Stress and Endocrine Disruptors

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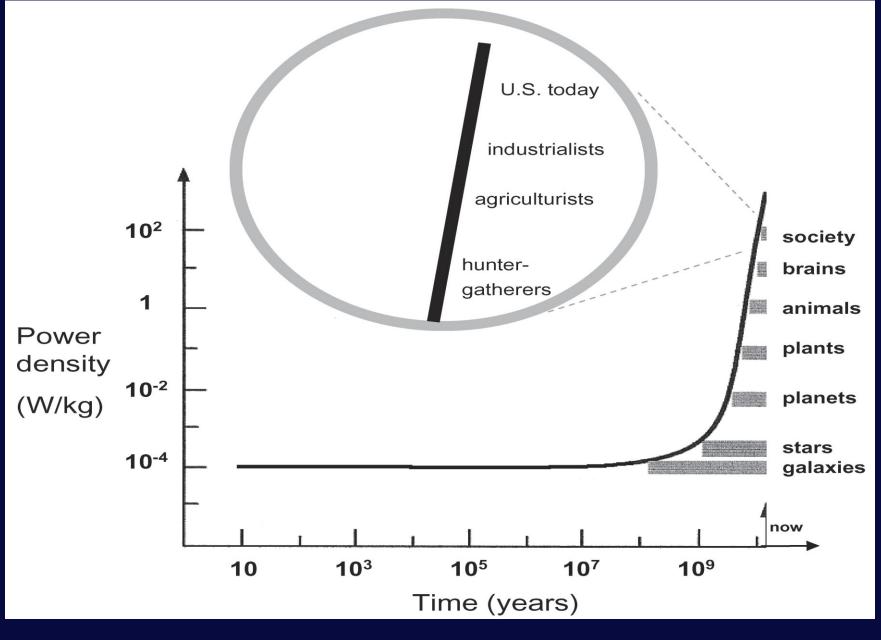
HUMAN COMPLEXITY: POST(EPI)GENOMIC ERA

Human genome:

About 3+3 billion bases ("Non-junk" DNA 98 vs. 2 %) **About 60% retroviral origin About 20 thousand protein-coding genes About 22 thousand ncRNA-coding genes About 200 thousand transcripts** (mRNA, ncRNA) **About 200-260 thousand proteins** Single nucleotide polymorphisms (snp' s or snv' s), microsatellites or copy number variants : (0.9% difference) About >25 million snp' s (snv' s), 1.5 million indels **About 20 million microsatellites** >5000 cnv's (many million bases) > 100 k disease-related mutations >1 million reg sequences >60% of promoters have CpG islands, **EPIGENETICS/EPIMUTATIONS**

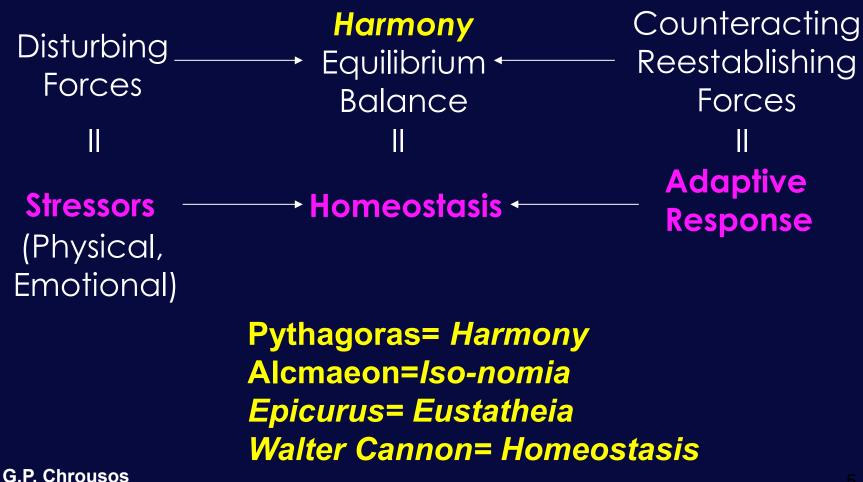
HUMAN COMPLEXITY: SOME HUMAN BRAIN NUMBERS

- ~ 100 billion neurons (100x10¹²) x >10.000 synapses per neuron = >10¹⁸ synapses)
- ~ 100.000 km of fibers
- ~ 1 trillion or more glial cells
- ~ 1.25 terabytes
- ~ 15 Watt lamp (2% of BW uses 20% energy) **Plasticity**



Chaisson E, New Scientist 2009

Complex Systems Theory



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

STRESS ETYMOLOGY

Indoeuropean root:

Gk: *Strangaleuin* = to strangle, also *Catastrophe*, and *Strabismus*

Lt: **Stringere** = to draw tight, to press

Homeostasis over Time

Improved Homeostasis=Hyperstasis

Elevated mood

Baseline Homeostasis=*Eustasis*

Constitutional mood

Deteriorated Homeostasis=Cacostasis

Suppressed mood

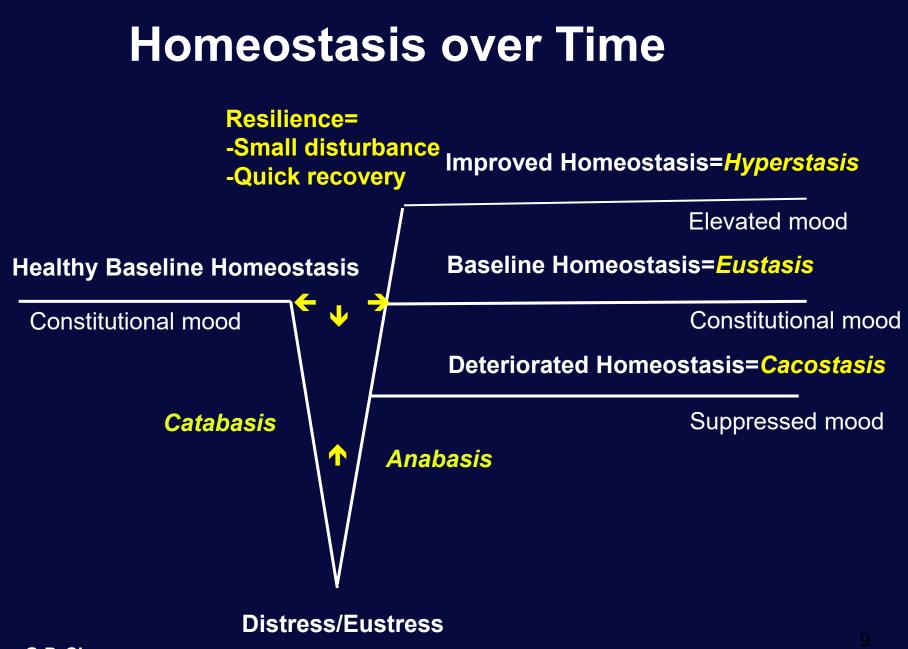
Anabasis

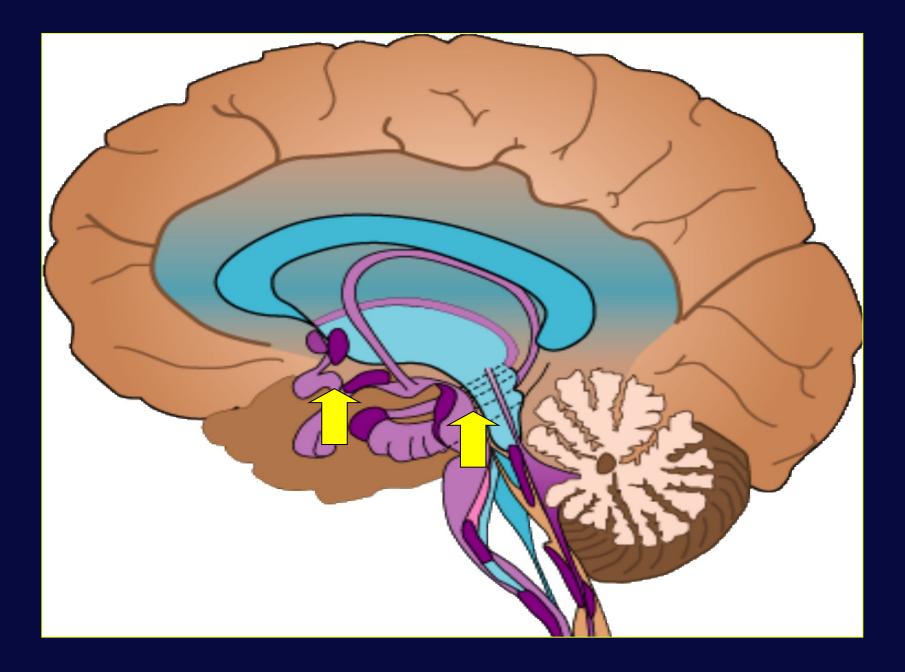
Distress (Eustress)

Healthy Baseline Homeostasis

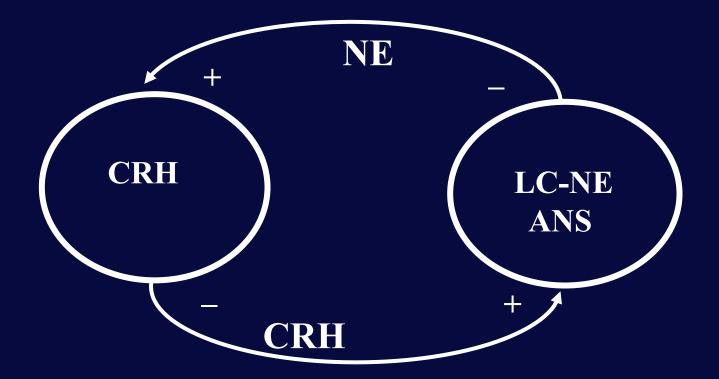
Catabasis

Constitutional mood

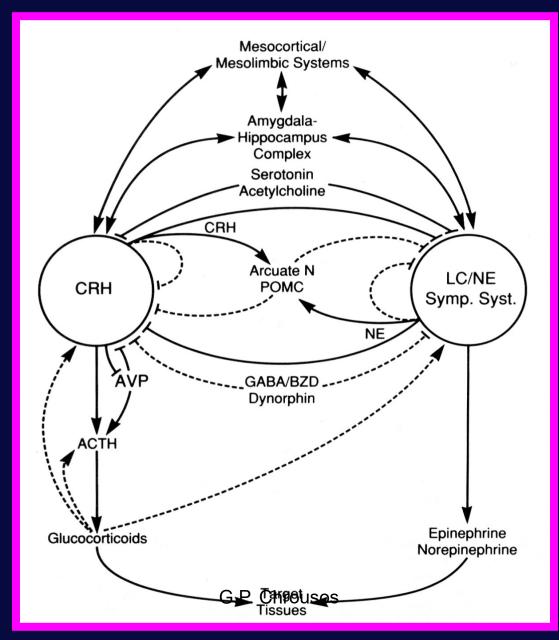




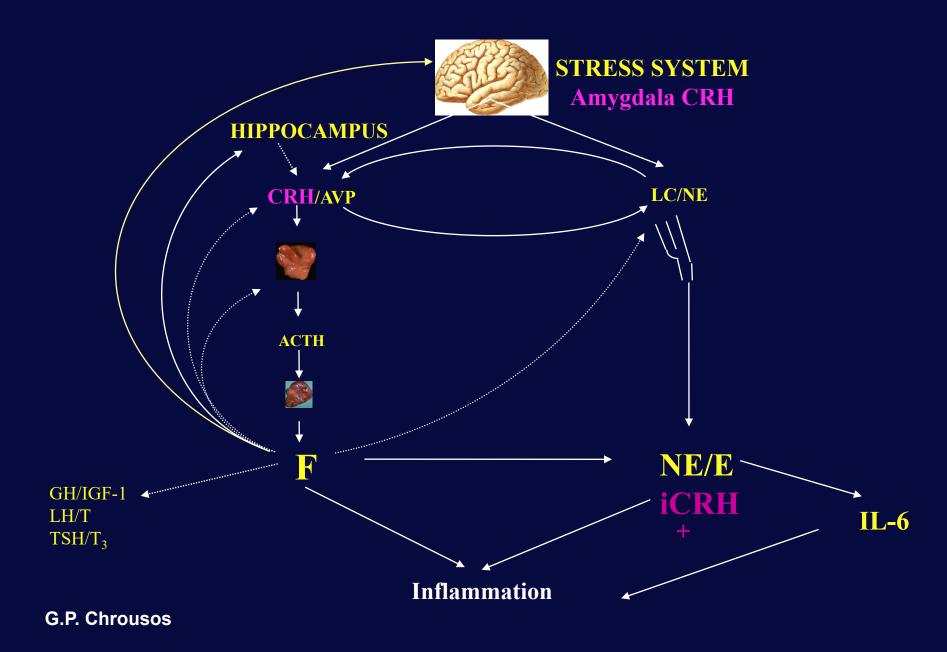
Stress System

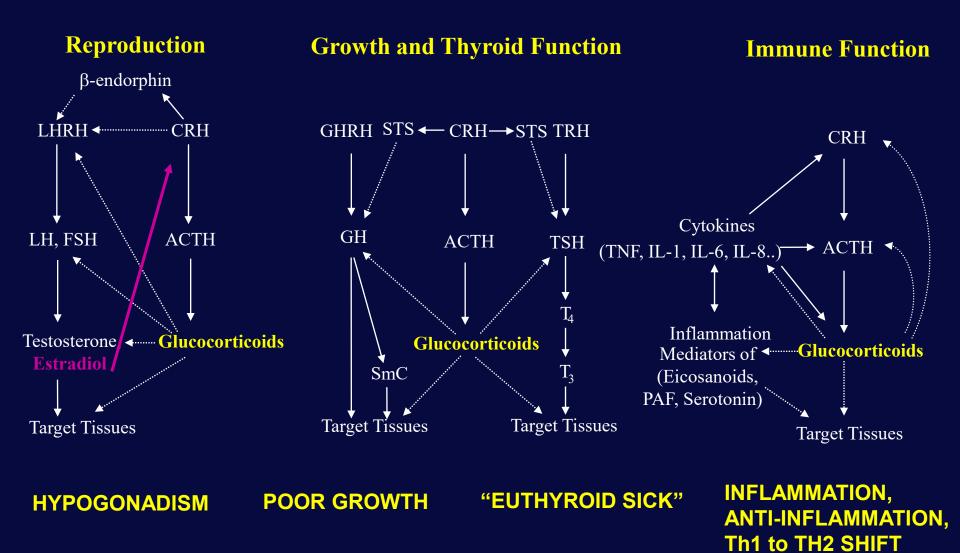


Stress System



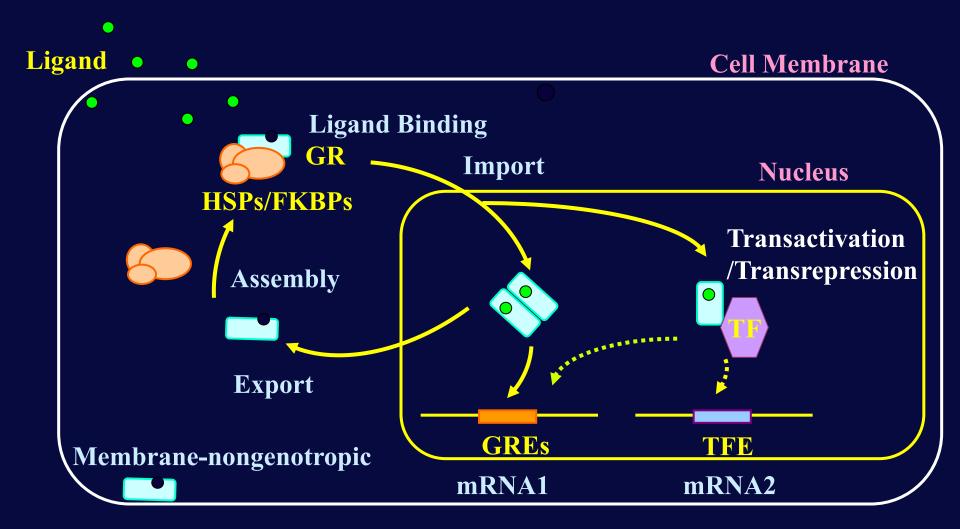
Chrousos JAMA 1992

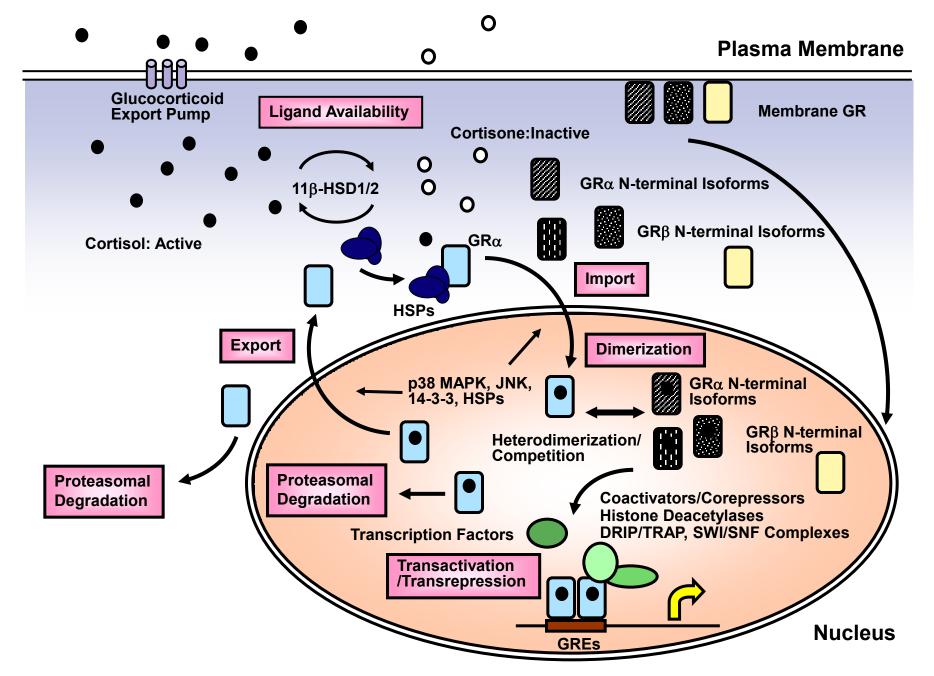




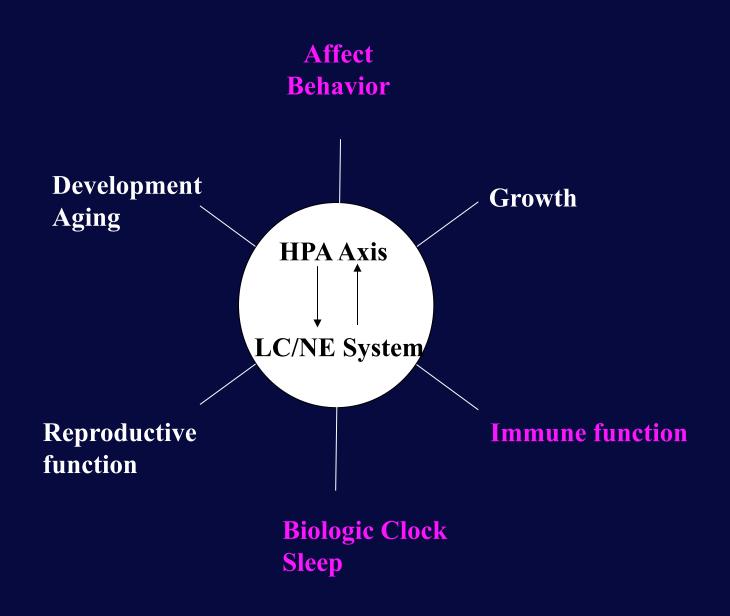
PARAINFLAMMATION

Glucocorticoid Receptor Signaling





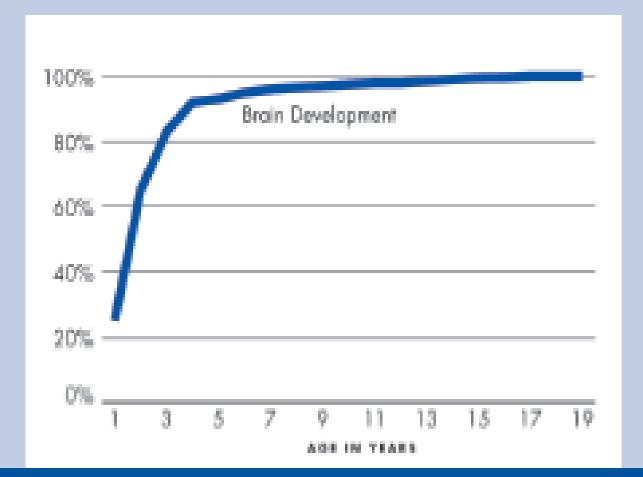
Chrousos & Kino Science STKE 2005



Physical and Emotional Stress Pathophysiology

- Timing (Critical periods=prenatal, first 5 y and adolescence)
- Acuity
- Chronicity

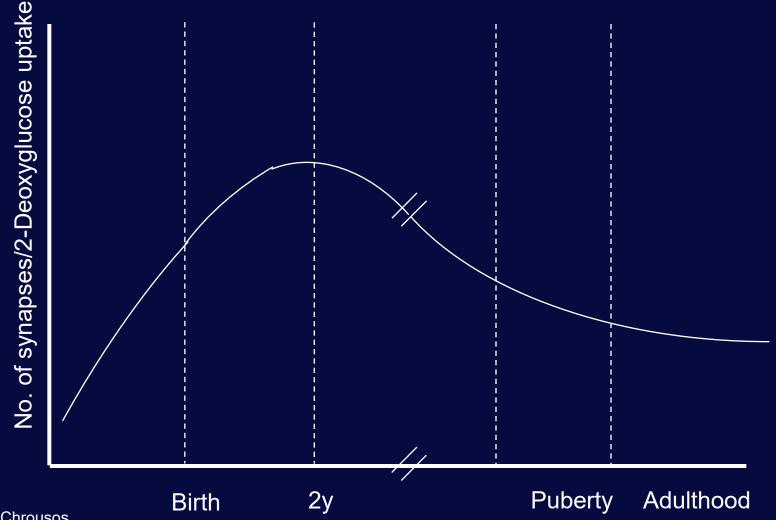
Brain Growth and Child Age



Source: RAND Corporation

G.P. Chrousos

THE DEVELOPING BRAIN



Cognitive and Language Development

Prefrontal/Frontal Lobe "Higher Functions"

- Interpretation of the environment, social cues
- Problem solving
- Planning for the future
- Proper control of impulses (emotional auto-regulation)

"CRITICAL" PERIODS OF LIFE

Prenatal, Early Childhood, Puberty (Human brain ontogeny complete at 25-27 y)

"Organizational" Effects of Hormones, Epigenetics, "Predictive programming"

(CRH, glucocorticoids, sex steroids, cytokines)

Inflammatory Injurious Agents

- Microbial products
- Intracellular molecules (proteins, lipids, carbohydrates, nucleic acids)
- Denatured molecules (proteins, lipids, nutrients)
 Oxidation, nitrosylation, misfolding, etc.
- O and NO radicals
- Adducts
- Xenobiotics/Toxins

INNATE IMMUNITY-First Line of Defense-PRRs

Activated by engagement of germ-line encoded PRRs (Pattern Recognition Receptors)

PRRs recognize the presence of:

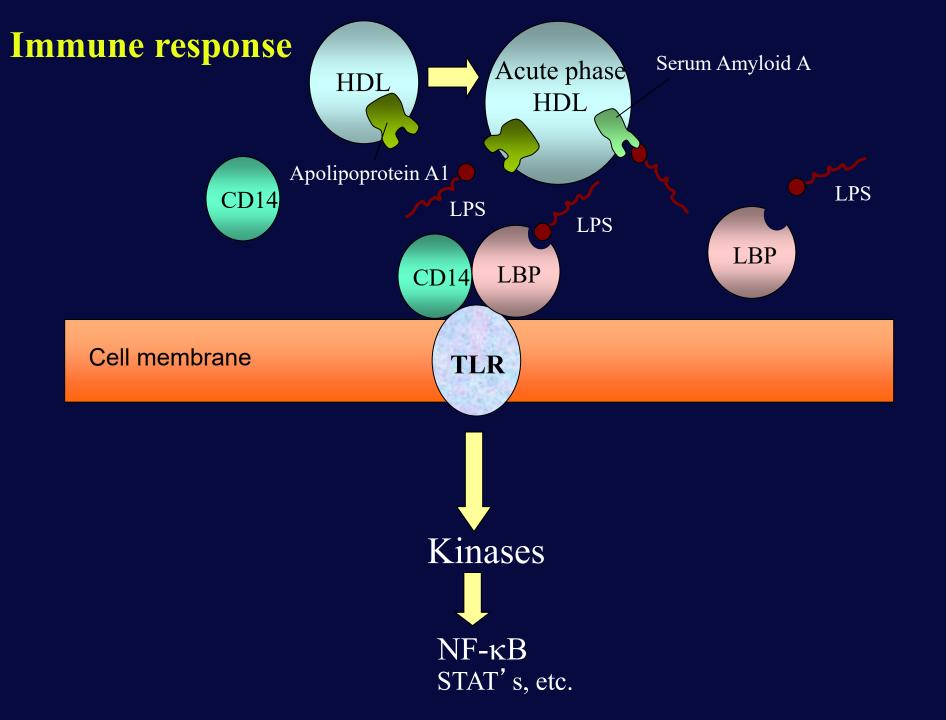
Microbial PAMPs (Pathogen-Associated Molecular Patterns)

Endogenous DAMPs (Danger-Associated Molecular Patterns)

Pattern Recognition Receptors

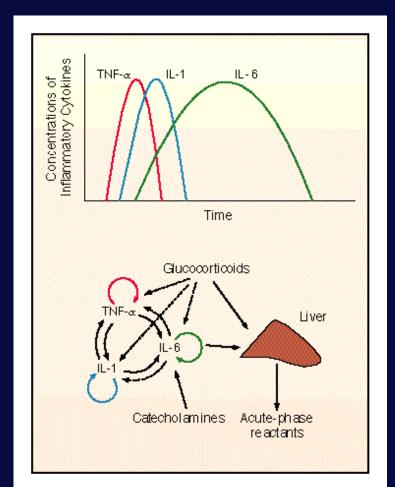
- Toll-like Receptors (TLRs)
- RIG-1-like Receptors
- C-Type Lectin Receptors
- Nod-like Danger Receptors (NLDRs)

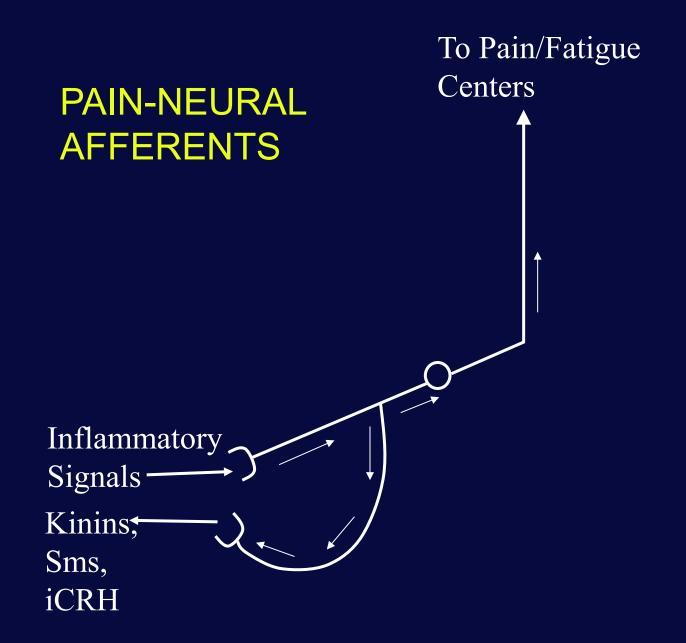
NF-KB NFAT STATs MAPKs Inflammasome, caspase1, IL-1beta, IL-18

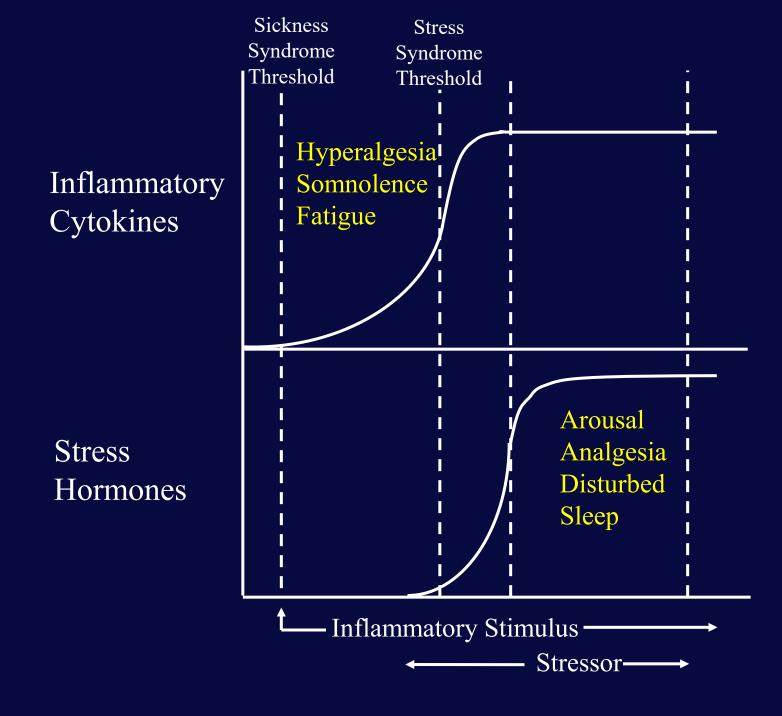


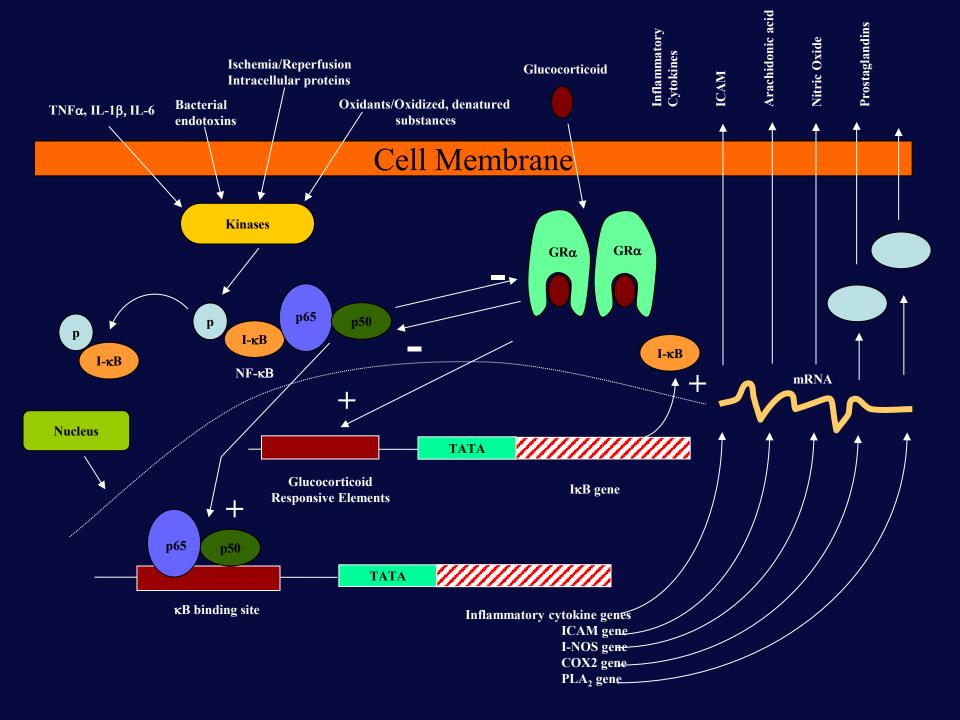
ACTIVATION OF IMMUNE AND IMMUNE ACCESSORY CELLS

- INFLAMMATORY CYTOKINES: TNFα, IL-1, IL-6, IL-8
- OTHER MEDIATORS OF INFLAMMATION:
- Prostanoids, PAF

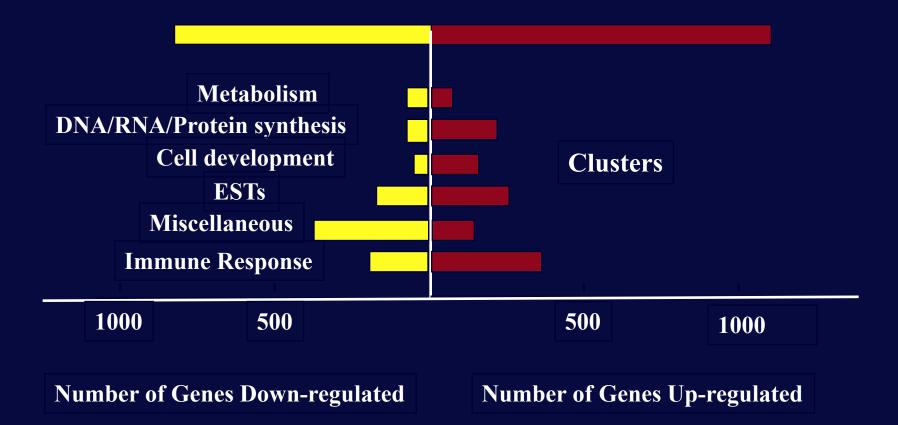






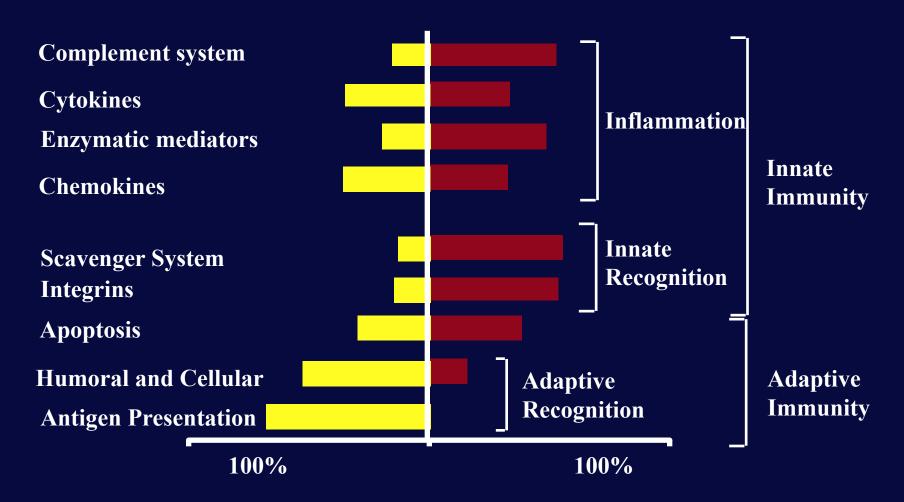


Regulated genes

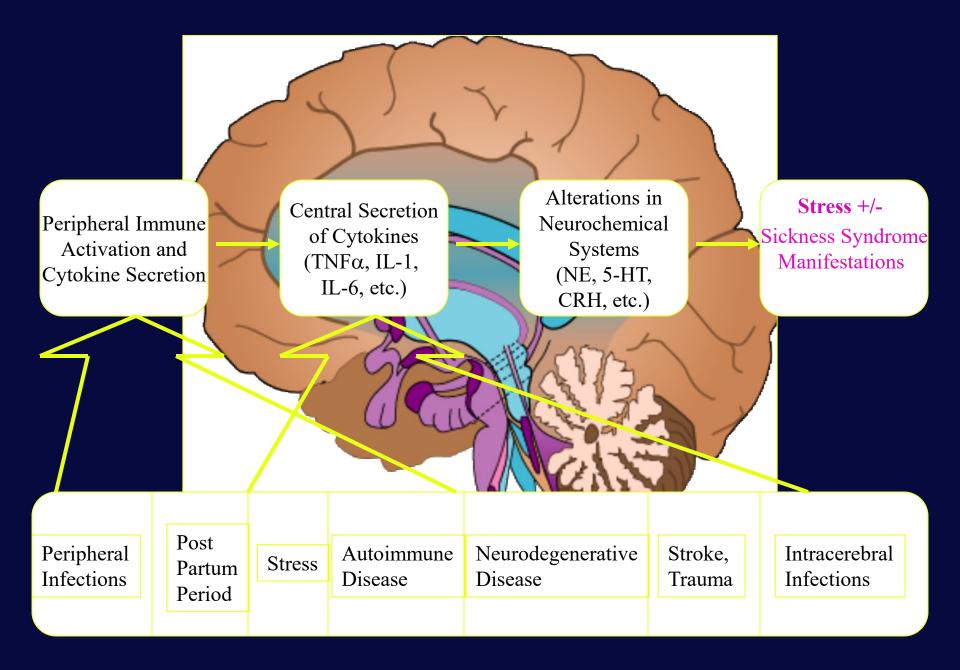


Subcategories

Subclusters



Galon .. Chrousos ... FASEB 2002



Epigenetics of Retrotransposons (Piwi protein-associated ncRNAs called piRNAs)

~60% of genome of retroviral origin
10% of genome consists of Alu repeats
10,000 HERV-K retrotransposons
3,000-5,000 SVA retrotransposons

The Piwi protein-piRNA pathway provides an adaptive defense in the transposon + viral arms race

Increasingly complex networks of small RNAs act through RNA-interference (RNAi) pathways to:

 restrain the spread of "selfish" genetic elements

mediate antiviral responses

regulate gene expression

organize chromosomal domains

Chronic effects of stress system activation:

fatigue, pain (sickness behavior,

stress behavior MUS)

smoldering para-inflammation, immune dysfunction, Th1 to Th2 shift, certain autoimmune disorders,

Vulnerability to certain infections and certain cancers

CHRONIC NONCOMMUNICABLE DISEASES

Chronic effects of stress system activation:

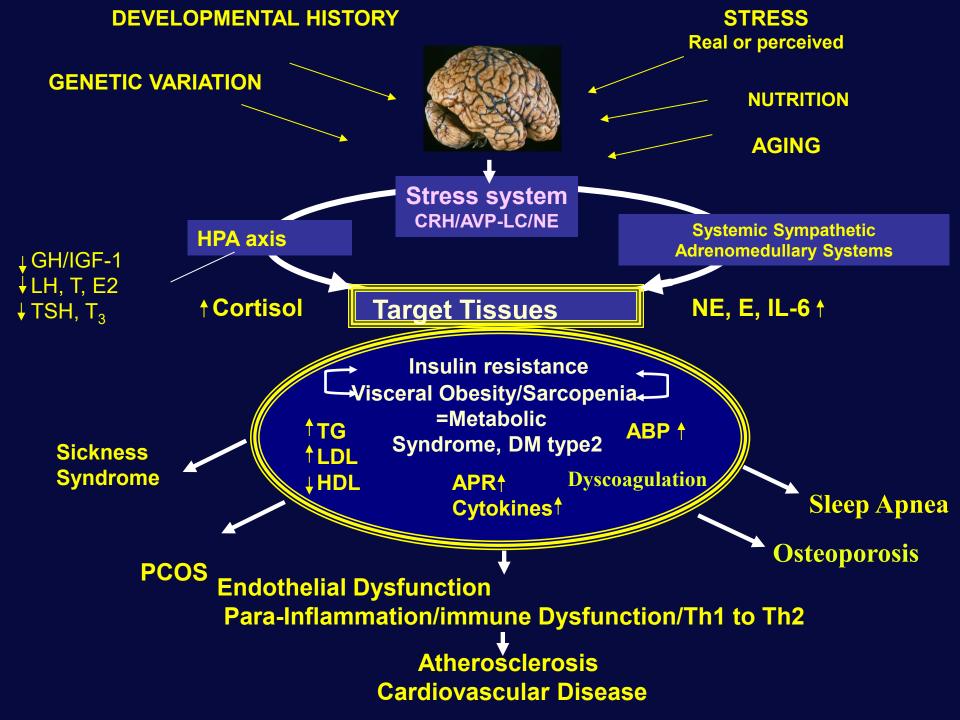
Vulnerability to certain infections

Viral: Common cold viruses

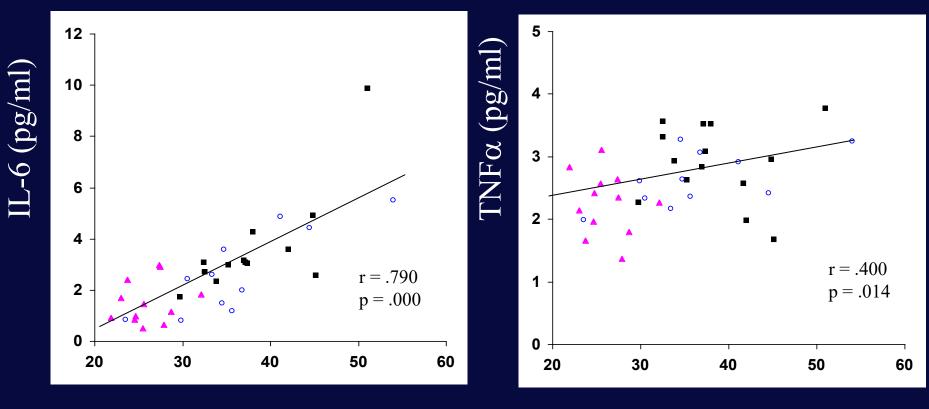
Bacterial: Tuberculosis, Leprosy

Saprophytic infections

Fungal



Both IL-6 and TNFa correlate with BMI



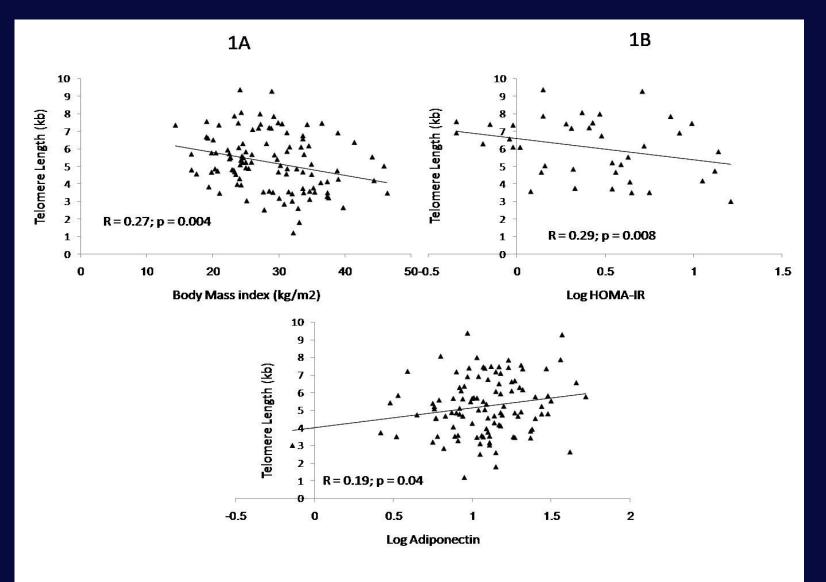
BMI

Vgontzas et al. JCEM 1997

HYPERCYTOKINEMIA

TRAUMA/ BURNS **INFECTIOUS ILLNESSES** AUTOIMMUNE INFLAMMATORY DISEASES ALLERGIC INFLAMMATIONS **CNS INFLAMMATIONS** NONINFLAMMATORY STRESS **OBESITY/VISCERAL OBESITY** AGING





Al-Attas et al. 2010

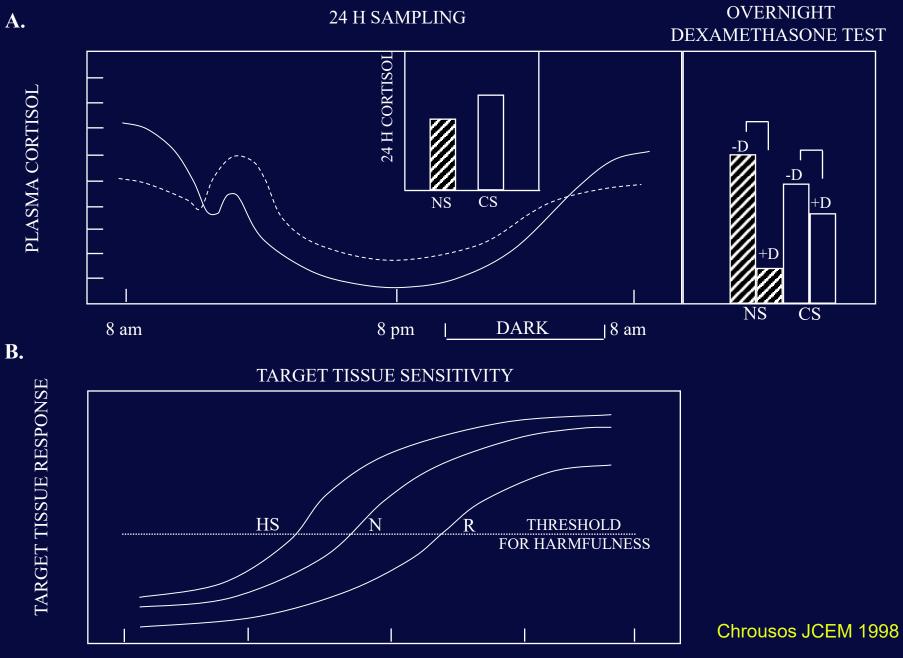
Cellular Stress

Nutritional — Inflammatory

Oxidative

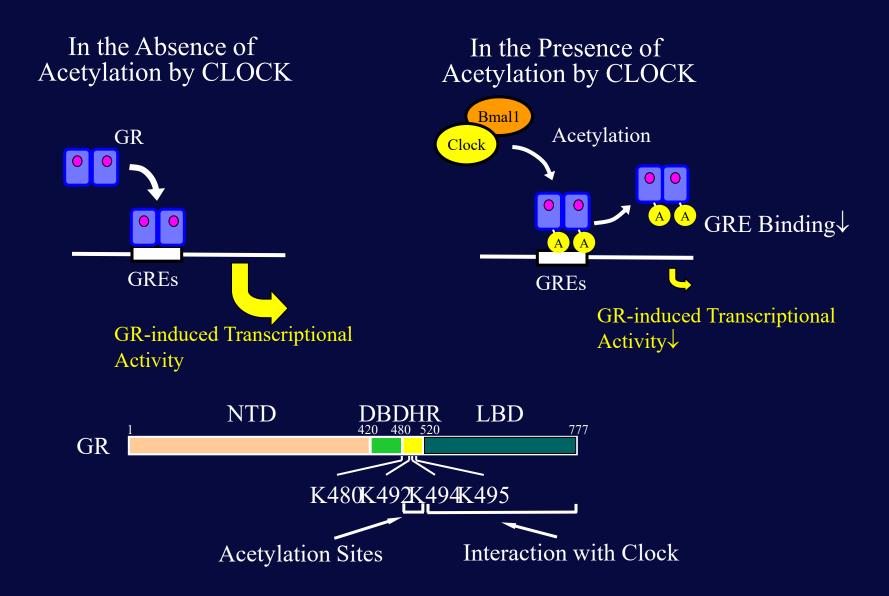
Cellular Stress

Oxidative Mitochondria Keap1 (SH sensor)-Nrf2-ARE



CORTISOL CONCENTRATION

Results #1: Clock/Bmal1 Represses GR Transcriptional Activity through Acetylation



Uncoupling between Circadian Rhythm of Serum Cortisol and Tissue Glucocorticoid Sensitivity

