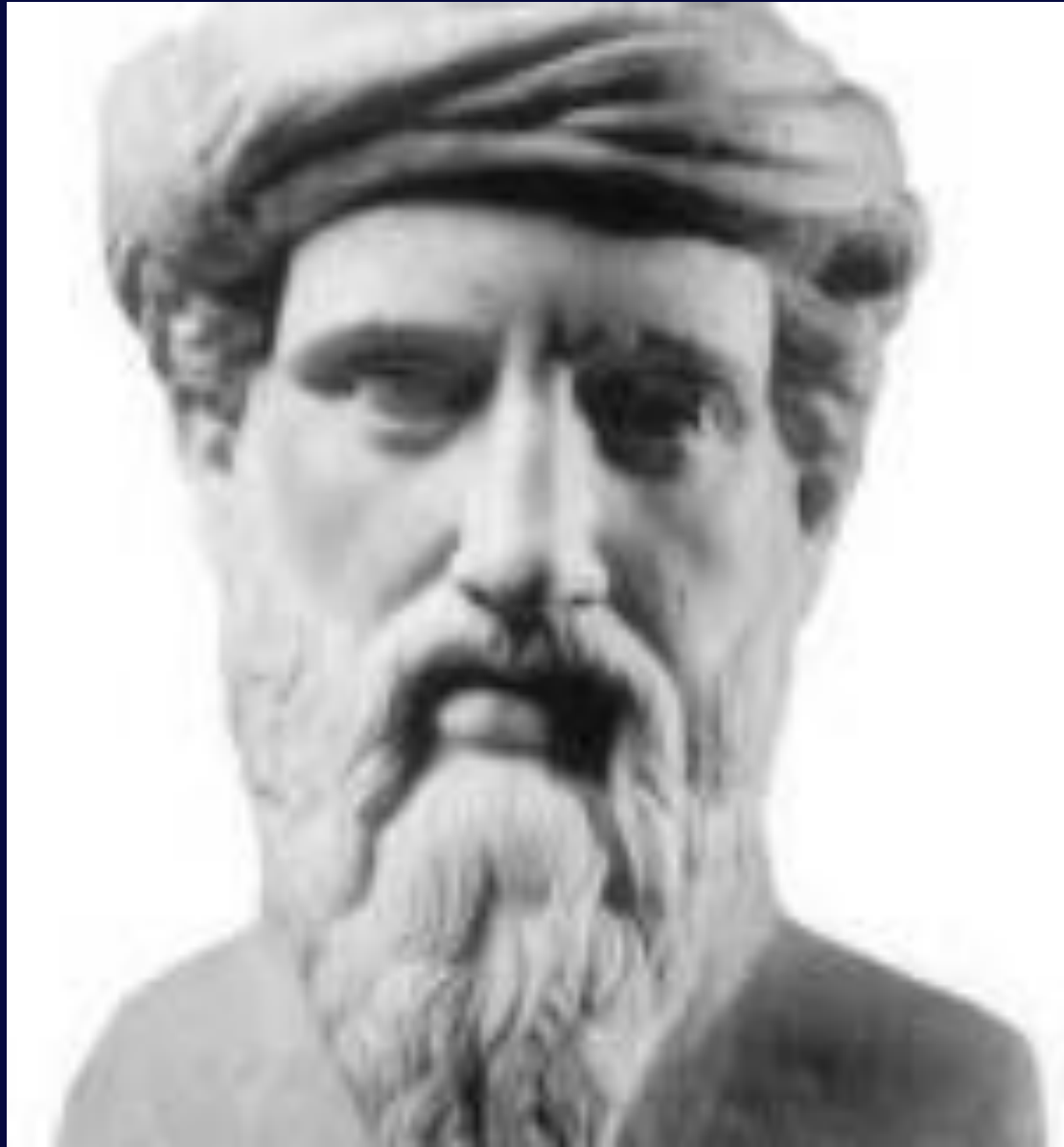
>5000 cnv' s (many million bases)

Over 10 k disease-related mutations

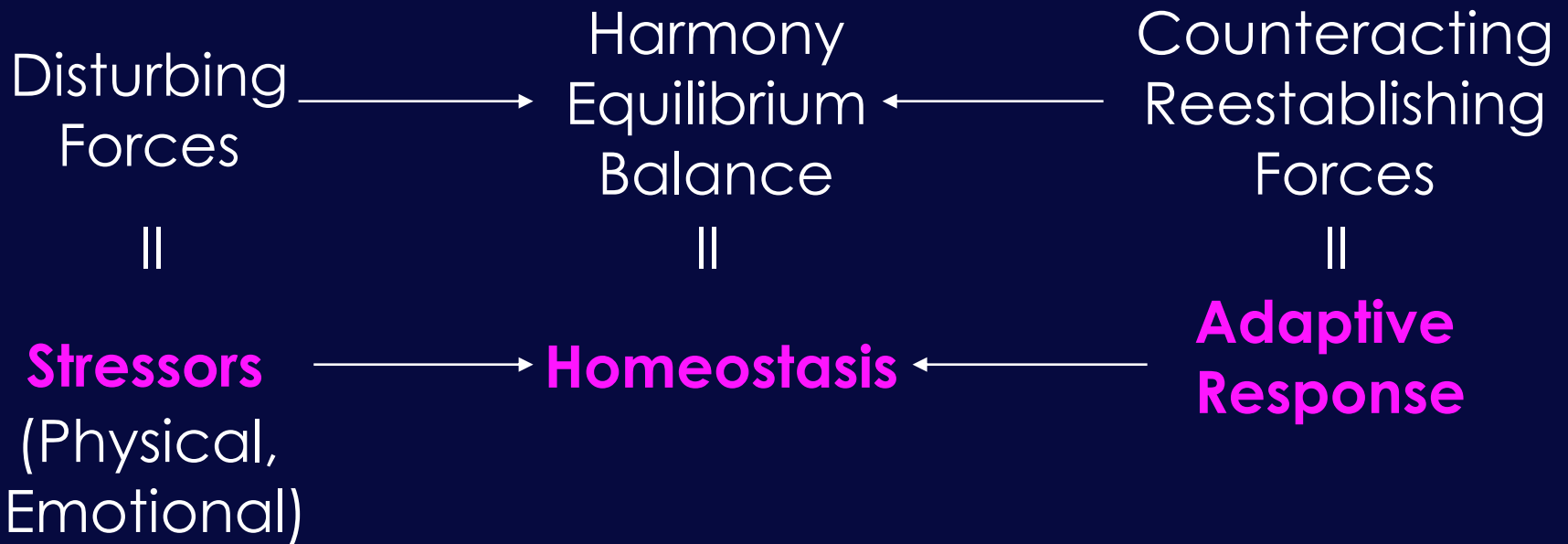
60% of promoters have CpG islands

HUMAN COMPLEXITY: **SOME HUMAN BRAIN NUMBERS**

- ~ 100 billion neurons (100×10^{12}) x >10.000 synapses per neuron = $>10^{18}$ synapses)
 - ~ 100.000 km of fibers
 - ~ 1 trillion or more glial cells
 - ~ 1.25 terabytes
 - ~ 15 Watt lamp (2% of BW uses 20% energy)
-



Pythagoras 6th century BCE



Pythagoras= *Harmony*

Alcmaeon= *Iso-nomia*

Walter Cannon= *Homeostasis*

**Stress is the State of
Threatened (or Perceived
Threatened) Homeostasis**

Genetics vs. Epigenetics

- **Complex Systems**
 - **Evolutionary and Developmental Stressors**
 - **Stress and Evo-Devo**
-

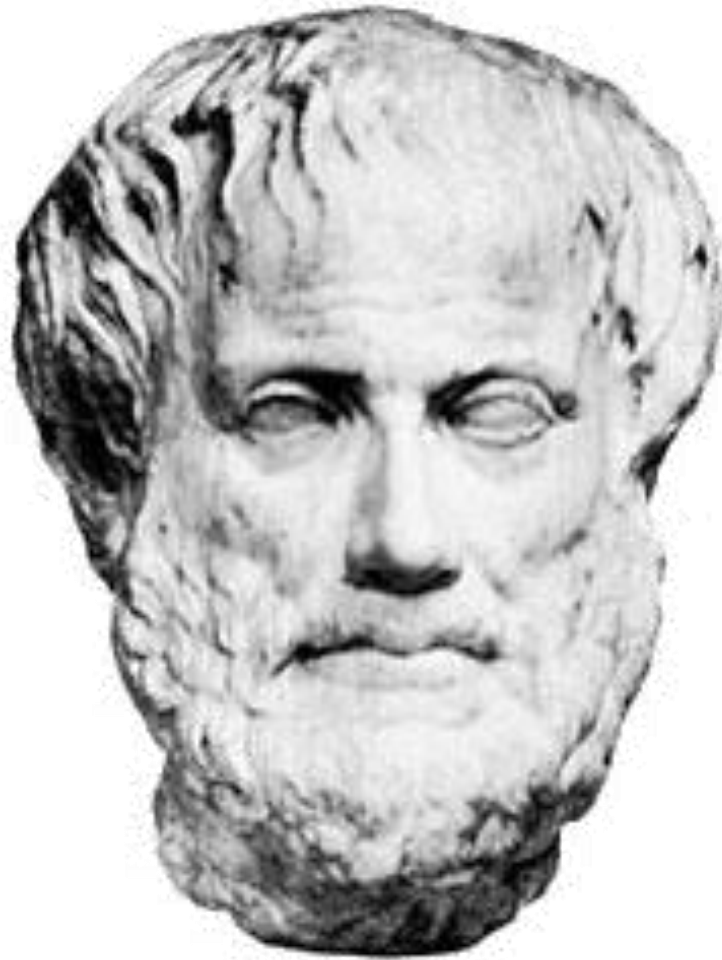
Evolutionary and Developmental Stressors

- **Starvation**
 - **Dehydration/hemorrhage (gastroenteritis, trauma)**
 - **Injurious agents (infections, toxic substances)**
 - **Adversaries (anticipation, minimization of exposure)**
 - **Tissue injury**
-



Raphael, 16th Century CE





Aristotle by Lysippos 4th Century BCE

The Human Brain

- Plato (*Meno*)

The innate preformation theory

(Genetic view)

- Aristotle

The blank state theory

(Epigenetic view)

Preformation (Plato)

(unfolding of preformed tissues)

Epigenesis (Aristotle)

(Embryology, Development)



Jean-Baptiste Lamarck (1744–1829)



Charles Robert Darwin (12 February 1809 – 19 April 1882)

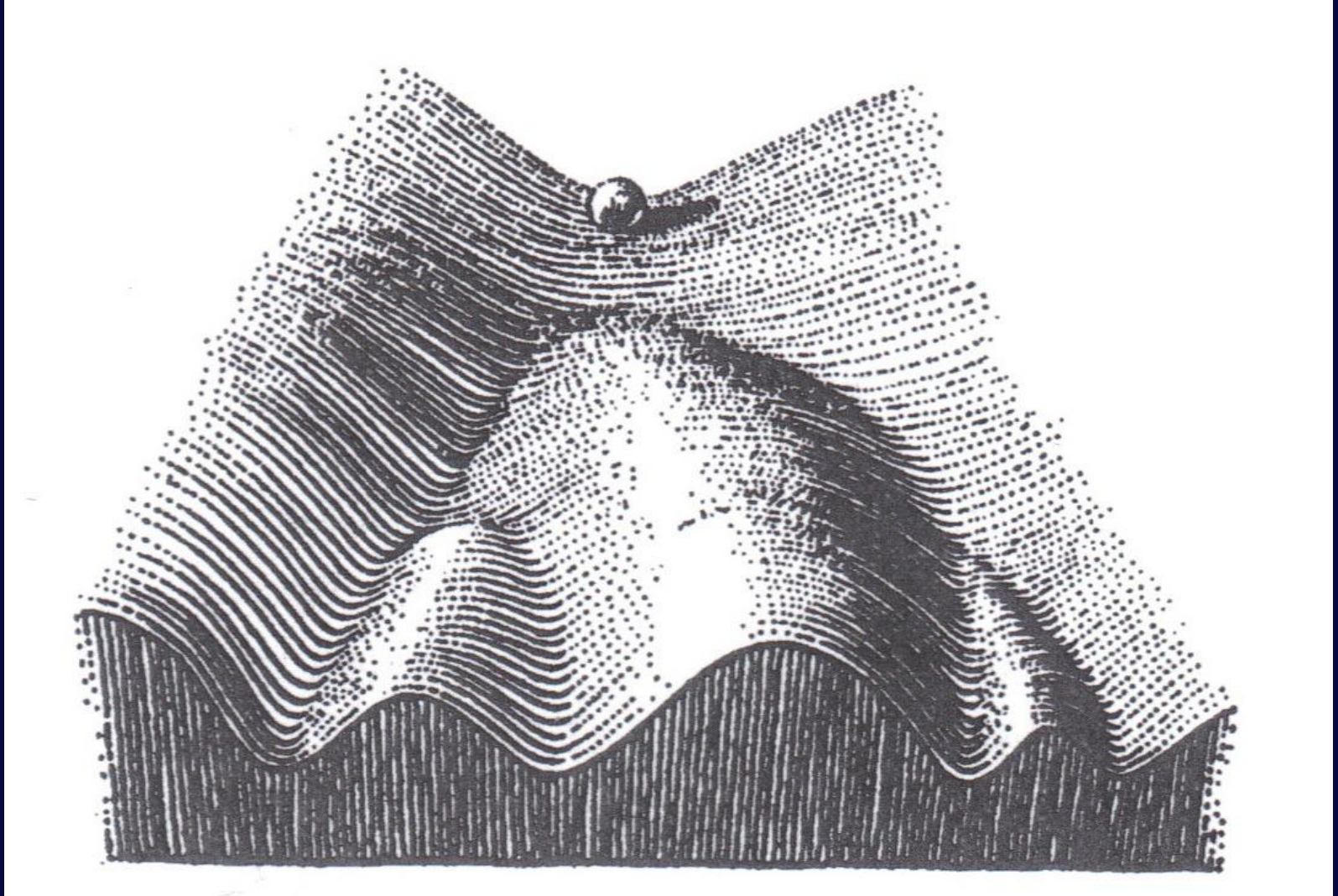
Modern definitions

“Epigenetics are the causal interactions between genes and their products which bring the phenotype into being” 1942

Conrad H Waddington 1900-1975

The Strategy of Genes, MacMillan 1957

“The Epigenetic Landscape”



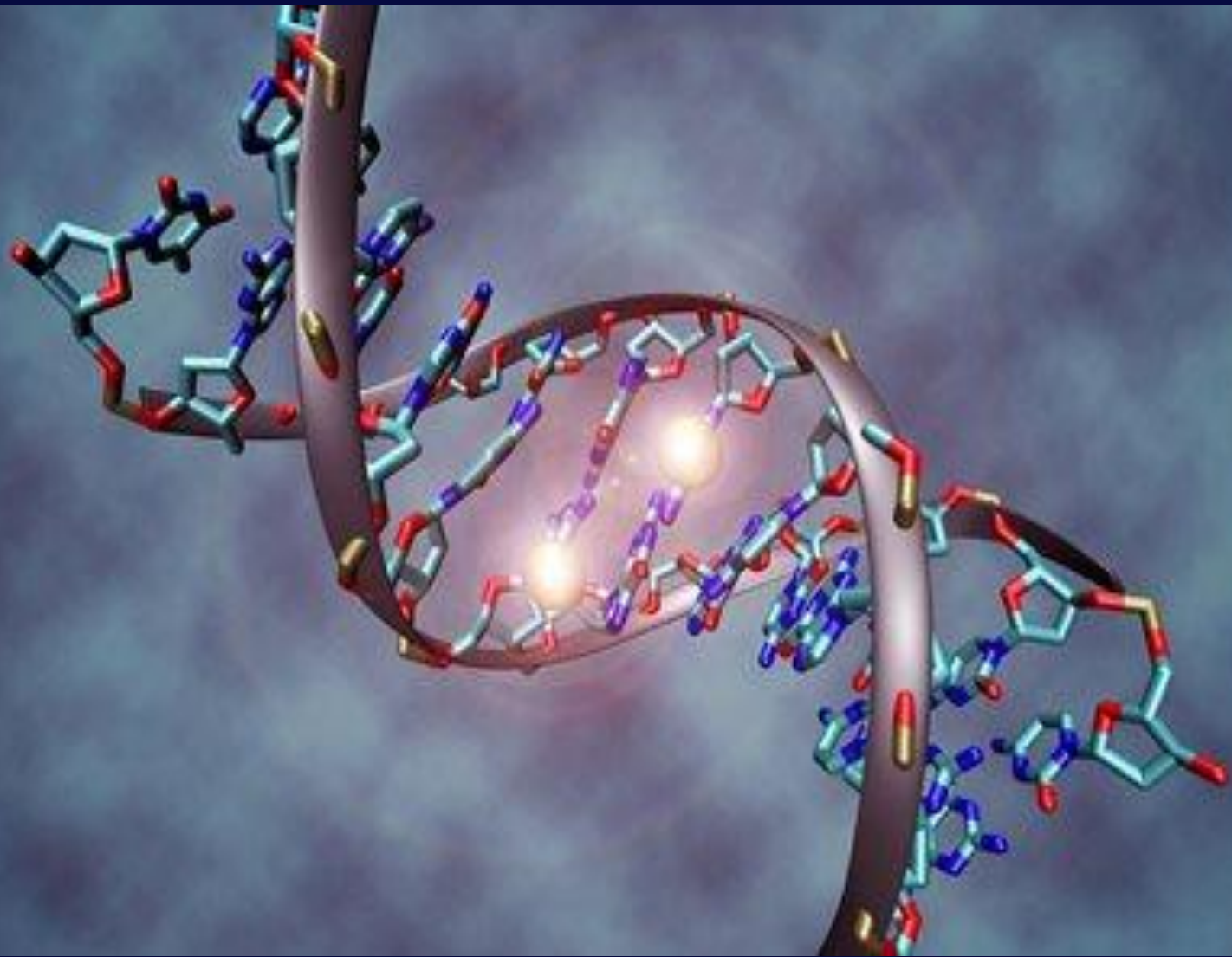
Conrad H Waddington 1900-1975

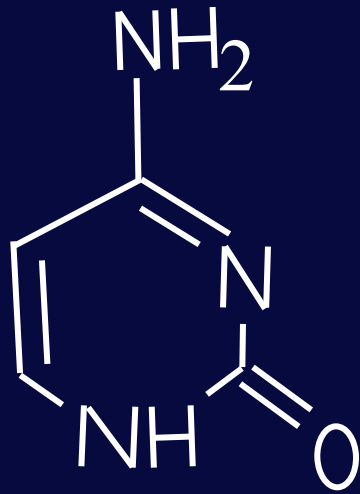
Modern definitions

***“Genetics proposes;
Epigenetics disposes”***

Medawar and Medawar 1983







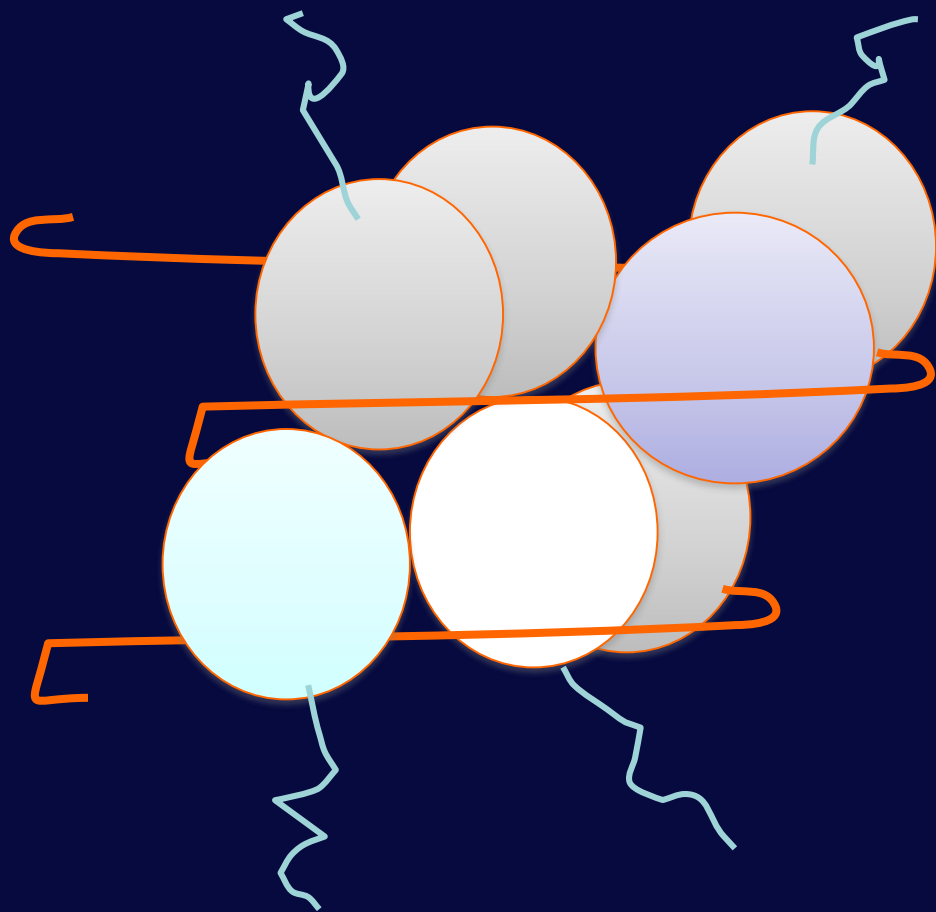
Cytosine

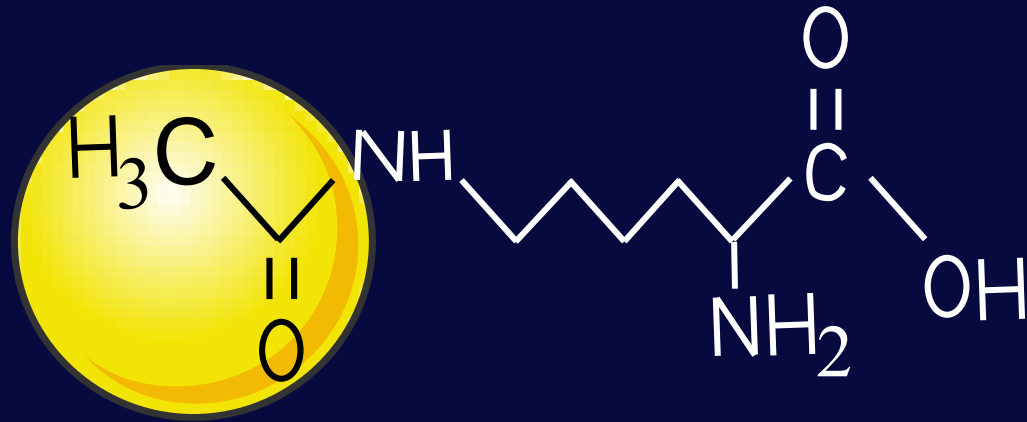


DNMT1
DNMT3A
DNMT3B

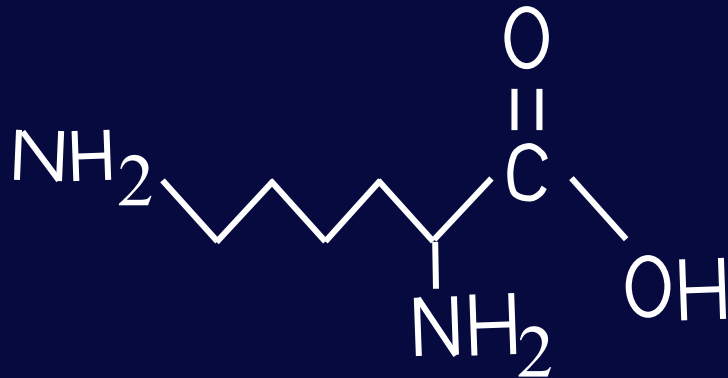


5-Methylcytosine





Acetyl-lysine



Lysine

Forms of Inheritance

- ***Genetic (blueprint)***
 - ***Structural***
 - ***Steady state***
 - ***Epigenetic***
 - ***Behavioral/Symbolic (memes)***
-

Components of Epigenetic Processes

- *Covalent bonds on DNA*
 - *Post-translational modification of proteins*
 - *DNA-binding proteins or complexes (Polycomb/Trithorax complexes (-/+ hox genes), Panoramix complex – other genes*
 - *miRNAs, other ncRNAs*
-

Epigenetic Mechanisms

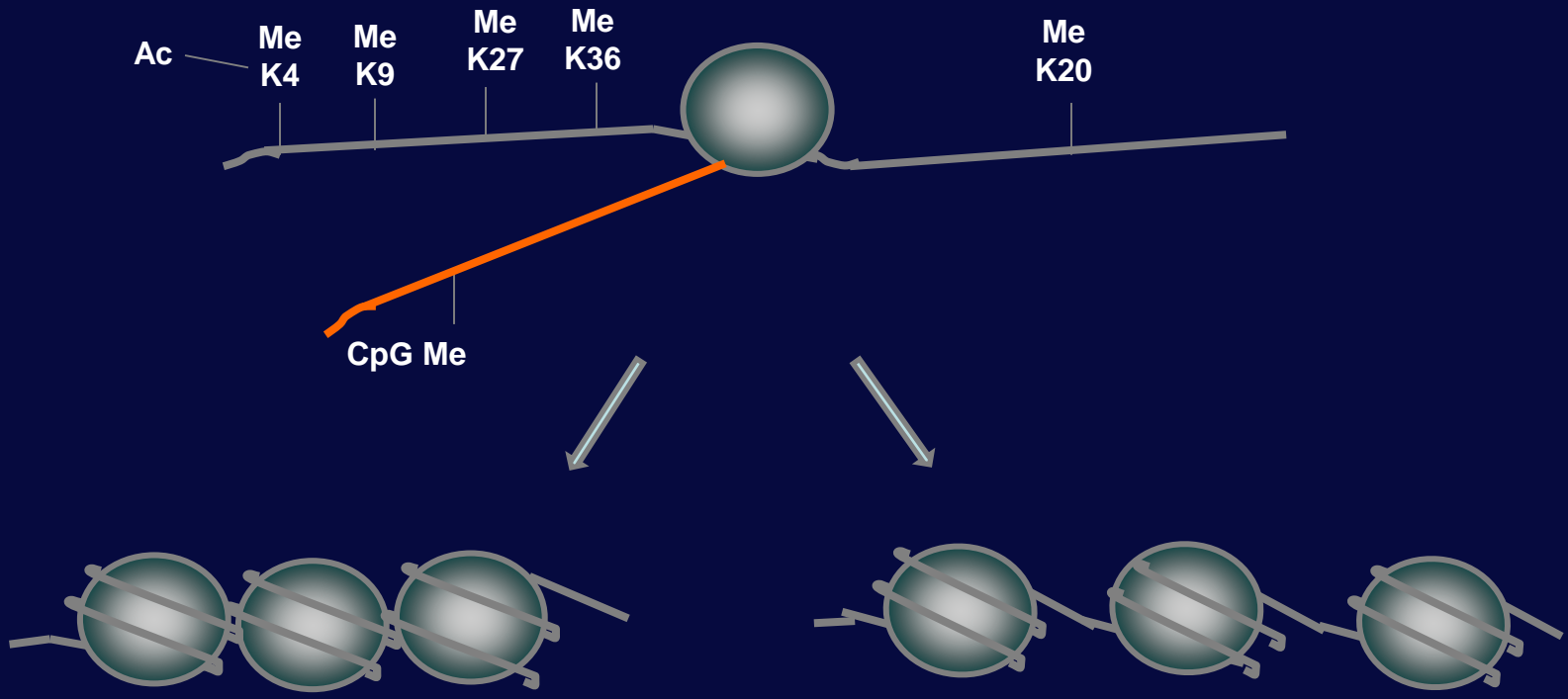
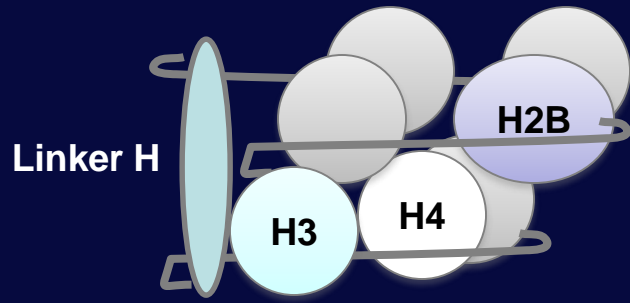
- **DNA Methylation/demethylation, Acetylation/deacetylation**

- **Covalent histone modifications**

(methylation, acetylation, phosphorylation, polyADP-ribosylation)

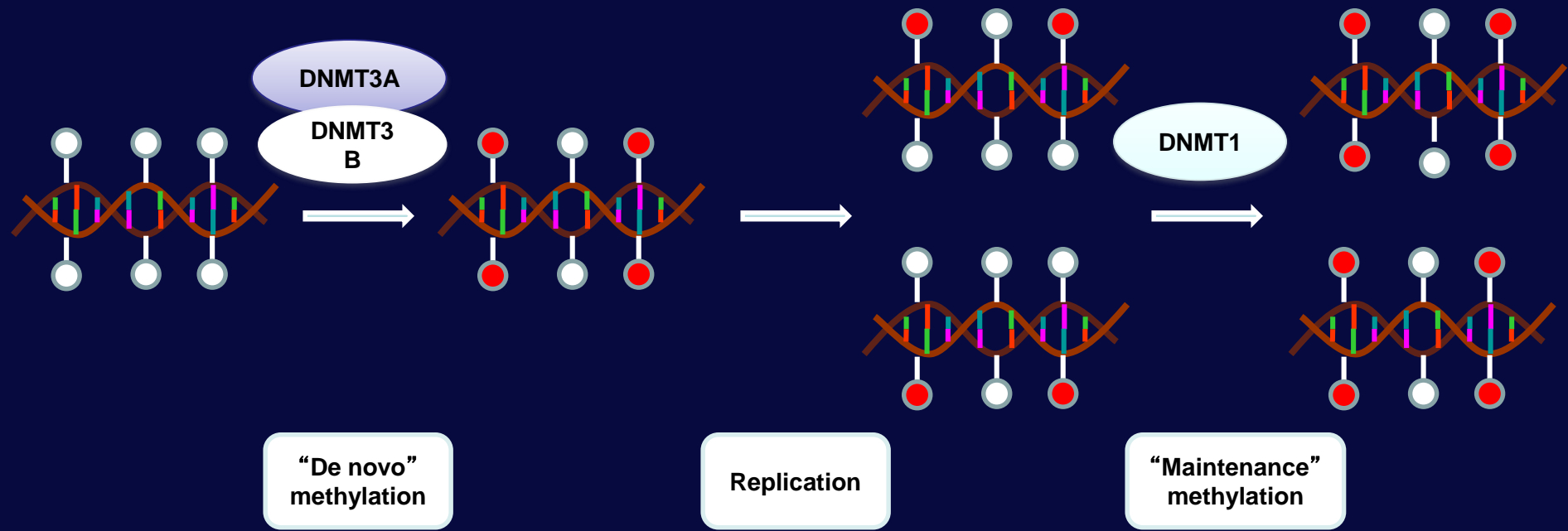
- **Methyl-CpG domain-binding proteins**

- **Chromatin compacting or unwinding complexes (polycomb, trithorax groups)**

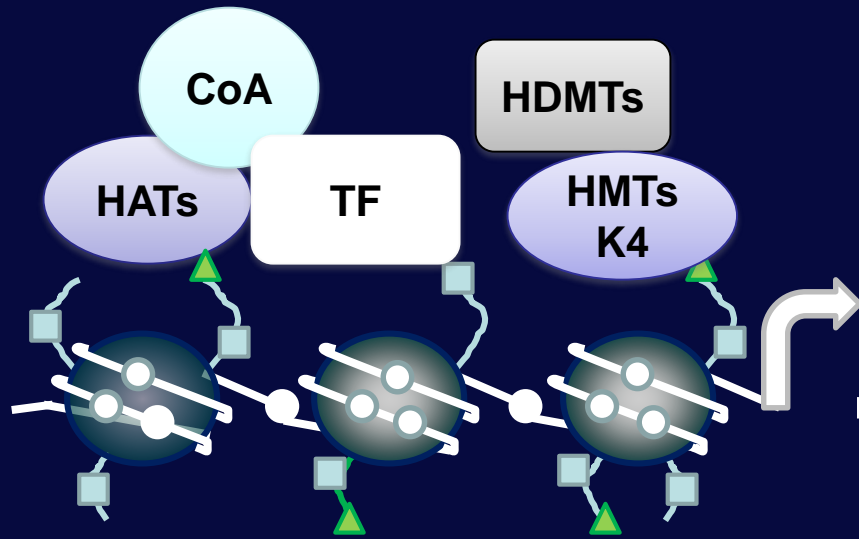


Gene Silencing

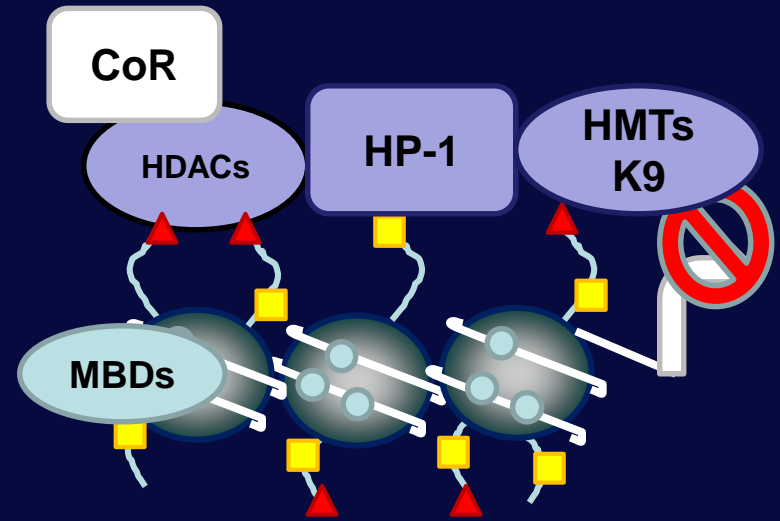
Gene Activation



Transcription



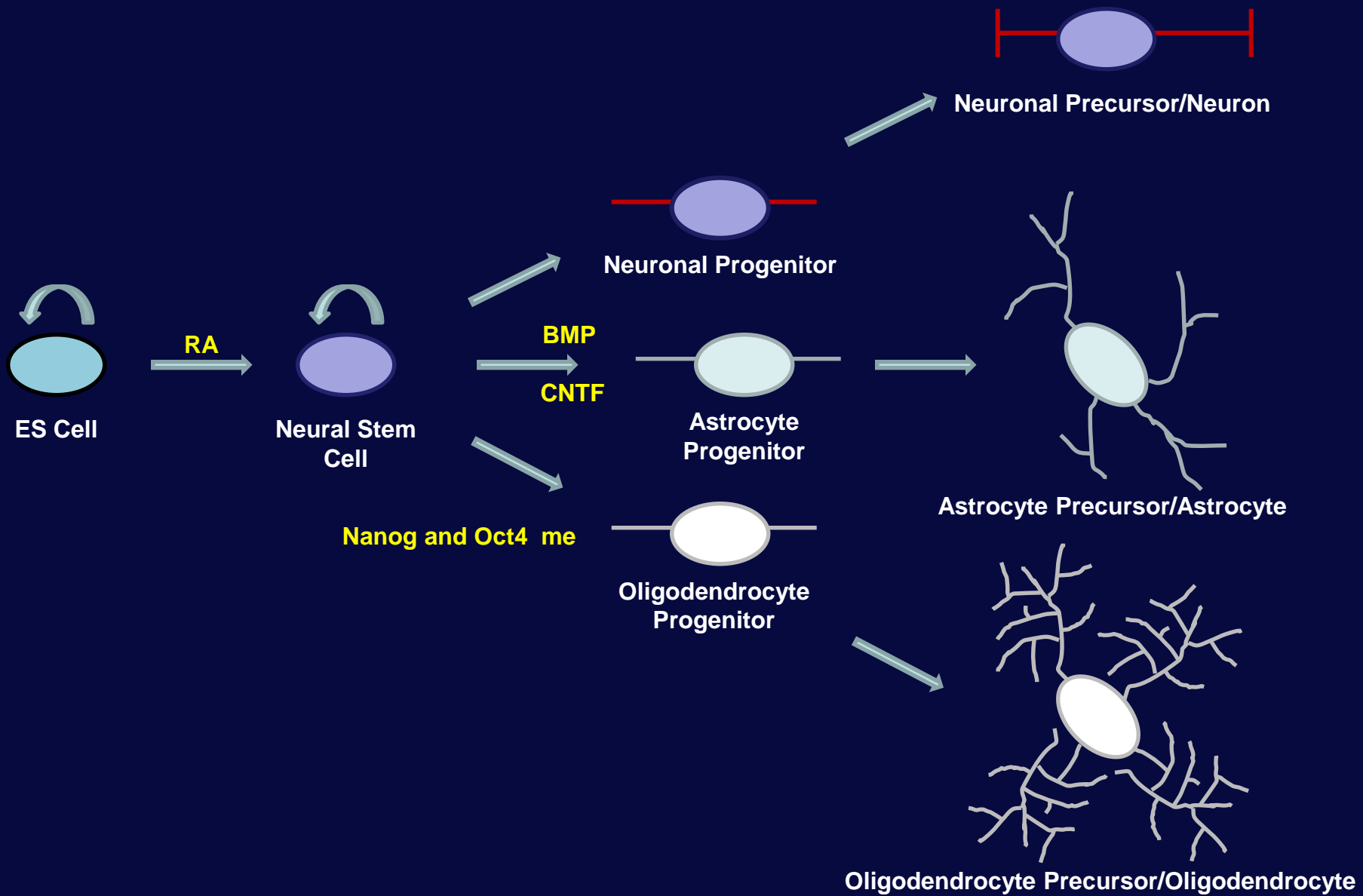
Silencing



- CpG
- ▲ Acetylation
- H3-K4me
- Me-CpG
- ▲ Deacetylation
- H3-K9me

Epigenetic Functions

- ***Cell differentiation***
 - ***Genomic imprinting***
 - ***X-chromosome inactivation***
 - ***Retrotransposon repression***
 - ***Puberty***
 - ***Sexual orientation***
 - ***Right/left handedness***
 - ***Labor and delivery***
 - ***Immune cell differentiation***
-



Genetic Imprinting



**Both copies
from mother:
Mouse
smaller than
normal**



**One copy
from each
parent:
Normal
mouse**



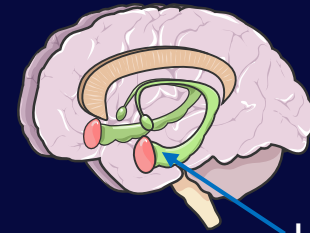
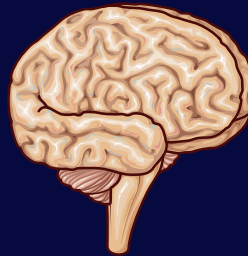
**Both copies
from father:
Mouse larger
than normal**

**Source of
chromosome
11**

Lots of licking and grooming of baby rat

Increased production of serotonin, the "happiness" neurotransmitter in the brain

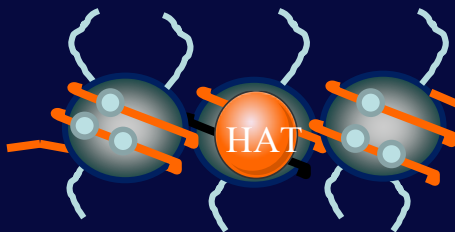
Serotonin signals to the hippocampus to increase production of an enzyme that acetylates histones (HAT)



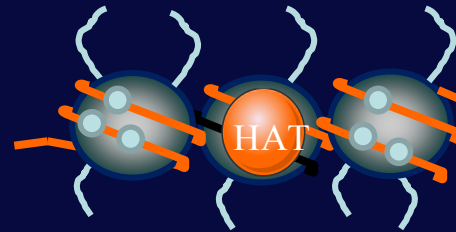
Hippocampus



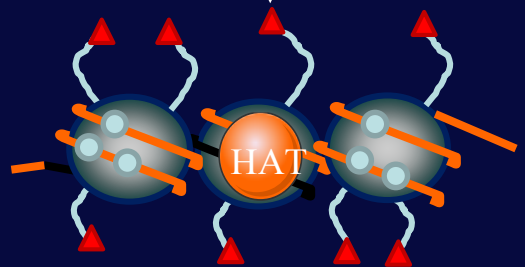
'Chilled out rat'



Decreased DNA methylation leads to higher expression of glucocorticoid receptor

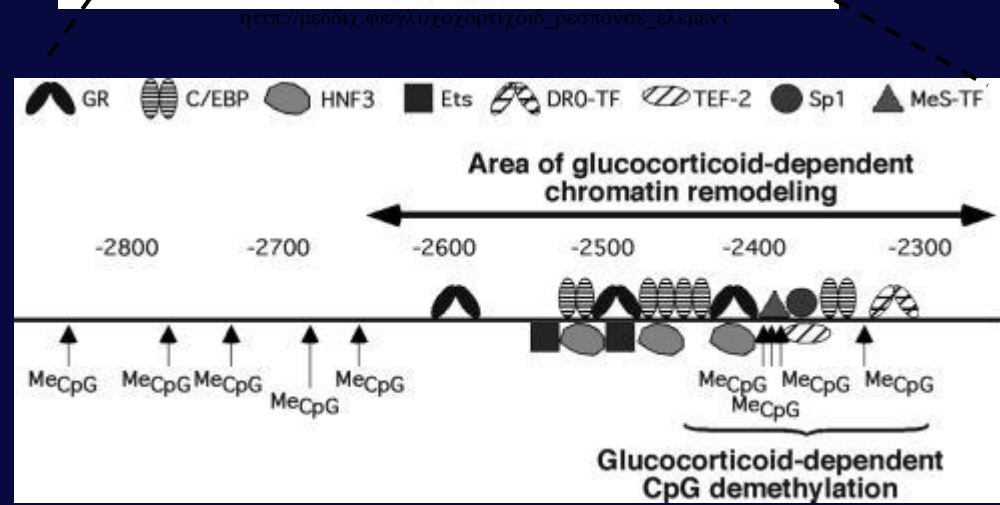
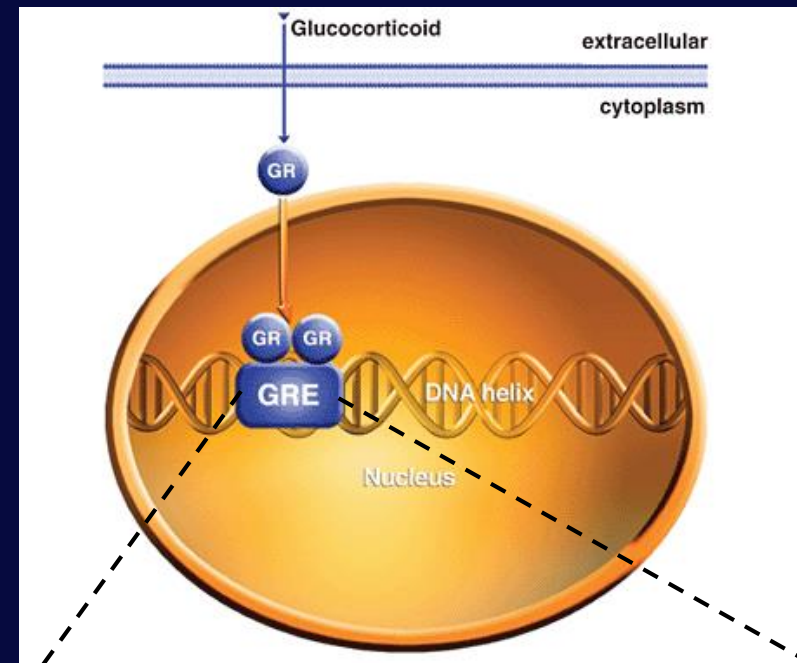
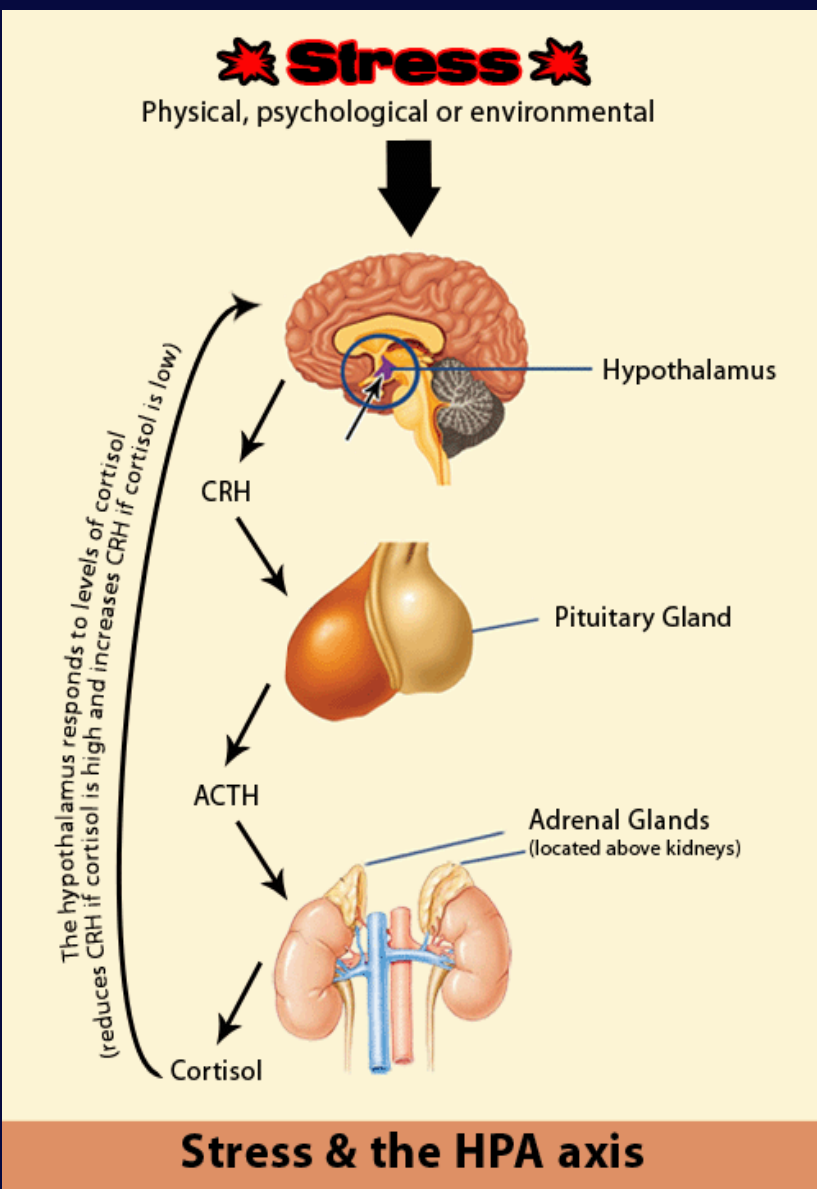


Histone acetylation creates a more relaxed chromatin environment. DNA methylation is removed



HAT binds to glucocorticoid receptor gene and adds acetyl groups to histone proteins

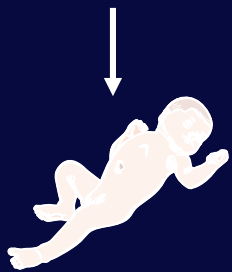
'Stressing' the epigenome: glucocorticoids



**Malnourished in
FIRST trimester**



Mary



Baby normal
birthweight/MS



Helen



Baby likely to be
heavier than
average/ MS

**Malnourished in
THIRD trimester**



Vicky



Baby reduced
birthweight/MS

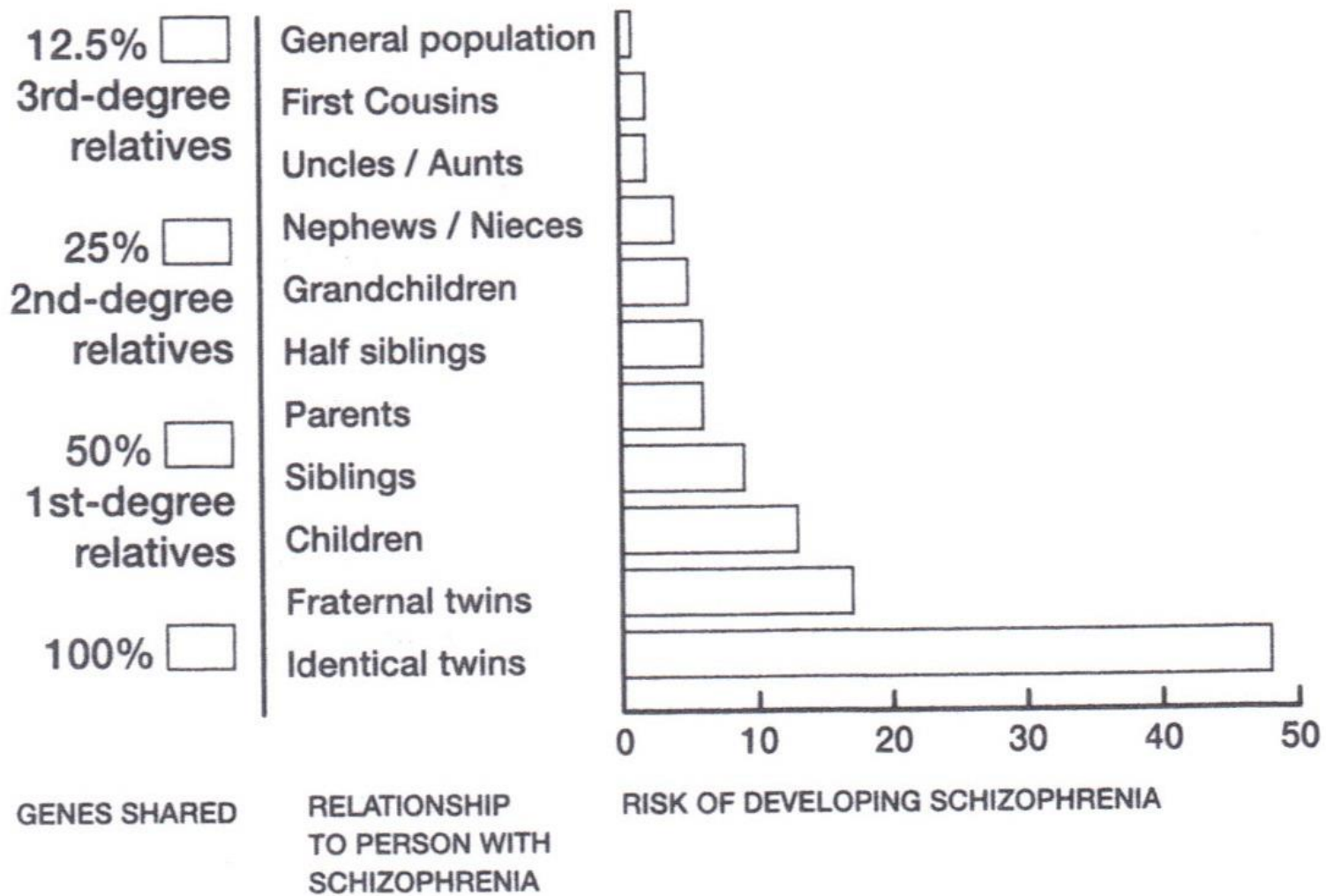


Kelly



Baby normal
birthweight/
no MS

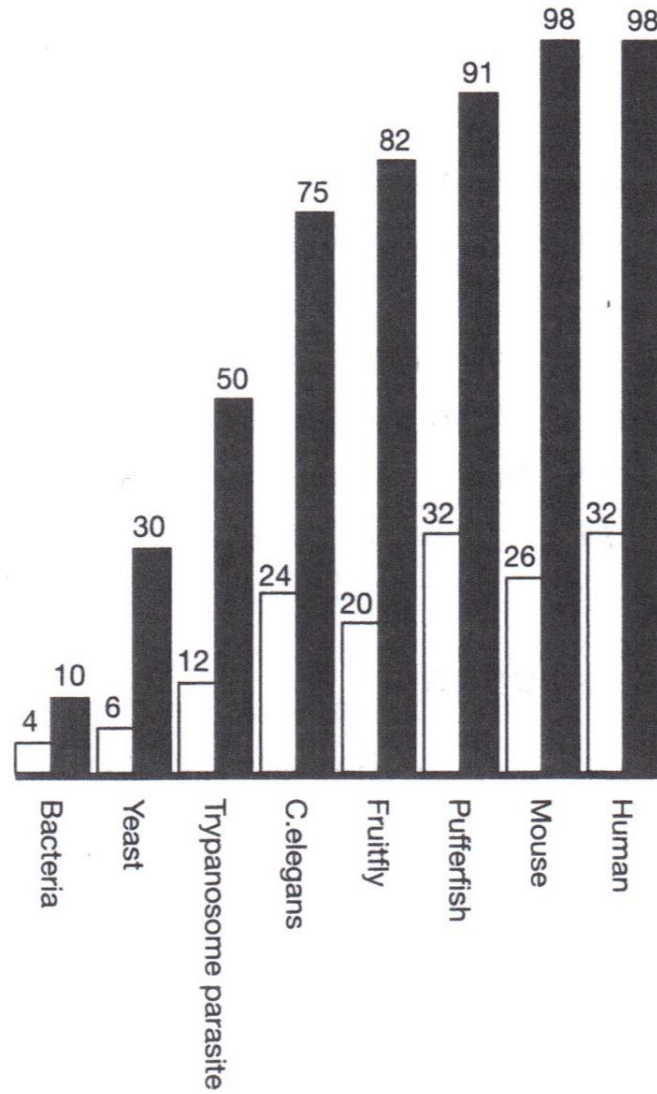
George



US Surgeon General's Report

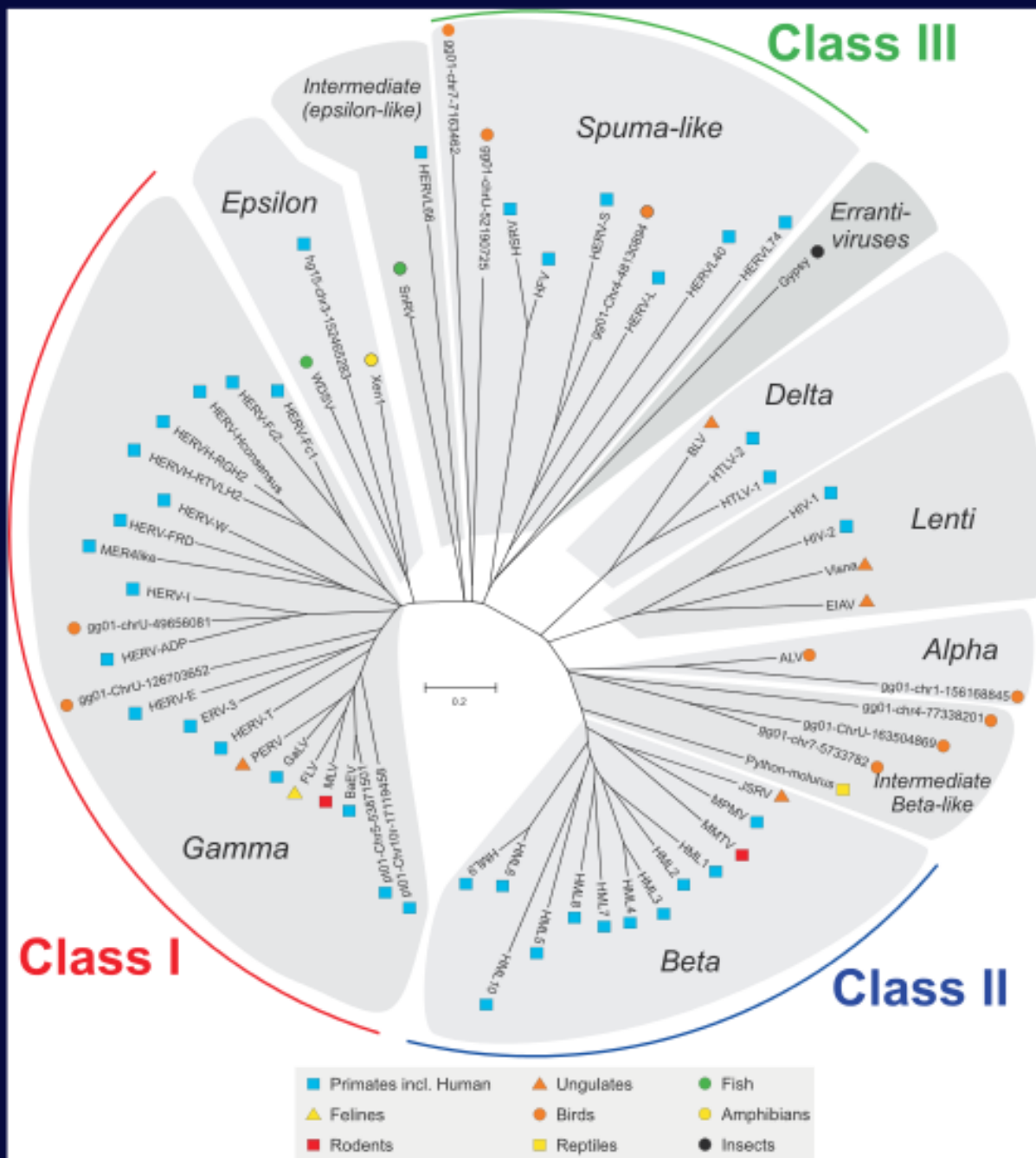
□ Amount of genome coding for protein in millions of basepairs

■ Percentage of genome that doesn't code for protein



Epigenetics of Retrotransposons (piRNAs)

- ***40-60% of genome of retroviral origin***
 - ***10% of genome Alu repeats***
 - ***10,000 HERV-K retrotransposons***
 - ***5,000 SVA retrotransposons***
-



tative unrooted Pol neighbor joining (NJ) dendrogram

Genetics vs. Epigenetics

- **Complex Systems –Stress Concepts**
 - **Evolutionary and Developmental Stressors**
 - **Stress and Evo-Devo**
-

Genotype + Environment = Phenotype/Disease Phenotype

- *Epigenetic control mechanisms evolve*
 - *There is a Lamarckian dimension in evolution*
 - *Imprints and methylation marks are erased and reestablished de novo stochastically in each generation*
-

Epigenetic Regulation of Pediatric “Endocrine”-related Genes

GRalpha, hippocampus, NGF-IP (stress, sexual abuse)

GRalpha, liver (obesity)

PPARalpha, liver (obesity)

Pdx1, islets of Langerhans (diabetes mellitus type 2)

ERalpha, hypothalamus (female behavior)

AR brain (male behavior), skin (hirsutism)

FTO (obesity) -> IRX3 homeobox transcription factor

Nanog, Oct 4 (neural stem cell differentiation)

DLK1-MEG3 (obesity, diabetes type 1)

Lxralpha (obesity, carbohydrate intolerance)

Kisspeptin –puberty

CRH- labor and delivery

FKBP5-Depression Rx

**Human endogenous retroviral element
K10 (HERV-K10)
is altered in *in vitro* handled blastocysts**

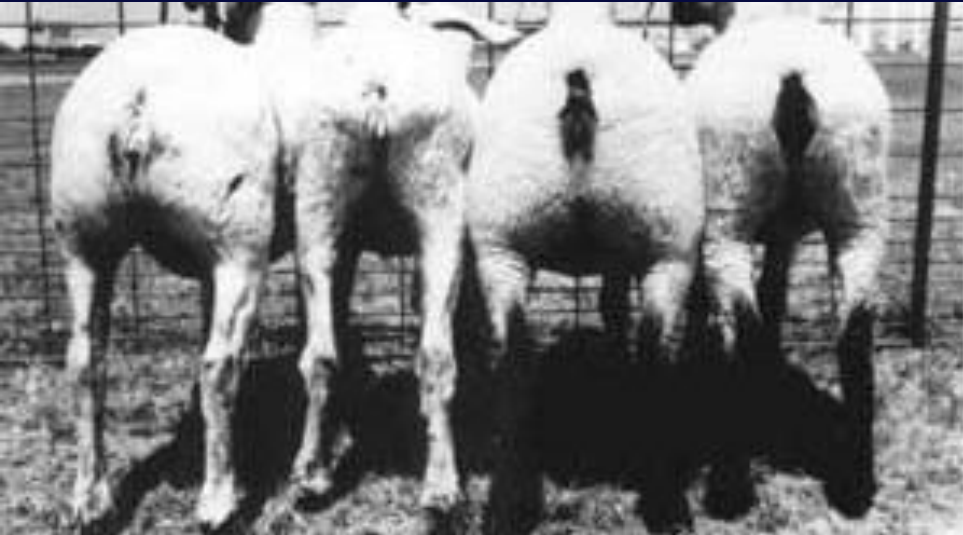


**Decreases methylation of the imprinted
DLK1-MEG3 gene region on chromosome
14q32.2.**

Dimitriadou et al., Stress 2013

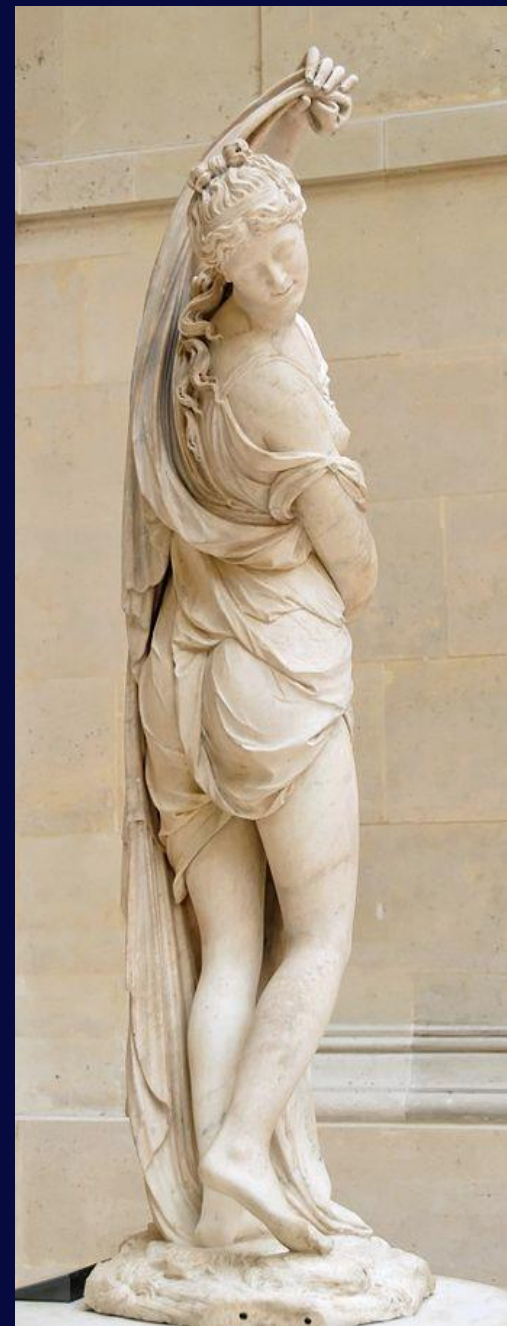
Venus Callipyge

Callipyge sheep



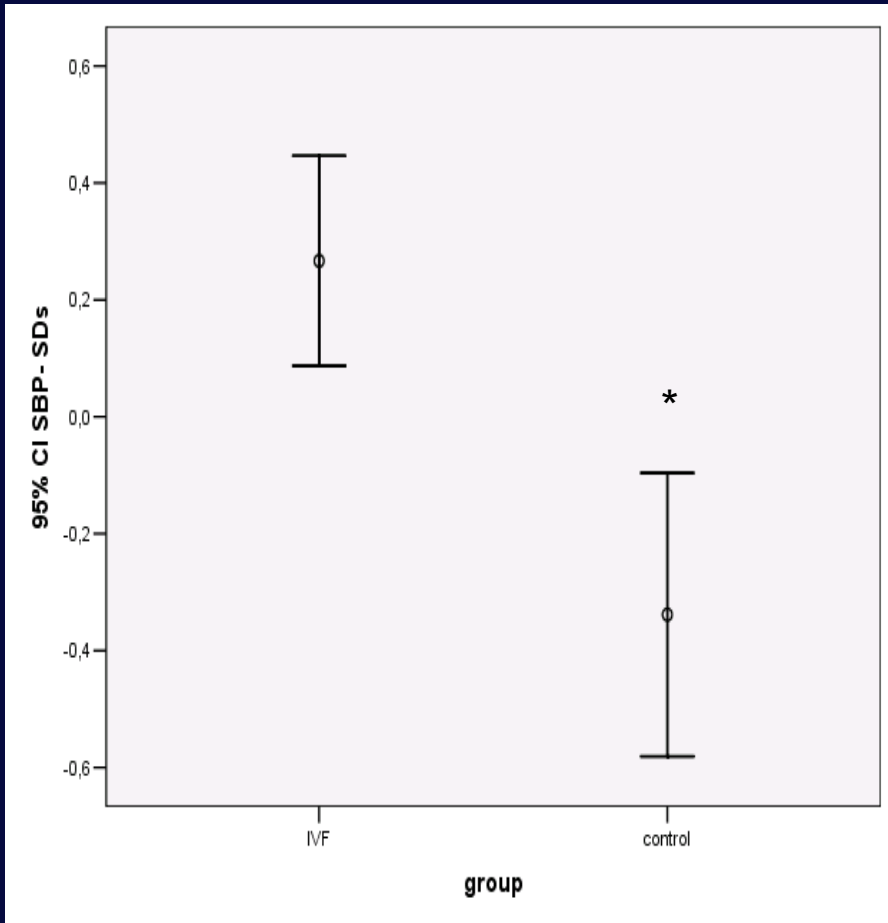
\> |™↓ || ≡LL↓□J©|S)

Callipyge

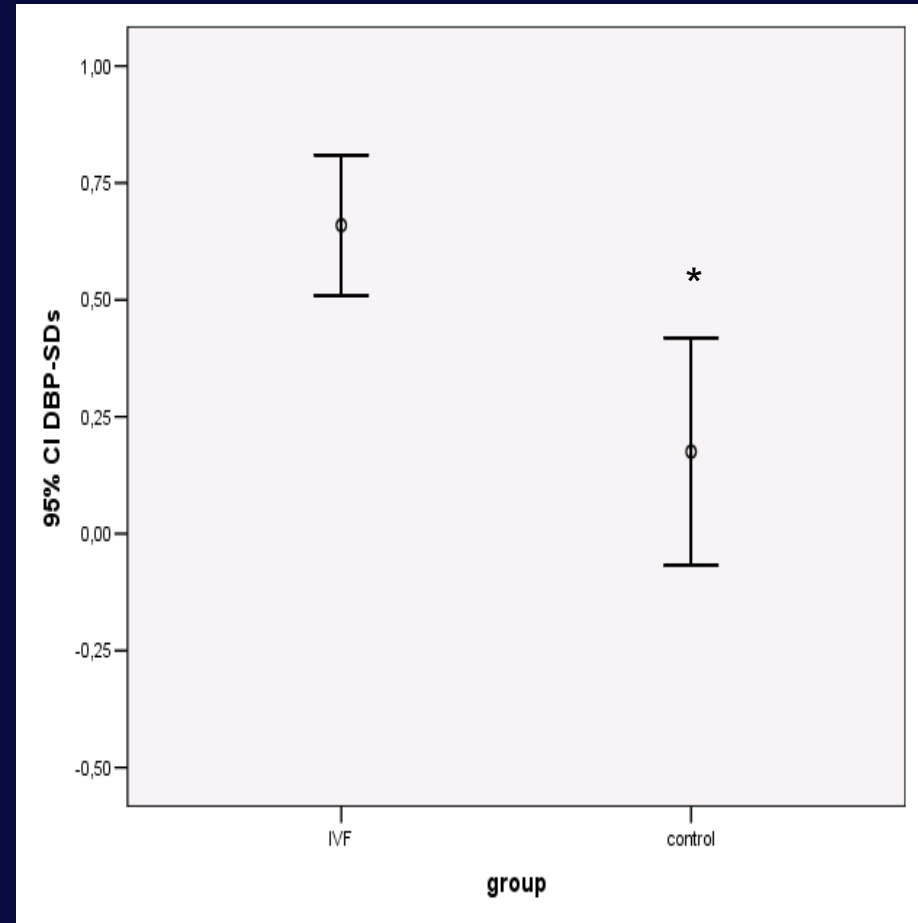




Systolic and Diastolic Blood Pressure-SDs



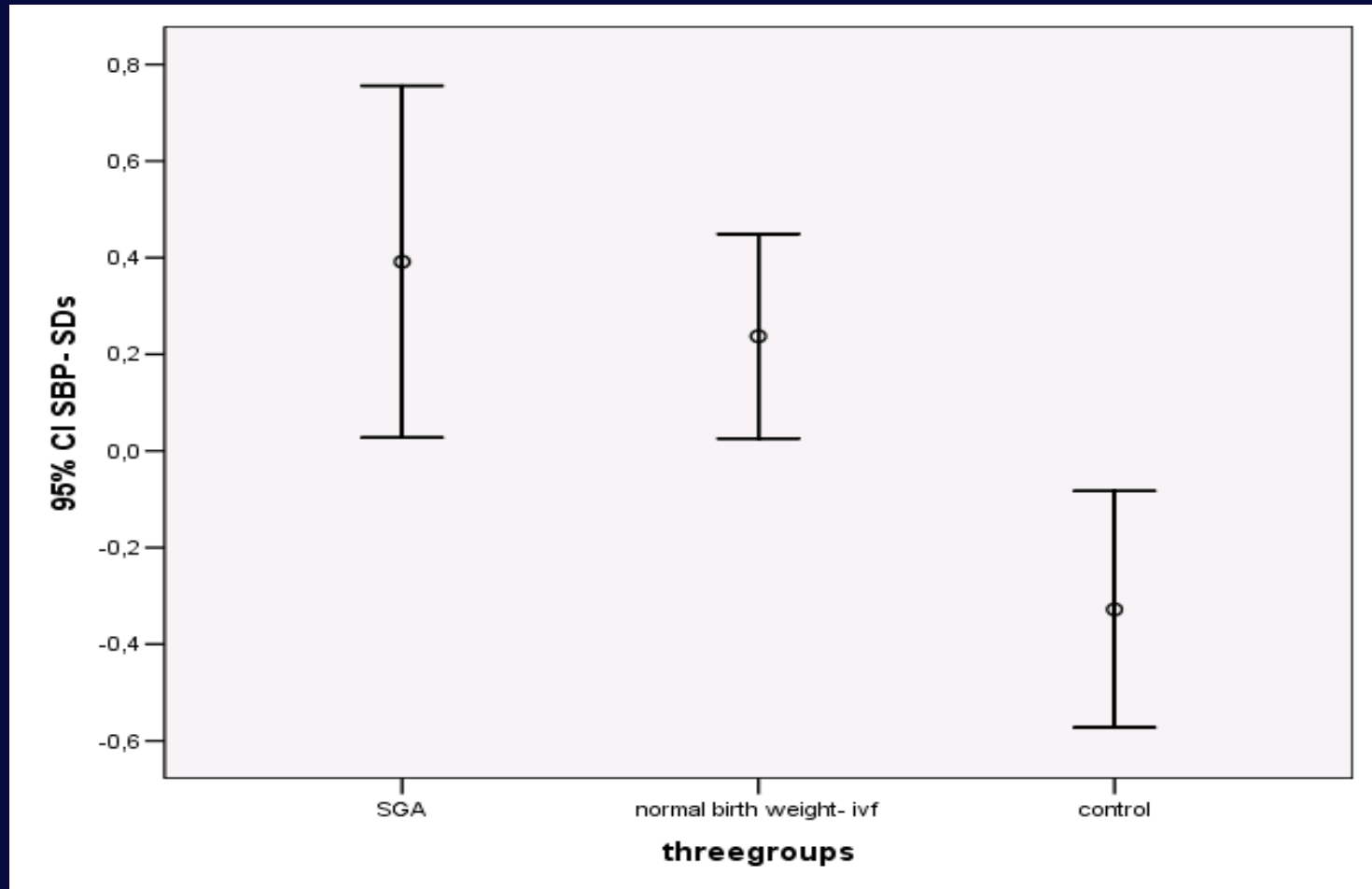
p=<0.001



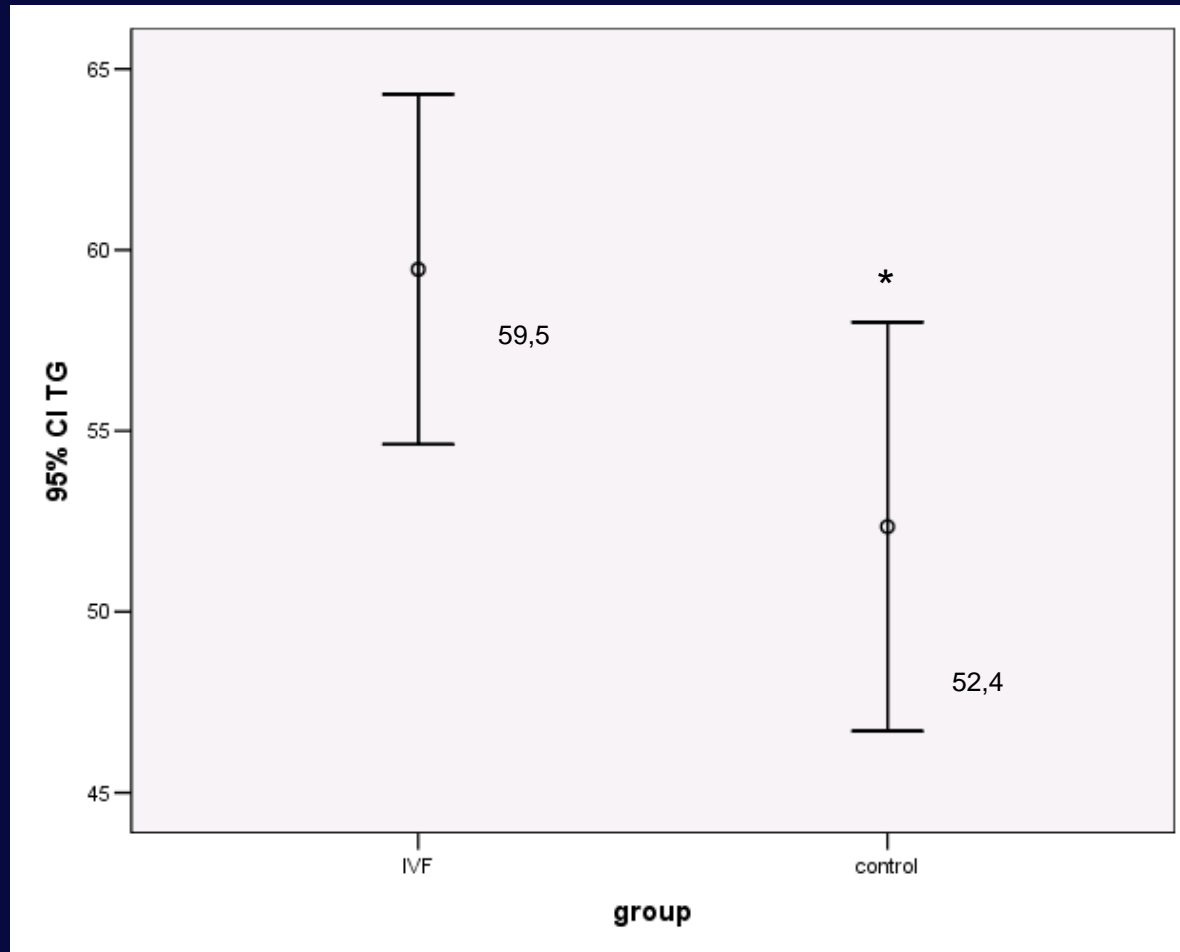
p=<0.001

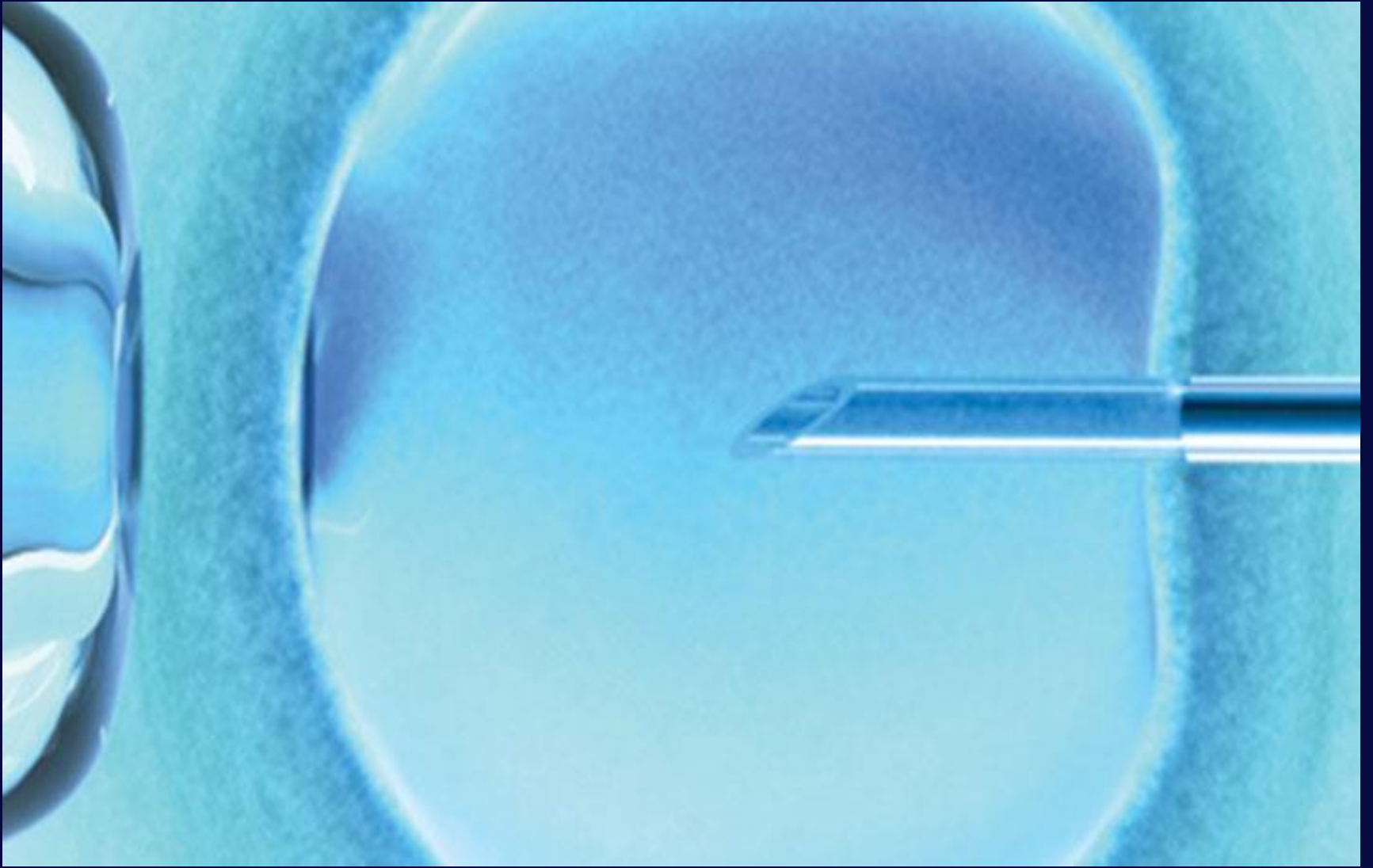
Results

Comparison between SGA-IVF, AGA-IVF and controls

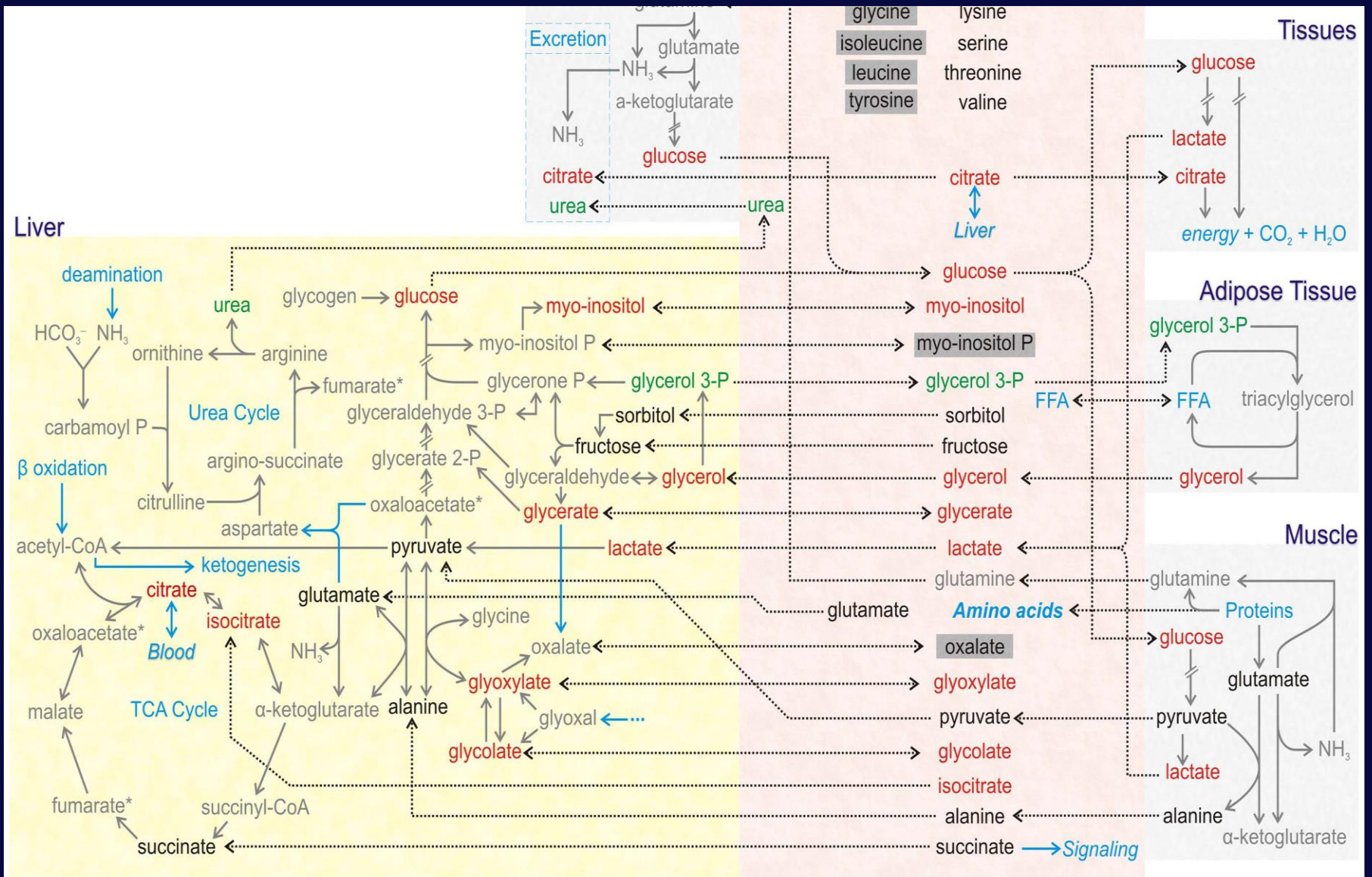


Triglycerides

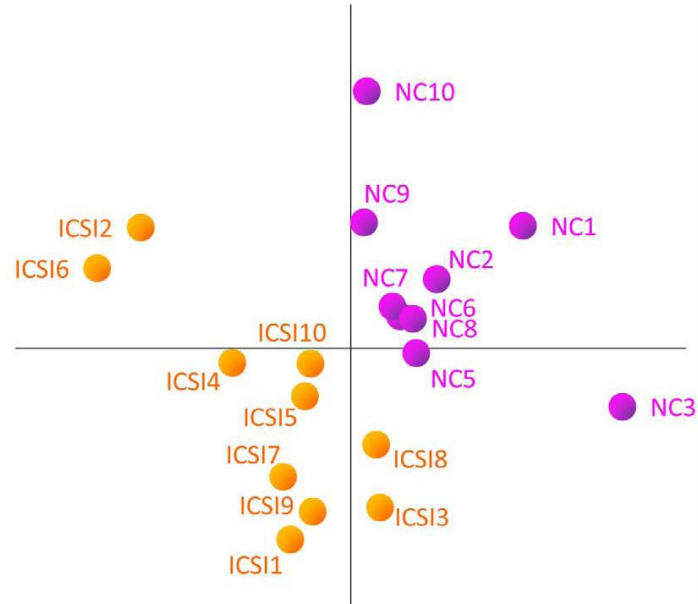




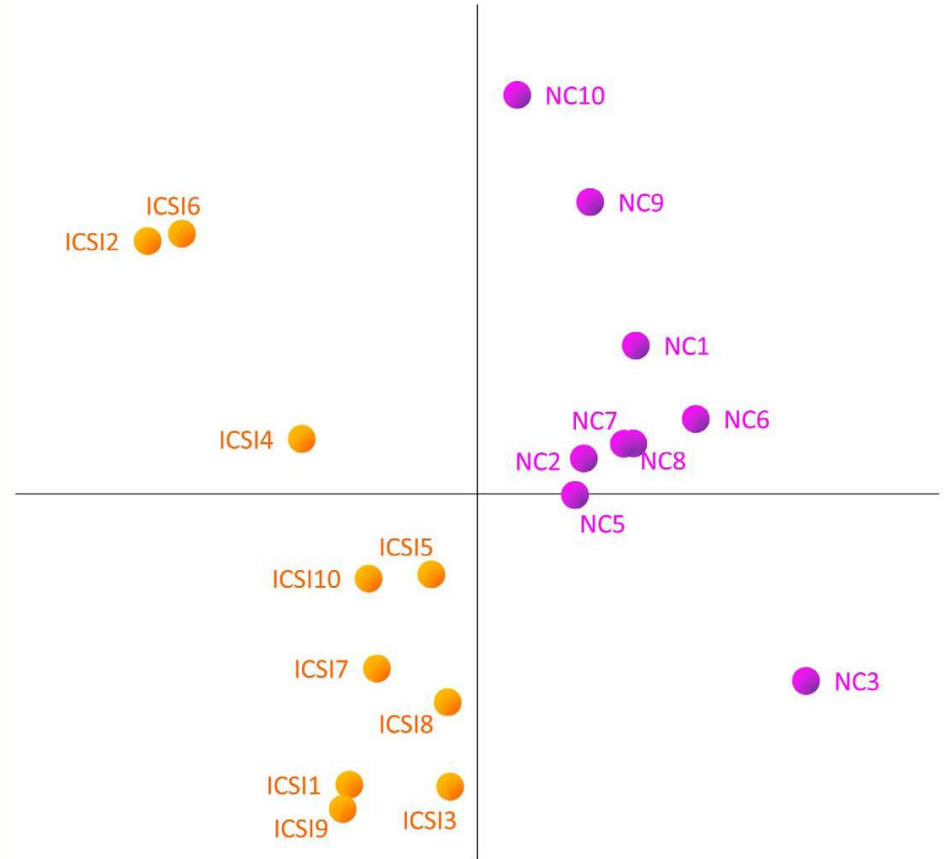
Parameters	Control N=42	ICSI N=42	Difference
Glucose (mg/dL)	83.7± 9.3	81±7.7	0.36
Insulin (mU/L)	6.9± 6.7	5.5± 2.5	0.9
HOMA index	1.5± 1.8	1.1±0.5	0.9
Total Chol (mg/dL)	172.7±24.5	167.7±25.3	0.35
Triglycerides (mg/dL)	54.2±22.6	45.4± 16.5	0.07
HDL-C (mg/dL)	60.8±12	63.9±8.9	0.17
LDL-C (mg/dL)	100.9±22	94.6± 21.2	0.18
ApoA1 (mg/dL)	156.1±19.8	153±21.1	0.5
ApoB (mg/dL)	74.2± 14.9	75.7±14.7	0.65
Lp(a) (mg/dL)	15.4± 20.3	11.8± 14.8	0.16
IGF-1 (ng/mL)	190.5±92.5	193.2± 115.3	0.58
YKL-40 (ng/mL)	27.08±15.5	15.45± 8.9	0.0002
hs-IL6 (pg/mL)	1.6± 1.5	2.3±4	0.38
hs-CRP (mg/L)	0.78± 0.87	0.44±0.3	0.022



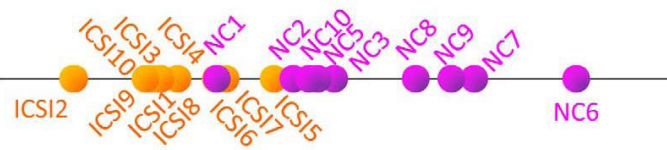
A. Mean Metabolomic Data



C. Biochemical & Mean Metabolomic Data



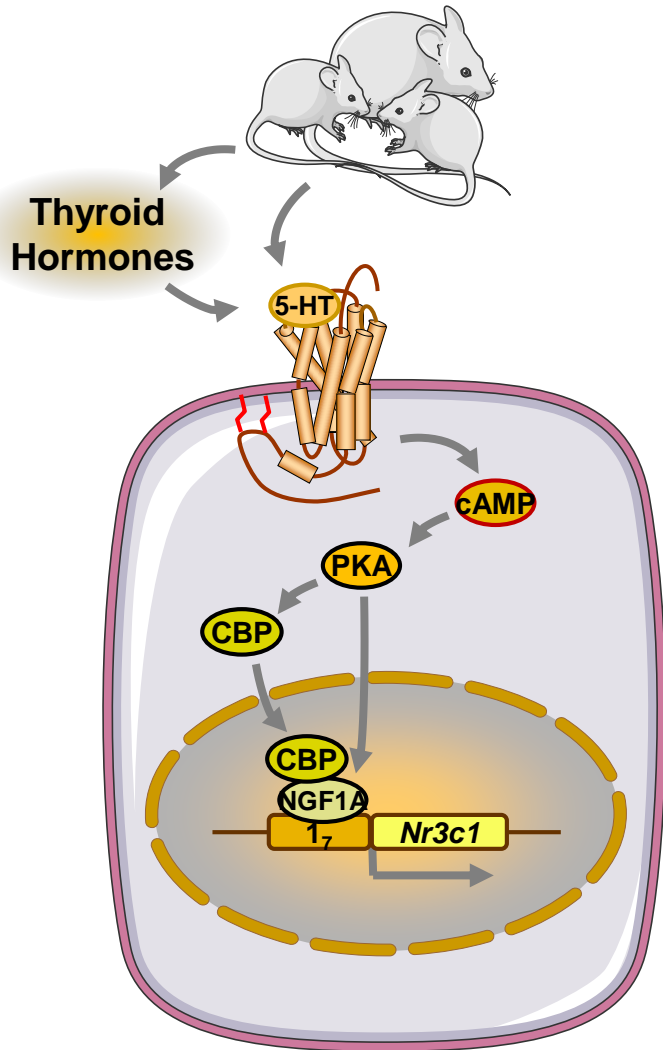
B. Biochemical Data



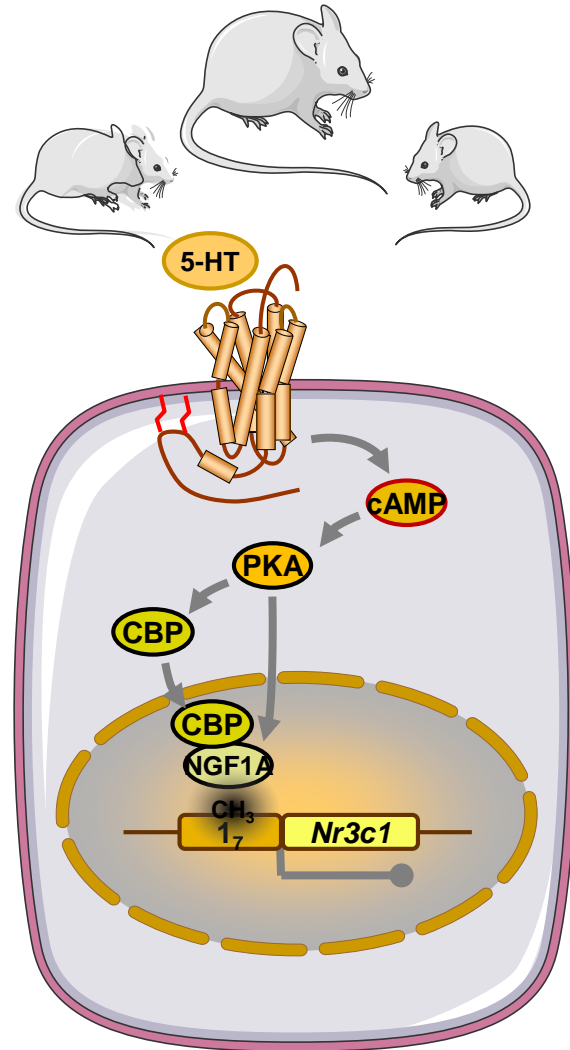


EPIGENETICS AND SEPARATION

High maternal care



Low maternal care



Champagne FA. *Behavioral Neuroscience* 2013; 127(5): 628–636

EPIGENETICS AND CHILDHOOD ABUSE \Rightarrow

Epigenetic regulation of the glucocorticoid receptor in human brain associates with childhood abuse

Patrick O McGowan^{1,2}, Aya Sasaki^{1,2}, Ana C D'Alessio³, Sergiy Dymov³, Benoit Labonté^{1,4}, Moshe Szyf^{2,3}, Gustavo Turecki^{1,4} & Michael J Meaney^{1,2,5}



McGoan PO et al. *Nature Neuroscience* 2009; 12(3): 342-348

EPIGENETICS AND CHILDHOOD ABUSE (33)

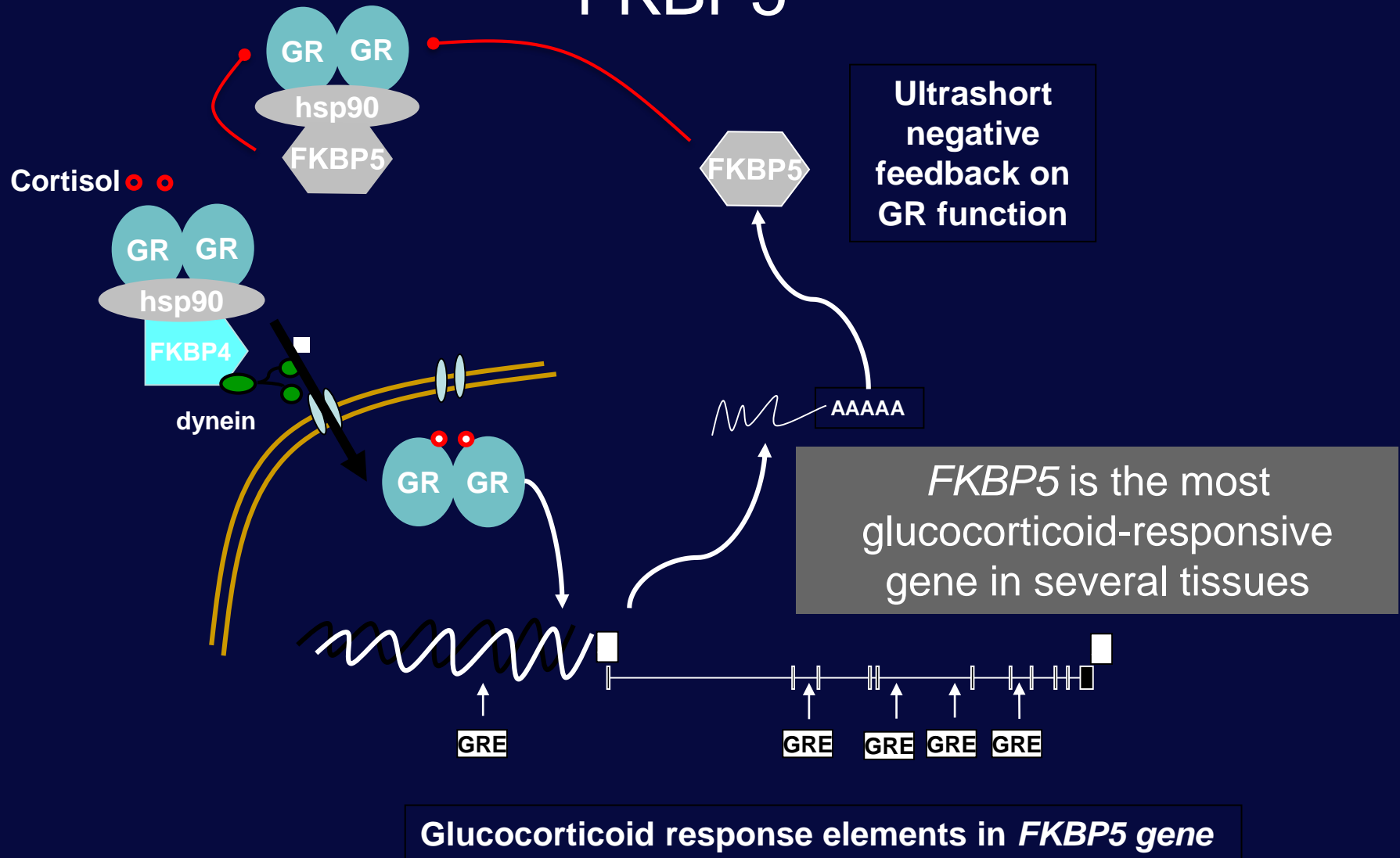
Increased methylation of glucocorticoid receptor gene (*NR3C1*) in adults with a history of childhood maltreatment: a link with the severity and type of trauma

N Perroud¹, A Paoloni-Giacobino², P Prada³, E Olié^{4,5,6}, A Salzmann¹, R Nicastro³, S Guillaume^{4,5,6}, D Mouthon², C Stouder², K Dieben³, P Huguelet³, P Courtet^{4,5,6} and A Malafosse^{1,2}

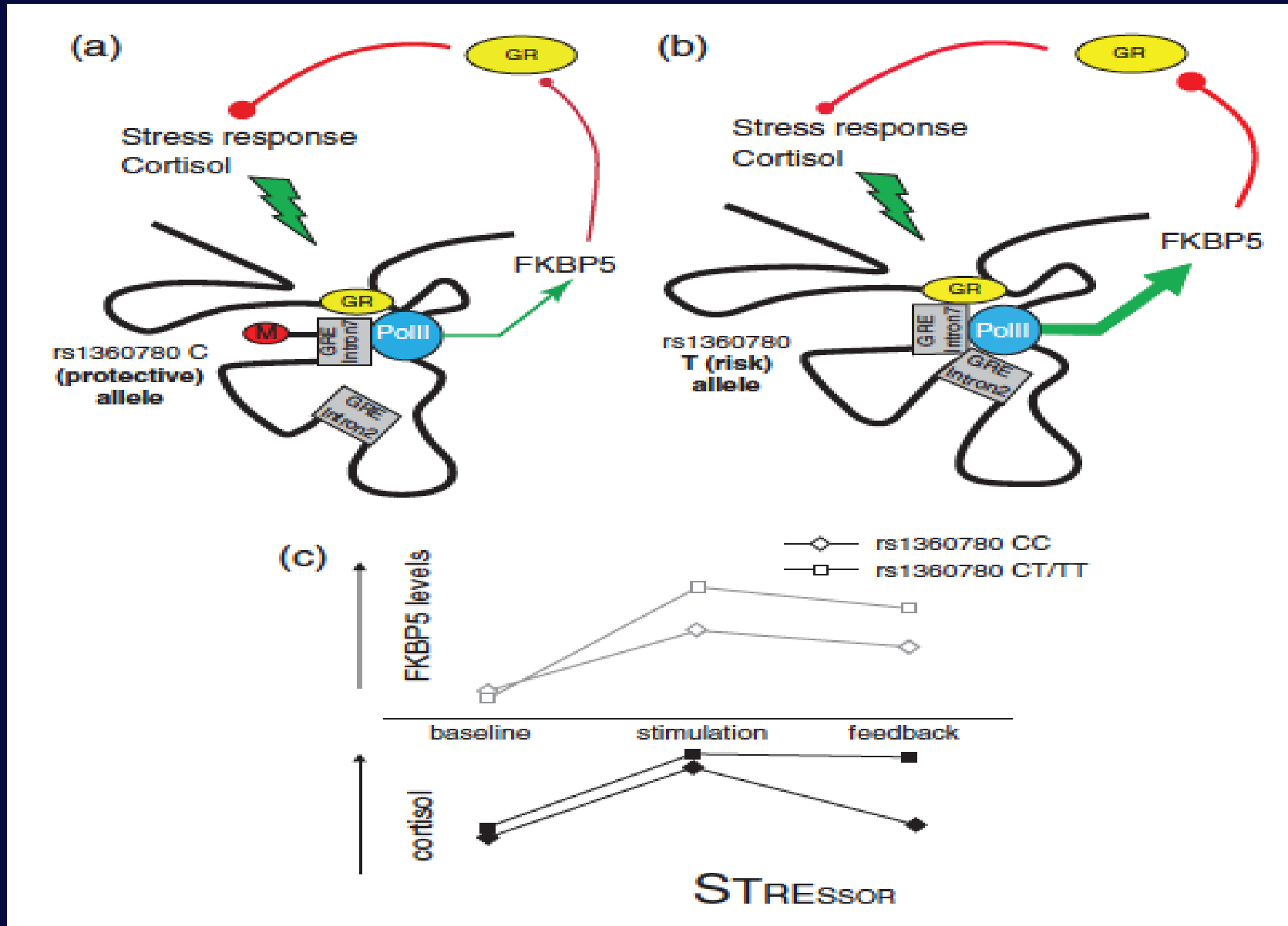


Perroud N et al. *Transl Psychiatry* 2011; (1): e59.

Molecular modulators of the stress response: FKBP5



FKBP5 disinhibition by gene-environment-epigenetic interactions



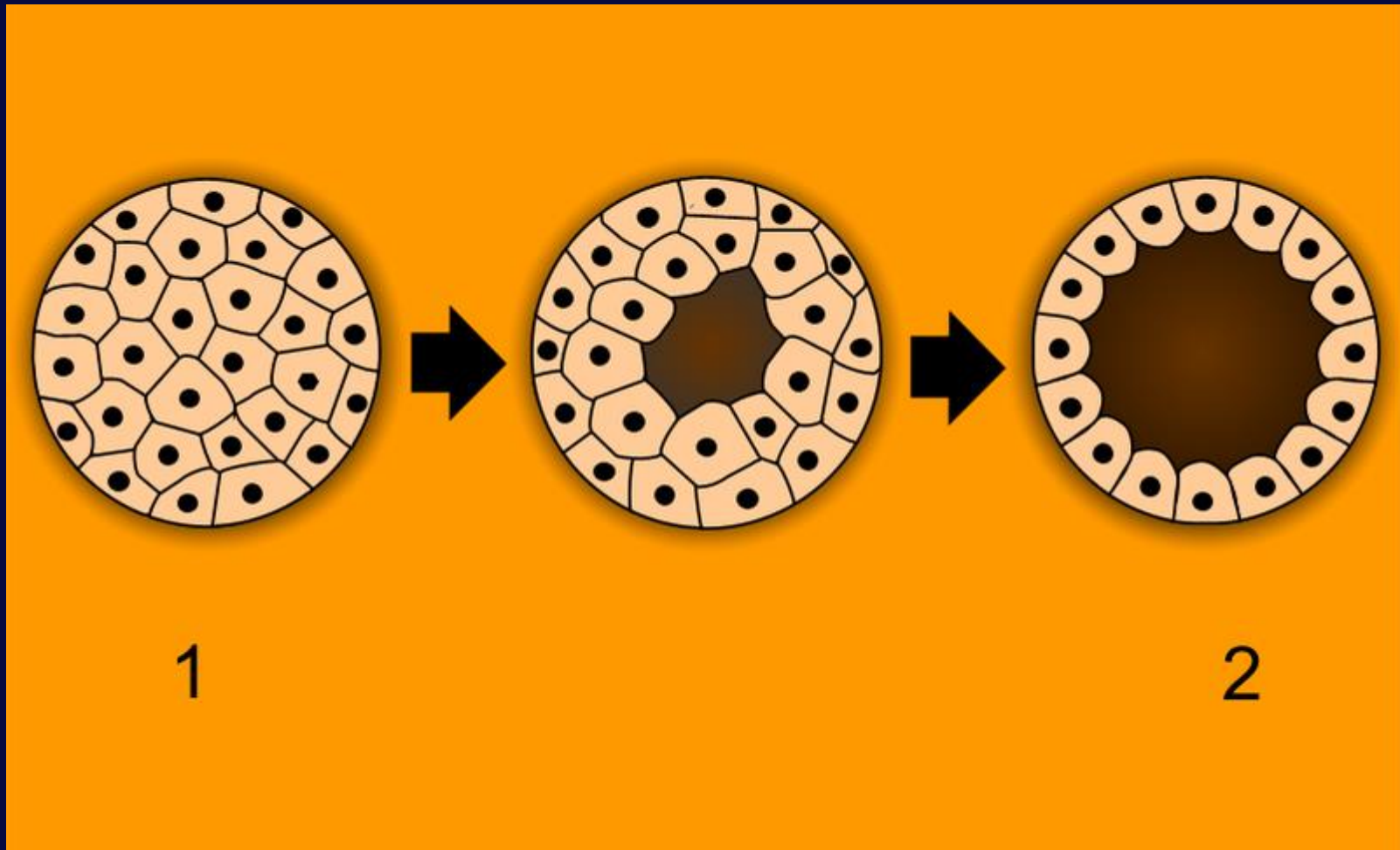
Epigenome

- ***CpG islands in 60% of gene promoters***
 - ***25% of methylation in stem cells is in non-CpG context***
 - ***Global demethylation of the trophoblast***
 - ***Global demethylation in aging***
 - ***Global demethylation in cancer***
-

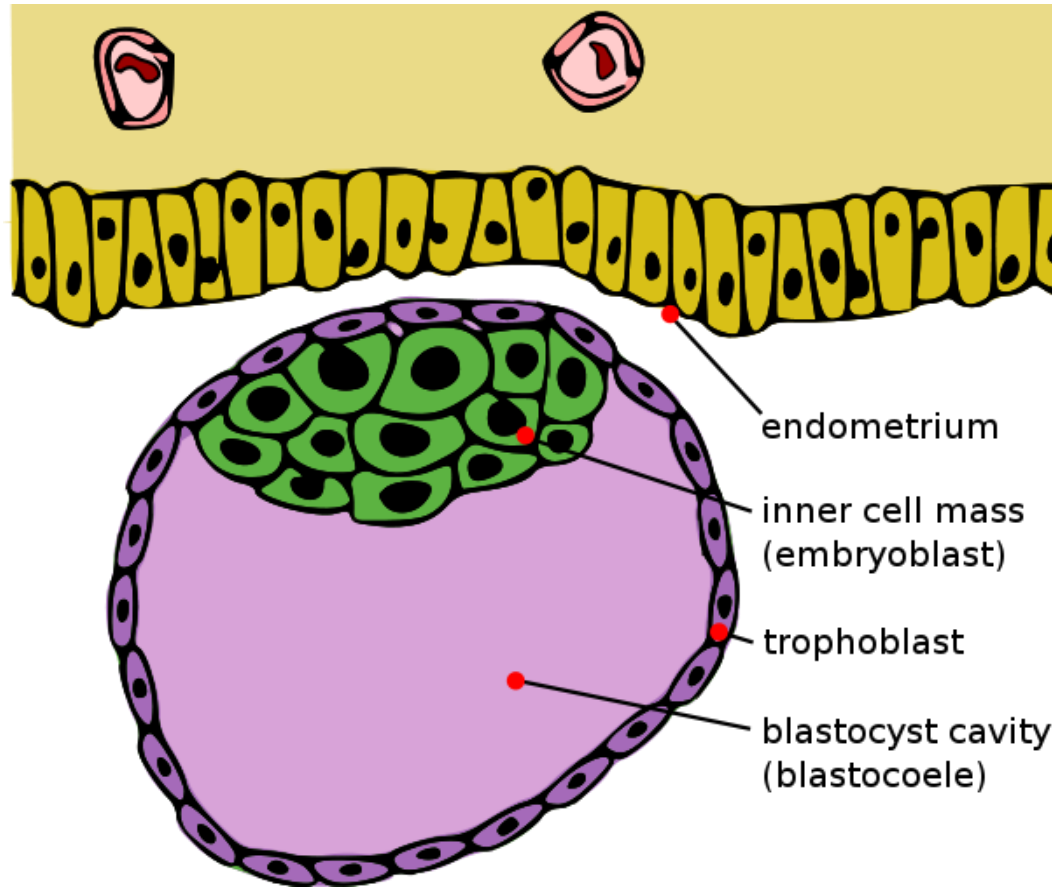
Epigenome

- ***Preimplantation: Methylome erasure***
Paternal first, maternal ensues
 - ***Remethylation: Morula-Blastocyst***
 - ***Inner cell mass: hypermethylation***
 - ***Trophoblast: hypomethylation***
 - ***Gametogenesis: epigenetic reprogramming/ erasure-remethylation***
-

Blastulation: 1 - morula, 2 - blastula.



- Oocyte to 2 cells 32 genes
- Two to 4 cells 129 genes



Blastocyst





ENVIRONMENTAL STRESSORS

Starvation
Dehydration
Injurious agents-inflammations
Adversaries-anticipation
Adversaries-avoidance
Injury-minimization

Maternal Stress,
Embryogenesis

Species

vs.

Individual

Evolution

Genetics

CNS complexity

Development

Epigenetics

CNS plasticity

Genotype

Epigenotype

Phenotype

The Economist

DECEMBER 13TH-19TH 2003

www.economist.com

Gore anoints Dean

PAGES 12 AND 33

America's Taiwan test

PAGES 12 AND 29

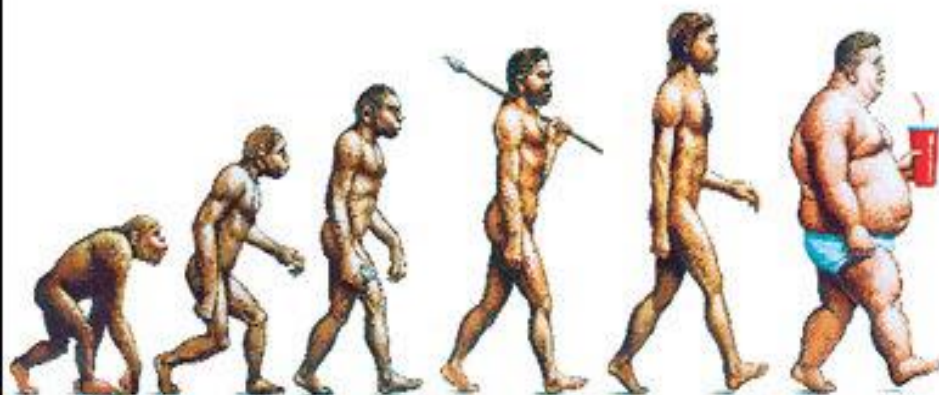
The future of flight

PAGES 79-81

A SURVEY OF FOOD

AFTER PAGE 52

The shape of things to come



US\$4.95 • C\$6.95



Argentina \$4.50
Australia \$4.95
Brazil \$5.95

Canada \$5.95
Cuba \$12.00
France \$5.95
Germany \$5.95

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Korea \$5.95
Mexico \$5.95

Netherlands \$5.95
New Zealand \$5.95
Norway \$5.95
Pakistan \$5.95
Singapore \$5.95
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Spain \$5.95
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Switzerland \$5.95
Taiwan \$5.95
Thailand \$5.95
UK \$5.95
USA \$4.95
Vietnam \$5.95

Selections of Gene and Epigenetic Networks Participating in Functions Important for Human Survival and Species Preservation

RESPONSE TO SURVIVAL THREAT

SELECTIVE ADVANTAGE

CONTEMPORARY DISEASE

Combat starvation

Energy conservation

Obesity/metabolic syndrome

Combat dehydration

Fluid and electrolyte conserve

Hypertension

Combat injurious agents

Potent immune reaction

Autoimmunity/Allergy

Anticipate adversaries

Arousal/fear

Anxiety/insomnia

Minimize exposure to danger

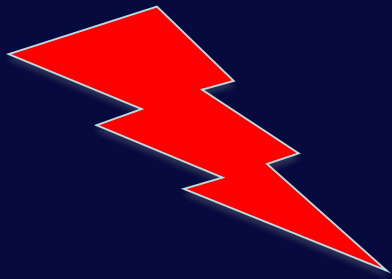
Withdrawal

Depression

Prevent tissue strain/damage

Retain tissue integrity

Pain and fatigue syndromes



Fetal Life

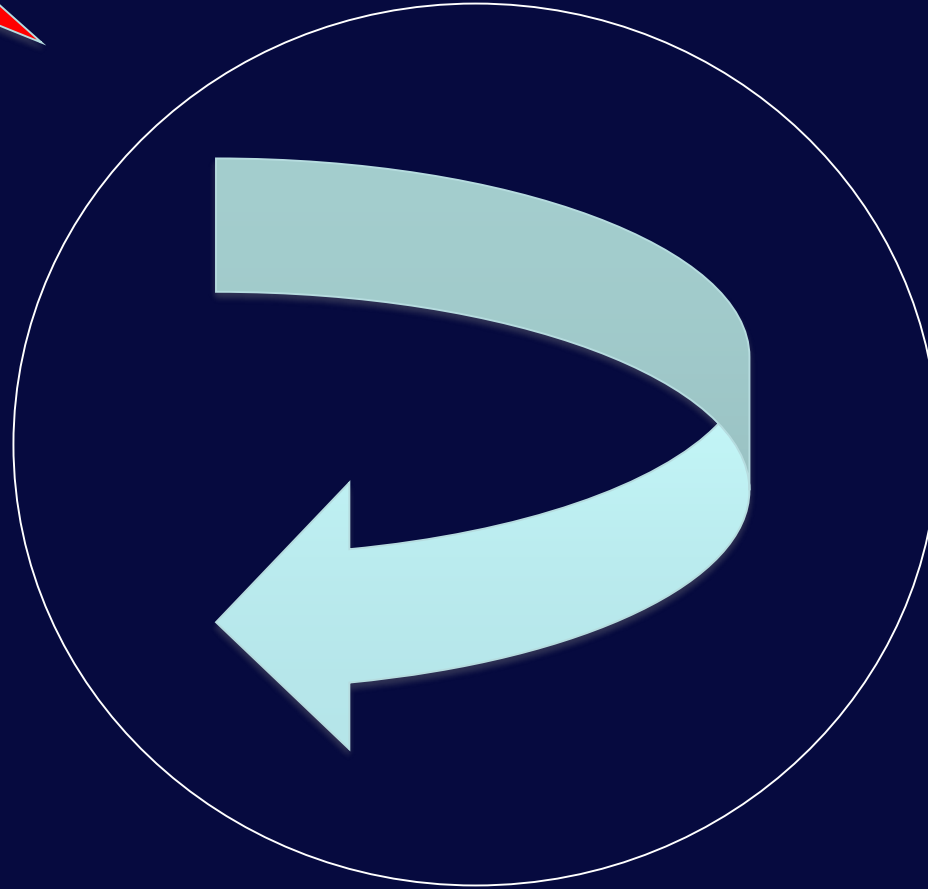
Conception

Childhood

Fertility

Puberty

Adulthood





1
 ΕΠΙΚΤΗΤΟΥ
 ΕΓΧΕΙΡΙΔΙΟΝ.
 ΕΡΙΣΤΕΤΙ
 ENCHIRIDION.

ΚΕΦ. α.
CAP. I.

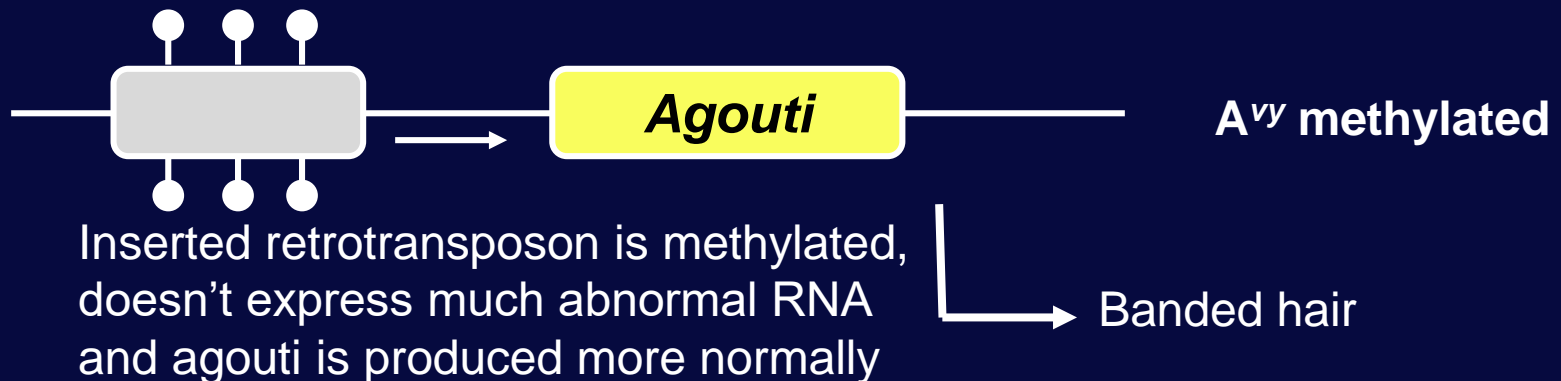
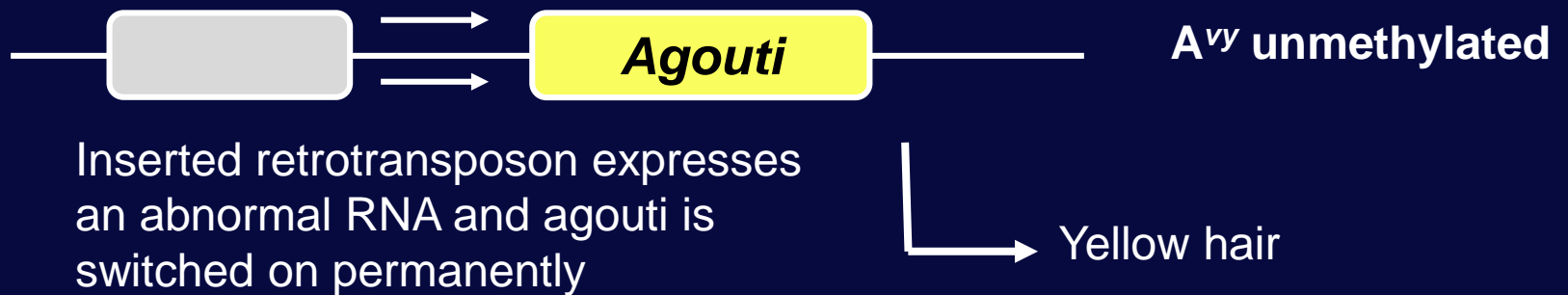
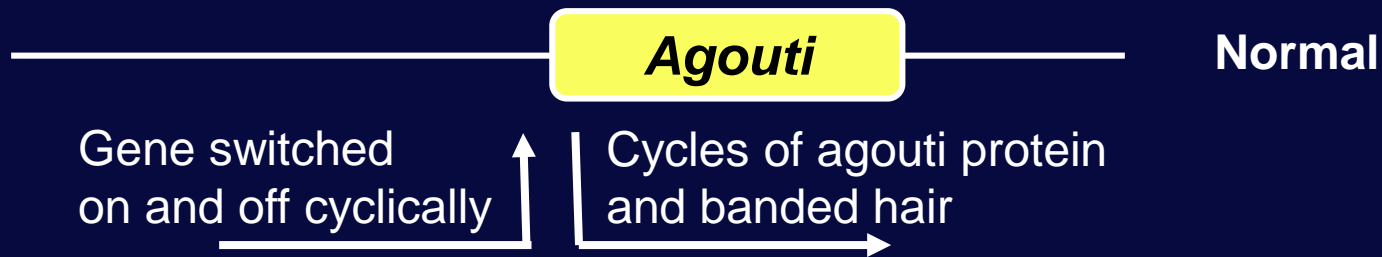
<p> Πῶν ὄντων πᾶ- μὲν ἐσὶν ἐφ' ἡ- μῖν, τὰ δὲ σὺκ- ἐφ' ἡμῖν. * ἐφ' ἡμῖν μὲν ὑποβολή, ὄρ- μη, ὄρεξις, ἐκκλησις, κῆ- ἐνὶ λόγῳ, ὅσα ἡμέτερα ἔργα. ὅσα ἐφ' ἡμῖν δεῖ, τὸ σῶμα, * ἡ κτῆσις, δόξα, δόχαι. ἢ ἐνὶ λόγῳ, ὅσα εἰς ἡμέτερα ἔργα. </p>	<p> Res quaedam in potestate no- stra sunt, quae- dam non sunt. In nostra potestate est opi- nio, appetitio, desiderium, avaritio; & ut uno com- plectar verbo, quaelibet nostrae actiones. Nostri arbitrii non sunt corpus, pecunia, gloria, imperia: ad summam, ea quae ipsi non agimus, omnia. </p>
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1 Tribuitur hoc Enchiridion Epi-
 ceto, quoniam ipse id non feci-
 sset, sed Actianus, qui de uberioribus
 in id commentarium edidit, quo dis-
 putationes Epicteti plenius profequi-
 tur. Testatur id Simplicius in prefa-
 tione commentarii ad hunc libellum
 hisce verbis: τὸ βιβλίον τῶν τῶ
 Ἐπικλήτου ἐστὶν ἐπισημοῦ καὶ
 ῥῆμον, ἢ τὸ πᾶσι σοφιστικῶν ὁ
 Ἀρριανὸς, τὰ καὶ ἐπισημοῦ καὶ ἀνα-
 κείτου ἐστὶν Φιλοσοφίας, καὶ κενή-
 κείτου ἐστὶν ἡλικῶν ἐπισημοῦ καὶ
 ὡς τὸ πᾶσι λέγει. Hoc est:

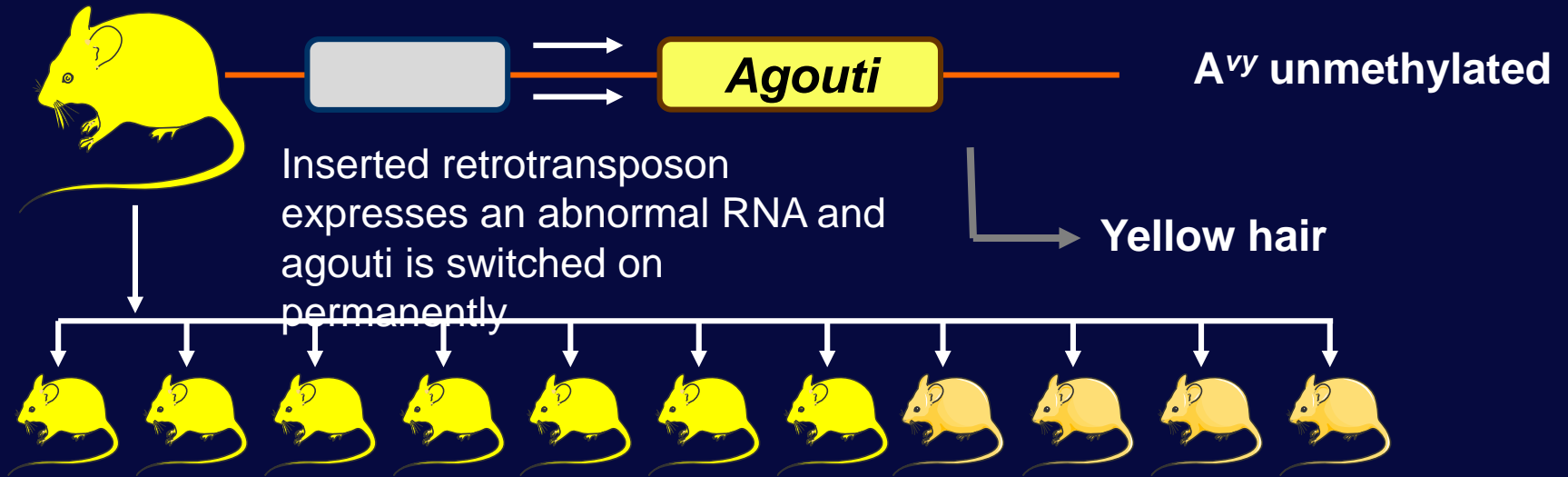
Idem Arrianus & hunc libellum, quod
 Enchiridion inscribitur composuit, de-
 ditque ex Epicteti discipulorum phi-
 losophia locis maxime idoneis ac neces-
 sariis, & animas vehementissime per-
 moventibus. Seneanus.
 2 Non solum pugio Graecis hoc
 nomine vocatur, sed etiam quid-
 quid ad manum est, & in usum
 promptum & obvium, ut instru-
 menta quaelibet ac vasa hygiea, &c.
 quae parata & in manu semper
 gestanda. Hesychius, Ἐγχειρίδιον,
 σκεῦος ἐργασίας σκεῦος, ἢ γυναικεί-
 ον.

□ ⊂ ⊄ ⊅ ⊆ ⊇ ⊈ ⊉ ⊊ ⊋ ⊌ ⊍ ⊎ ⊏ ⊐ ⊑ ⊒ ⊓ ⊔ ⊕ ⊖ ⊗ ⊘ ⊙ ⊚ ⊛ ⊜ ⊝ ⊞ ⊟ ⊠ ⊡ ⊢ ⊣ ⊤ ⊥ ⊦ ⊧ ⊨ ⊩ ⊪ ⊫ ⊬ ⊭ ⊮ ⊯ ⊰ ⊱ ⊲ ⊳ ⊴ ⊵ ⊶ ⊷ ⊸ ⊹ ⊺ ⊻ ⊼ ⊽ ⊾ ⊿ ⊿ ⊿ □

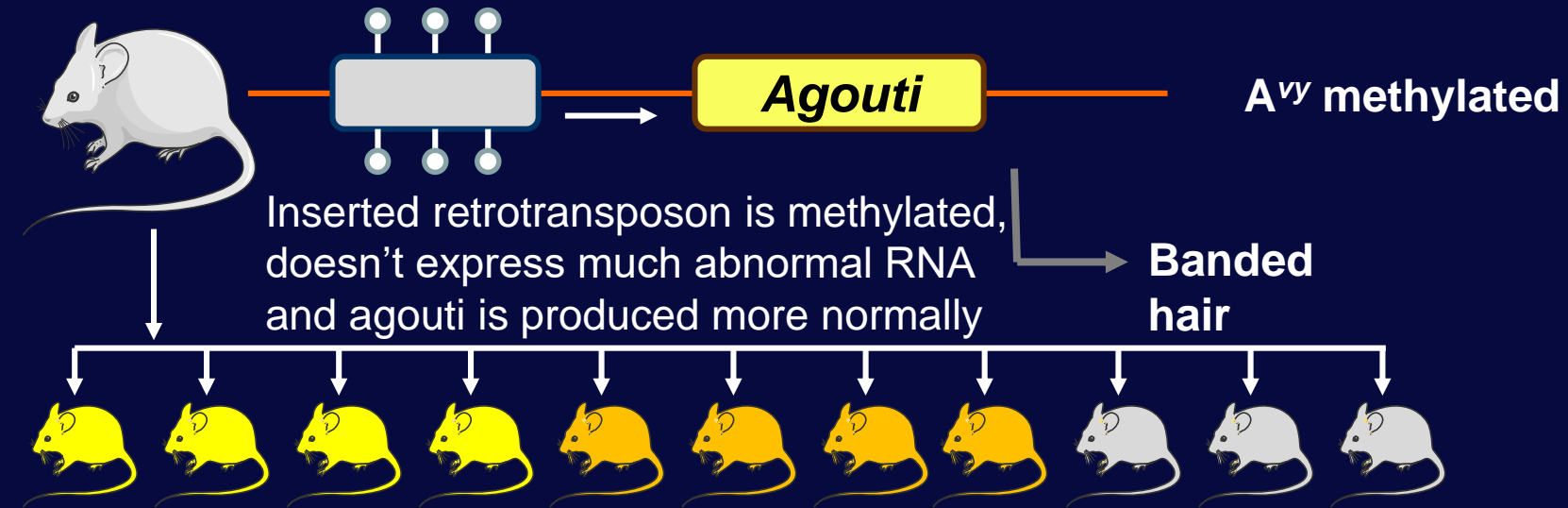
'Be equanimous and remember
 not to believe easily'

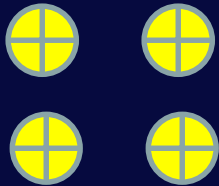
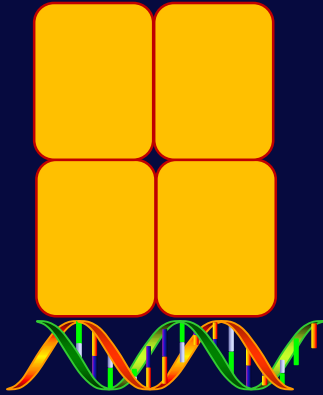


A yellow mother only gives birth to yellow or lightly colored pups

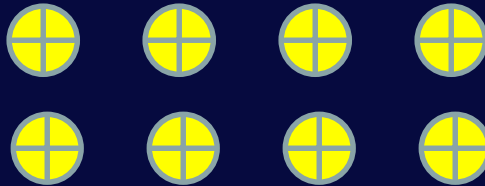


A dark mother gives birth to yellow, lightly colored and dark pups

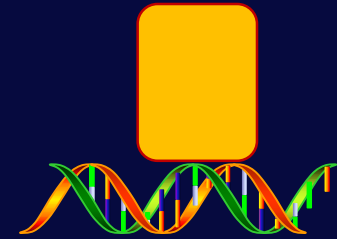




Hypothesis:
ncRNA represses
expression of
target gene

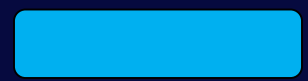


Prediction:
Decreasing levels of
ncRNA leads to
increased expression
of target gene



Actual outcome:
Decreasing levels of
ncRNA leads to
decreased expression
of target gene

Embryo and first larval stage




LIN-14 mRNA is translated into protein

Second larval stage



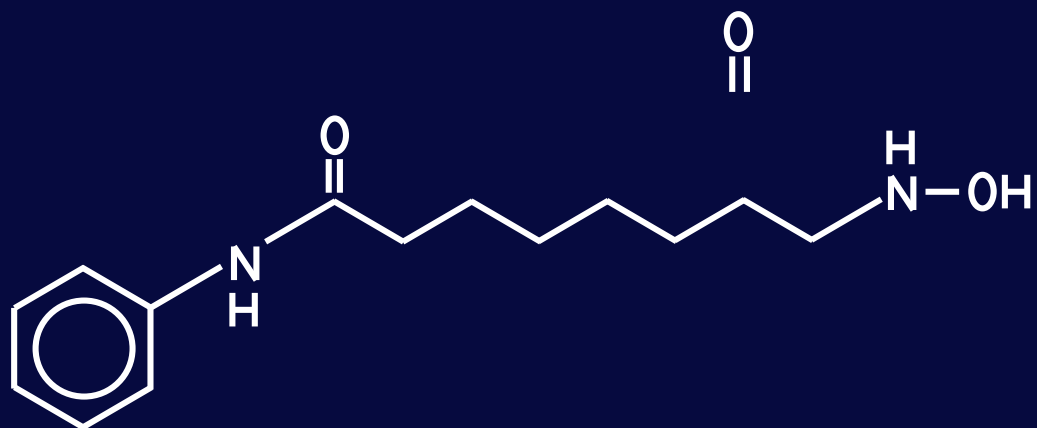
Binding of Lin-4 ncRNA to 3' UTR of LIN-14 mRNA prevents translation of protein

 Region of LIN-14 mRNA that gets translated into protein

 Region of LIN-14 mRNA that DOES NOT get translated into protein
= 3' untranslated region
= 3' UTR

 LIN-14 protein

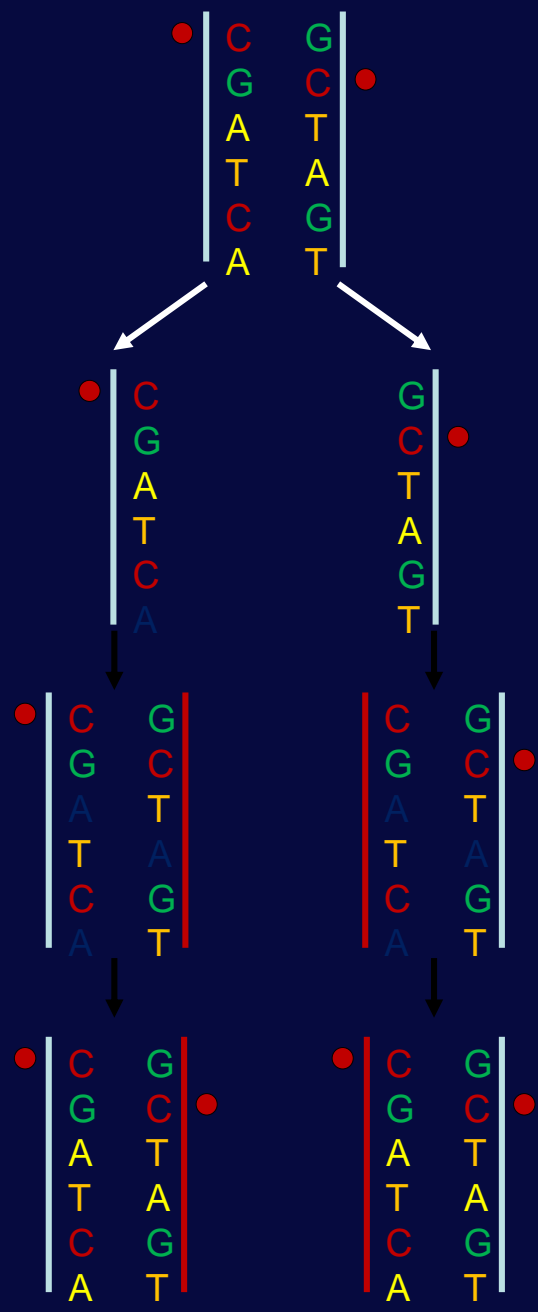
 Lin-4 non-coding RNA



SAHA

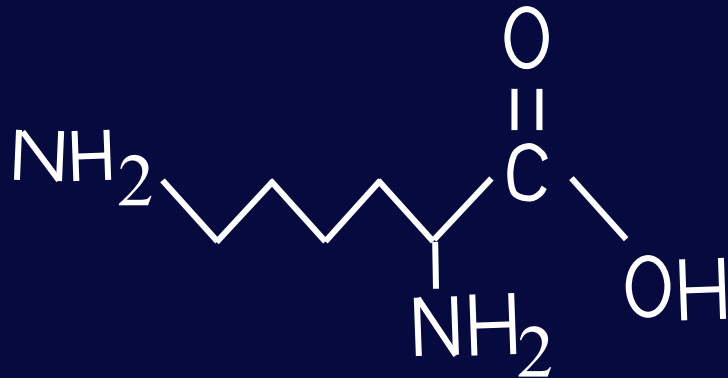


10HDA



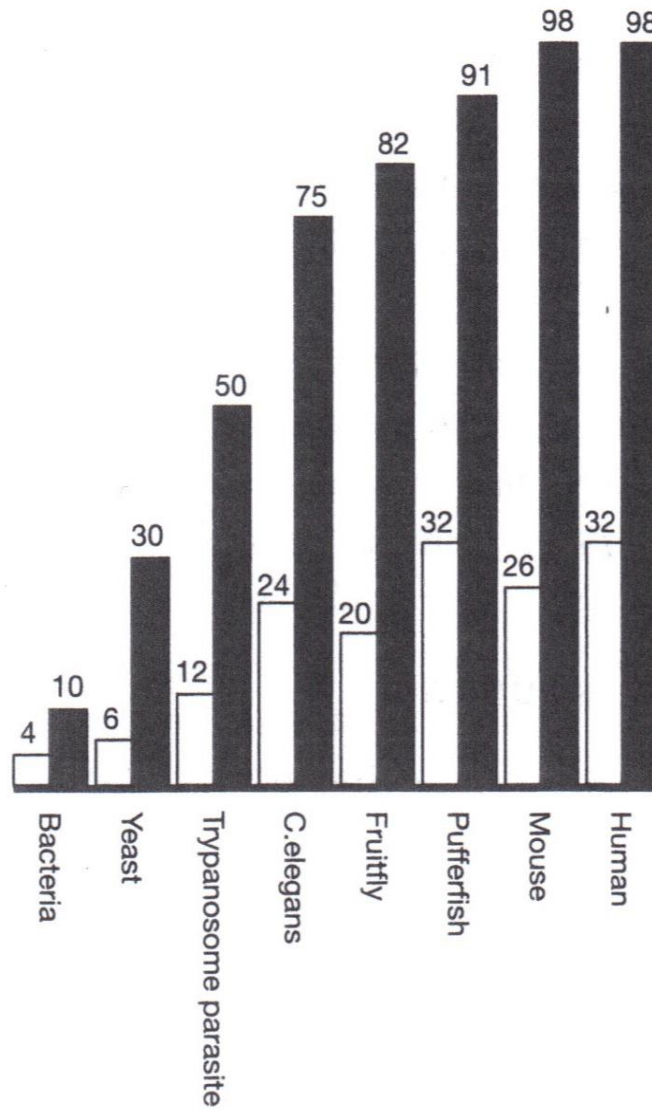


Acetyl-Lysine

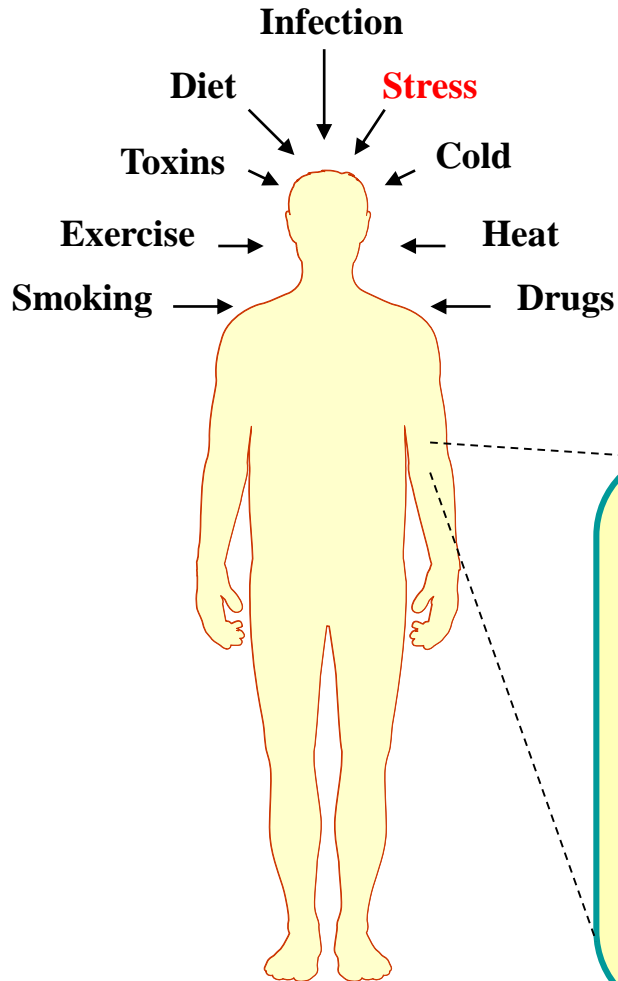


Lysine

□ Amount of genome coding for protein in millions of basepairs
■ Percentage of genome that doesn't code for protein



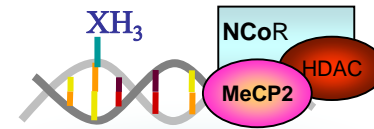
ENVIRONMENTAL STRESSORS



GENETIC MACHINERY

Binding of methyl-CpG binding proteins

Recruitment of HDACs & corepressors



DNA methylation

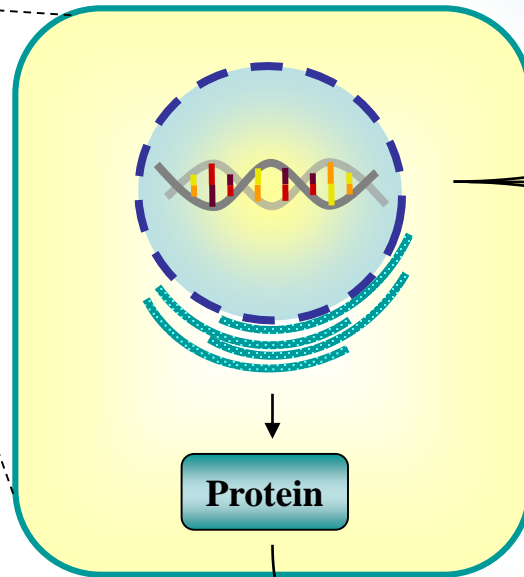


DNA variation



Histone modification

Phenotype



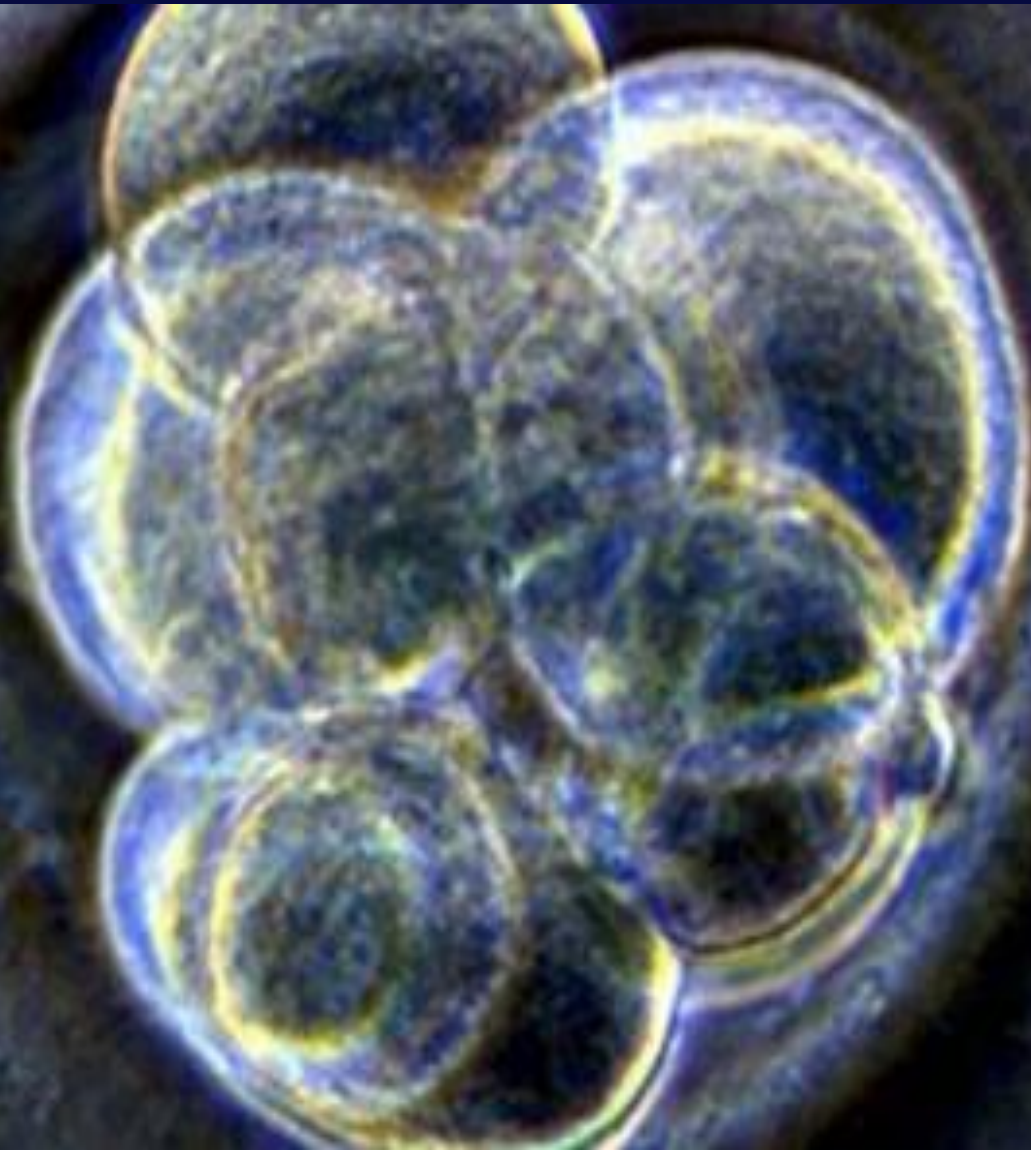
Epigenetics at the Early Stages of Childhood

Evo Devo: Genetics vs. Epigenetics ,

**George P. Chrousos, MD,
University of Athens,
Athens, Greece**

(No Disclosures)

Early Embryon



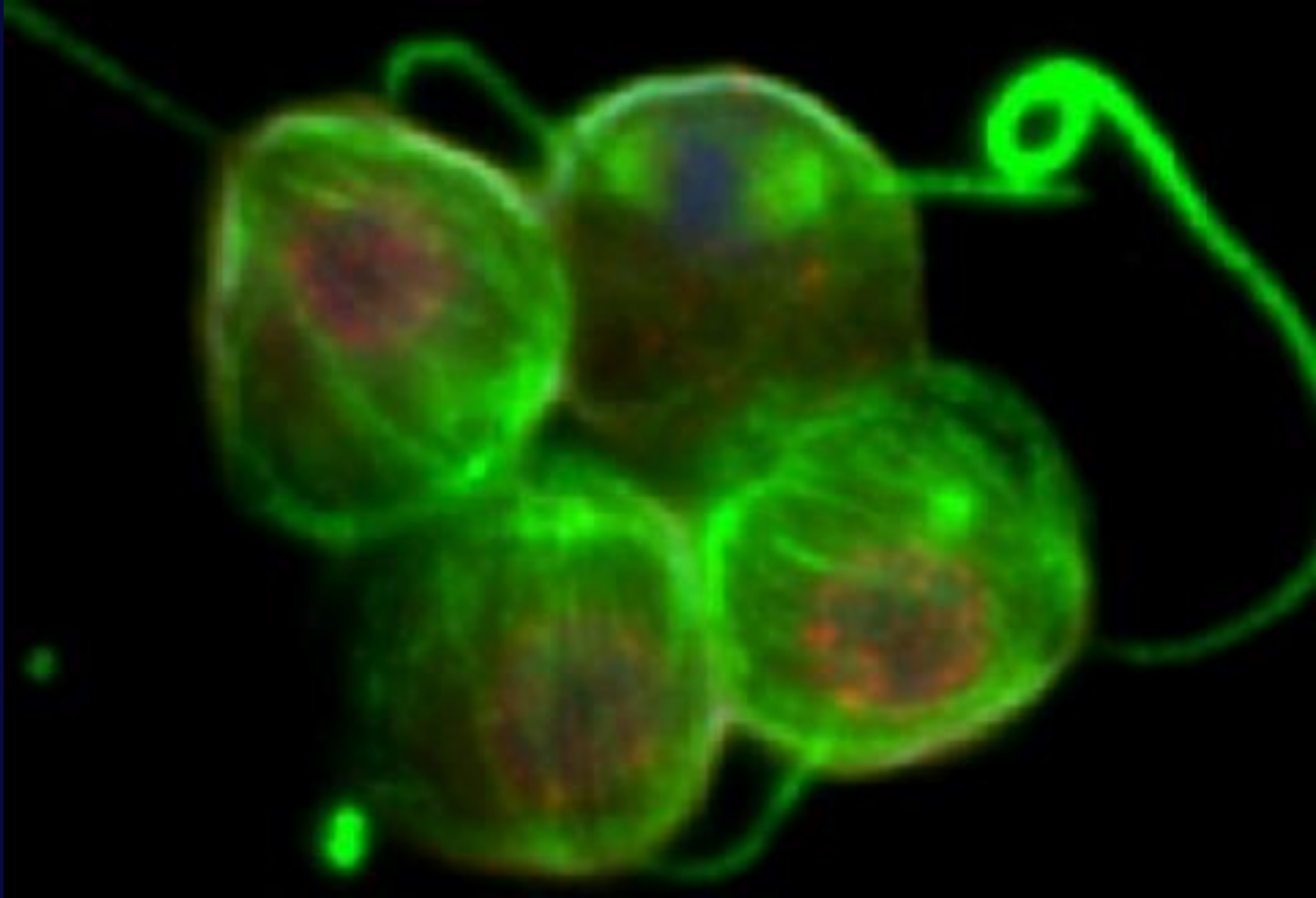


1

2

Blastulation: 1 - morula, 2 - blastula.

Choanoflagellates



ALL I REALLY
NEED TO KNOW
I LEARNED IN
KINDERGARTEN

THE CLASSIC
INTERNATIONAL BESTSELLER

25TH
ANNIVERSARY
EDITION

UNCOMMON THOUGHTS
ON COMMON THINGS

MORE THAN 7 MILLION
COPIES SOLD

ROBERT FULGHUM

Tandem Zinc Finger Proteins

