



Microbiote, Obésité et atteintes Rénales

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Journée Mondiale du Rein

Obesity & Cardiometabolic disorders

- ❑ 192 Billion Euros / year EU (57% direct cost 21% loss of productivity , 22% indirect cost) *Leal J. et al, 2012, Economic Costs In: European Cardiovascular Disease Statistics).*
- ❑ Worldwide progression obesity & diabetes & CVD incl. Children
- ❑ Expected Tsunami
- ❑ Socio economic disparities
- ❑ Attrition in CVD & Metabolic drugs
- ❑ Need for new approaches !



6.5 MILLION
OBESE PEOPLE IN FRANCE

500 MILLION
OBESE PEOPLE WORLDWIDE

3 MILLION
PEOPLE WITH DIABETES IN FRANCE

17.3 MILLION
ANNUAL DEATHS WORLDWIDE
DUE TO CARDIOVASCULAR DISEASE

LEADING
CAUSE OF DEATH
WORLDWIDE

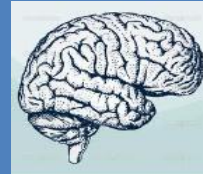
Obesity and kidney disease: Hidden consequences of the epidemic

Csaba P. Kovesdy, Susan L. Furth, Carmine Zoccali,
on behalf of the World Kidney Day Steering Committee ✉

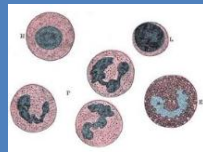
First published: 16 February 2017 [Full publication history](#)

.....Obésités : Maladie d'organes, Maladie des systèmes....

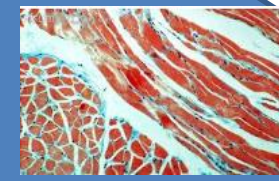
Modes de vie



Cerveau
Système endocrin
humeur



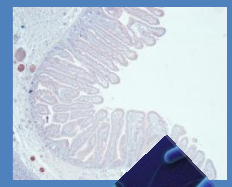
Système immunitaire
Production cytokine



Muscle
Fonction
Endocrine



Coeur

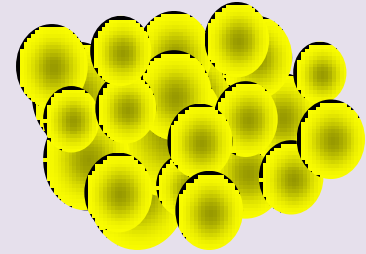


Intestin
Microbiote

Hypertrophie

Fonctions
Endocrines
perturbées

altérations
Immuno
inflammatoires



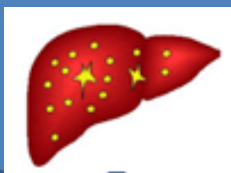
Fibrose

Angiogenese

Dysfonction
Organelle



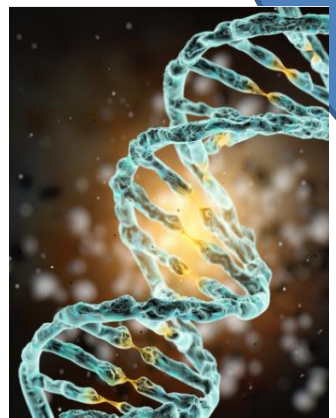
Système
Vasculaire
Atherothrombo



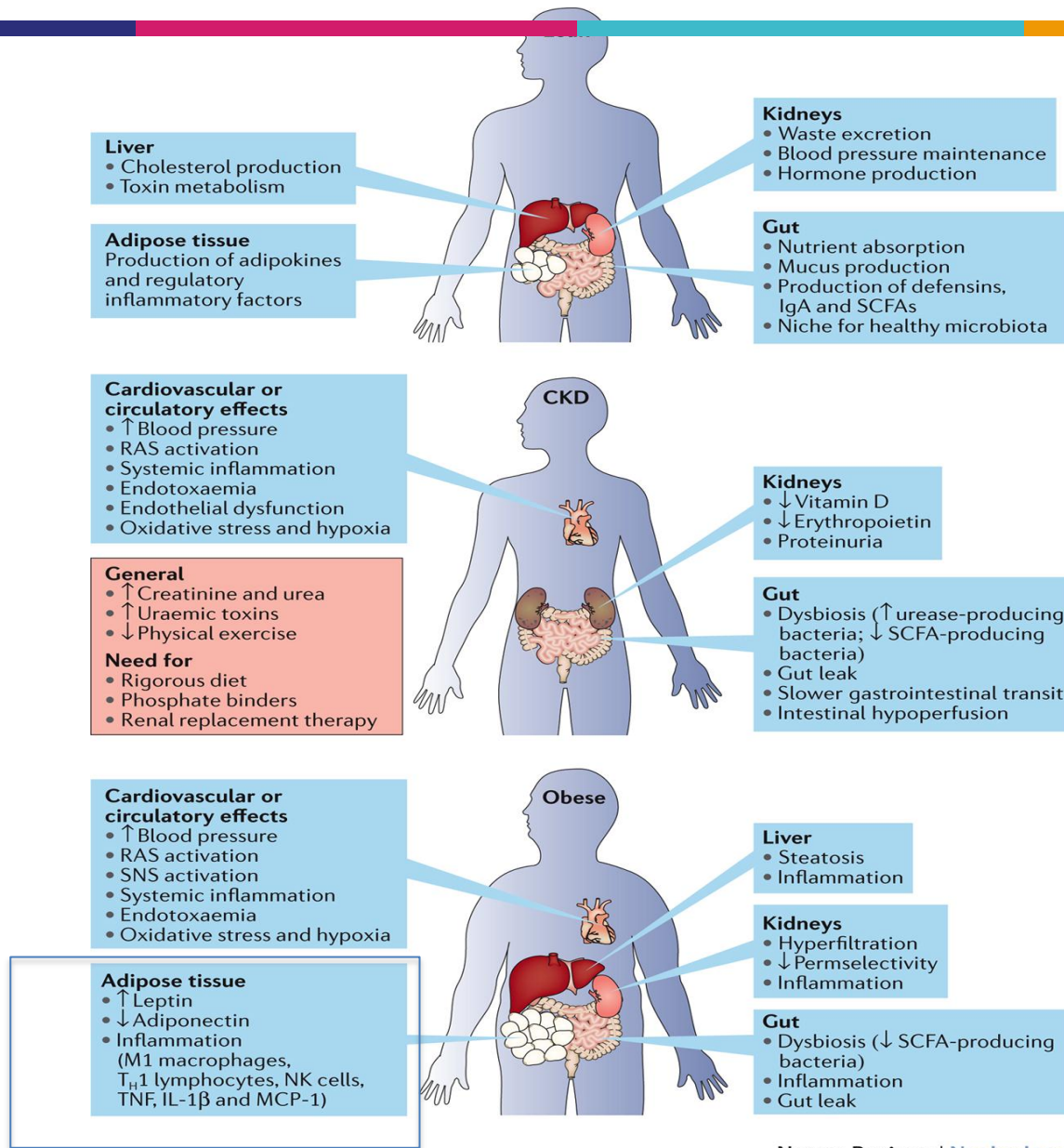
Foie
NAFLD/NASH



Génome/Epigénome

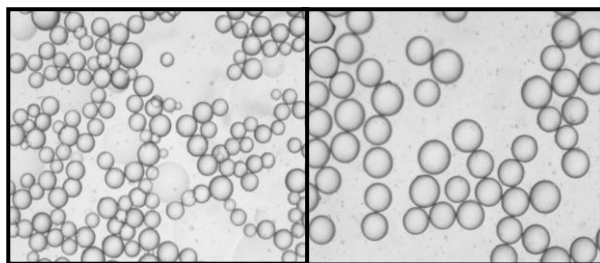


The fat–intestine–kidney axis



Nature Reviews | Nephrology

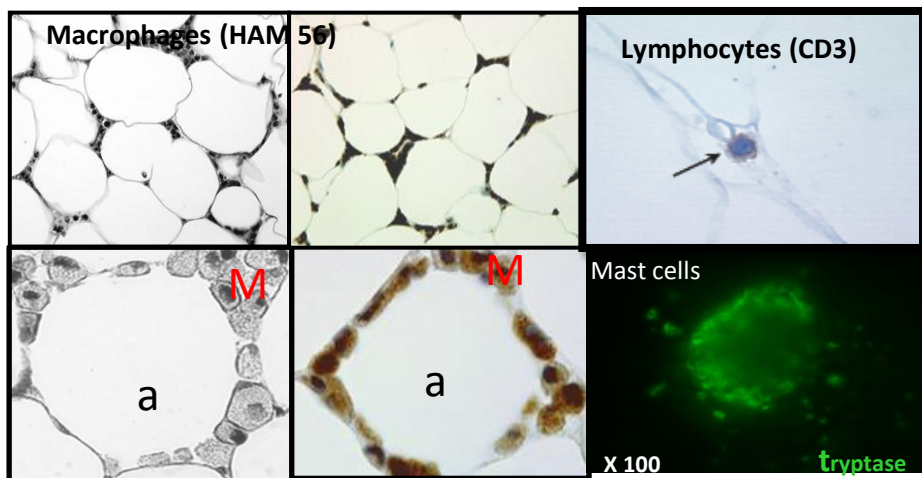
Adipocyte hypertrophy



Lean

Obese

Inflammatory cell accumulation

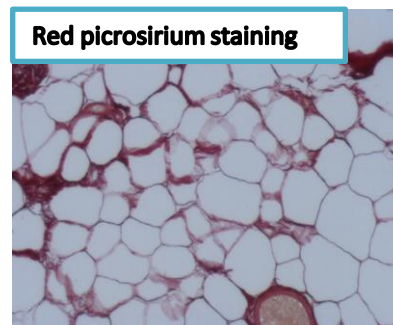


Macrophage (M) accumulation Other immune cells

Cancello, Tordjman, Diabetes 2010 U872

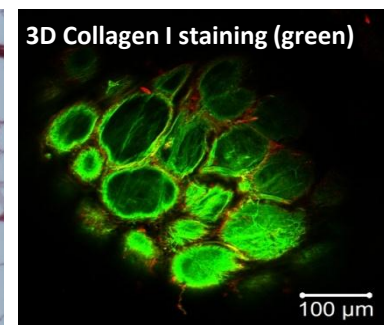
Liu, Divoux Nature Medicine, 2009

Fibrosis (pericellular)



Red picrosirium staining

Divoux, Diabetes 2010

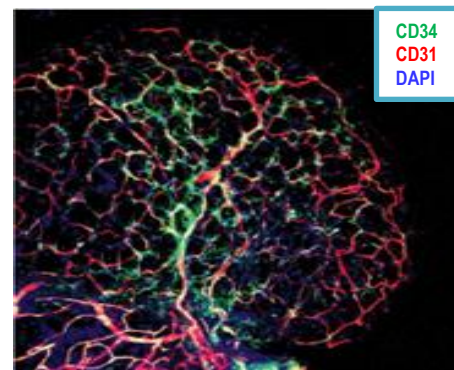


3D Collagen I staining (green)

Pellegrinelli, U872

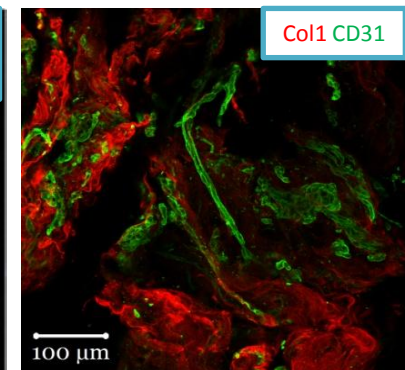
100 μm

Vascular alterations (inflammation & senescence)



CD34
CD31
DAPI

Villaret, Diabetes 2010, U



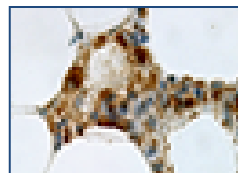
Col1 CD31

100 μm

Pellegrinelli, U872

Local interactions

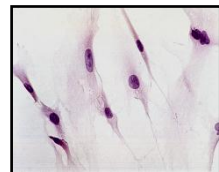
Macrophages
M1/M2 state
Polarization
Ex. *KLF4* (Coll)
J Clin Invest 2011



TNF α , IL-6
CCL5/Rantes
CXCL2

Keophipath, ATVB. 2010
Rouault, Endocrinology, 2013

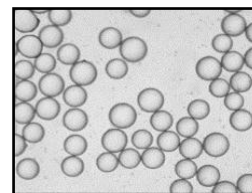
NEFA?



Proliferation,
Migration,
Inflammation
Profibrotic

Profibrotic progenitors

Lacasa, Endocrinology 2007
Keophipath, Mol Endoc 2009
Marcellin Cell Metab 2017



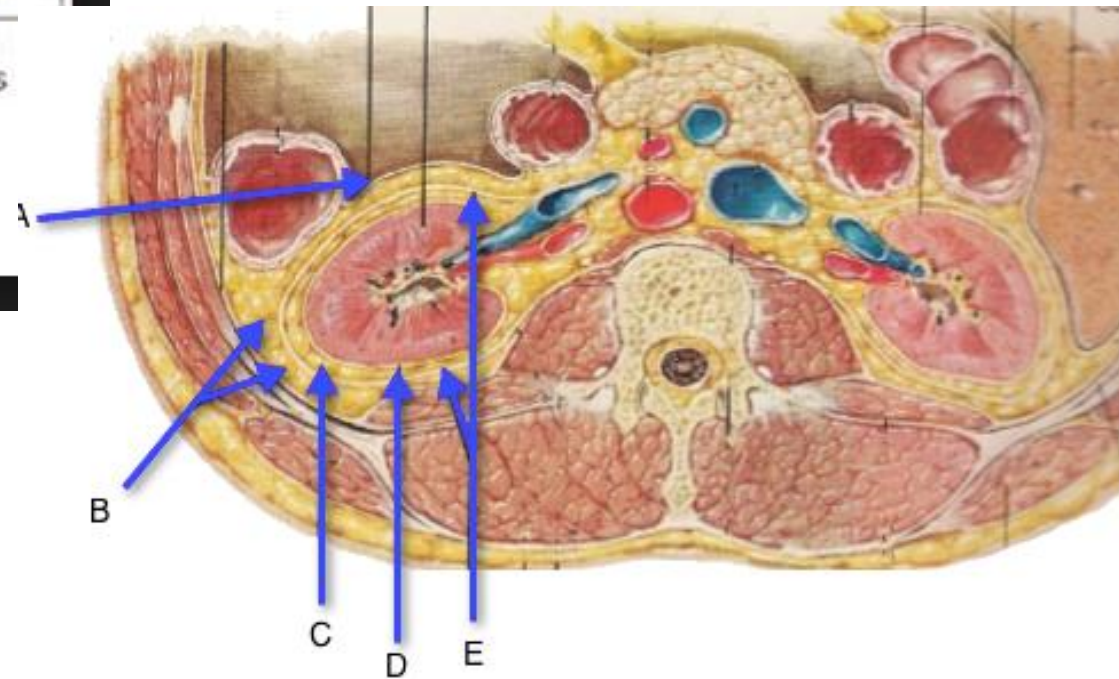
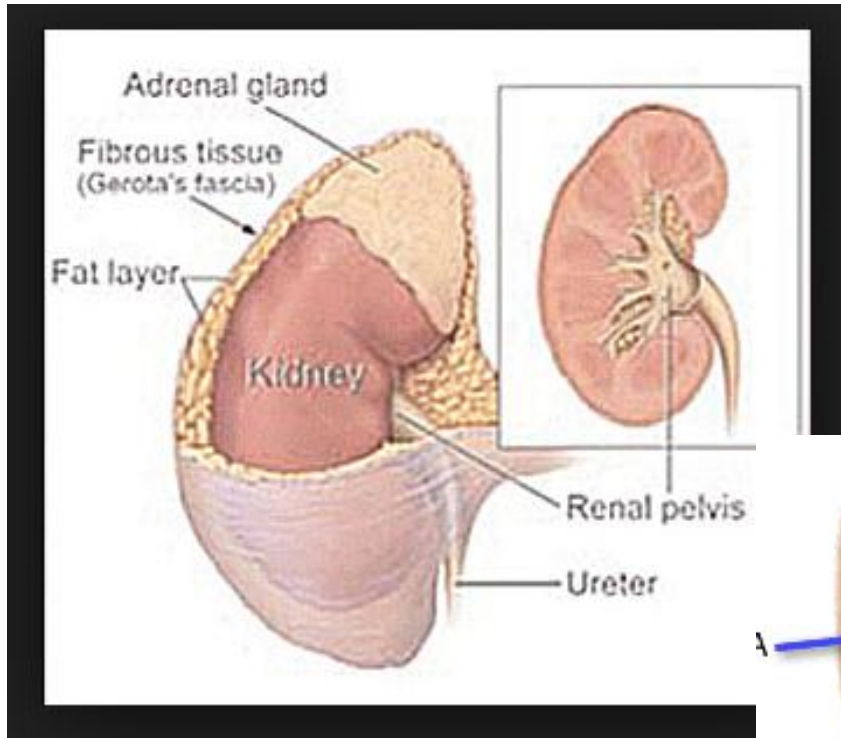
Organelle Dysfunctions
Lysosomes etc..
Soussi (Diabetes, 2016)

Inflammation
GPS2 (A Toubal, JCI, 2013)

Metabolic perturbations
Glucose uptake
Insulin sensitivity

Pellegrinelli, J Pathol 2014

Fat Layer around Kidney



Obesity

Insulino-resistance

Accumulation of macrophages in omental AT & deep scWAT

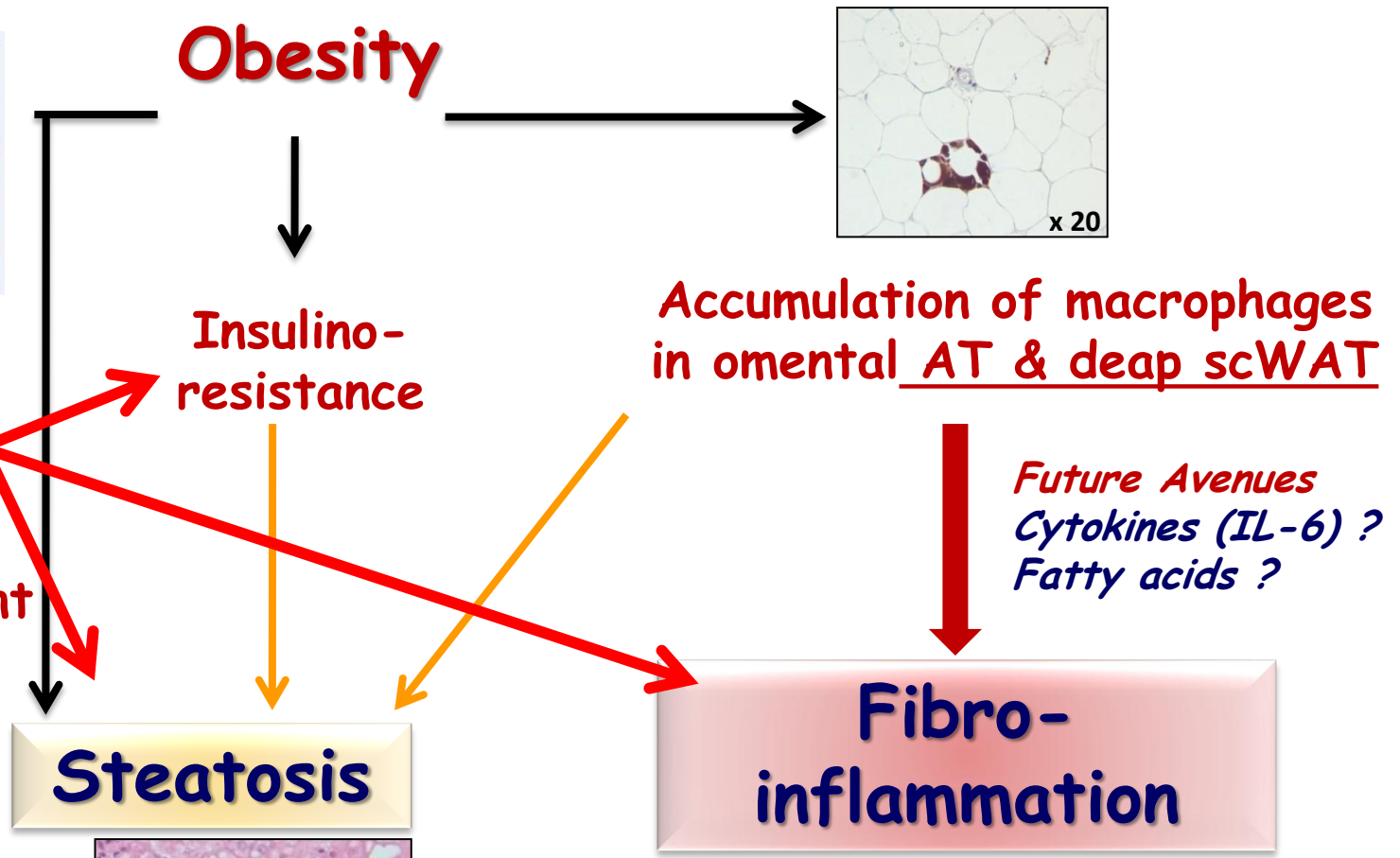
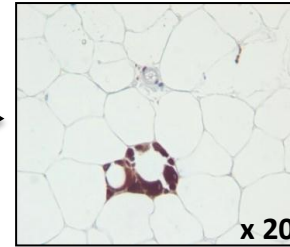
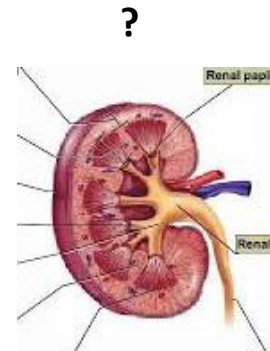
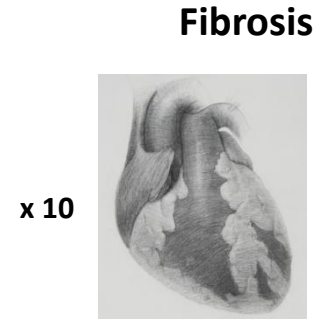
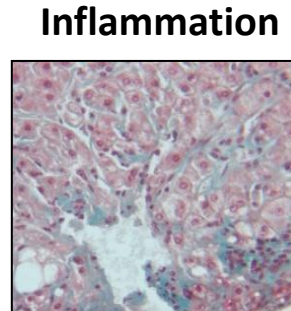
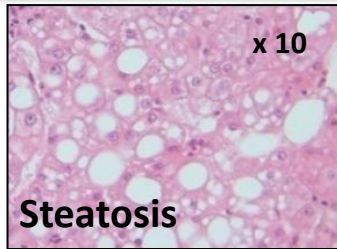
*Future Avenues
Cytokines (IL-6) ?
Fatty acids ?*

Chronic & intermittent Hypoxia

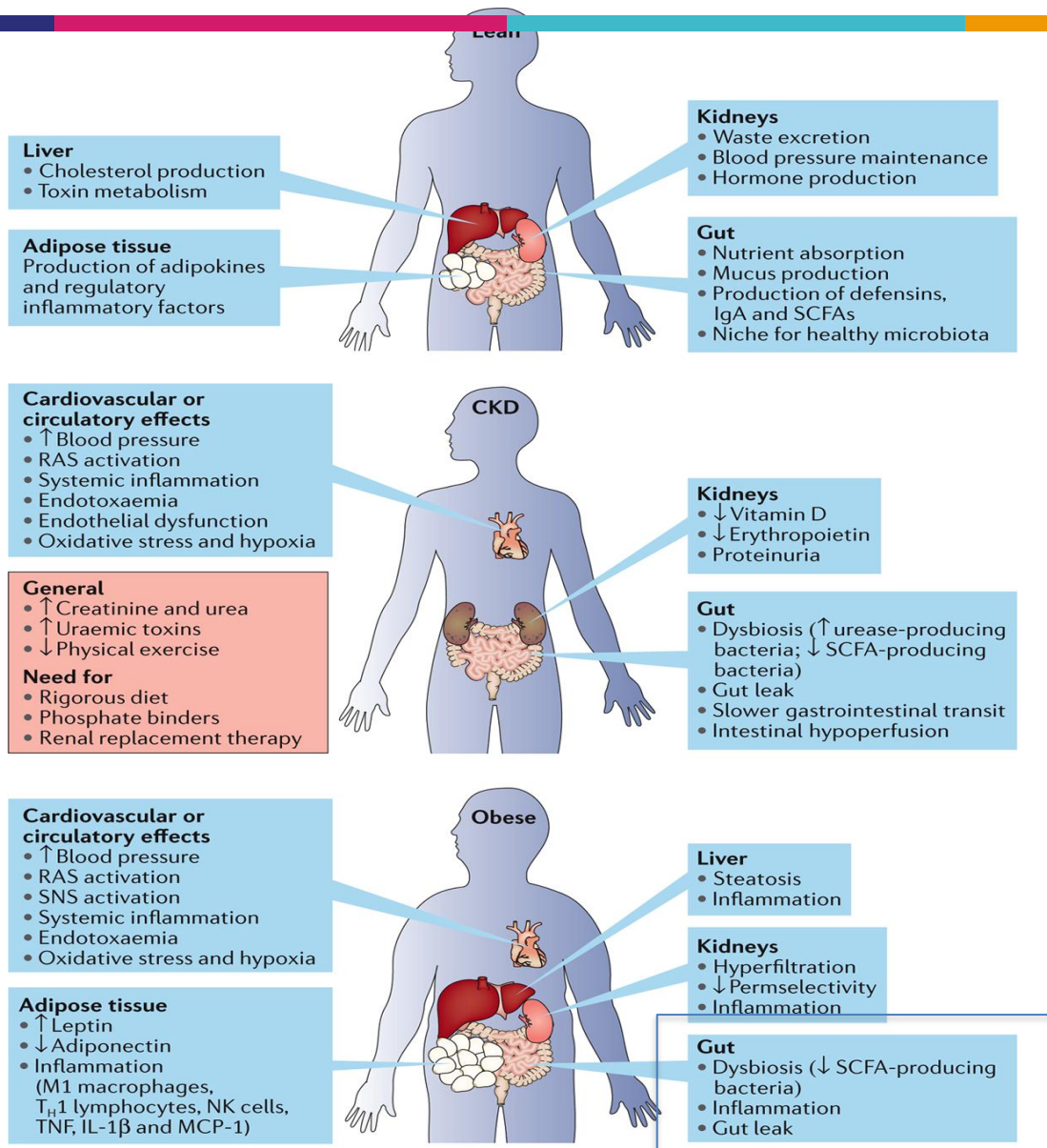
Steatosis

Fibro-inflammation

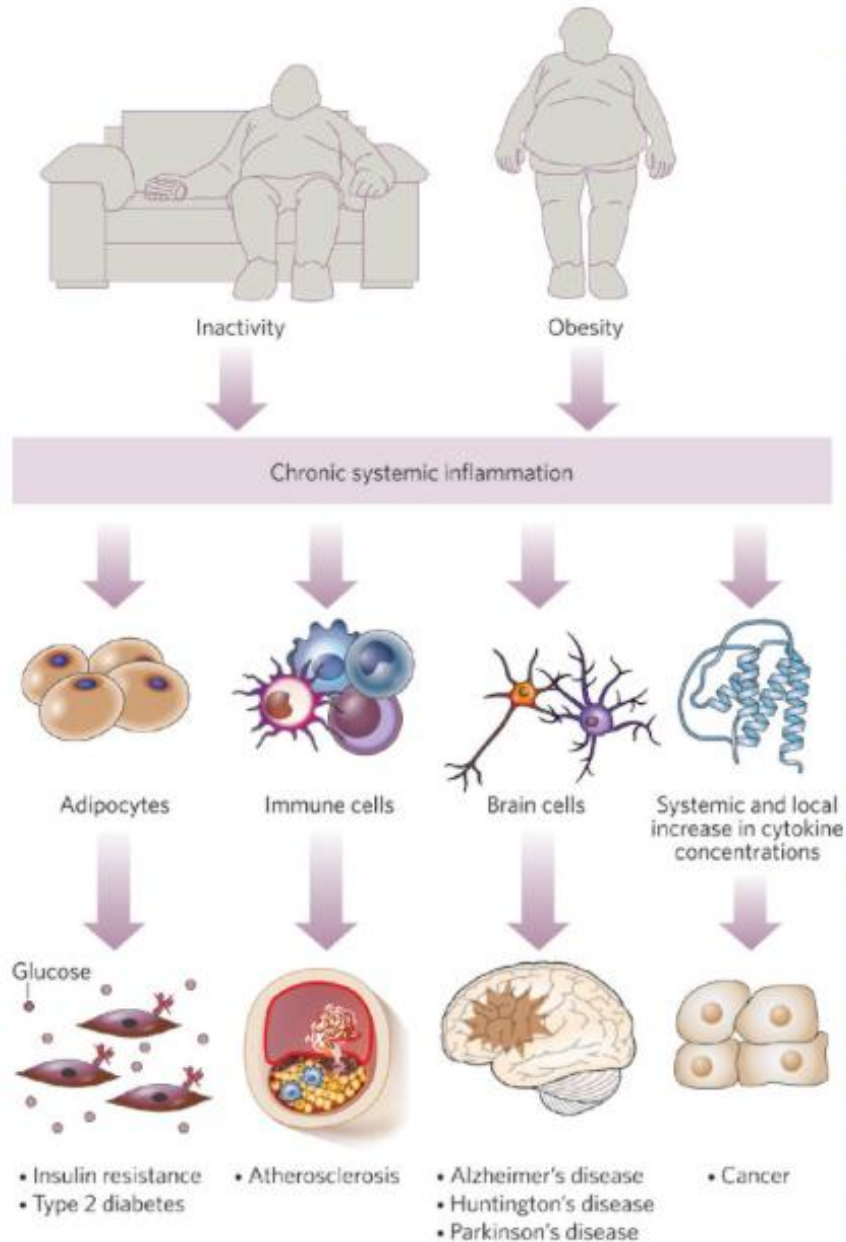
- Cancello Diabetes 2006
- Tordjman J Hepatol 2009
- TAM Diabetes 2011
- Tordjman J Hepatol 2012
- Bedossa Hepatology 2012
- Aron J Hepatol 2012
- Venteclef Eur. Heart J 2015
- Bedossa Gut, 2016



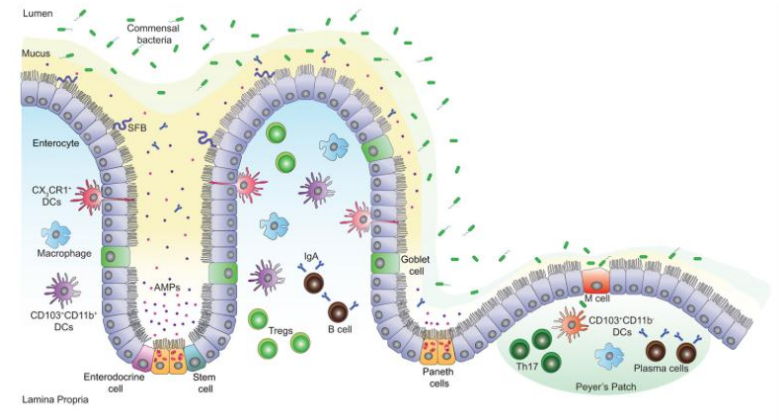
The fat–intestine–kidney axis



Nature Reviews | Nephrology



What about intestinal barrier?



- Largest lymphoid tissue in size of the body: complex immune system
- Dysbiosis of gut microbiota in obesity (*Cotillard et al., Nature 2014*).
- **Gaps in knowledge in obesity-related inflammation of the jejunum**

Summary of findings

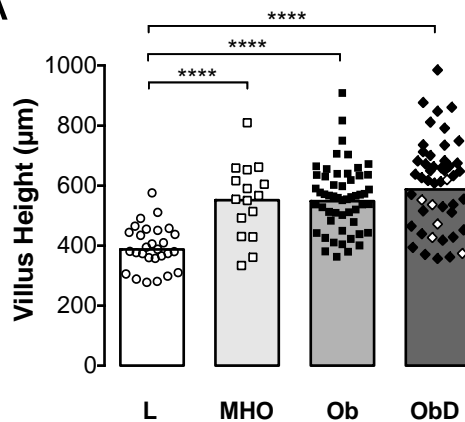


Increased surface of absorption

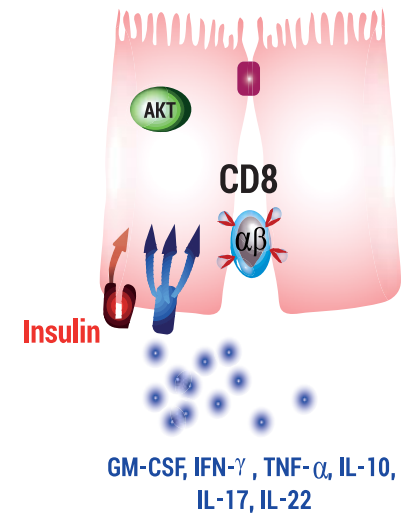
More inflammation (jejunum)

Altered function (insulin resistance)

A

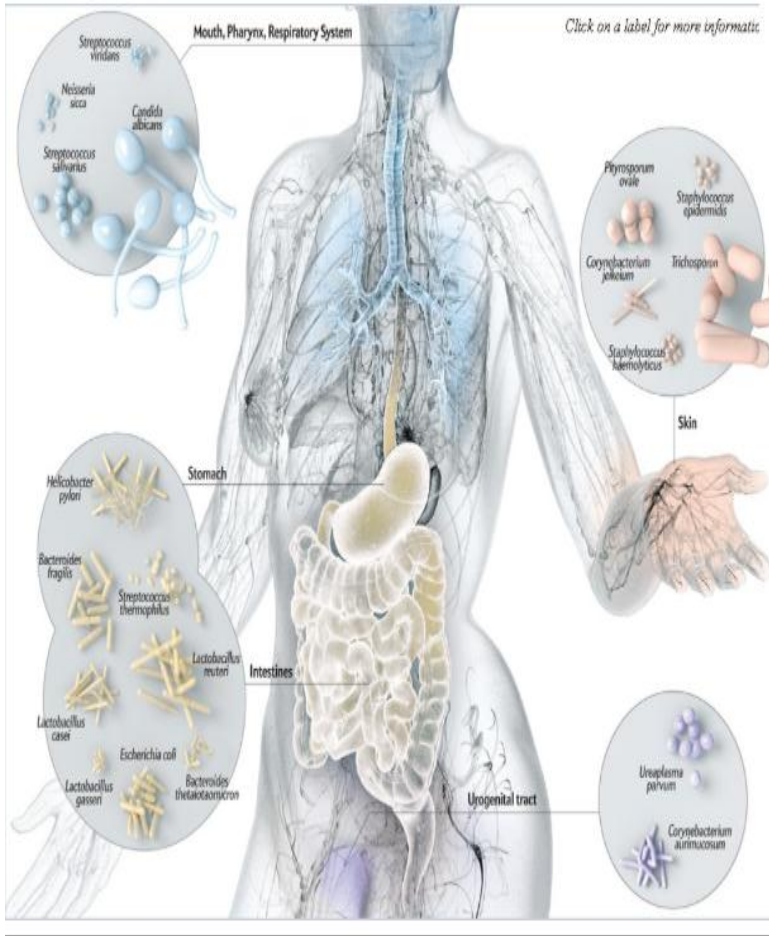


	n/mm ²	Ob vs. Lean
Mature DC	5	x 5
NK cells	12	x 6
Macrophage	150	x 1.5
CD8αβ		
n/mm ²	Ob vs. Lean	
Lamina propria	350	x 1.75
Epithelium	1200	x 1.5



Links with obesity & complications

Corpulence, liver disease and dyslipidemia



Microbiota as an asset

- Defense - bacterial antagonism
- Priming of mucosal immunity
- Peristalsis
- Metabolism of dietary carcinogens
- Synthesis of B & K vitamins
- Epithelial nutrients (e.g. SCFAs)
- **Degradation of Dietary Oxalate++**
- Conversion of prodrugs
- Utilisation of indigestible (CH_2O)_n

Microbiota as a liability

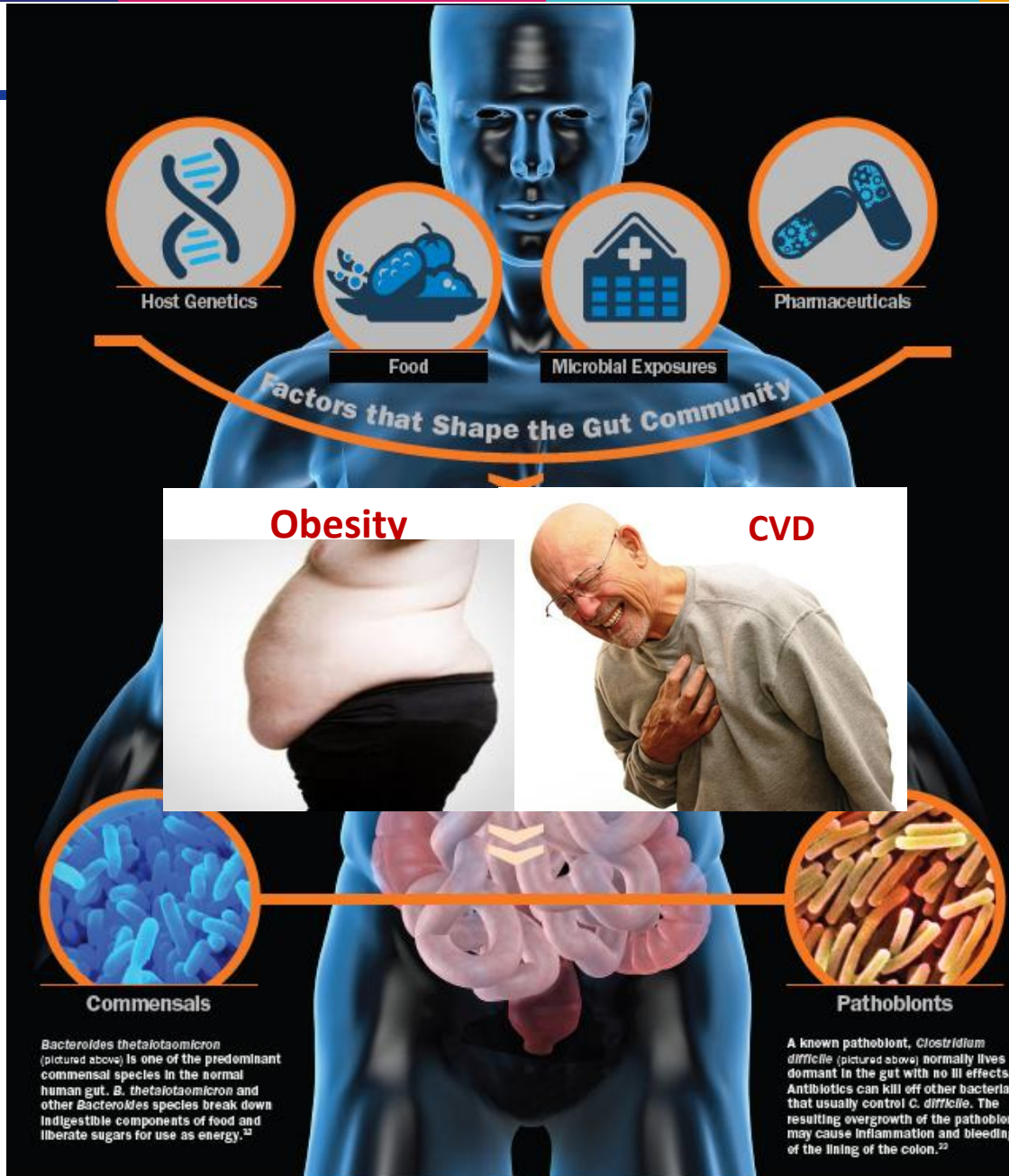
- Procarcinogens → carcinogens
- Overgrowth syndromes
- Opportunism – Translocation
- Implicated in obesity, metabolic syndrome and colorectal cancer, IBD and other diseases (CKD...)



Kidaura *et al.*
Science 2013

➤ **Gut Microbiome** could be one of the causes of **Obesity**

Equilibre entre santé et maladie



Bacteroides thetaiotaomicron (pictured above) is one of the predominant commensal species in the normal human gut. *B. thetaiotaomicron* and other *Bacteroides* species break down indigestible components of food and liberate sugars for use as energy.²²

A known pathobiont, *Clostridium difficile* (pictured above) normally lives dormant in the gut with no ill effects. Antibiotics can kill off other bacteria that usually control *C. difficile*. The resulting overgrowth of the pathobiont may cause inflammation and bleeding of the lining of the colon.²²

Fibromyalgia

IBD

Diabetes

Colon Cancer

T1 & T2

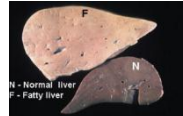
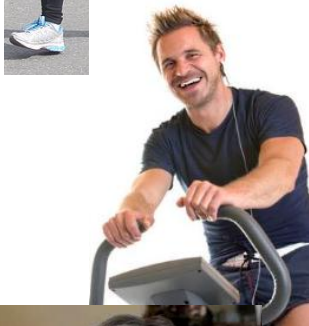
NASH

CVD

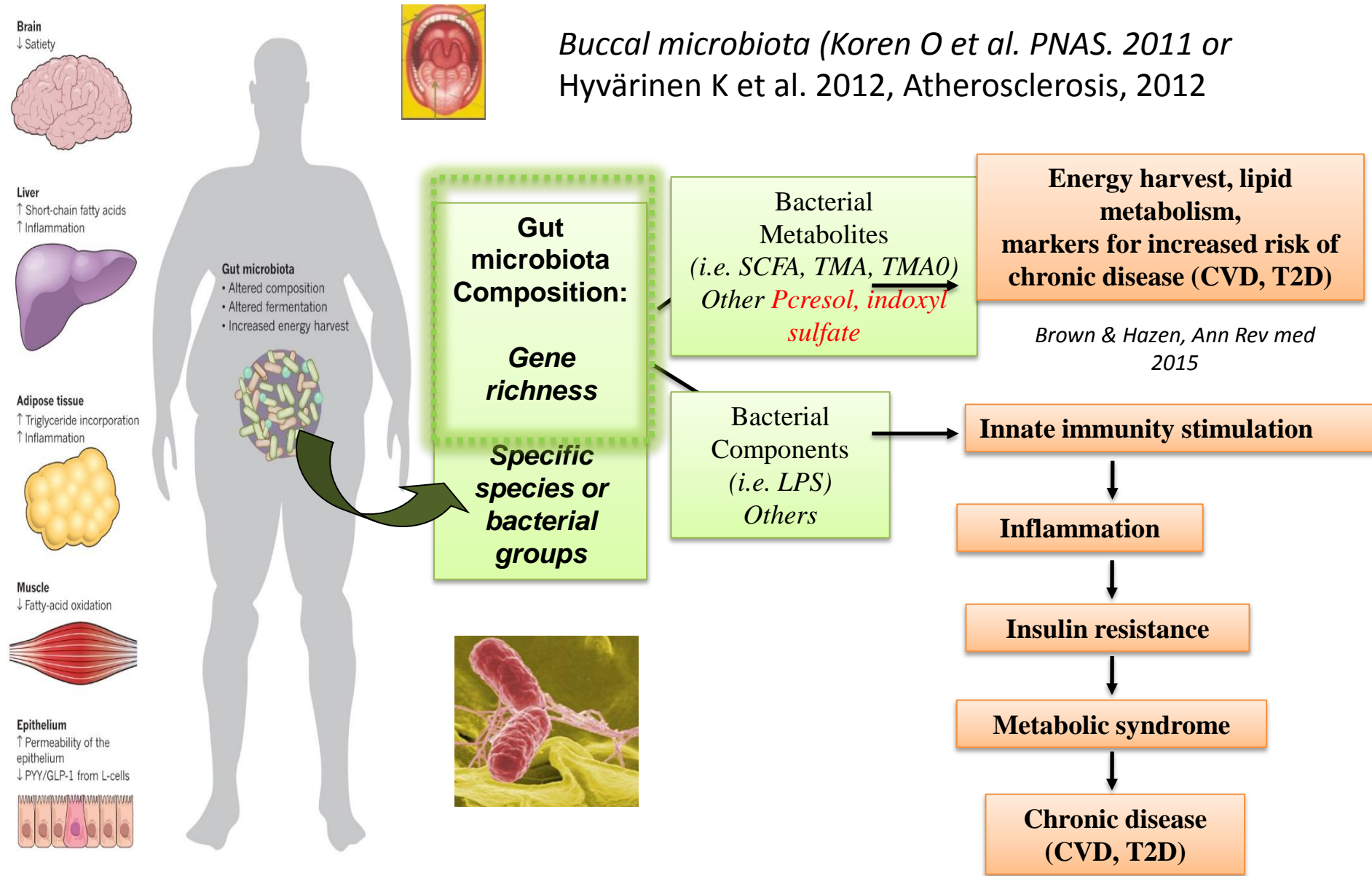
Depression

Kidney stones

Atopic disease



Potential role for microbiota in the development of cardiometabolic (&Kidney) diseases



Buccal microbiota (Koren O et al. PNAS. 2011 or Hyvärinen K et al. 2012, Atherosclerosis, 2012)

Gut Microbiome and Obesity: Diversity matters

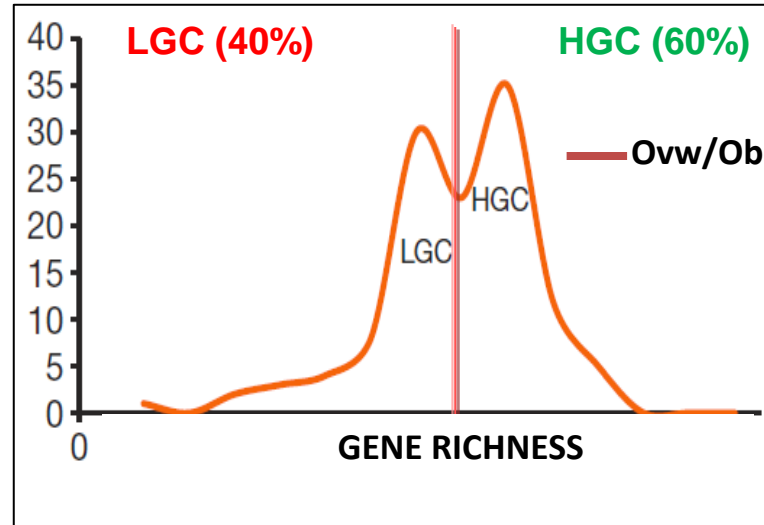
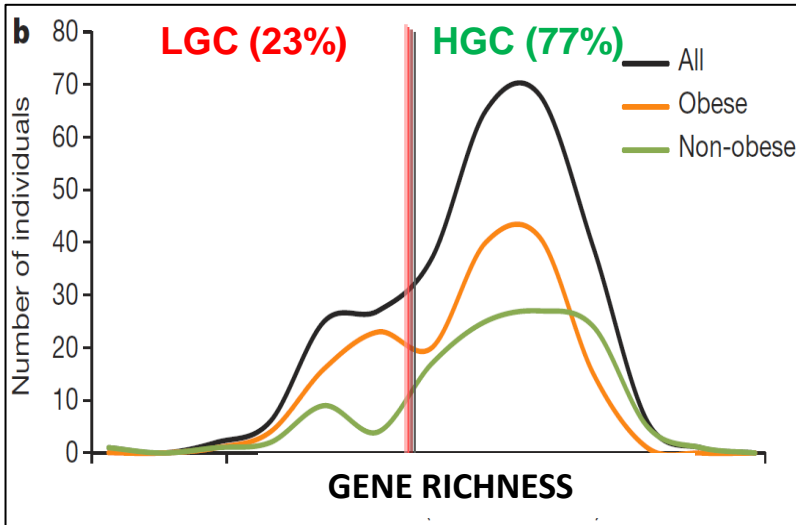


METAHIT

292 subjects

MicroObes

49 subjects



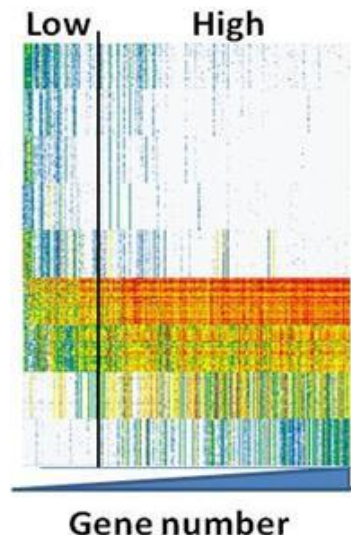
LGC = Low gene count

HGC = High gene count

LGC: ↗ Pro-inflammatory

LGC: ↘ Anti-inflammatory

- Cl. bolteae*
- Cl. symbiosum*
- Cl. clostridioforme*
- Cl. ramosum*
- R. gnavus*
- F. prausnitzii*
- R. inulinivorans*
- Co. eutactus*
- M. smithii*



LGC associates with CMD risks

- ↑↑ dyslipidemia
- ↑ adiposity
- ↑ insulin resistance (surrogates)
- ↑ inflammation (circulating and adipose tissue)



49 obese or overweight patients

Diet: High fiber and protein, low carbohydrate index

49 subjects

Energy-restricted diet
(1200-1500 kcal)

Body weight
maintaining diet

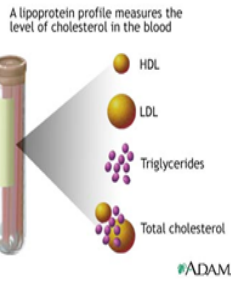
0wk

6wks

12wks



Diet



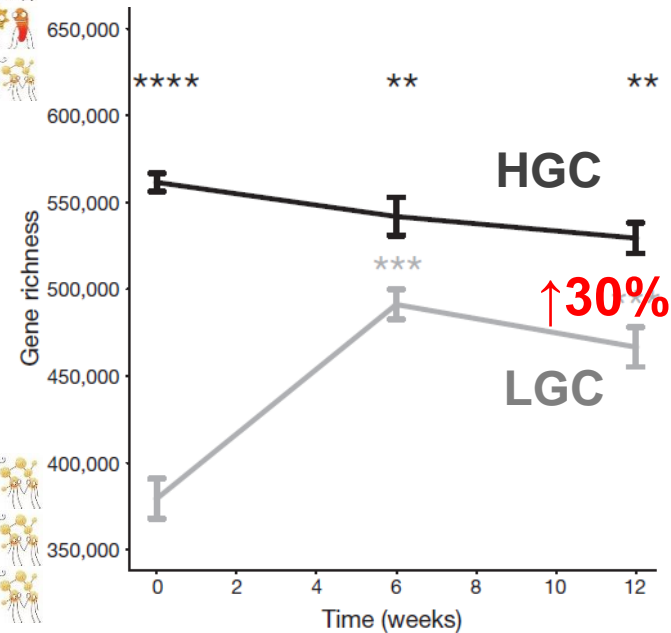
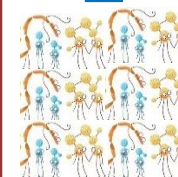
Phenotype



Gut flora*

* SOLiD™ reads were mapped on a 3.3 million genes catalog [1]

[1] Qin J. et al., g. Nature, 464 7285:59-65, 2010.



Cotillard *et al.* Nature 2013

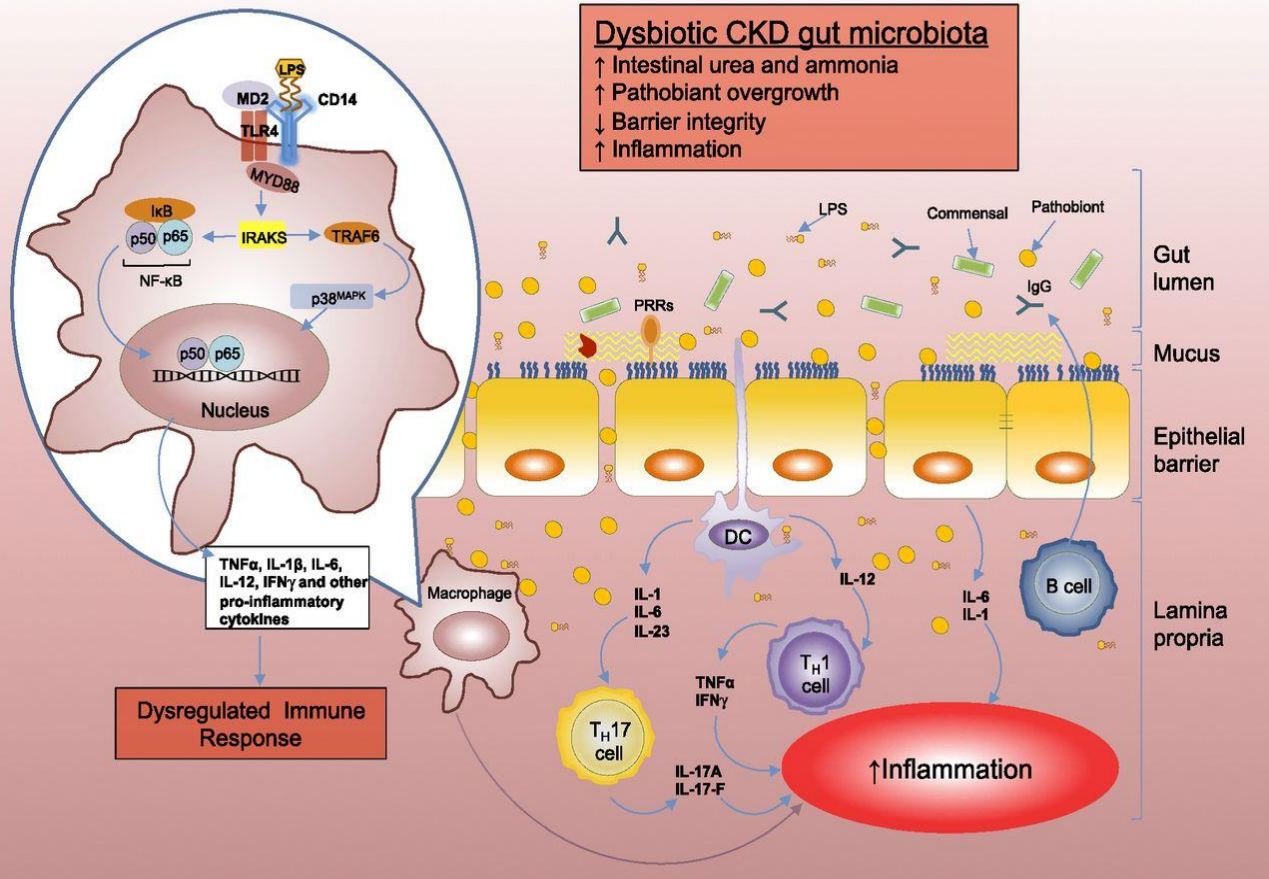
Qualitative and quantitative alterations in CKD ?

- ESRD patients: 190 OTU phyla Firmicutes (subphylum Clostridia), Actinobacteria & Proteobacteria (**Gammaproteobacteria**) Vaziri ND, Kidney Int (2013)
- Nephrectomized rats: 175 OTU. Dec Bacteroidetes & Firmicutes, *especially Lactobacillaceae and Prevotellaceae*
- ESRD vs. healthy : enrichment in **bacteria producing urease and uricase and less SCFA** (Wong J, Am J Nephrol. 2014)
- CKD : more Firmicutes > bacteroidetes (Barros AF, 2015)
- ESRD patients (Pyrosequencing) > *Klebsiella, Proteus, Escherichia, and Pseudomonas* (Wang et al,)
- **Lower richness species; lactobacillus & prevotellaceae** (Vaz
- **Bacterial translocation ?** / LPS (Wang et al, 2012)



Bidirectional dialogue

B



Bacteria release uremic toxins (cleared by the Kiney)

.....

Others
 Dietary fibers
 Antibiotics
 Slow transit
 Acidocetosis
 Oral iron

.....

Ali Ramezani, and Dominic S. Raj JASN 2014;25:657-670





Gut microbiota

Composition

Microbial richness

Specific species (i.e. *A. muciniphila*)

Metabolites

TMAO

Bacterial components

Cell wall, *LPS*

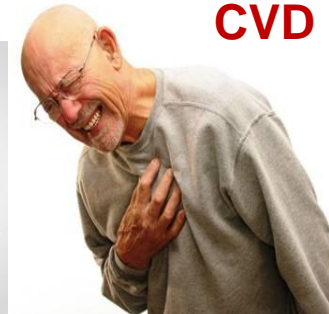


Obesity and co-morbidities

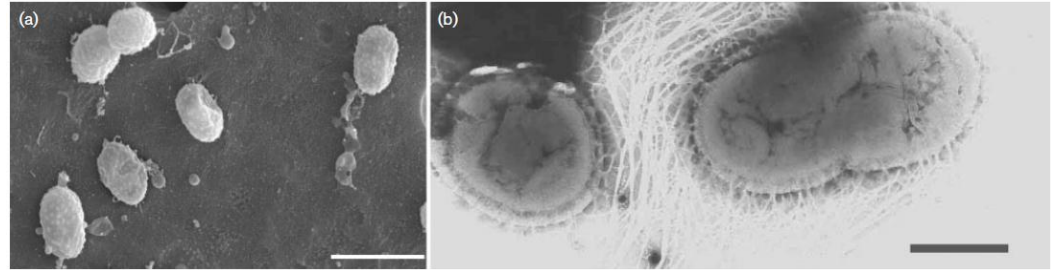
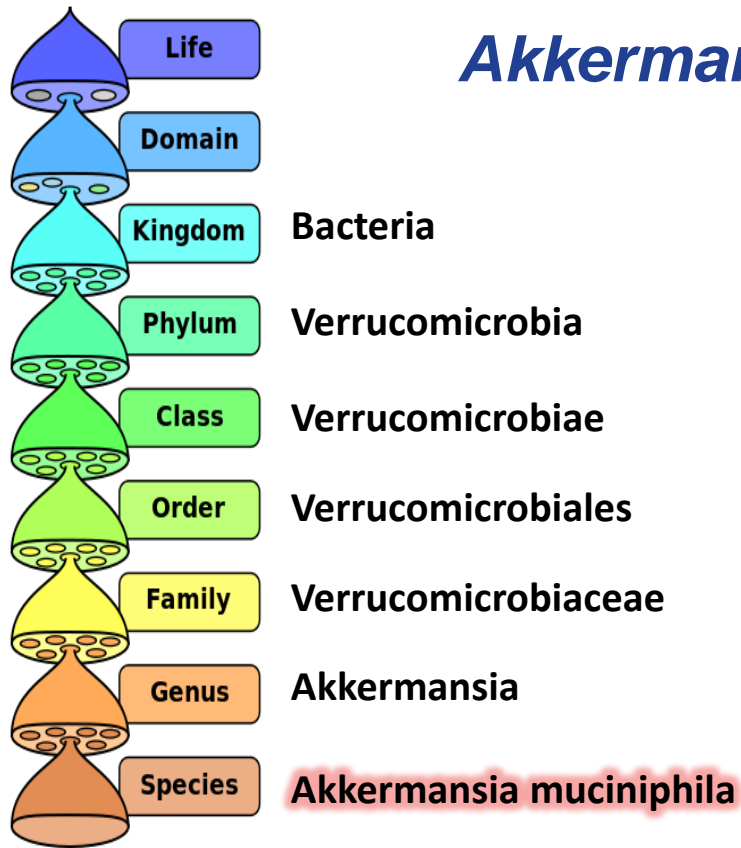
Obesity



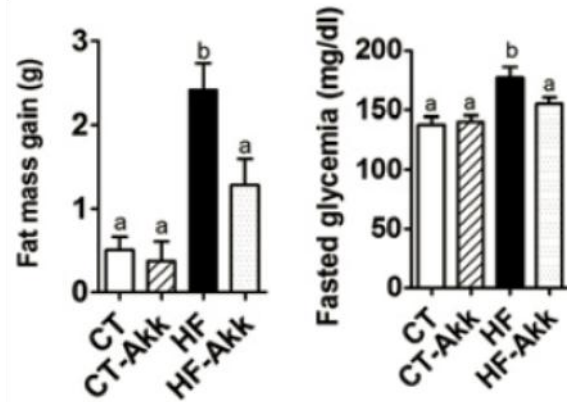
CVD



Akkermansia muciniphila (Akk)

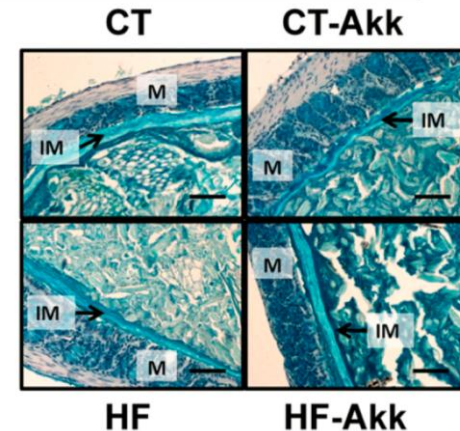


Derrien *et al.* IJSEM 2004



Health implications

- **Mouse studies: maintenance of glucose homeostasis.** *Everard et al. PNAS 2013*
- **Links with Metformin.** *Shin et al. Gut 2013*
- **Inverse association between Akk and iR humans (more clear in mice).**



Everard *et al.* PNAS, 2013

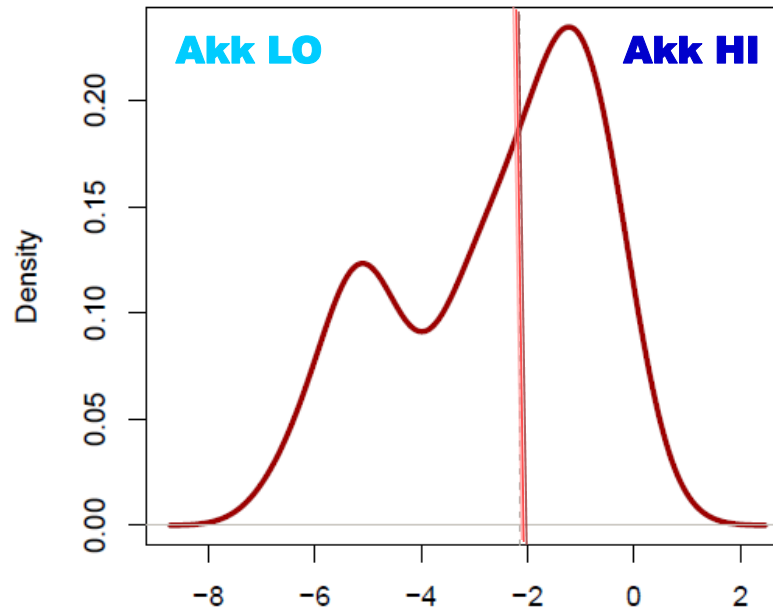
AIM: Study the association between *A. muciniphila* abundance, metabolic status, and microbial richness



T0

49 overweight
and obese adults

Akk abundance distribution



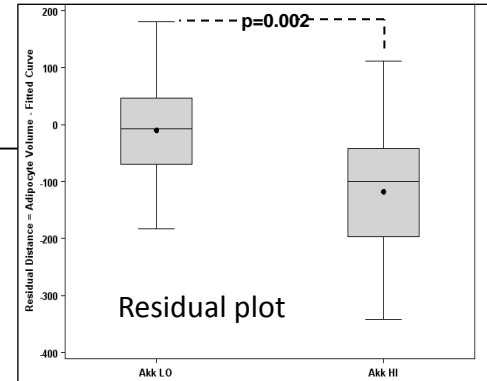
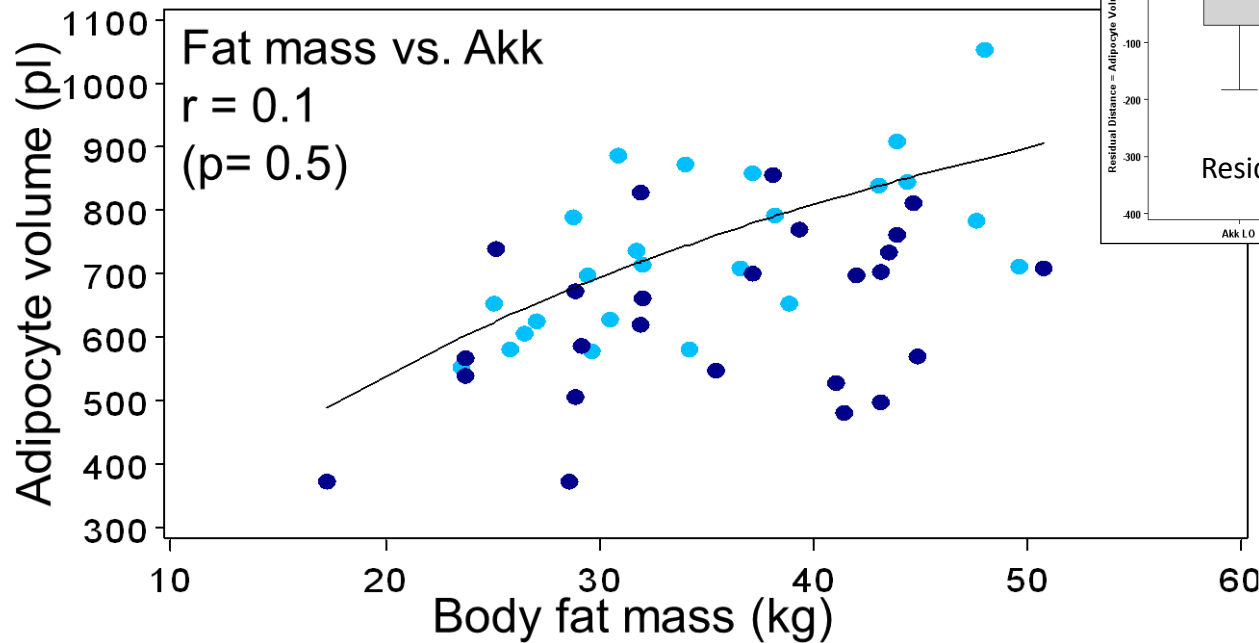


Higher **baseline** *A. muciniphila* is associated with a healthier metabolic status





Subjects with higher *A. muciniphila* have smaller adipocytes



 Akk LO

 Akk HI

$$\text{Theoretical Adipocyte volume (pl)} = \frac{(40.7 * \text{Kg Fat Mass})}{(1 + (0.025 * \text{Kg Fat Mass}))}$$

Methodology:





Hirsch et al. Lipid Res, 1968

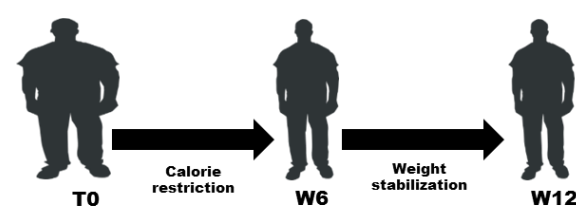
Spalding et al. Nature, 2008

Dao et al. Gut 2015

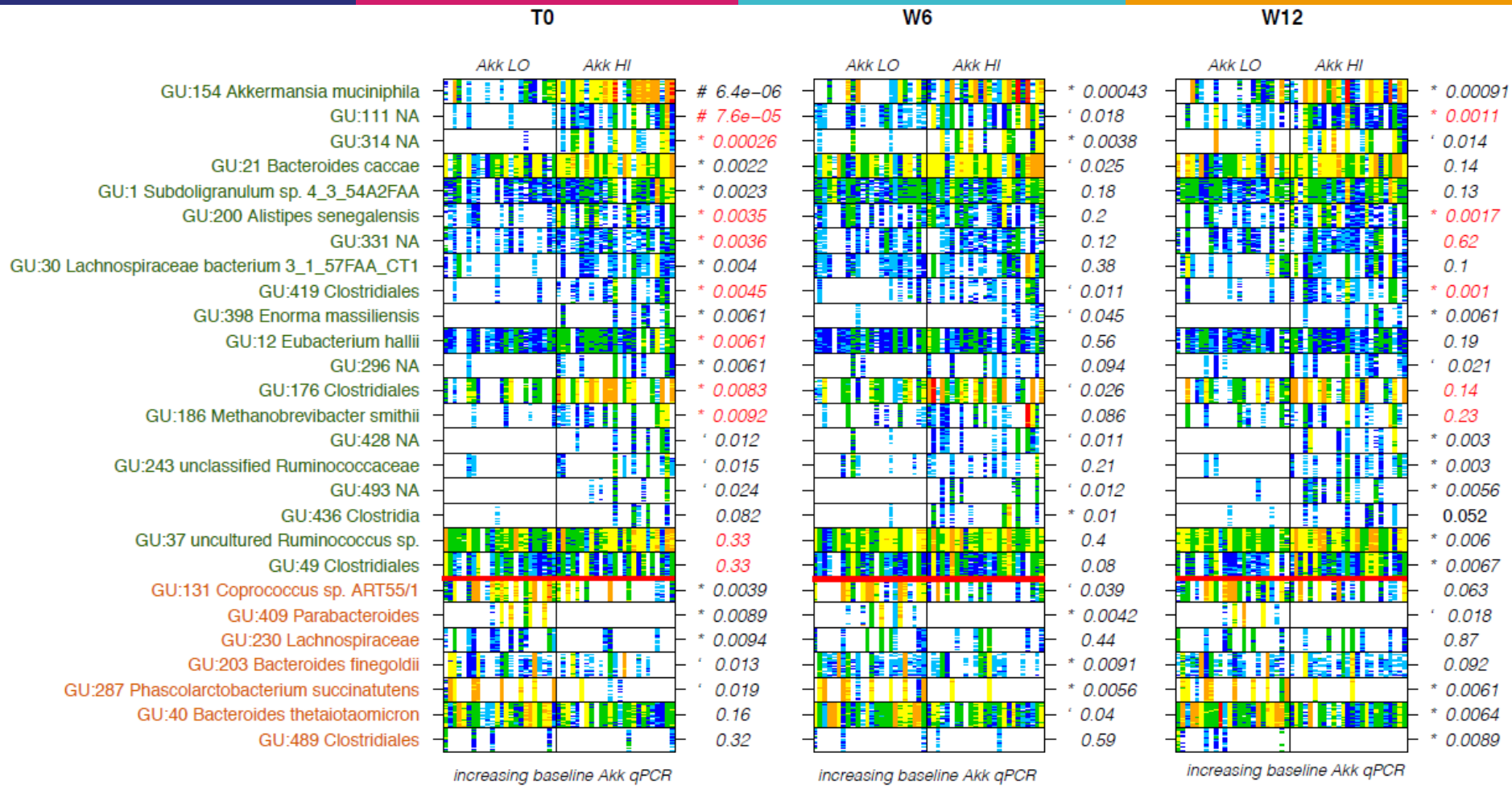


Subjects with **higher** *A. muciniphila* abundance and gene richness have **healthier metabolic profile**

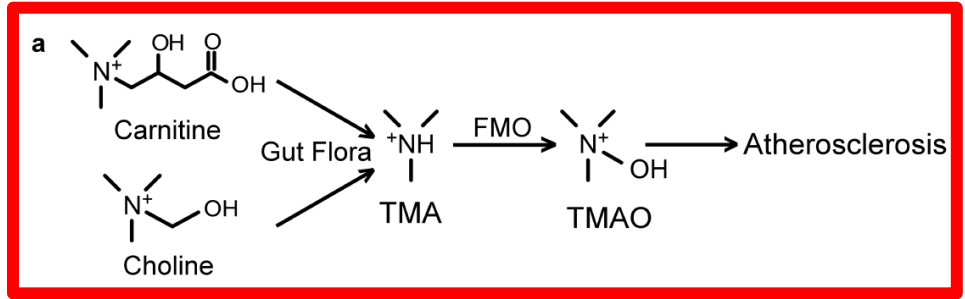
-  Akk LO, LGC (N = 9)
-  Akk HI, LGC (N = 9)
-  Akk LO, HGC (N = 11)
-  Akk HI, HGC (N = 16)



A core of MGS associated with Akk over time in MicroObes



13 Firmicutes, 5 Bacteroidetes, 1 Actinobacteria and 1 Euryarchaeota

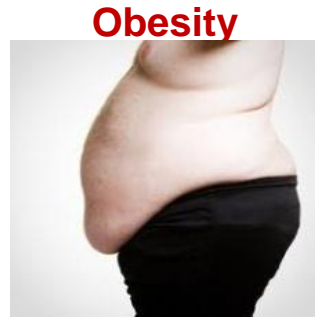


Gut microbiota
 Composition
 Microbial richness
 Specific species (i.e. *A. muciniphila*)

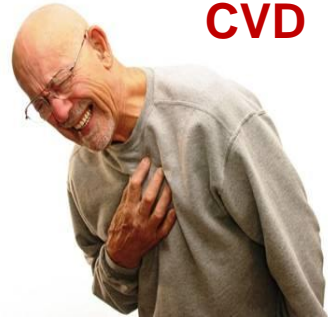
Metabolites
 TMAO

Bacterial components
 Cell wall, LPS

.....▶ **Co-morbidities**

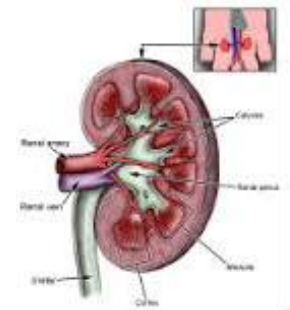


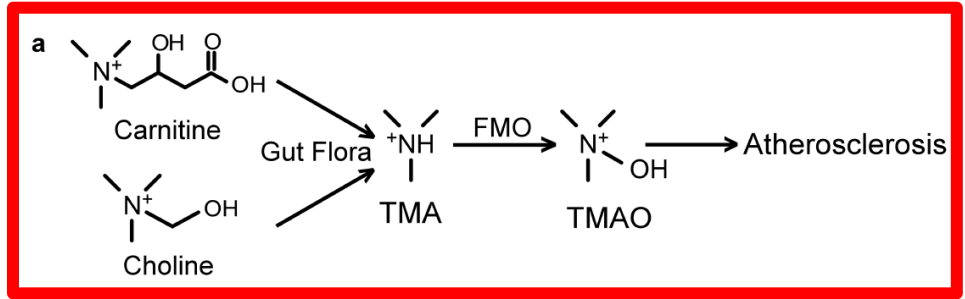
Obesity



CVD

CKD?





Gut microbiota
 Composition
 Microbial richness
 Specific species (i.e. *A. muciniphila*)

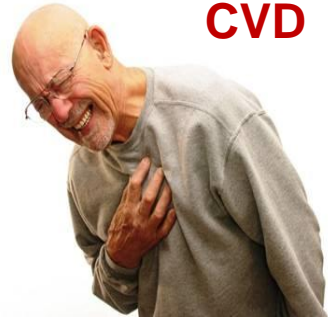
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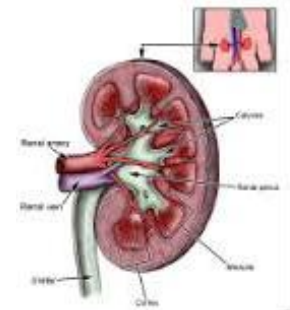


Obesity

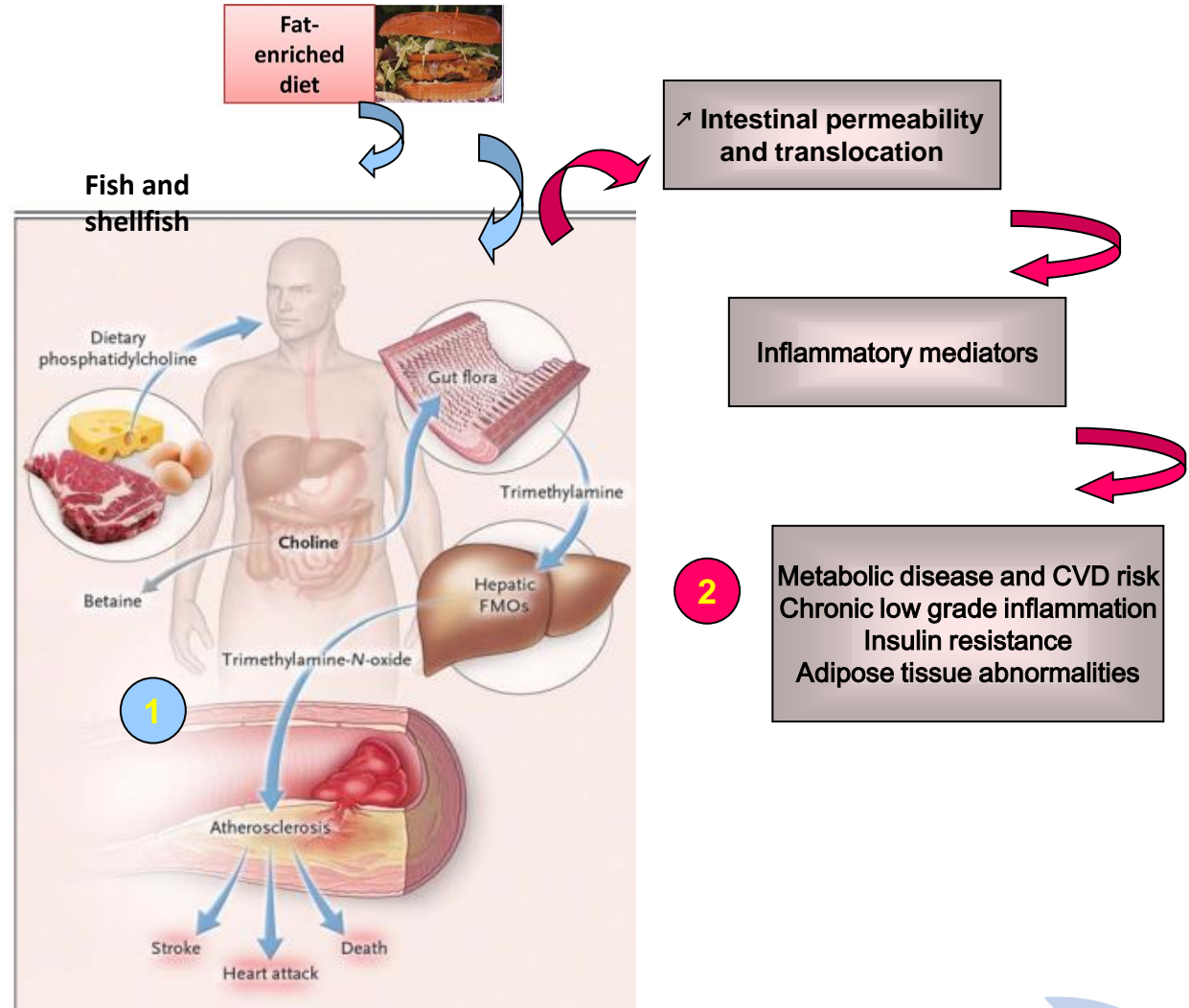


CVD

CKD?



Gut Microbiota, TMAO & CVD risks



Adapted from R. Burcelin et al *Seminars in Immunology* 2012 et Wilson Tang et al, *NEJM* 2013 Cani P.D. et al. *Diabetes* 2007, Cani P.D. et al. *Diabetologia* 2007, Tsukumo et al. *Diabetes* 2007, Cani P.D. et al. *Diabetes* 2008, Kim et al. *Circ. Res.* 2007



4007 patients
undergoing
elective coronary
angiography

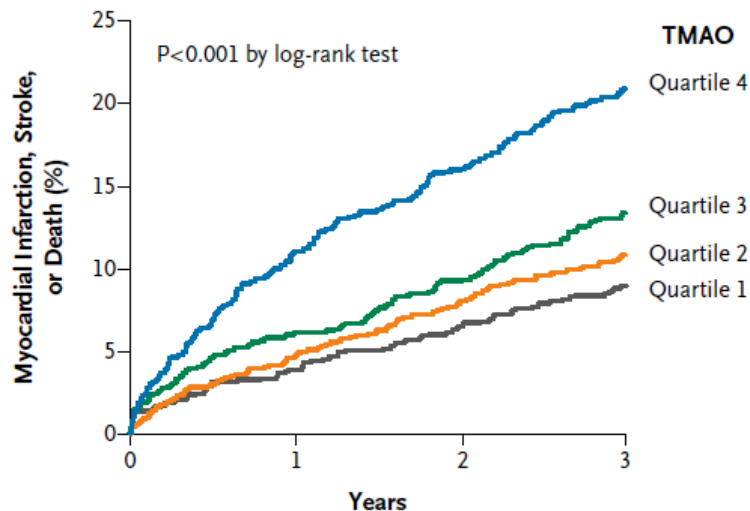
3yr follow up

Table 1. Baseline Characteristics of the Participants in the Clinical-Outcomes Study, According to Status with Respect to Major Adverse Cardiovascular Events at 3 Years.*

Characteristic	All Participants (N=4007)	Participants without Events (N=3494)	Participants with Events (N=513)	P Value†
Age — yr	63±11	62±11	68±10	<0.001
Male sex — %	64	65	62	0.16
Median TMAO (interquartile range) — μM	3.7 (2.4–6.2)	3.5 (2.4–5.9)	5.0 (3.0–8.8)	<0.001

Older, higher glycemia, % T2D ↗ % AHT ↗, MI history

2.5-fold increase in risk



↗ basal TMAO remains an independent factor of CV events after adjusting for traditional risk factors.

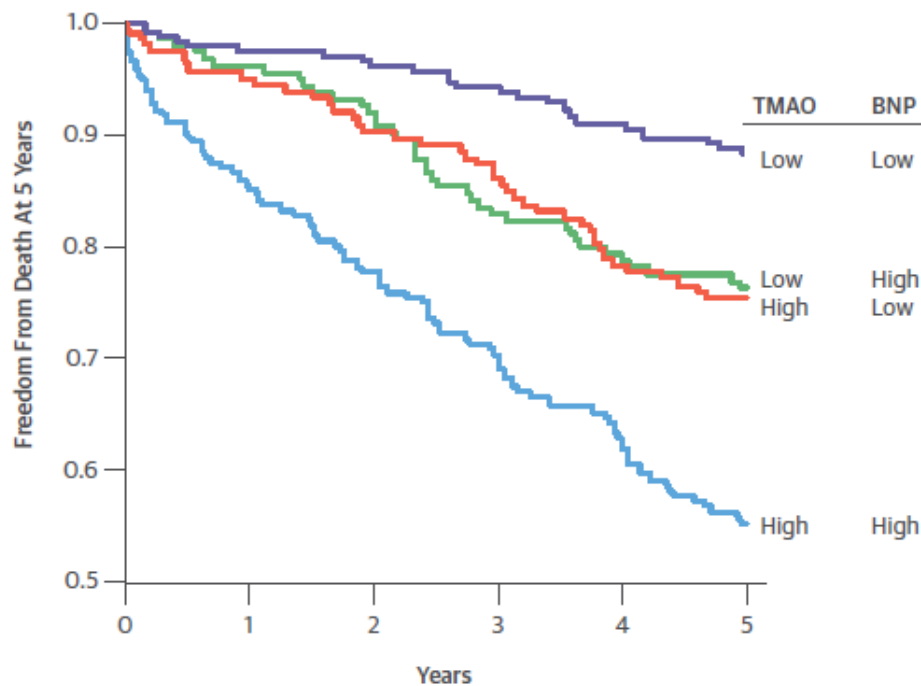
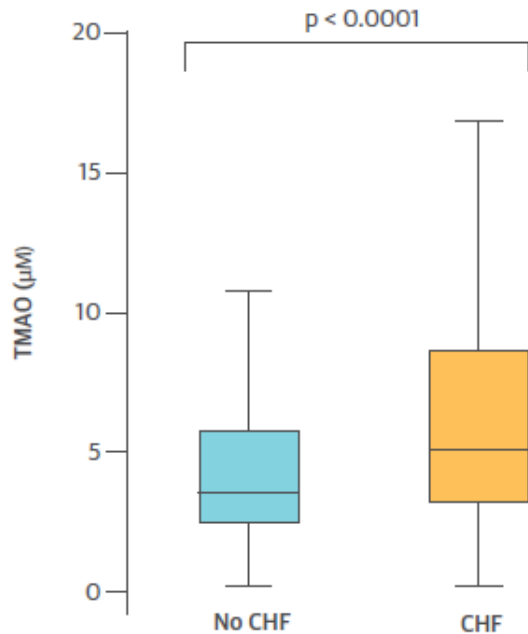


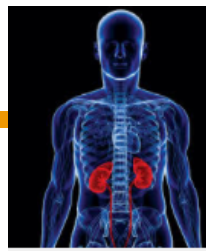
720 patients with stable HF

5yr follow up

TABLE 1 Baseline Characteristics of Heart Failure Cohort (n = 720)

	Overall	TMAO <5 μM	TMAO ≥5 μM	p Value
Age, yrs	66 ± 10	64 ± 11	68 ± 10	<0.001
Male	59	59	59	1.000
Diabetes mellitus	41	31	51	<0.001
Hypertension	78	76	79	0.316
Ischemic etiology	64	63	65	0.673
LV ejection fraction	35 (25-50)	35 (25-51)	40 (25-50)	0.567
Body mass index, kg/m ²	28.4 (25.1-33.1)	28.7 (25.2-33.3)	28.1 (24.8-33.0)	0.298



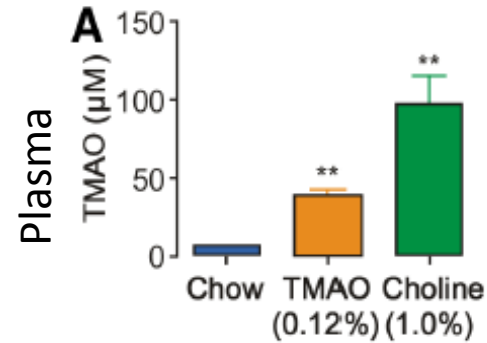


- CKD patients: incr. TMAO (Tang WHW et al, J Card Fail, 2015)
 - Hemodialysis (1 session): Dec. TMAO (Bain MA, 2006)
 - Kidney transplantation: Dec. TMAO (Stubbs JR, JASN. 2015)
 - High TMAO: Predictor of all cause mortality (Onopiuk A 2015)
or long-term cardiac death or coronary atherosclerosis
(Stubbs JR, JASN. 2015)
- No information on CKD stage of progression
 - No information on food intake consumption

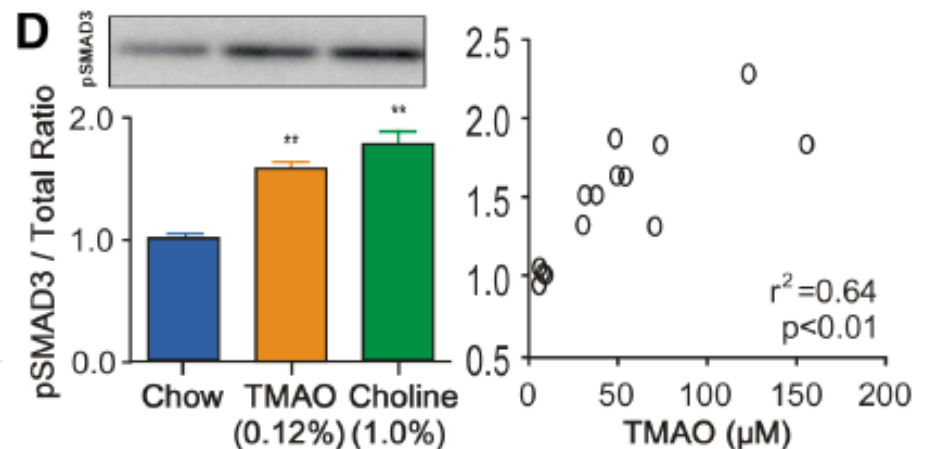
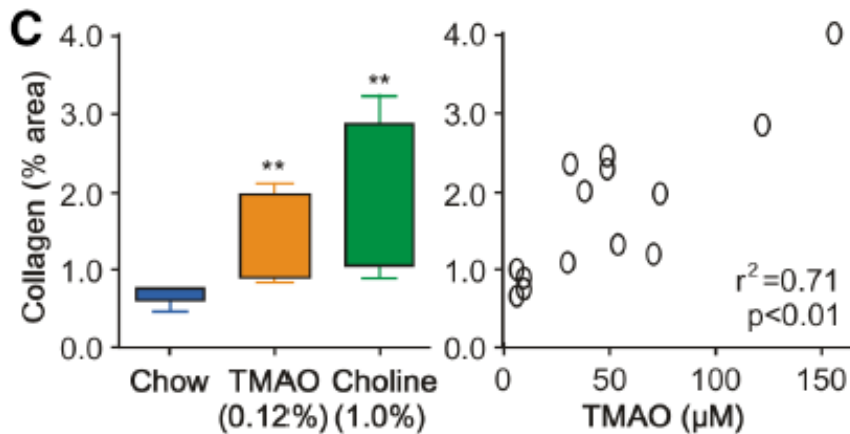
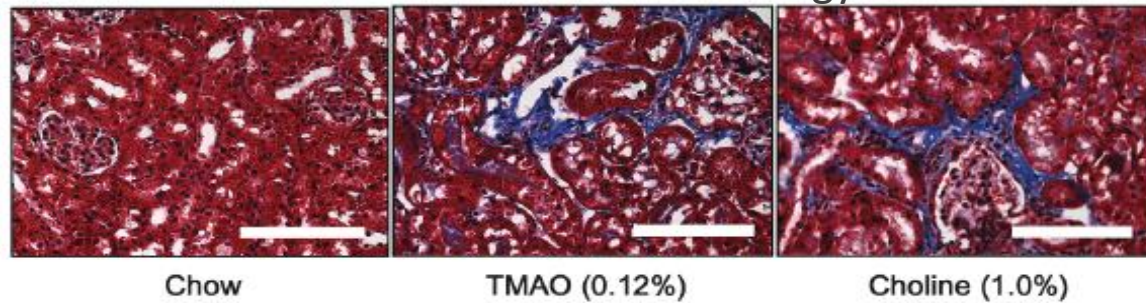
6 weeks

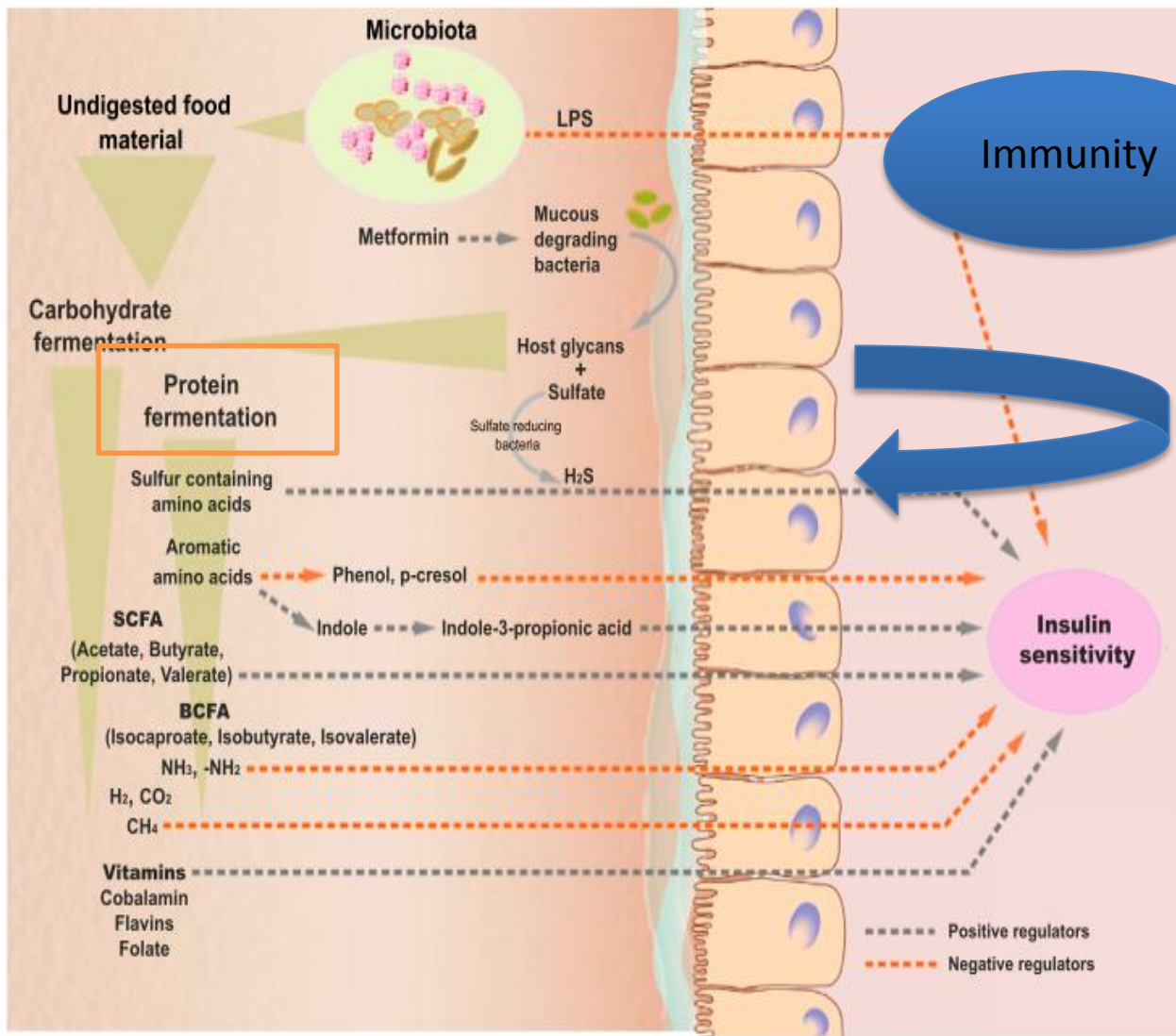


- Chow (choline 0,08%)
- Chow + **TMAO** (0,12%)
- Chow + **Choline** (1%)



Mason trichrome histology





Intestinal barrier

Two candidates

Indoxyl Sulfate

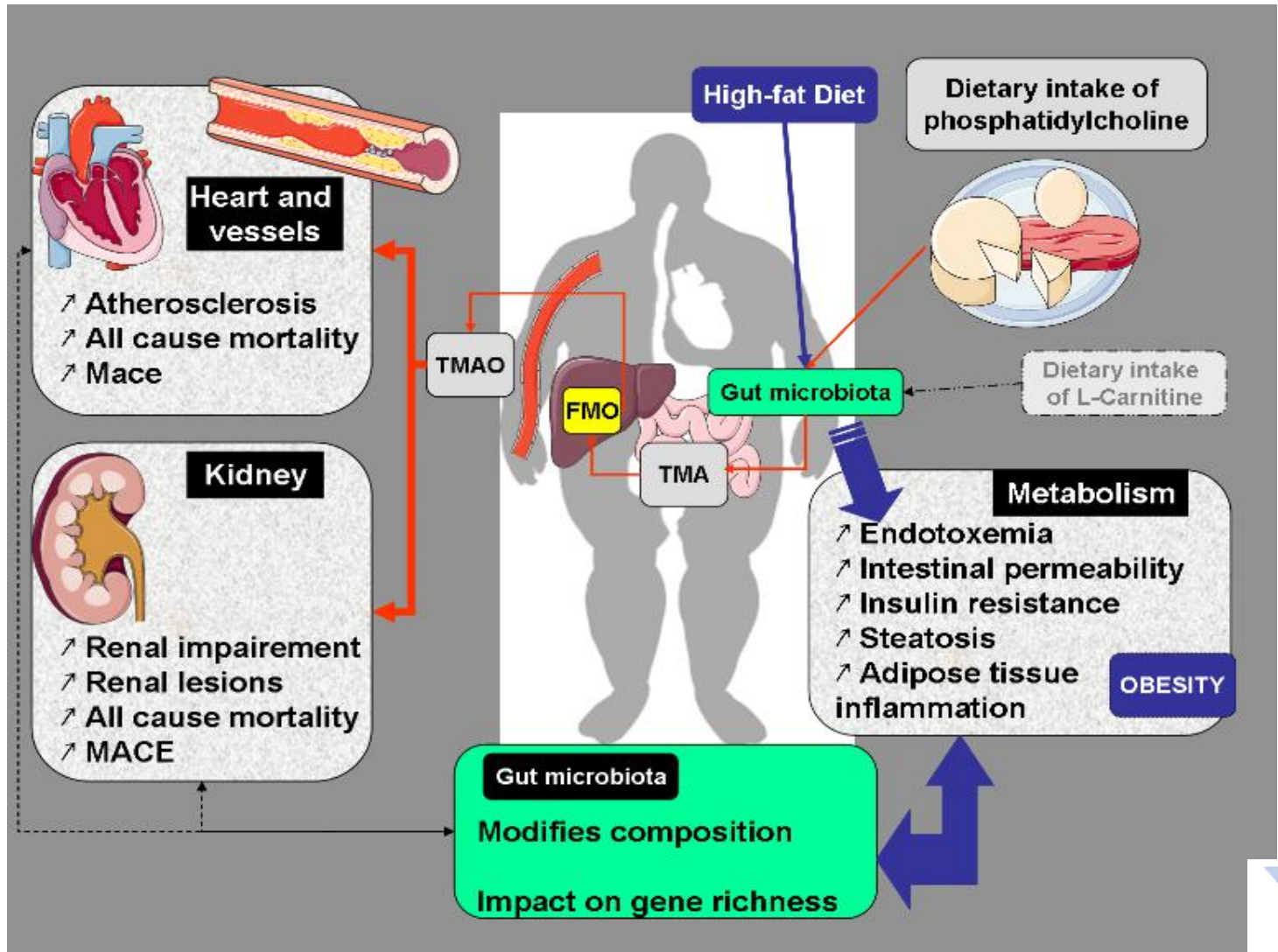
P Cresol



Linked to Kidney function
CKD progression
Mortality (dialysis)

Vascular stiffness & CVD

Summary : shared pathway?

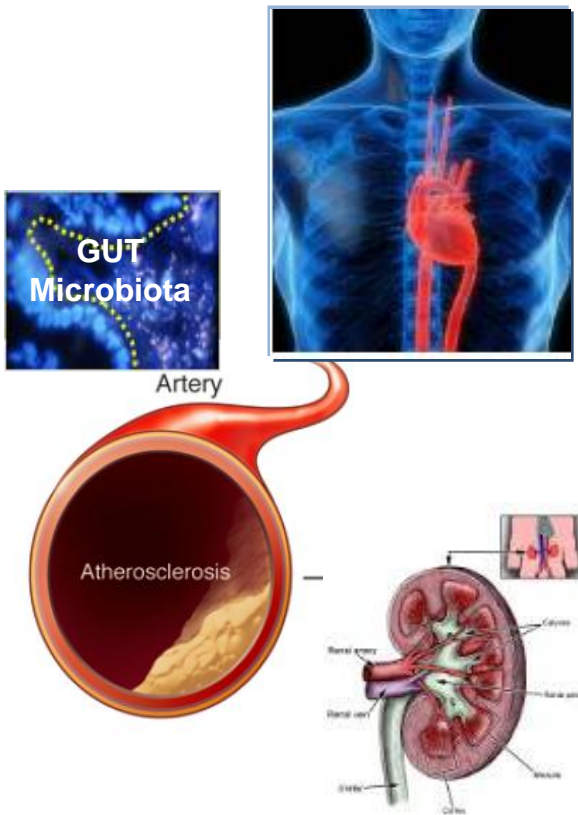


Aron-Wisnewsky J and Clément K, Nature Nephrol. In press

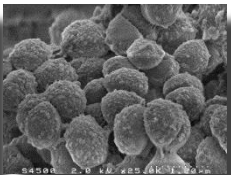


General conclusions

- ❖ Direct link shown between diet, microbiota and CVD, especially in animal models.
- ❖ Although mechanisms remain to be explored, different factors of gut microbiota (composition and richness, bacterial function and structural components) are linked with the health of the host.
- ❖ Gut microbiota is modifiable, and this can serve as target for improvement of disease.
- ❖ THERAPY ? (PRE, PRO BIOTICS, TRANSFER..)



Faecalibacterium prausnitzii
Photos INRA



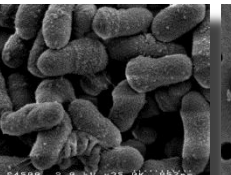
Ruminococcus
spp



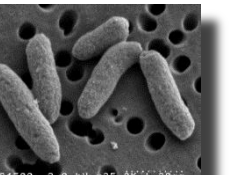
Clostridium difficile
en caecum souris



Bactéries ancrées
dans une Plaque de
Peyer,
Intestin de souris



Bacteroides dorei



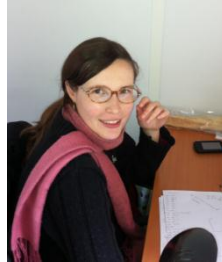
Escherichia coli



**Christine
Poitou**



**Salwa
Rizkalla**



**Aurelie
Cotillard**



**Jean-Daniel
Zucker**



Edi Prifti

the Musketeers



**Carlota
Dao**



**Judith
Aron**



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