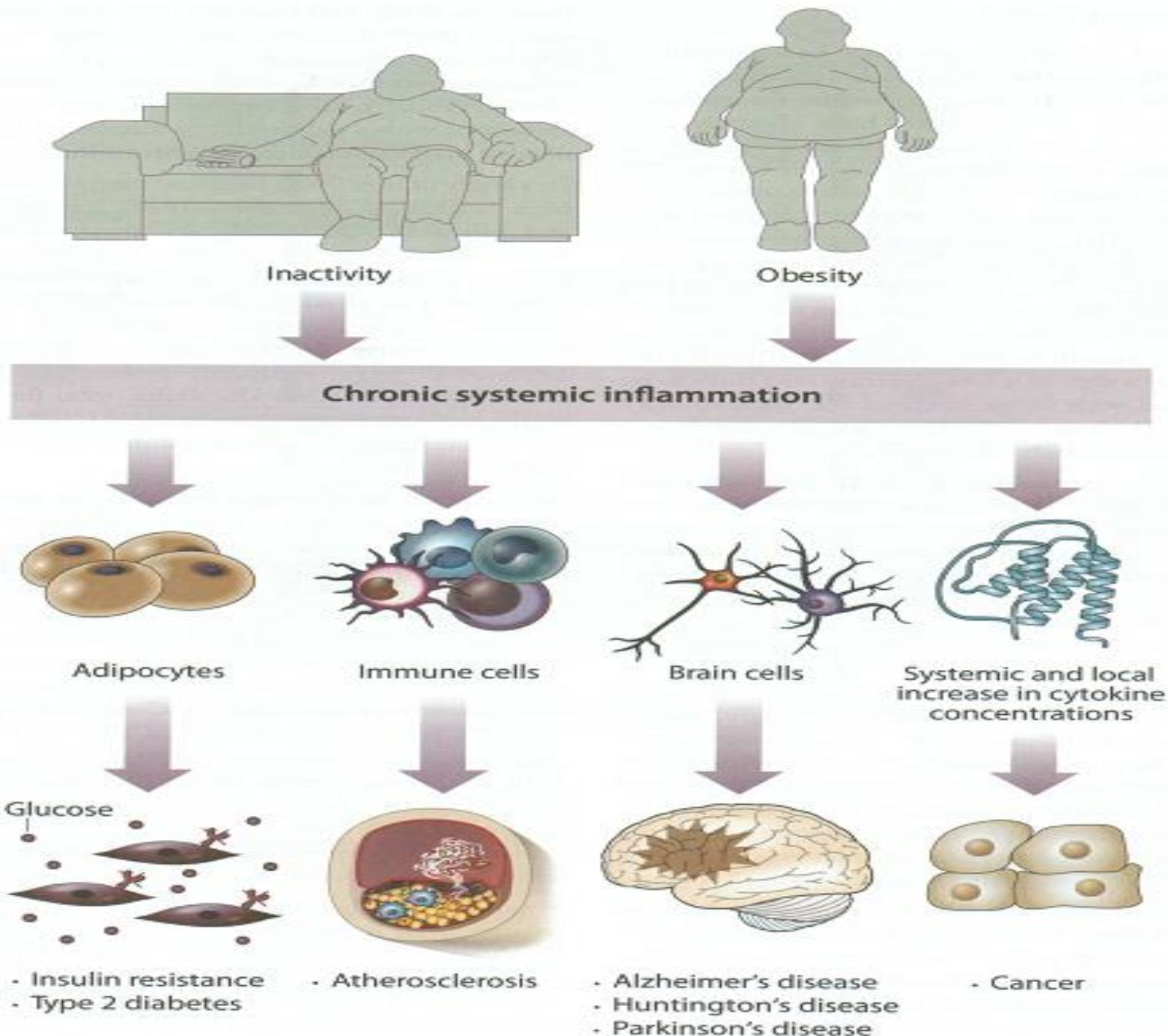


# The Nutrition/Neuroendocrineimmune Network

Jeffrey Bland, Ph.D., FACN, FACB  
President  
Personalized Lifestyle Medicine Institute  
[www.plminstitute.org](http://www.plminstitute.org)

# Connecting the Immune, Endocrine and Nervous Systems

*Network Biology and Functional Medicine*



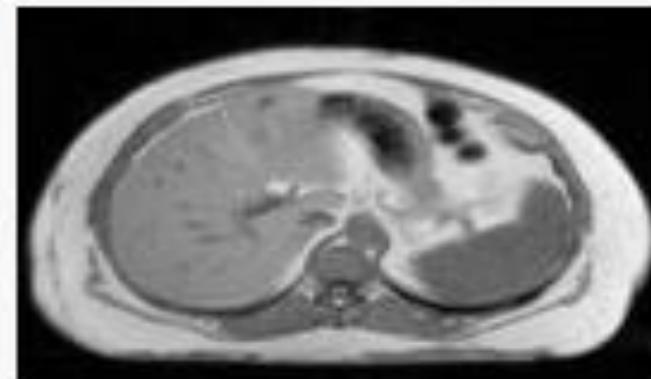
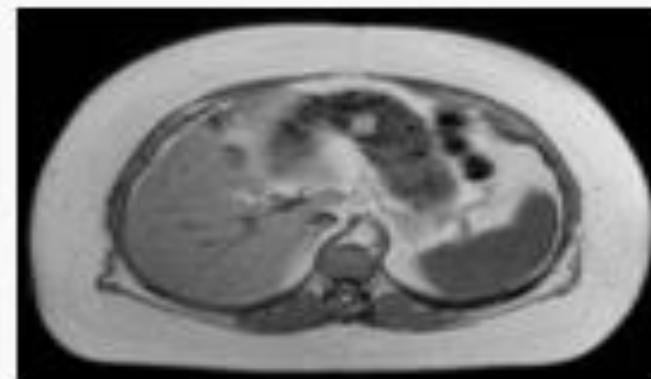
# Absence of an Effect of Liposuction on Insulin Action and Risk Factors for Coronary Heart Disease

Samuel Klein, M.D., Luigi Fontana, M.D., Ph.D., V. Leroy Young, M.D., Andrew R. Coggan, Ph.D., Charles Kilo, M.D., Bruce W. Patterson, Ph.D., and B. Selma Mohammed, M.D., Ph.D.

Before  
Liposuction

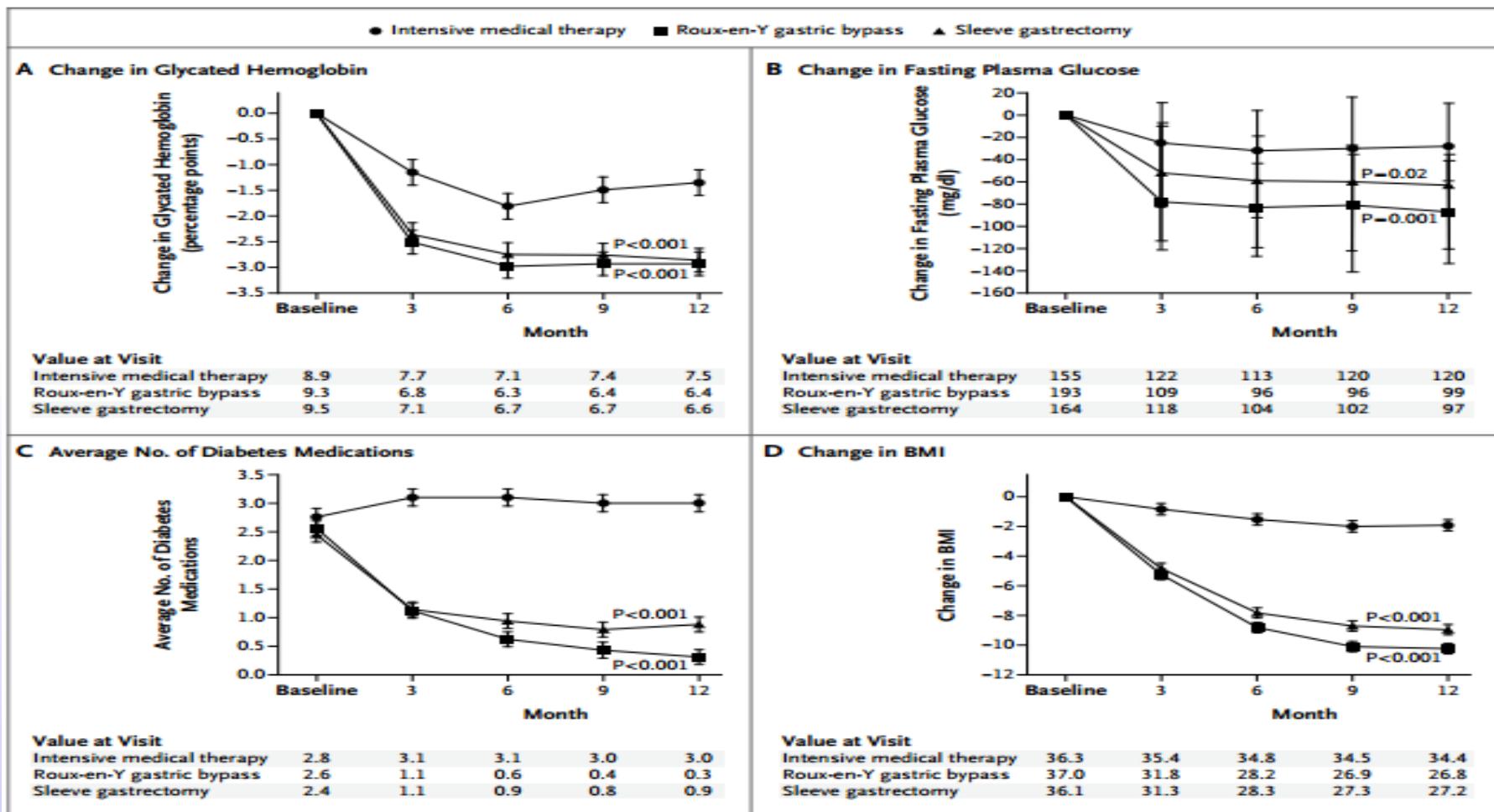


After  
Liposuction

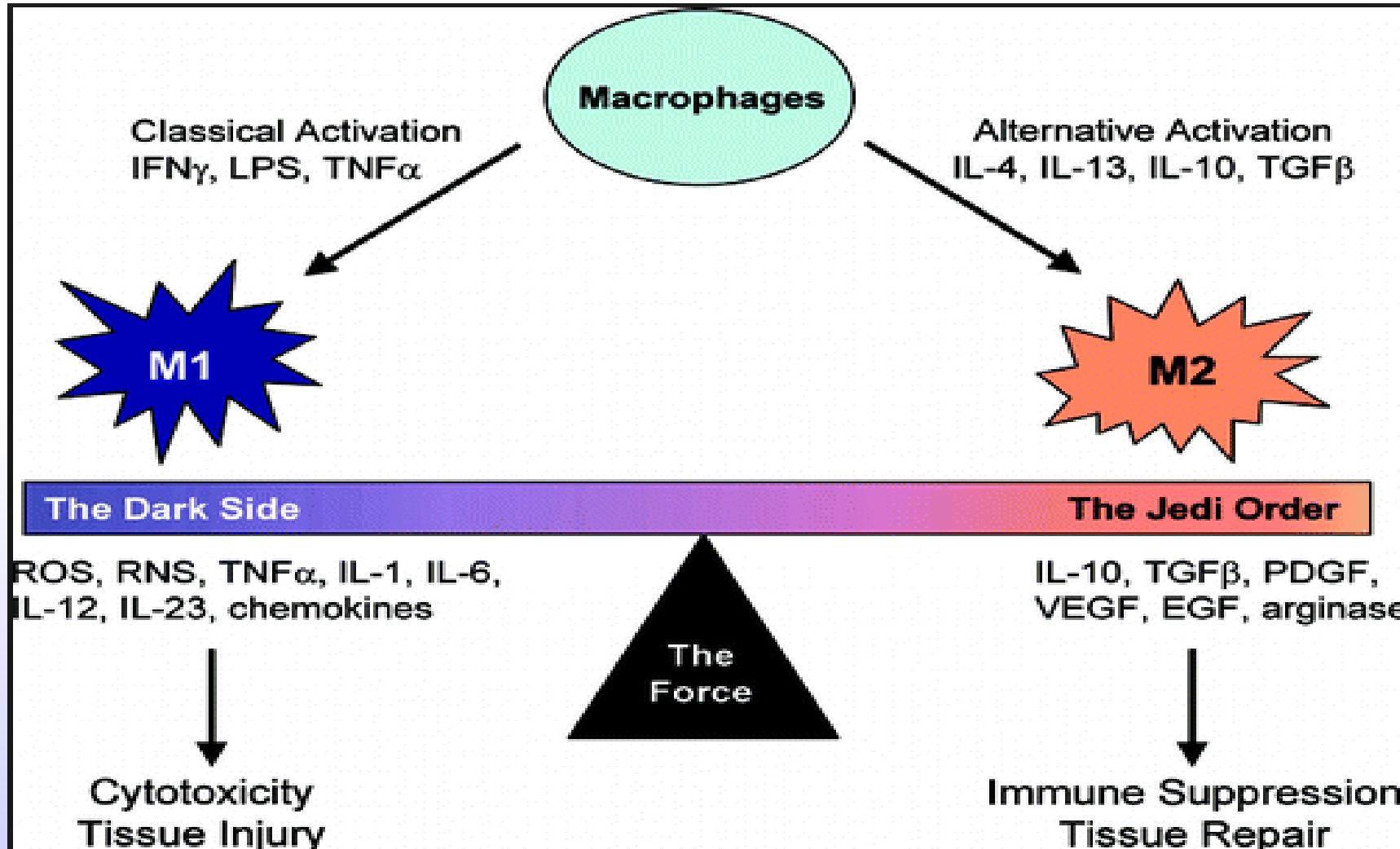


# Bariatric Surgery versus Intensive Medical Therapy in Obese Patients with Diabetes

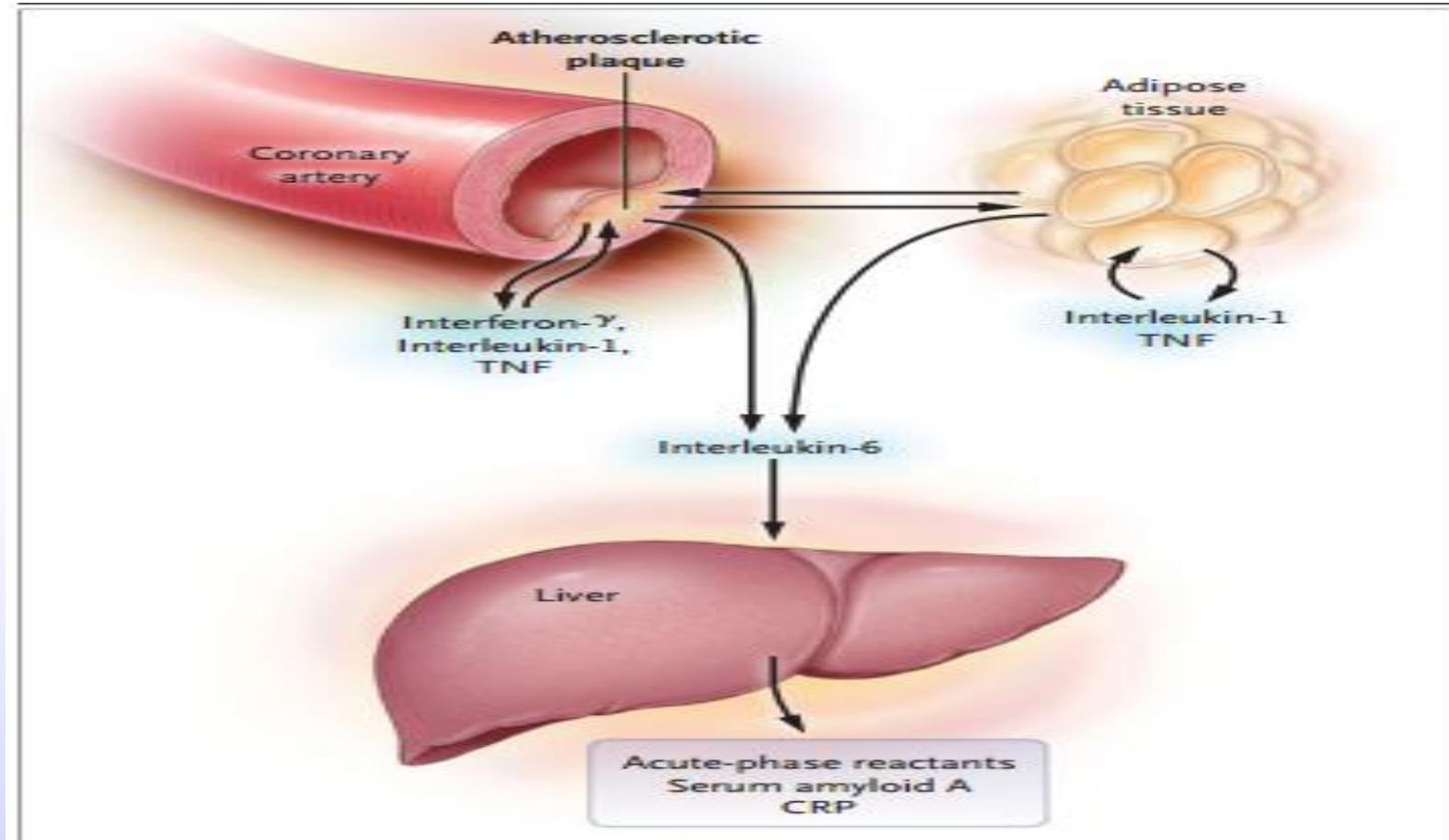
Philip R. Schauer, M.D., Sangeeta R. Kashyap, M.D., Kathy Wolski, M.P.H., Stacy A. Brethauer, M.D., John P. Kirwan, Ph.D., Claire E. Pothier, M.P.H., Susan Thomas, R.N., Beth Aboot, R.N., Steven E. Nissen, M.D., and Deepak L. Bhatt, M.D., M.P.H.



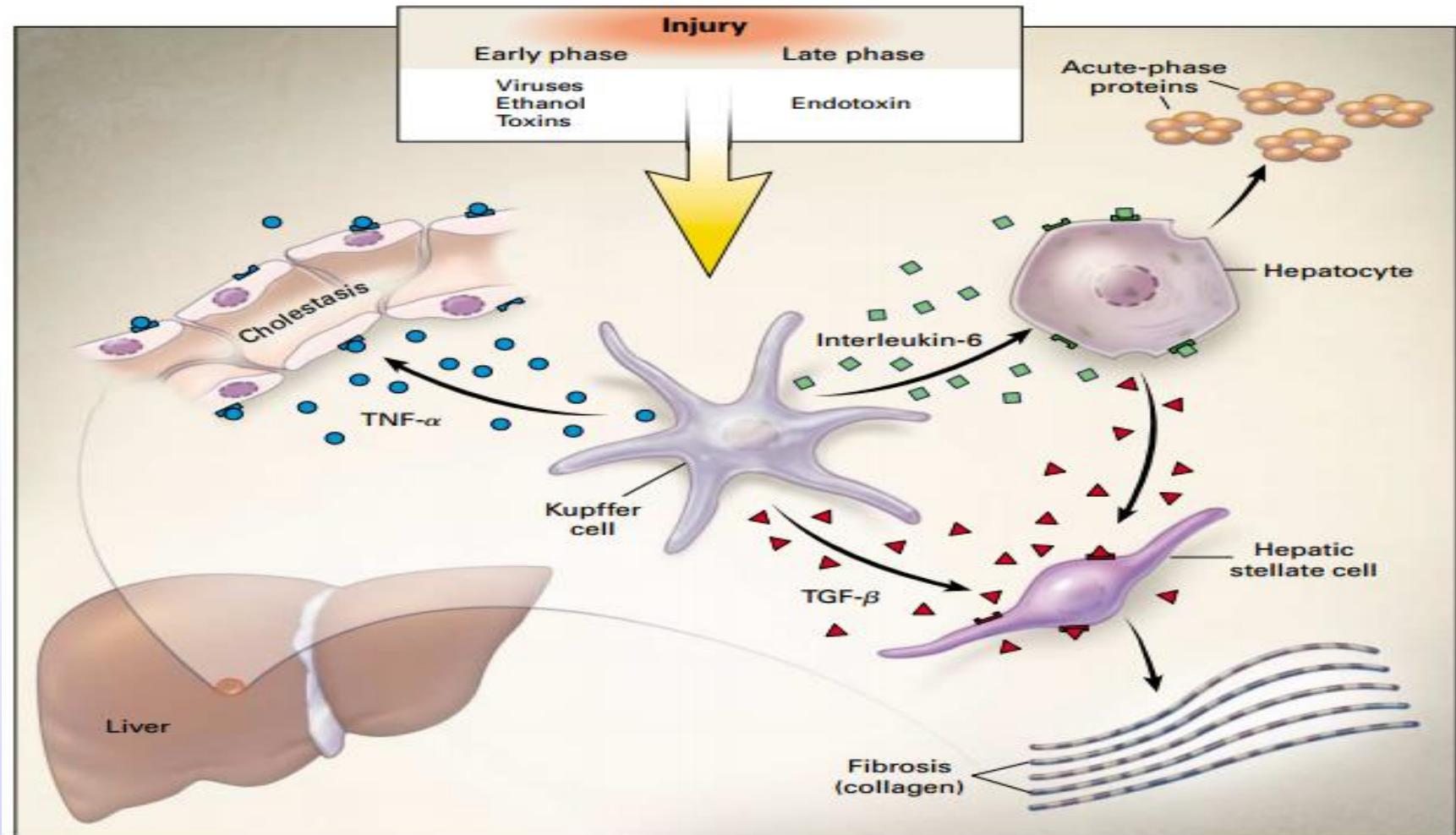
# M1 and M2 Macrophages and Immune Balance



# Metabolic Inflammation and the Cytokine Cascade



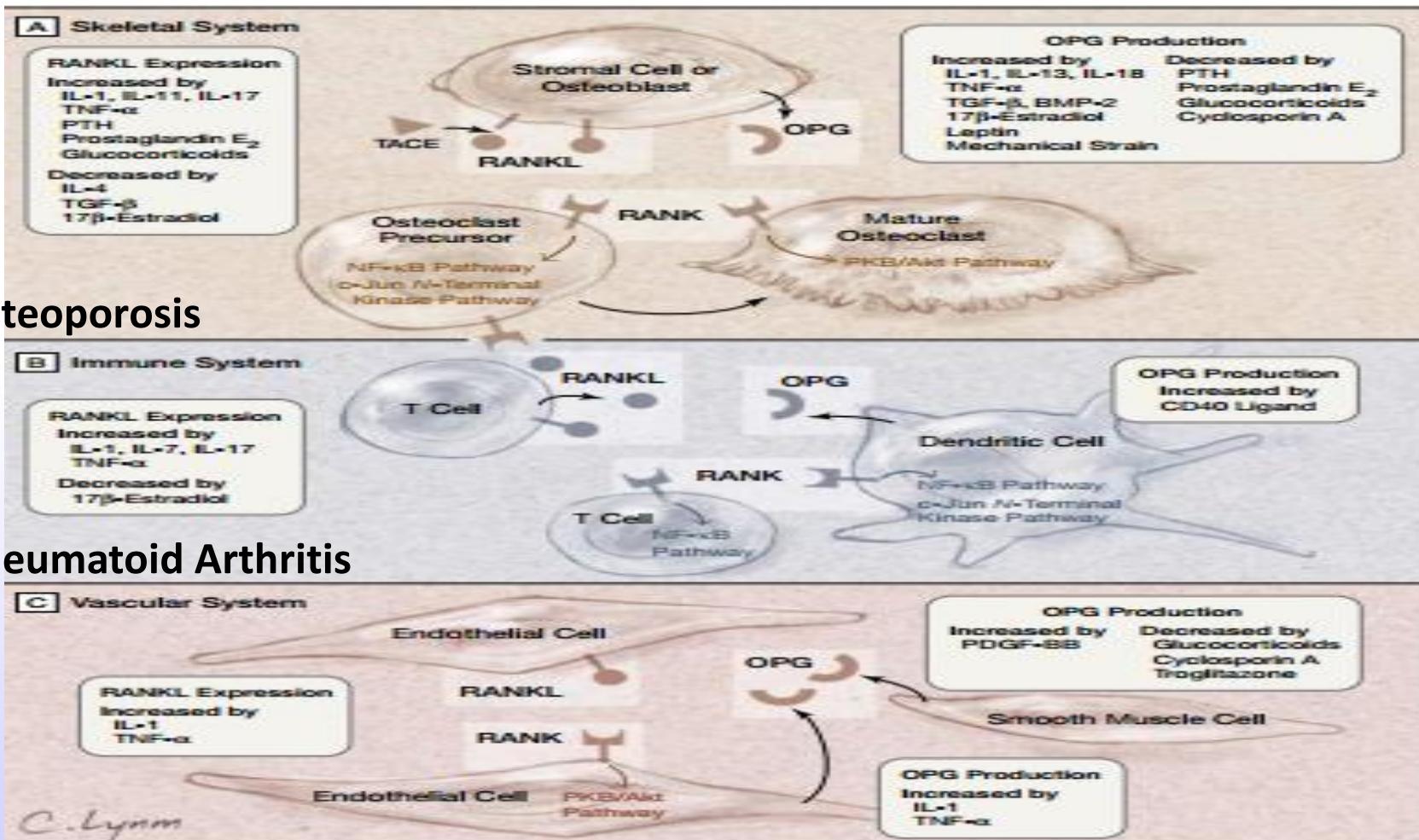
# Hepatic Kupffer Cell Activation, Cytokines and Metabolic Inflammation



# Clinical Implications of the Osteoprotegerin/RANKL/RANK System for Bone and Vascular Diseases

JAMA. 2004;292:490-495

Think Shared Common Mechanism



Coronary Heart Disease

What is the Source of the  
Chronic Inflammation?

# Endotoxin, Inflammation, and Mitochondrial Uncoupling

## Clues to Gene Activity in Inflammation Found

Tracy Hampton, PhD

**D**NA MICROARRAY TESTS THAT REVEAL patterns of gene activity show potential for better understanding human responses to injury and infection, according to new research findings published online in *Nature* on August 31 (Calvano et al. Available at: <http://www.nature.com>).

The study authors, a multi-institutional research team from the National Institute of General Medical Sciences' Inflammation and Host Response to Injury program, are investigating how the systemic inflammation that can occur in patients with injury or infection alters the expression of genes within white blood cells. Such gene expression patterns could help reveal underlying regulatory mechanisms of the inflammatory response, suggest therapeutic interventions, and (because patients respond differently to therapies) may allow physicians to tailor therapies specifically for individual patients.

The investigators injected healthy participants with endotoxin, a bacterial component that can cause sepsis in susceptible burn and trauma patients. (When

injected into healthy individuals, endotoxin produces a widespread but controlled inflammatory response.) The researchers then collected blood samples at subsequent time points over 24 hours, analyzed gene expression patterns in circulating white blood cells over time, and compared the results with those of control participants. The laboratory technologies used tested nearly 45 000 probes, representing all human genes.

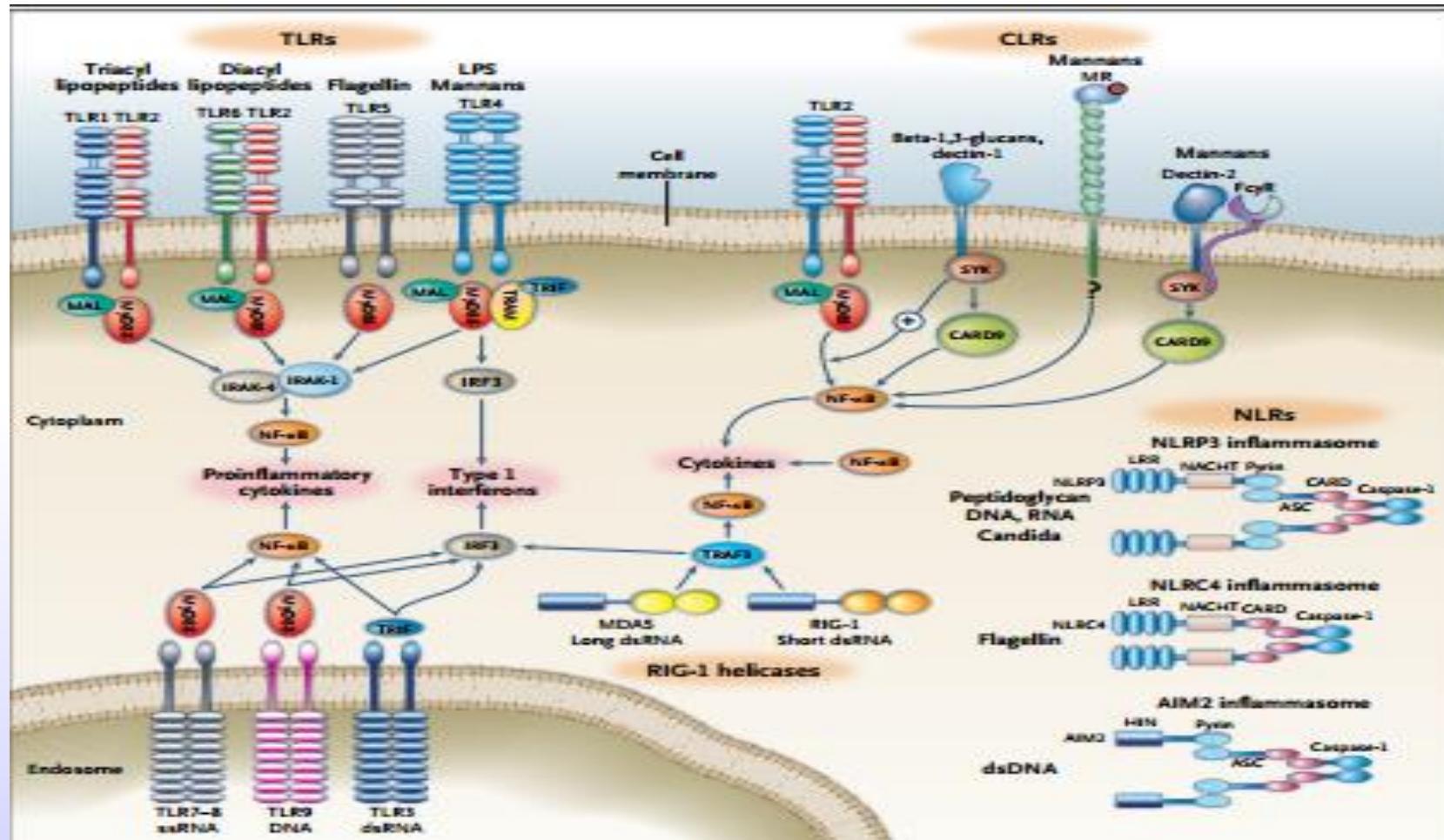
Analyses of the data revealed that activity of more than 3700 genes in white blood cells changed significantly in individuals who received endotoxin, while activity of those genes in control participants was unchanged. Over time, more than half of the genes were expressed at reduced levels, including several genes involved in the function of cells' energy-producing organelle, the mitochondria. After these changes in gene expression occurred, almost all returned to their baseline level of expression by 24 hours, indicating resolution of the inflammatory response.

To understand how these changes in gene expression might affect other genes and biological processes, the team turned to a database of information on thou-

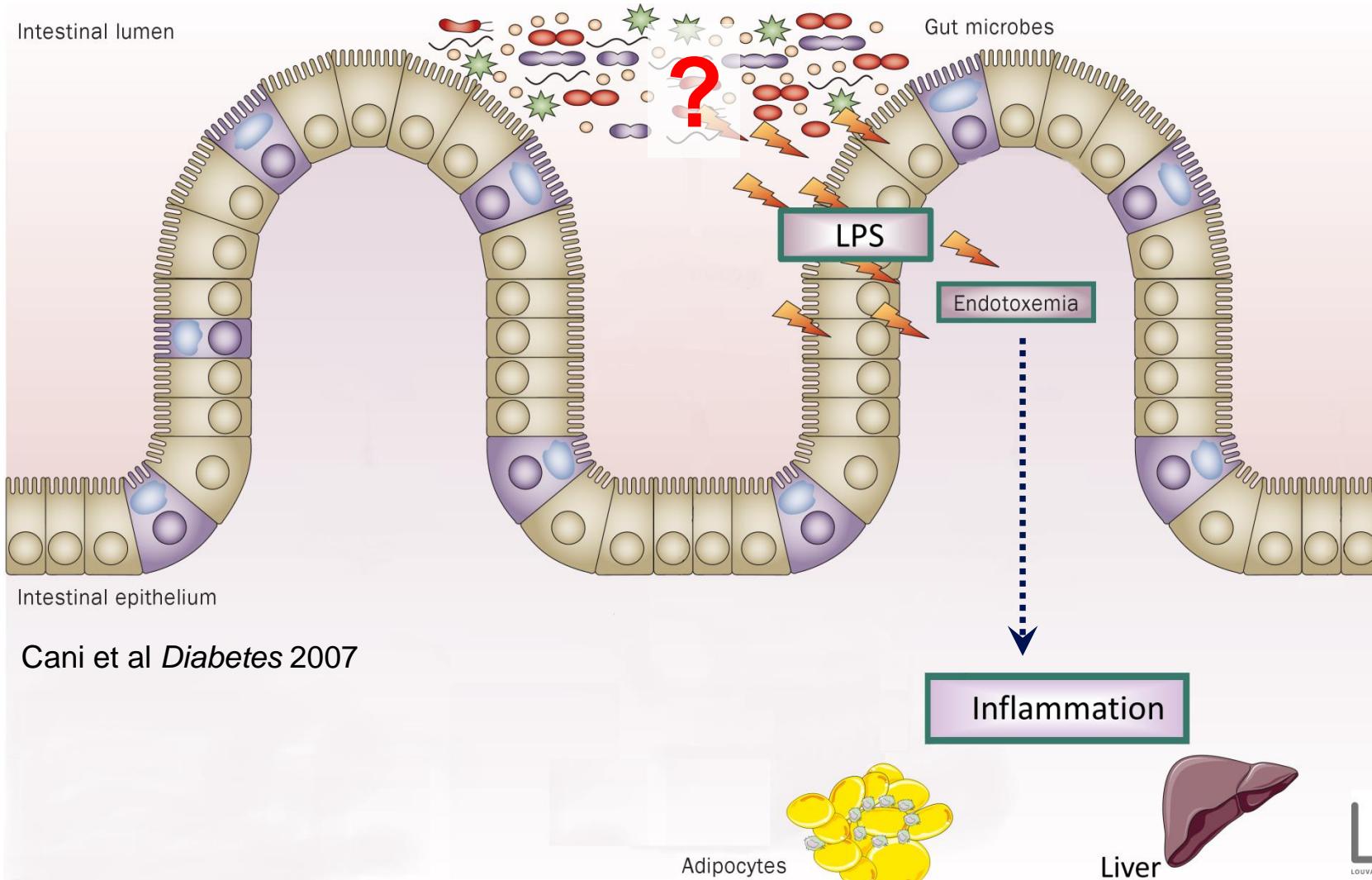
sands of human, mouse, and rat genes compiled from more than 200 000 scientific articles. From the information there, they constructed inflammation-associated molecular networks that showed how the genes altered in the endotoxin study could interact with more than 4000 additional genes. For example, the investigators reported abnormalities in gene networks responsible for energy production and protein synthesis and degradation in white blood cells. Moreover, alterations in the expression of hundreds of other genes and pathways not previously known to be associated with inflammation were discovered.

The investigators' work may lead to a better understanding of how some individuals recover quickly from traumatic injuries while others can develop inflammatory complications long after the initial injury. The techniques used in the effort are now being applied to samples collected from burn and trauma patients for a new study. Nearly 300 patients to date have been enrolled for this study; a preliminary analysis of the first 100 or so patients is currently under way. □

# 4 Types of Pattern Recognition Receptors



# Specific gut microbiota derived compounds are able to trigger metabolic inflammation



Cani et al *Diabetes* 2007

Adapted from

**nature**  
REVIEWS

ENDOCRINOLOGY

Nathalie M. Delzenne, Audrey M. Neyrinck, Fredrik Bäckhed and Patrice D. Cani

# Gut Microbiota from Twins

## Gut Microbiota from Twins Discordant for Obesity Modulate Metabolism in Mice

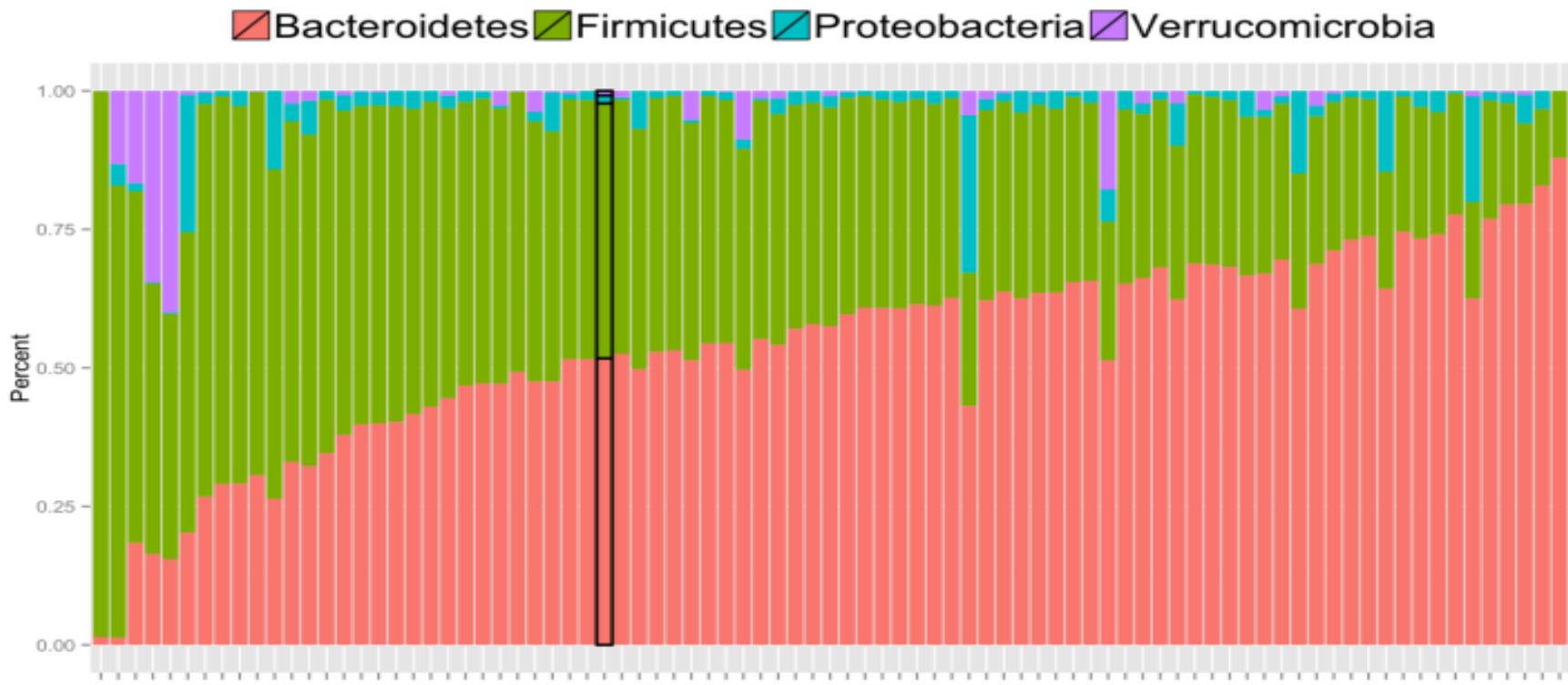
**Interspecific Disease:** Antimicrobial resistance genes, phage and plasmid load from bacterial communities of all human gut microbiomes are causally related to a significant change in disease phenotypes in children. Below, we illustrate how causality generally are represented by the disease heterogeneity, differences between and within disease-associated gut microbiomes, disease, gut flora and antimicrobials. Translating this into a direct causation of antibiotic resistance genes into antibiotic resistance genes of gut microbiomes is a complex process of many factors. The impact of disease heterogeneity on the causality of antibiotic resistance genes in the microbiome, and thus influence of antibiotic resistance genes is not discussed. In addition, comparing children with antibiotic resistance genes to healthy children, we observed that gut microbiomes are representative of disease status. In contrast, most disease-associated genes showed through communication of sugar intake, Aspergillus, and heterogeneity with the gut microbiome, were not causally related to the disease heterogeneity. The causality of antibiotic resistance genes in the gut microbiome is not discussed.



- Gut microbiota from lean animals transplanted to fat animals confer altered gene expression associated with lean phenotype of a low fat, high fiber diet
- Effect is NOT due to decreased calorie release from lean microbiota, but metabolic influence
- Science 2013; 341: 1079

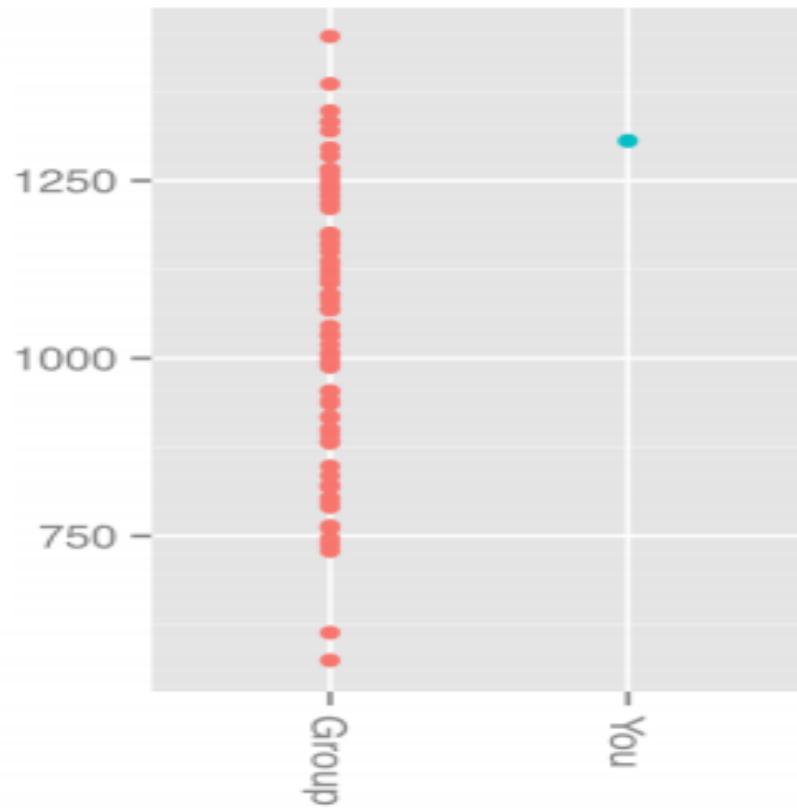
# Example of Species Diversity in the Gut Microbiome

We found a broad spectrum of **gut microbiome** composition among the Pioneers.

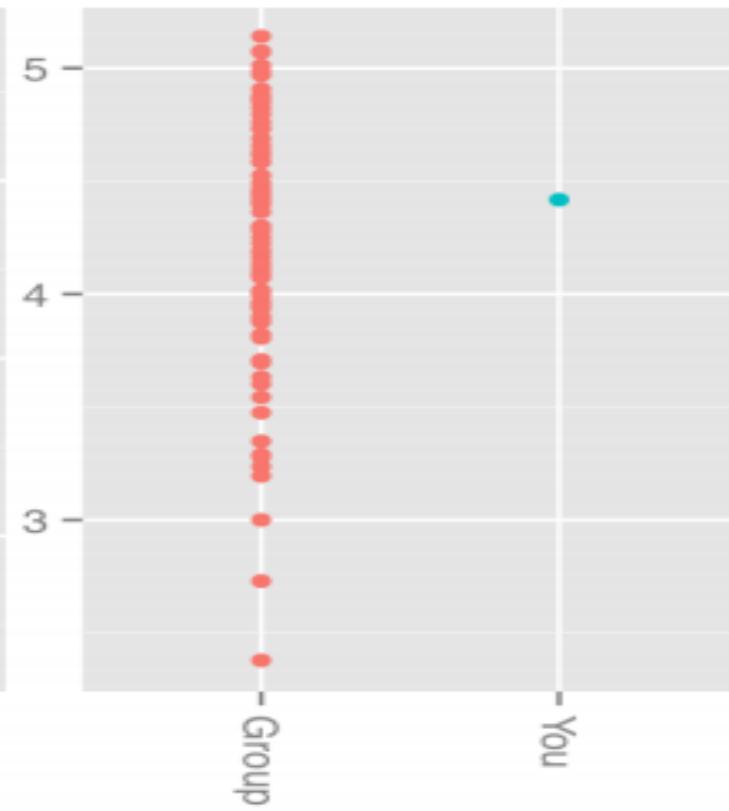


# Relative Evaluation of Microbiome Species Diversity

**Number of Total Species**



**Diversity Score**

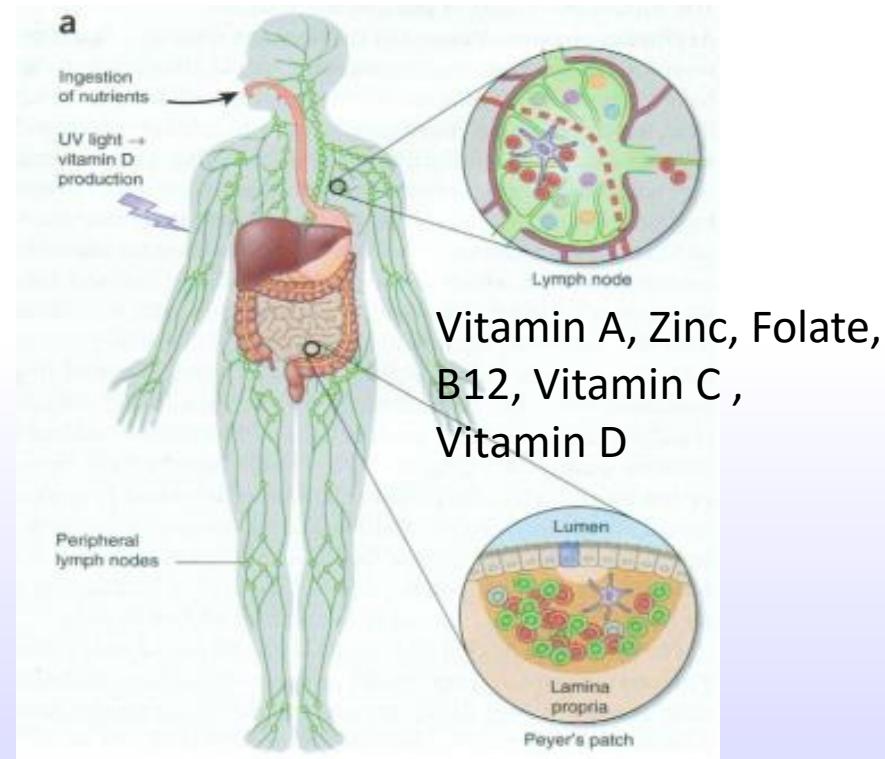


# Role of Specific Nutrients on Humoral Immunity

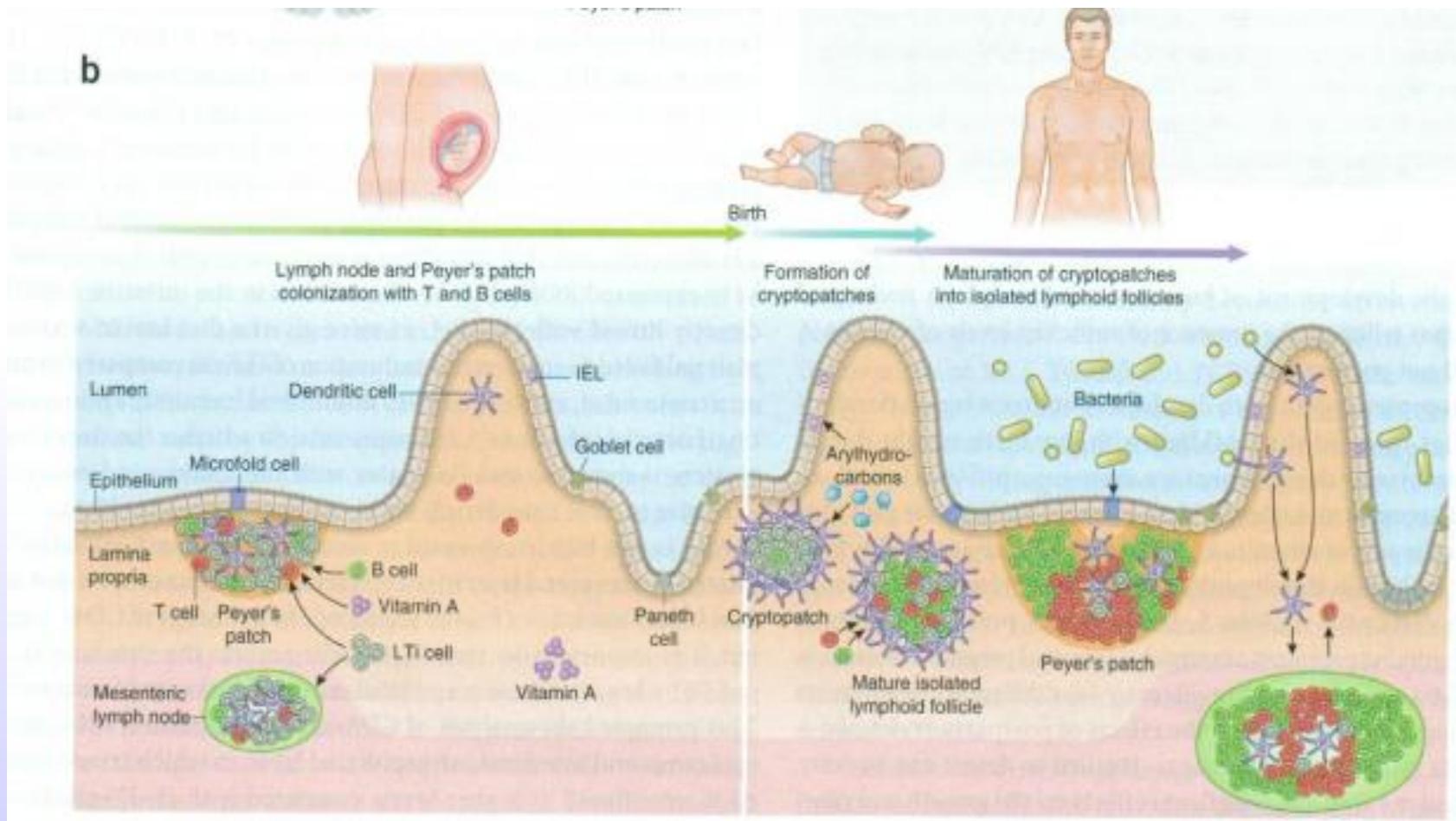
## Influence of nutrient-derived metabolites on lymphocyte immunity

Marc Veldhoven & Cristina Ferreira

Organisms need to protect themselves against potential dangers from their surroundings, yet they require constant and intimate interactions with the same environment for their survival. The immune system is instrumental for protection against invading organisms and their toxins. The immune system consists of many cell types and is highly integrated within other tissues. Immune activity is particularly enriched at surfaces that separate the host from its environment, such as the skin and the gastrointestinal tract. This enables protection at sites directly at risk but also enables environmental factors to influence the maturation and function of immune structures and cells. Recent work has indicated that the diet in particular is able to influence the immune system and thus affect the development of inflammatory disease. This review aims to highlight recent work on how external factors, with a focus on those derived from the diet such as vitamin A, can have a direct or indirect deterministic influence on the activity and function of immunity.



# Development of GALT immune Function

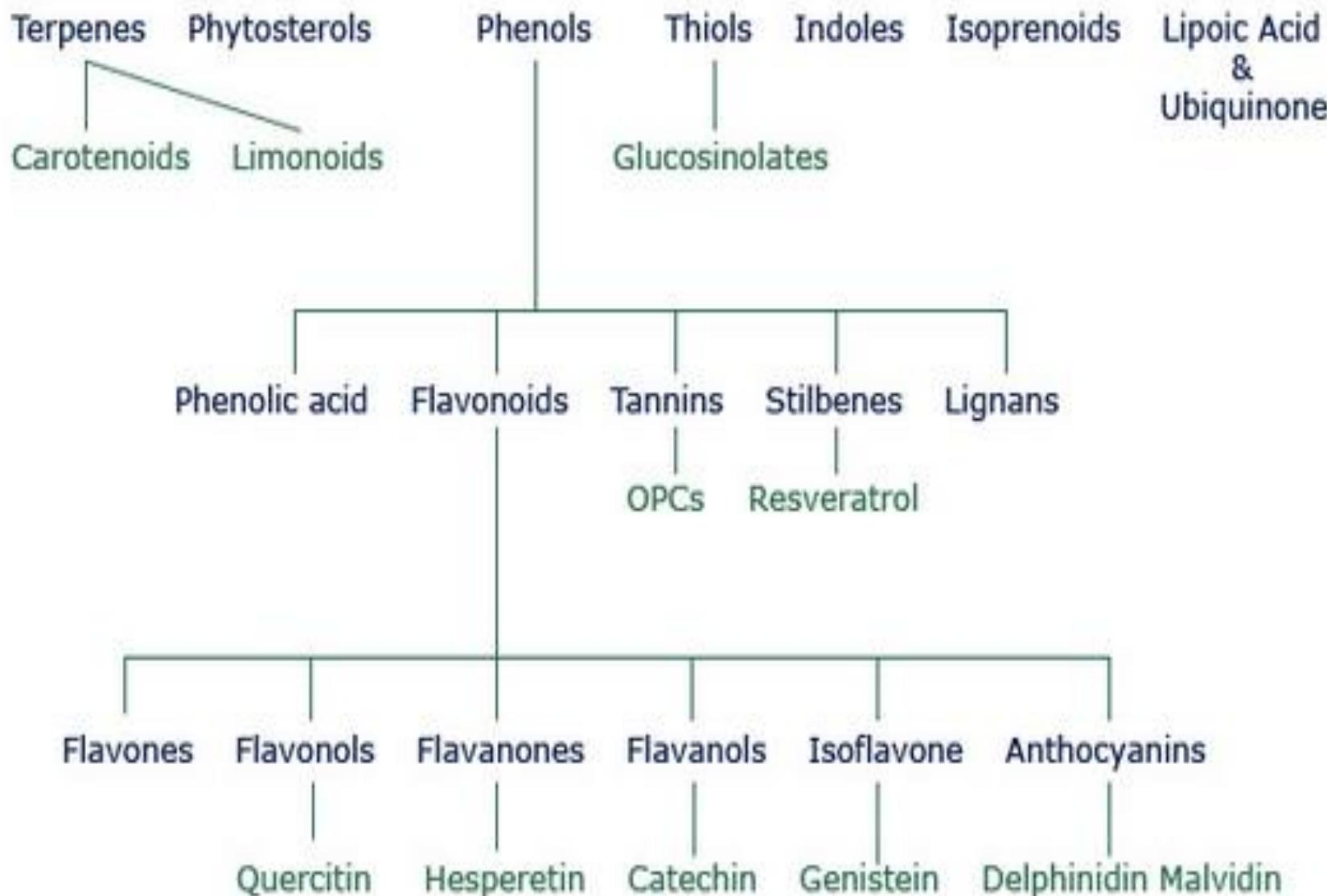


# Phytochemical Regulation of Neuroendocrineimmune Networks



**Food is Much More than We Thought for the Previous 100 years in the Western World**

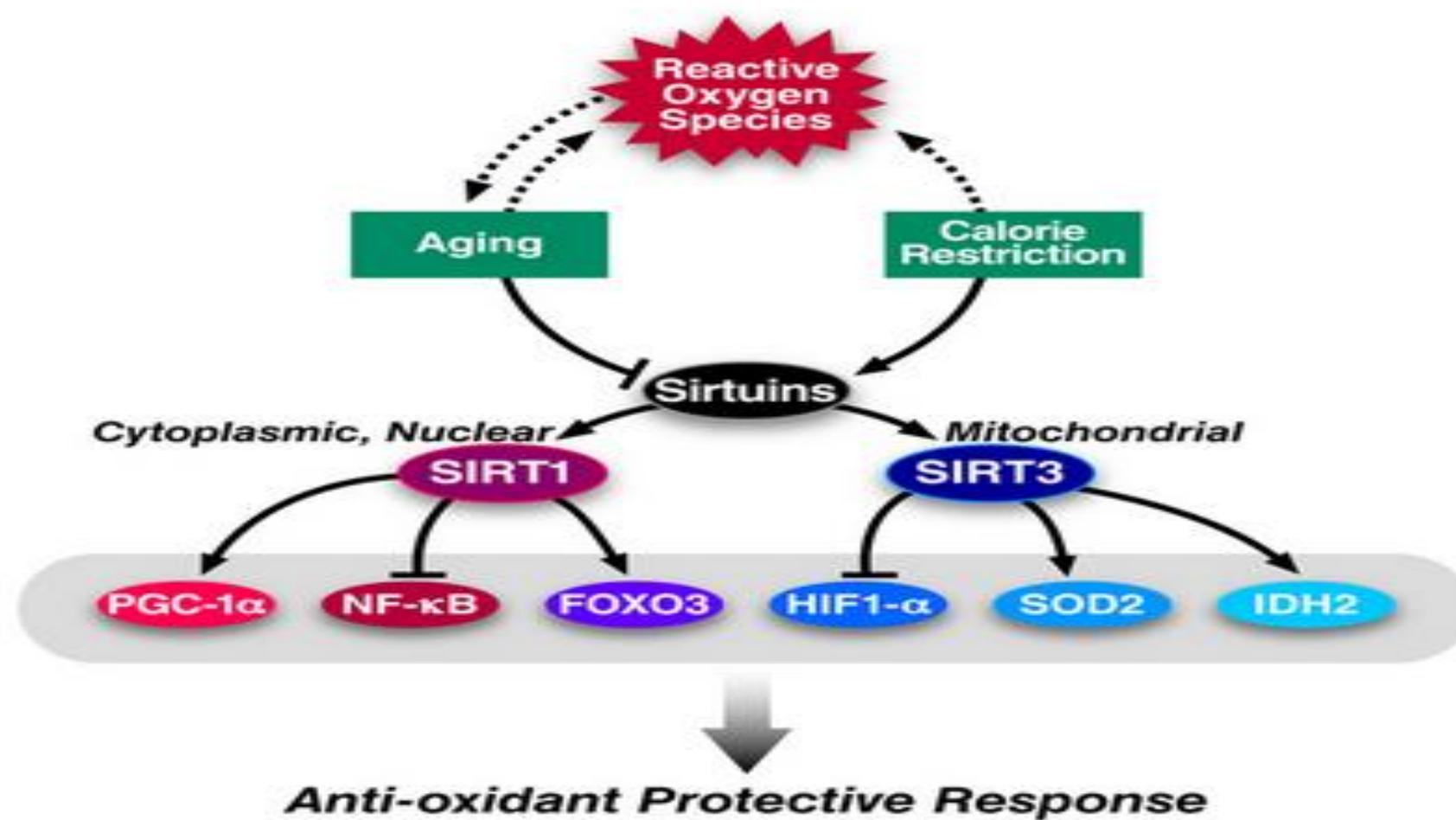
# Phytochemicals



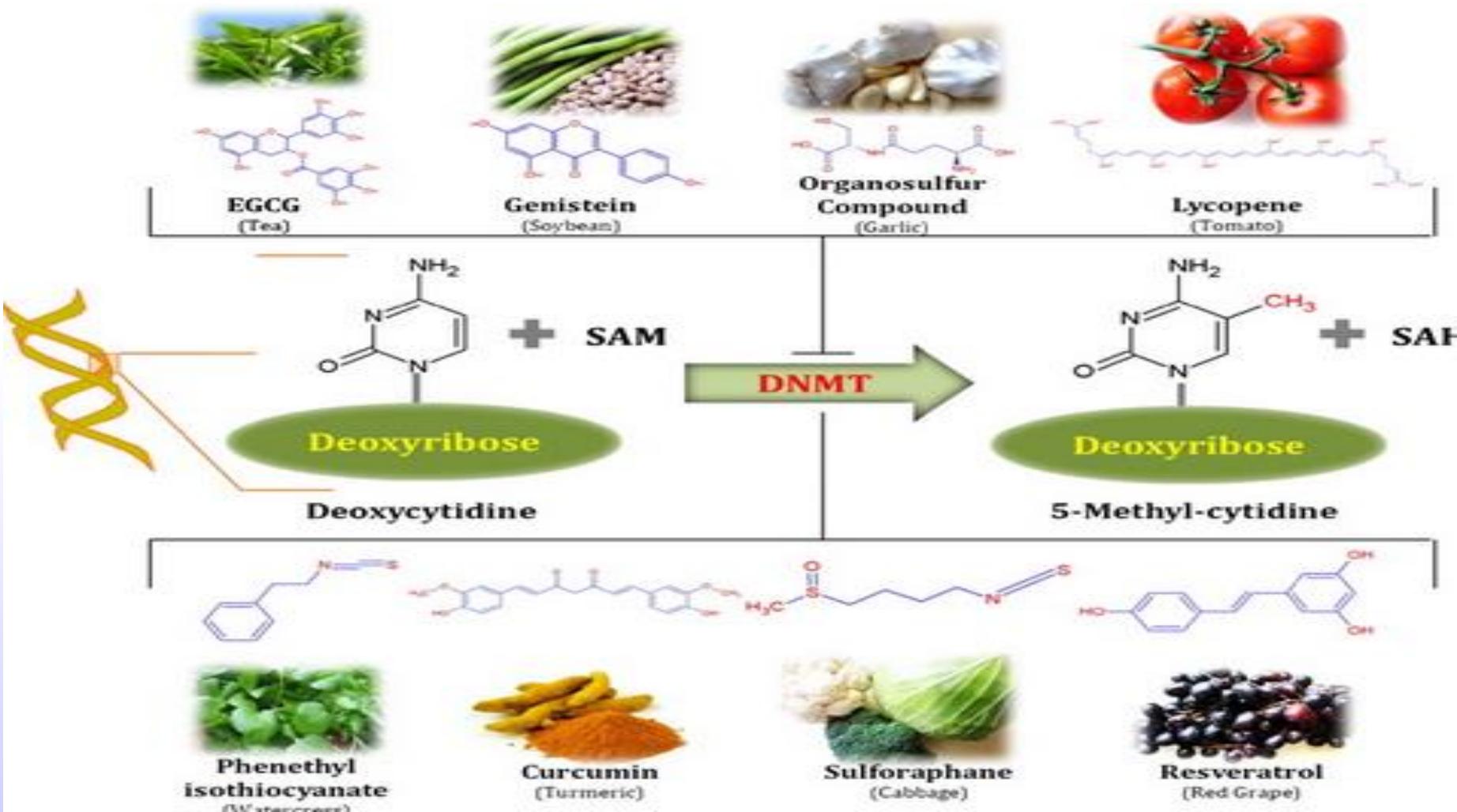
# The Sirtuin-Histone Deacetylase Story

**Upregulation of Sirtuin Activity by  
Specific Phytochemicals**

# Sirtuins as Regulators of Gene Expression



# Foods and Phytochemicals that Influence Genome Methylation and Sirutins



# Example of How Phytochemicals Influence the Endocrine System

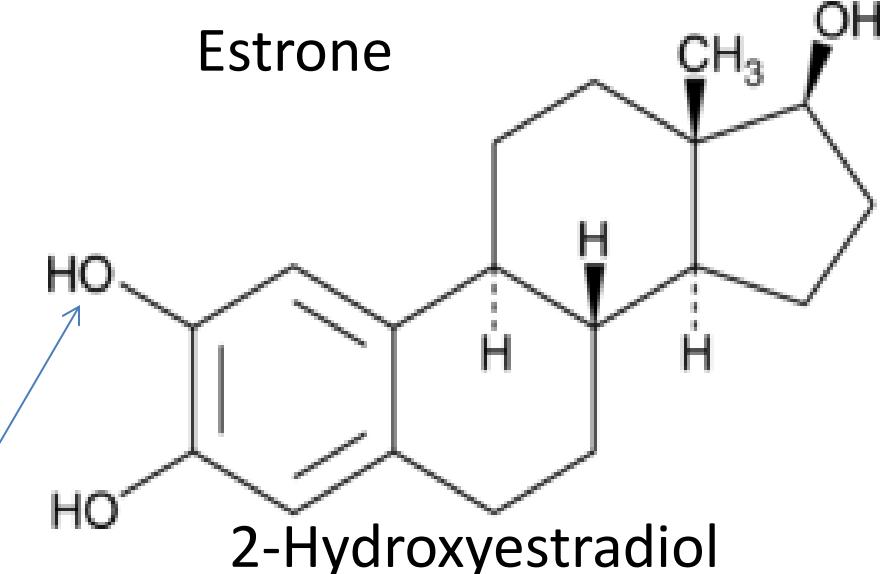
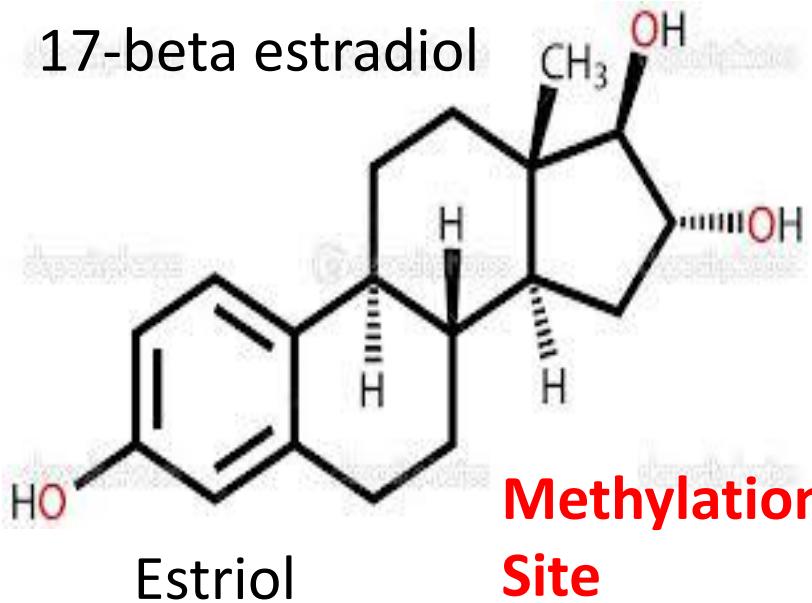
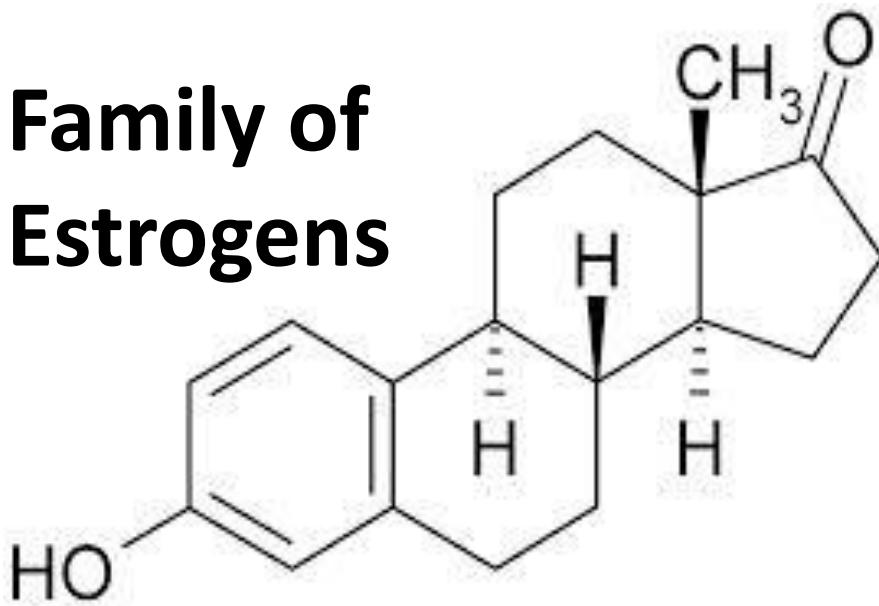
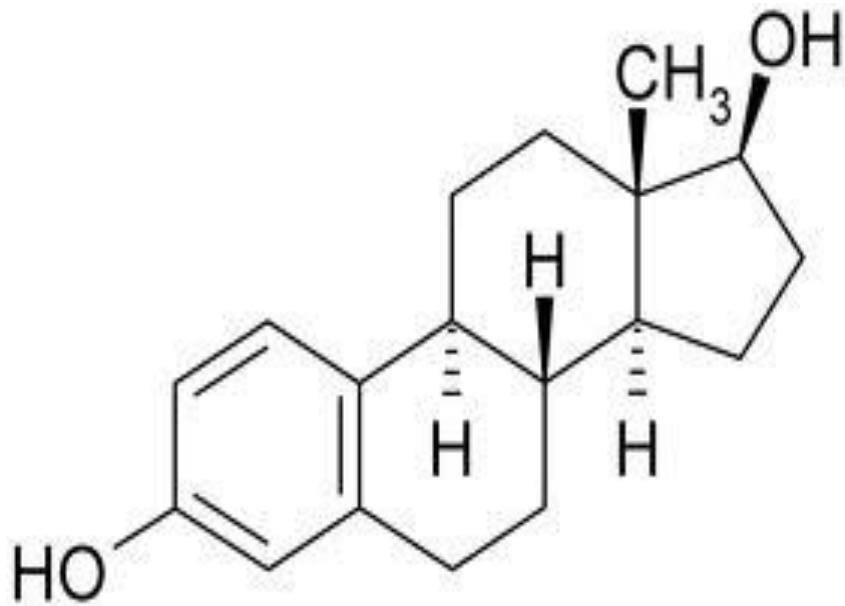
# Elizabeth Rogan, Ph.D.

## Phytochemicals and Estrogen Metabolism

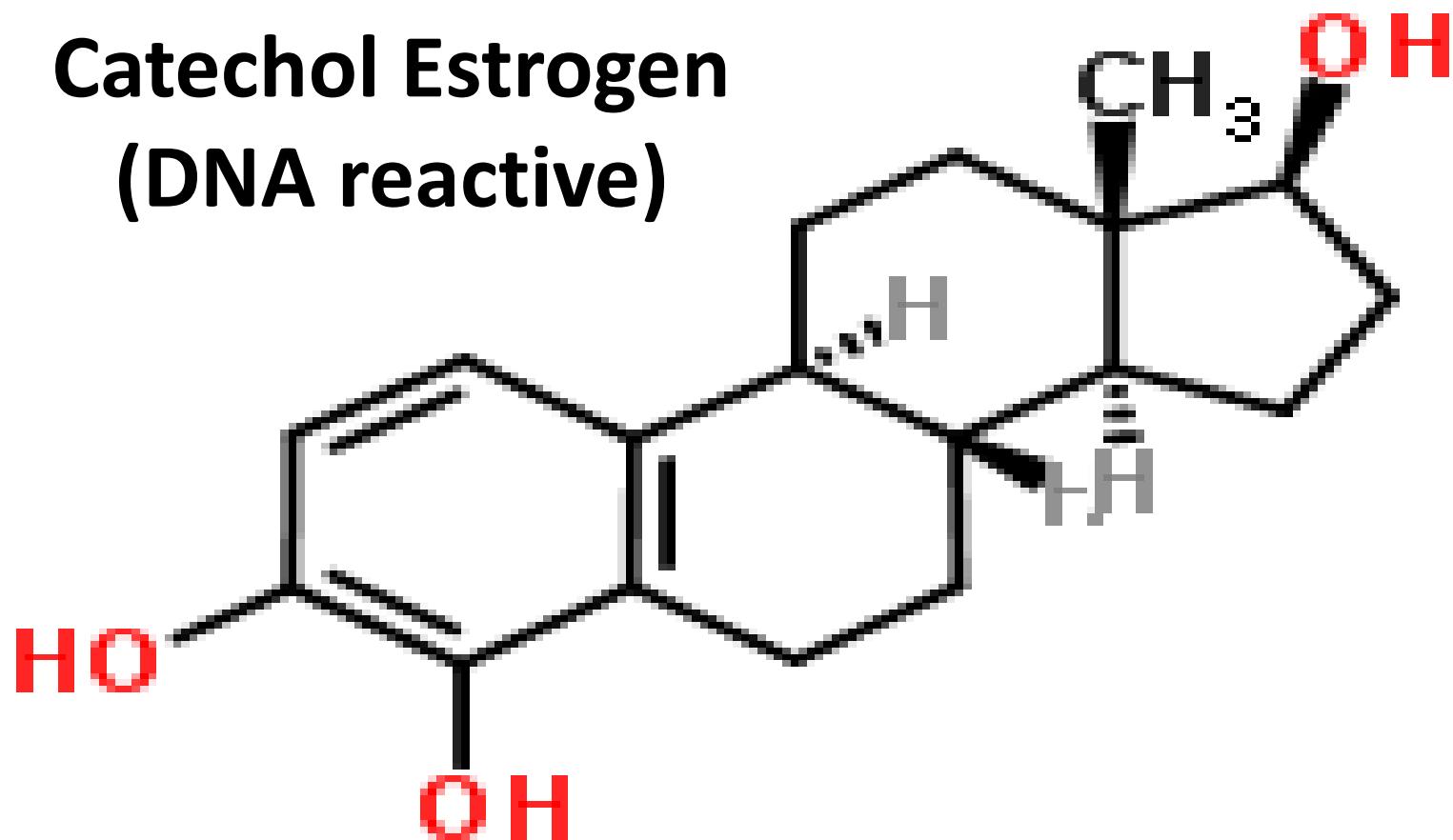


- PROFESSOR AND CHAIR
- Department of Environmental, Agricultural & Occupational Health
- College of Public Health
- Professor
- Eppley Institute for Research in Cancer and Allied Diseases
- University of Nebraska Medical Center

# Family of Estrogens

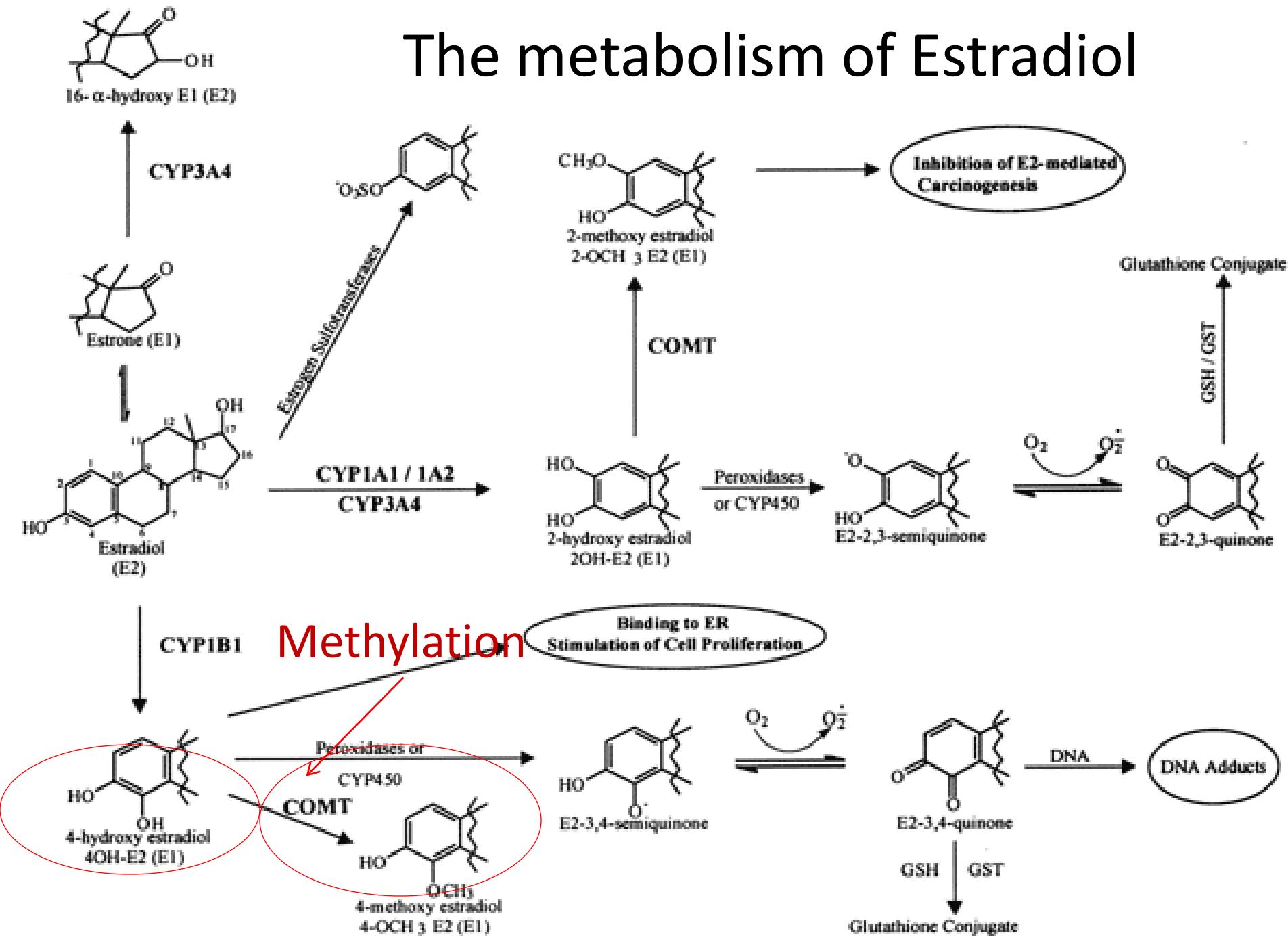


# Catechol Estrogen (DNA reactive)

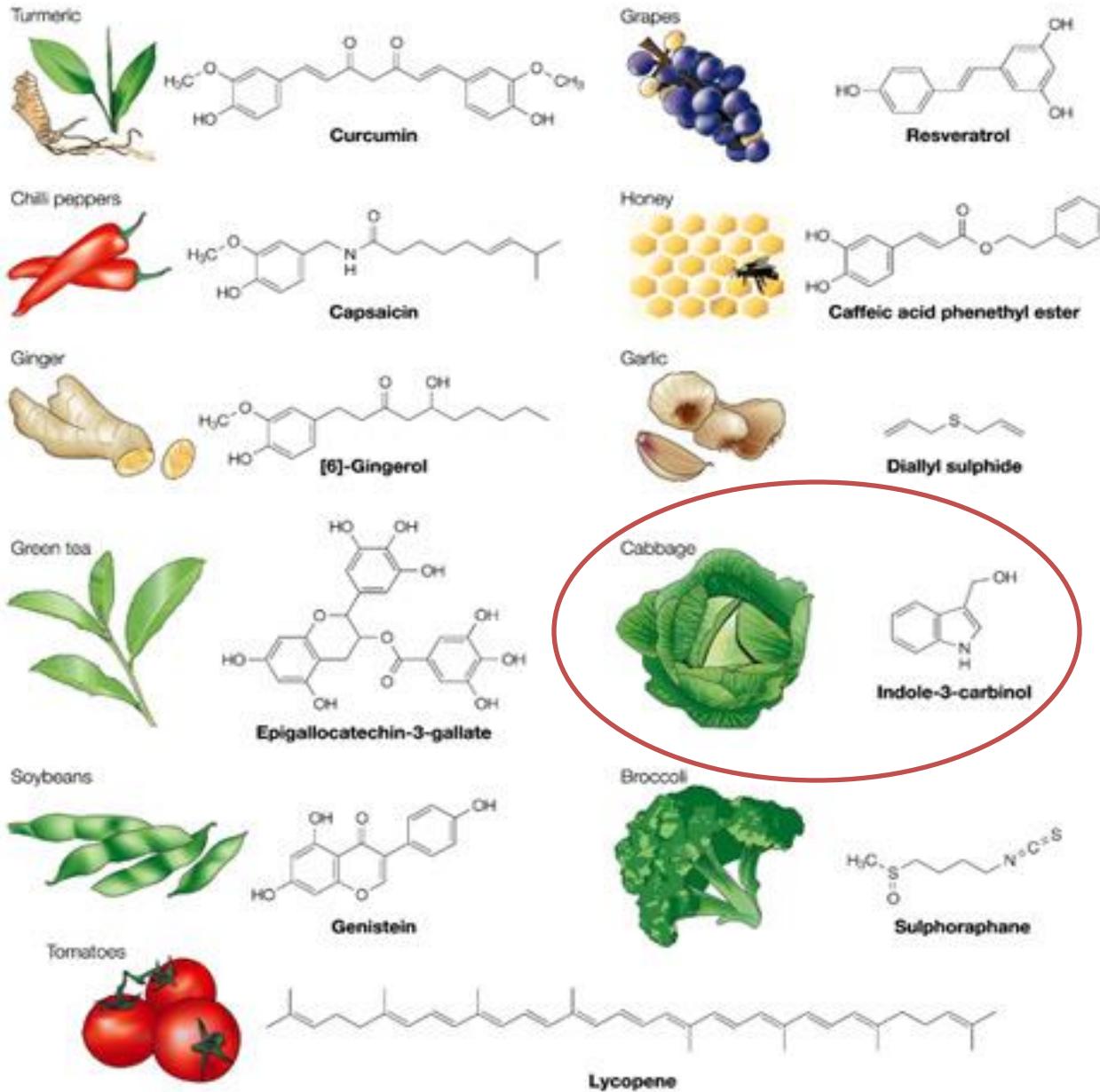


# 4-Hydroxyestradiol

# The metabolism of Estradiol



# Bioactive Phytochemical Components Modulating Estrogen



# Indole-3 Carbinol Intervention and the Risk to Cervical Cancer

- Gynecol Oncol. 2000 Aug;78(2):123-9.
  - **Placebo-controlled trial of indole-3-carbinol in the treatment of CIN**
  - “Thirty patients with biopsy proven CIN II-III were randomized to receive placebo or 200, or 400 mg/day I-3-C administered orally for 12 weeks”.
  - “None (0 of 10) of the patients in the placebo group had complete regression of CIN. In contrast 4 of 8 patients in the 200 mg/day arm and 4 of 9 patients in the 400 mg/day arm had complete regression based on their 12-week biopsy”.

# Nutrition and Phytochemicals for Dementia Neuroendocrineimmune Treatment

www.impactaging.com

AGING, September 2014, Vol 6 N 9

Review

## Reversal of cognitive decline: A novel therapeutic program

Dale E. Bredesen<sup>1,2</sup>

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<sup>2</sup> Buck Institute for Research on Aging, Novato, CA 94945.

**Key words:** Alzheimer's, dementia, mild cognitive impairment, neurobehavioral disorders, neuroinflammation, neurodegeneration, systems biology

Received: 9/15/14; Accepted: 9/26/14; Published: 9/27/14

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**Abstract:** This report describes a novel, comprehensive, and personalized therapeutic program that is based on the underlying pathogenesis of Alzheimer's disease, and which involves multiple modalities designed to achieve metabolic enhancement for neurodegeneration (MEND). The first 10 patients who have utilized this program include patients with memory loss associated with Alzheimer's disease (AD), amnestic mild cognitive impairment (aMCI), or subjective cognitive impairment (SCI). Nine of the 10 displayed subjective or objective improvement in cognition beginning within 3-6 months, with the one failure being a patient with very late stage AD. Six of the patients had had to discontinue working or were struggling with their jobs at the time of presentation, and all were able to return to work or continue working with improved performance. Improvements have been sustained, and at this time the longest patient follow-up is two and one-half years from initial treatment, with sustained and marked improvement. These results suggest that a larger, more extensive trial of this therapeutic program is warranted. The results also suggest that, at least early in the course, cognitive decline may be driven in large part by metabolic processes. Furthermore, given the failure of monotherapeutics in AD to date, the results raise the possibility that such a therapeutic system may be useful as a platform on which drugs that would fail as monotherapeutics may succeed as key components of a therapeutic system.



Alzheimer's & Dementia (2015) 1-8

Alzheimer's  
&  
Dementia

## MIND diet associated with reduced incidence of Alzheimer's disease

Martha Clare Morris<sup>a,\*</sup>, Christy C. Tangney<sup>b</sup>, Yamin Wang<sup>a</sup>, Frank M. Sacks<sup>c</sup>,  
David A. Bennett<sup>d,e</sup>, Neelum T. Aggarwal<sup>d,f</sup>

<sup>a</sup>Department of Internal Medicine and the Rush Alzheimer's Disease Center at Rush University Medical Center

<sup>b</sup>Department of Clinical Nutrition and the Rush Alzheimer's Disease Center at Rush University Medical Center

<sup>c</sup>Department of Nutrition, Harvard School of Public Health

<sup>d</sup>Department of Behavioral Sciences and the Rush Alzheimer's Disease Center at Rush University Medical Center

<sup>e</sup>Department of Neurology and the Rush Alzheimer's Disease Center at Rush University Medical Center

### Abstract

**Background:** In a previous study, higher concordance to the MIND diet, a hybrid Mediterranean-Dietary Approaches to Stop Hypertension diet, was associated with slower cognitive decline. In this study we related these three dietary patterns to incident Alzheimer's disease (AD).

**Methods:** We investigated the diet-AD relations in a prospective study of 923 participants, ages 58 to 98 years, followed on average 4.5 years. Diet was assessed by a semiquantitative food frequency questionnaire.

**Results:** In adjusted proportional hazards models, the second (hazards ratio or HR = 0.65, 95% confidence interval or CI 0.44, 0.98) and highest tertiles (HR = 0.47, 95% CI 0.26, 0.76) of MIND diet scores had lower rates of AD versus tertile 1, whereas only the third tertiles of the DASH (HR = 0.61, 95% CI 0.38, 0.97) and Mediterranean (HR = 0.46, 95% CI 0.26, 0.79) diets were associated with lower AD rates.

**Conclusion:** High adherence to all three diets may reduce AD risk. Moderate adherence to the MIND diet may also decrease AD risk.

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### Keywords:

Cognition; Alzheimer disease; Nutrition; diet; Epidemiological study; Aging

# The Nutrition Connection to the Neuroendocrineimmune Network

