

# “Towards Microbial Fermentation Metabolites as Biomarkers for Health Benefits of Prebiotics”

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for working group on Microbial Metabolism and Fermentation

SUPPORTED BY THE ILSI EUROPE PREBIOTICS TASK FORCE



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# Background

The ILSI Europe Prebiotic Task Force identified a particular need to explore available evidence for positive or negative physiological effects of microbial (faecal) metabolites on the host.

# Aim

- Exploring how to define and impact digestive health
- Looking at the different gut functions that contribute to a healthy gastrointestinal tract and thereby to overall health

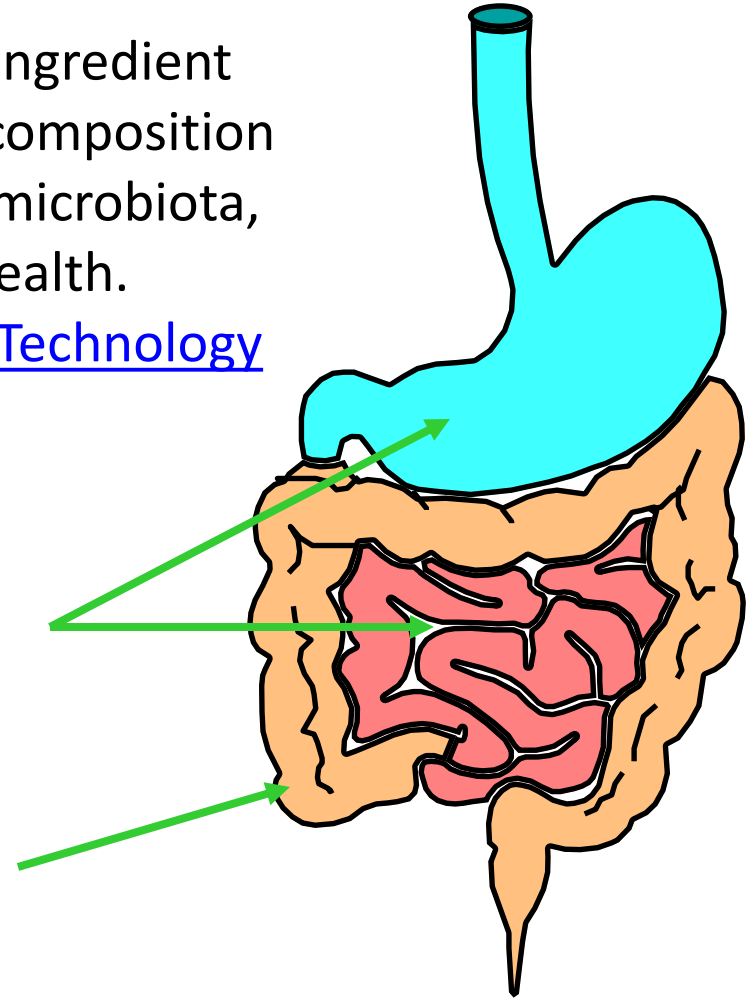
# Prebiotic definition:

A **prebiotic** is a selectively fermented ingredient that results in specific changes in the composition and/or activity of the gastrointestinal microbiota, thus conferring benefit(s) upon host health.

[\(Gibson et al. 2010. Food Science and Technology Bulletin: Functional Foods 7 \(1\) 1–19.\)](#)

**Persist until the colon**

**Selective fermentation in the colon**



# Objectives and Methods:

A literature search was performed to address:

- Available evidence for the beneficial or harmful effects of known microbial metabolites including short chain fatty acids and protein fermentation products.
- The potential for functional analysis of faecal water
- The applicability of metabolome signatures and systems biology

# Overview on different metabolites

## Summarising available evidence

### a. List of relevant metabolites

### b. Highlighted metabolites

Integrated perspectives (harm+benefit) including:

- Pharmacological perspectives
- Toxicological perspectives
- Nutrition perspectives
- Normal ranges

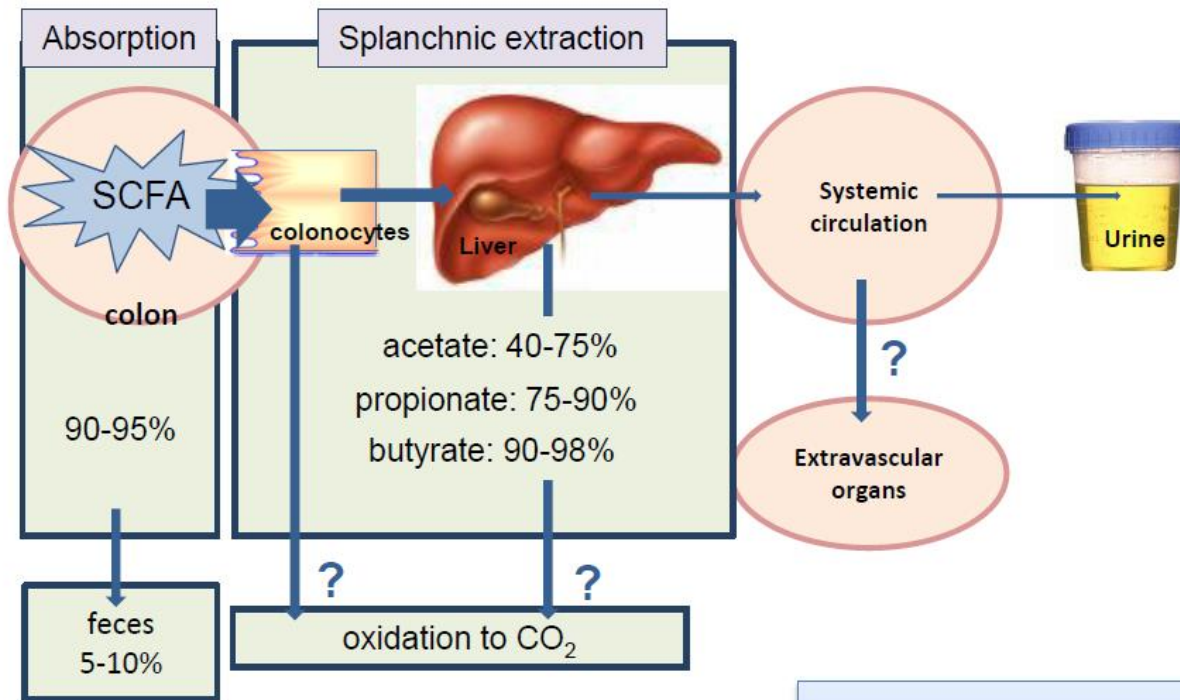
### c. More holistic approach: Profiles

- Functional analysis of faecal water (include genotoxicity)
- Metabolomics
- Metagenome analysis

# Colonic Microbial Metabolites

Carbohydrate (Dietary fibre)	Protein	Plant Polyphenolics	Fat and related bile acids
<ul style="list-style-type: none"> <li>• Short chain fatty acids               <ul style="list-style-type: none"> <li>○ Acetate</li> <li>○ Propionate</li> <li>○ Butyrate</li> </ul> </li> <li>• Lactate</li> <li>• Succinate</li> <li>• Alcohols</li> <li>• Gasses:               <ul style="list-style-type: none"> <li>○ Hydrogen</li> <li>○ Methane</li> <li>○ Carbondioxide</li> </ul> </li> </ul>	<ul style="list-style-type: none"> <li>• Ammonia</li> <li>• Hydrogen Sulphide</li> <li>• Phenols</li> <li>• p-cresol</li> <li>• Indoles</li> <li>• Branched chain fatty acids</li> </ul>	<ul style="list-style-type: none"> <li>• Large range of phenolic compounds and acids including:</li> <li>• Simple phenols</li> <li>• Glycinated benzoic acids</li> <li>• Derivatives of benzoic acid</li> <li>Derivatives of               <ul style="list-style-type: none"> <li>○ Phenyl acetic acid</li> <li>○ Phenylpropionic acid</li> <li>○ Mandelic acids</li> <li>○ Cinnamic acids</li> <li>○ Equols</li> </ul> </li> </ul>	<ul style="list-style-type: none"> <li>• Hydroxy fatty acids</li> <li>• Secondary bile acids</li> <li>• Long chain aldehydes</li> </ul>

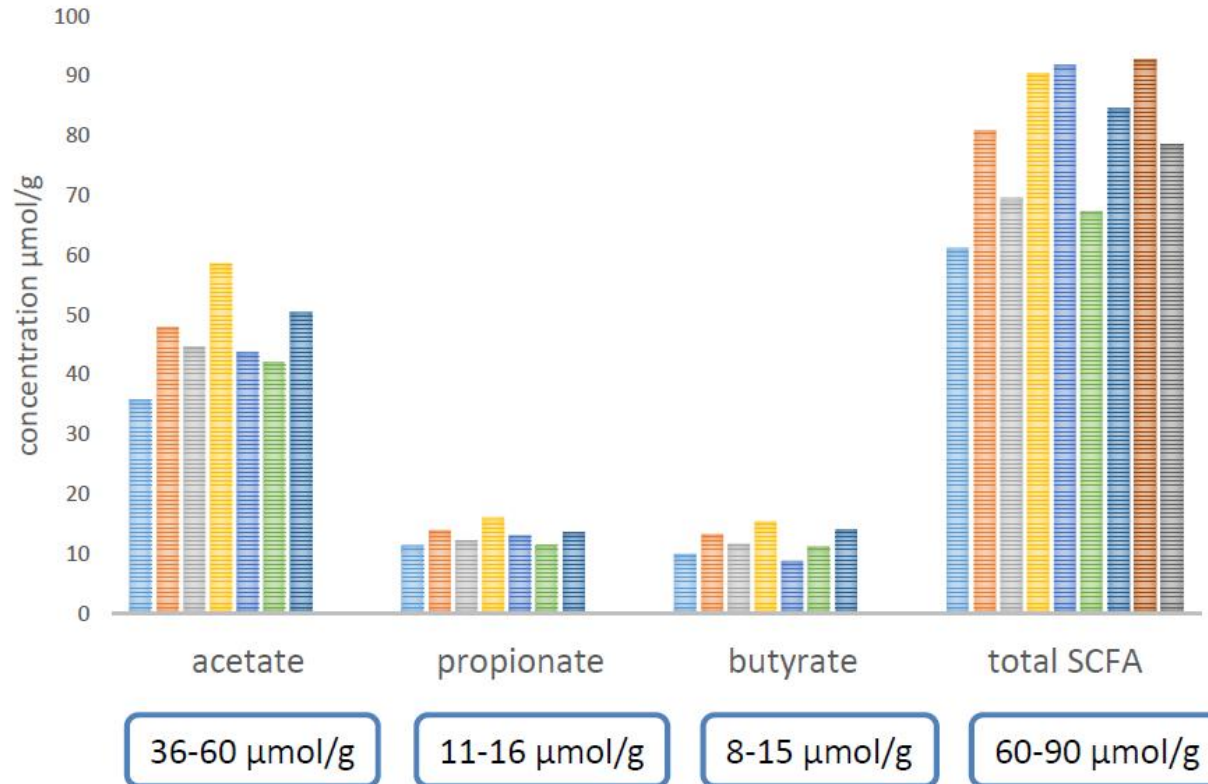
# Metabolism of SCFA



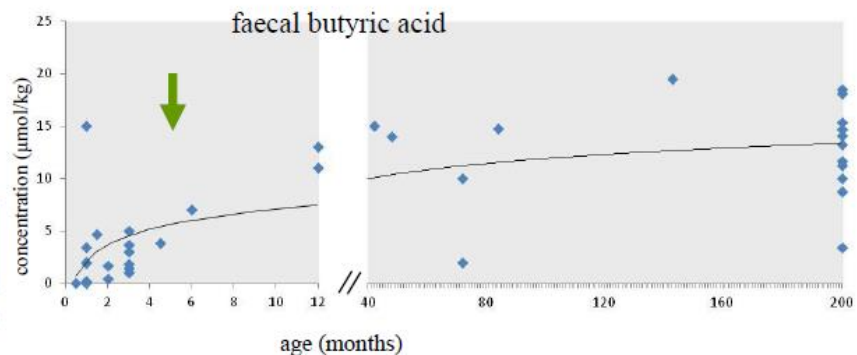
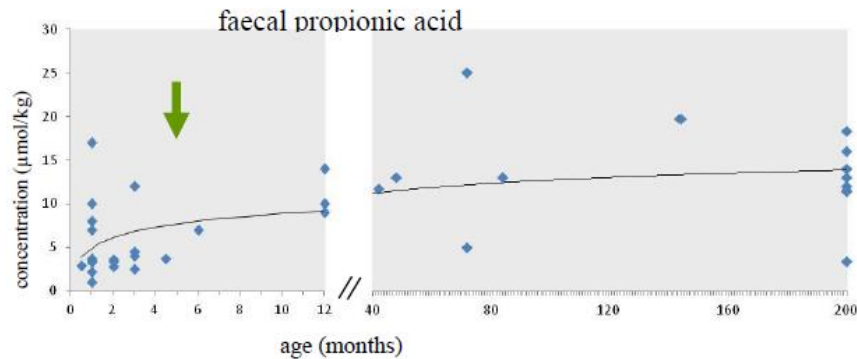
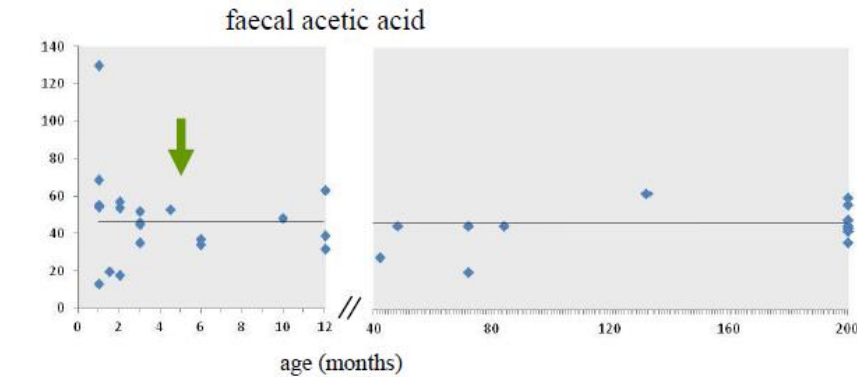
- Inaccessibility of appropriate body compartment
  - ⇒ leaves faeces, plasma, urine
  - ⇒ not representative
- Where / when to measure
- Large normal variations – diet and other factors



# Faecal SCFA concentrations



# Faecal [SCFA] depend on age



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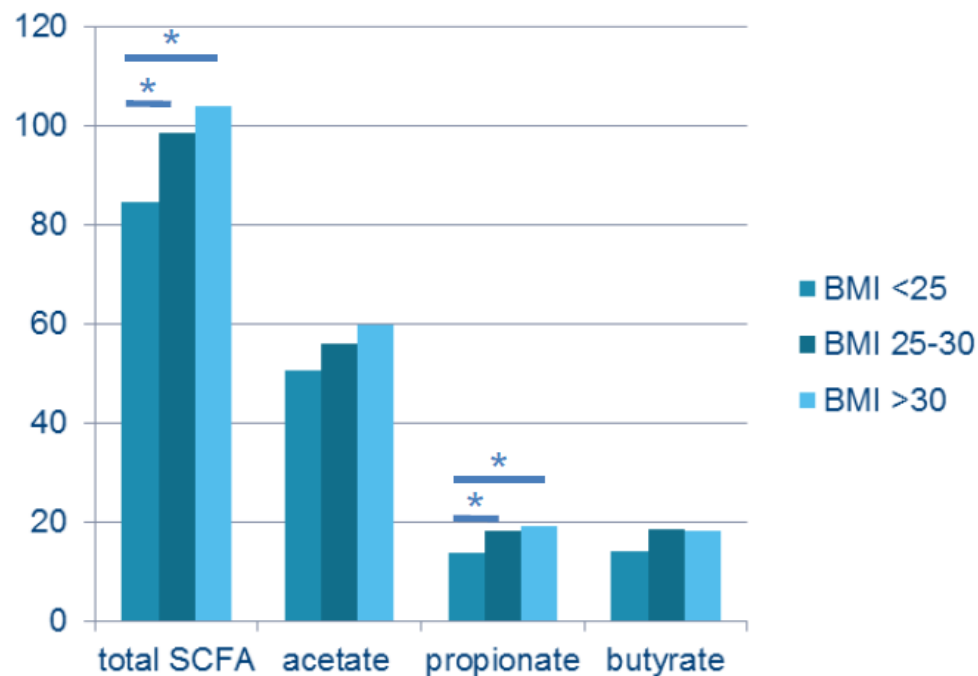
- Edwards *et al.* Acta Paediatr 1994, 83, 459-462  
 Midtvedt *et al.* J Pediatr Gastroenterol Nutr 1992, 15, 395-403  
 Parrett *et al.* Am J Clin Nutr 1997, 65, 927-933  
 Tjellstrom *et al.* Microb Ecol Health Dis 2013, 24, 20905  
 Wang *et al.* Dig Dis Sci 2012, 57, 2096-2102  
 De Filippo *et al.* Proc Natl Acad Sci U S A 2010, 107, 14691-14696  
 Guerin-Danan *et al.* J Pediatr Gastroenterol Nutr 1997, 25, 281-289  
 Siigur *et al.* Acta Paediatr 1993, 82, 536-538  
 Holscher *et al.* J Parenter Enteral Nutr 2012, 36, 95s-105s  
 Norin *et al.* Microb Ecol Health Dis 2004, 16, 8-12  
 Bakker-Zierikzee *et al.* Br J Nutr 2005, 94, 783-790  
 Samuelsson *et al.* Diabet Med 2004, 21, 64-67  
 Parrett *et al.* Arch Dis Child 1997, 76, 249-253

# Faecal [SCFA] differ in lean v obese

## Microbiota and SCFA in Lean and Overweight Healthy Subjects

Andreas Schwartz<sup>1</sup>, David Taras<sup>2</sup>, Klaus Schäfer<sup>2</sup>, Silvia Beijer<sup>3</sup>, Nicolaas A. Bos<sup>3</sup>, Christiane Donus<sup>4</sup> and Philip D. Hardt<sup>4</sup>

*Obesity* (2009) **18**, 190–195. doi:10.1038/oby.2009.167



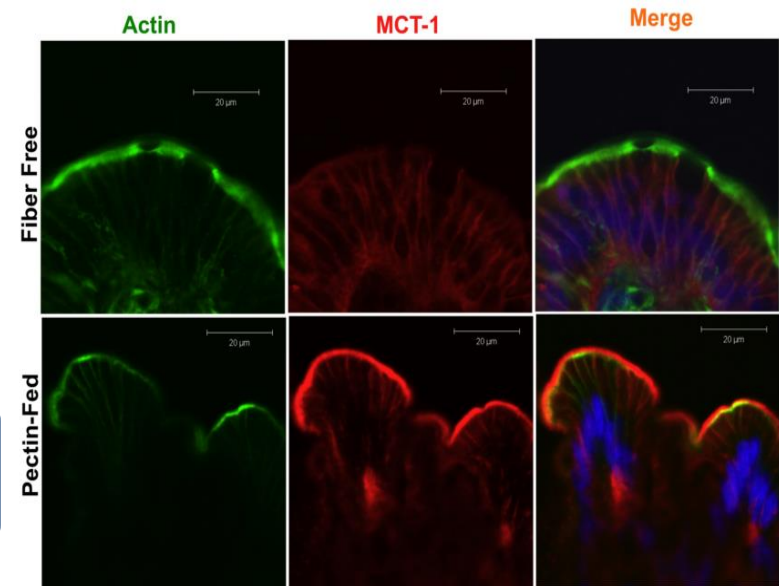
# Faecal [SCFA] poor biomarker of gut health

Increased faecal SCFA excretion due to decreased SCFA uptake?

MCT transporters  
expression and  
apical location is  
promoted by  
luminal SCFA

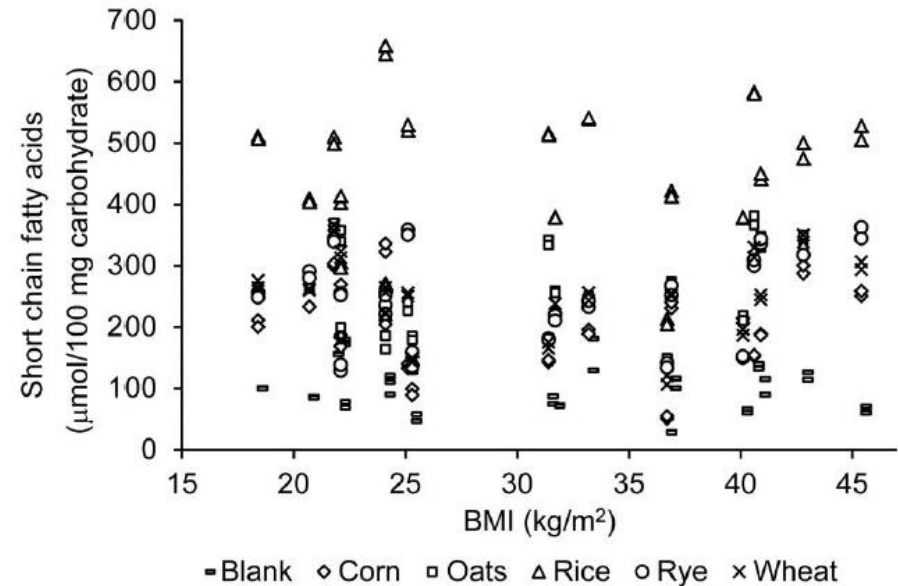
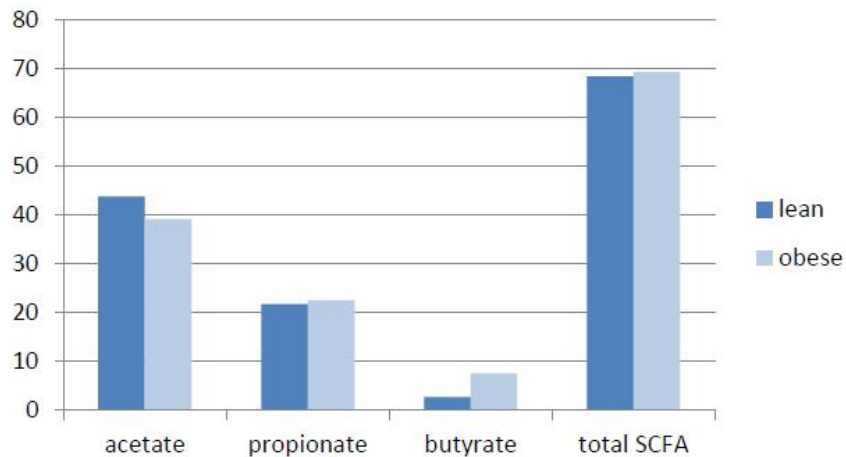
Increased faecal SCFA excretion could be due to  
decreased MCT active uptake in high fat/low CHO diets

Bile salt CDCA  
and *E. coli* EPEC  
inhibit butyrate  
uptake



Goncalves et al. J Cell Biochemistry 2012, 113, 2937-2947  
Borthakur et al. Am J Physiol Gastrointest Liver Physiol  
2012, 303, G1126-G1133

# Little difference in SCFA production *in vitro* between lean and obese microbiota



# Faecal [SCFA] in disease states

- IBD: lower levels of faecal SCFA
- celiac disease: increased levels of total SCFA and acetate
- allergy: lower faecal levels of propionate and butyrate

⇒ causative to disease?

⇒ markers of disease?

# SCFA may beneficially impact on the following mammalian processes

- Apoptosis
- Cellular proliferation in non-cancer cells
- Inflammation
- Recruitment of immune cells to intestine
- Neutrophil activity/oxidative burst
- Adipose tissue – adipokine production, fat storage
- Tight junction control
- Expression of incretins/gut peptides, and regulation of food intake
- Intestinal motility
- Cholesterol production/lipogenesis
- Glucogenesis
- Cellular energy metabolism
- Thermogenesis
- Inhibits tumorigenesis
- DNA miss-match repair

Note: most mechanistic data from in vitro or animal studies – rodents, pigs, chickens

# Faecal minor organic acids

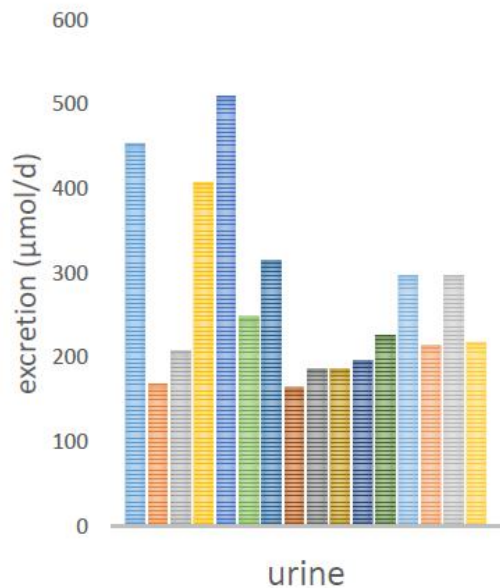
- Lactate, succinate
  - do not accumulate much in faeces in health
  - cross-feeding between bacteria leads to formation of main SCFA
  - Lactate:
    - considered as a marker of dysbiosis
    - cosubstrate for sulphate reducing bacteria ⇨ promote sulphide generation
  - Succinate:
    - may act as a signal of inflammation
    - Increased levels have been linked to inflammatory bowel diseases (IBD)



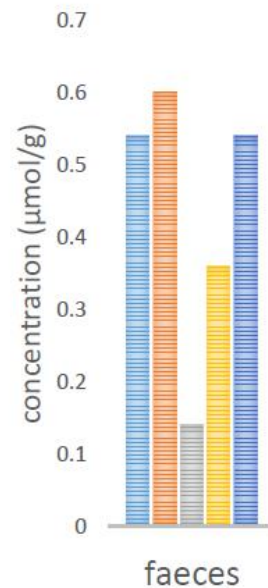
# Metabolites of protein fermentation

Metabolite	Origin
Ammonia	<ul style="list-style-type: none"><li>• Bacterial degradation of amino acids</li><li>• Hydrolysis of urea</li></ul>
Phenols	<ul style="list-style-type: none"><li>• Major metabolites of bacterial fermentation of aromatic amino acids</li><li>• Rapidly absorbed by colonic mucosa and excreted in urine</li><li>• Do not accumulate in healthy subjects</li></ul>
<ul style="list-style-type: none"><li>• p-cresol</li></ul>	< tyrosine
<ul style="list-style-type: none"><li>• phenol</li></ul>	< phenylalanine
<ul style="list-style-type: none"><li>• indole</li></ul>	< tryptophan

# Urinary and faecal levels of p-cresol



160-530 μmol/d

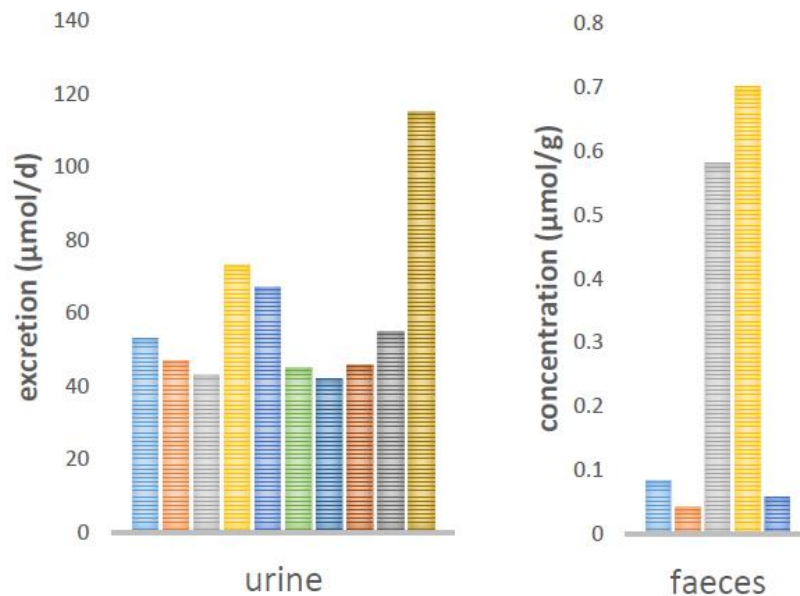


0.14-0.60 μmol/g

- Higher urine levels in obese than in normal weight subjects
- Urine p-cresol levels increase in very old subjects

Birkett *et al.* Am J Clin Nutr 1996, 63, 766-772  
Damen *et al.* J Nutr 2012, 142, 470-477  
Ling *et al.* J Nutr 1992, 122, 924-930  
Patel *et al.* Clin J Am Soc Nephrol 2012, 7, 982-988  
Renwick *et al.* Hum Toxicol 1988, 7, 267-272  
De Preter *et al.* Br J Nutr 2004, 92, 439-446  
De Preter *et al.* J Am Coll Nutr 2007, 25, 541-549  
De Preter *et al.* Am J Physiol Gastrointest Liver Physiol 2007, 292, G358-G368  
Cloetens *et al.* Br J Nutr 2010, 103, 703-713  
Cloetens *et al.* J Am Coll Nutr 2008, 27, 512-518  
Windey *et al.* PlosOne 2012, 7, Article Number: e52387  
Gostner *et al.* Br J Nutr 2006, 95, 40  
Adams *et al.* Lancet 1985, 2, 1313-1313  
Heavey *et al.* Br J Nutr 2003, 89, 509-515

# Urinary and faecal levels of phenol



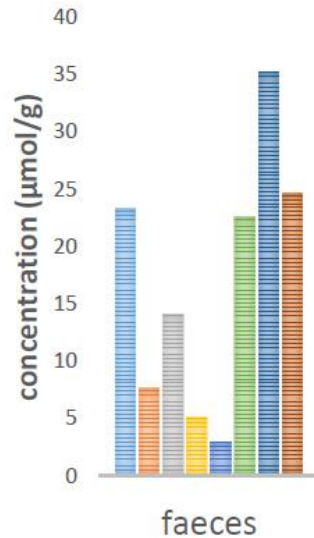
40-115 µmol/d

0.04-0.7 µmol/g

# Effects of phenolic derivatives of protein fermentation

- Effects on epithelial cells mainly determined using *in vitro* incubation tests
  - Decreased viability
  - Reduced epithelial barrier function
- No systemic toxicity in healthy subjects
- Accumulates in serum in chronic kidney disease - uremic toxin
  - Contributes to endothelial dysfunction
- Very limited data on toxicity of indole

# Faecal [ammonia]



3-35 µmol/g

- Comparable levels in overweight and normal weight subjects
- Higher levels in children with autism spectrum disorders (42 µmol/g)

# Effects of intestinal ammonia

- Effects on epithelial cells:
  - alters nucleic acid synthesis
  - changes the morphology and intermediary metabolism of intestinal cells
  - reduces the lifespan of cells
  
- No adverse effects upon oral administration
  
- Hepatic encephalopathy

**Table 6.** List of microbial catabolites of common plant polyphenols and their putative health effects<sup>(155)</sup>

Plant polyphenol	Microbial catabolite	Possible health effects	References
(-)-Epicatechin	4-Hydroxyphenylacetic acid	Antimicrobial/antimycotic activity <i>in vitro</i>	Alakomi (2007) <sup>(254)</sup> Ko (2009) <sup>(255)</sup> Roowi (2010) <sup>(256)</sup>
	3-(3-Hydroxyphenyl)propionic acid	Antimicrobial activity against Gram-negative enterobacteria via outer membrane destabilisation	
	5-(3,4-Dihydroxyphenyl)- $\gamma$ -valeric acid (-)-5-(3',4'-Dihydroxyphenyl)- $\gamma$ -valerolactone	? ?	
(-)-Epigallocatechin	4-Hydroxyphenylacetic acid (-)-5-(3',4'-Dihydroxyphenyl)- $\gamma$ -valerolactone	Antimicrobial/antimycotic activity <i>in vitro</i>	Roowi (2010) <sup>(256)</sup>
(-)-Epigallocatechin-3- <i>O</i> -gallate	Pyrocatechol		Ko (2009) <sup>(255)</sup> Roowi (2010) <sup>(256)</sup> Okello (2012) <sup>(257)</sup> Ni (2008) <sup>(258)</sup> Taguri (2006) <sup>(259)</sup>
	Pyrogallol	Antibacterial activity (especially against Gram-negative enterobacteria) An acetylcholinesterase inhibition greater than gallic acid parent Inhibition of <i>Vibrio</i> spp. quorum sensing	
Daidzein	4-Hydroxyphenylacetic acid (-)-5-(3',4'-Dihydroxyphenyl)- $\gamma$ -valerolactone Equol	Antimicrobial/antimycotic activity <i>in vitro</i> ? Phyto-oestrogen important for heart and bone health, and possible colon cancer protectants	Jackman (2007) <sup>(260)</sup> Ishimi (2009) <sup>(261)</sup> Davis (2009) <sup>(262)</sup> Selma (2009) <sup>(263)</sup>
	<i>O</i> -demethylangolensin	Oestrogenic and/or anti-oestrogenic activity	
Quercetin	2-(3,4-Dihydroxyphenyl)acetic acid 2-3-(3-Hydroxyphenyl)acetic acid 3,4-Dihydroxybenzoic acid Phloroglucinol 3-(3,4-Dihydroxyphenyl)propionic acid 3-(3-Hydroxyphenyl)propionic acid		Larrosa (2006) <sup>(264)</sup> Selma (2009) <sup>(263)</sup> Selma (2009) <sup>(263)</sup>
	2-(4-Hydroxyphenyl)acetic acid		
	3-(4-Hydroxyphenyl)propionic acid	Antimicrobial activity against Gram-negative enterobacteria via outer membrane destabilisation	
	Phloroglucinol 8-Prenylnaringenin	? ?	
		Antimicrobial activity against Gram-negative enterobacteria via outer membrane destabilisation	
Isoxanthohumol Catechin and epicatechin	3-(3-Hydroxyphenyl)propionic acid		Selma (2009) <sup>(263)</sup> Alakomi (2007) <sup>(254)</sup> Selma (2009) <sup>(263)</sup>
	5-(3',4'-Dihydroxyphenyl)- $\gamma$ -valerolactone 5-(3'-Hydroxyphenyl)- $\gamma$ -valerolactone 3-Hydroxyhippuric acid pyrogallol 5-(3,4-Dihydroxyphenyl)valeric acid 5-(3-Hydroxyphenyl)valeric acid 3-(3,4-Dihydroxyphenyl)propionic acid	Antimicrobial activity against Gram-negative enterobacteria via outer membrane destabilisation	
	5-(3-Methoxyphenyl)valeric acid 3-(3,4-Dihydroxyphenyl)propionic acid 5-(3-Methoxyphenyl)valeric acid 2,3-Dihydroxyphenoxy  3-(3',4'-dihydroxyphenyl)propionic acid		

# Possible solution

A more holistic approach



# Functional analysis of faecal water

- Functional analysis of faecal water provides an integrated measure of the overall contribution of the compounds present to a defined biological endpoint.
- Provides no information on compounds responsible for the functional effect

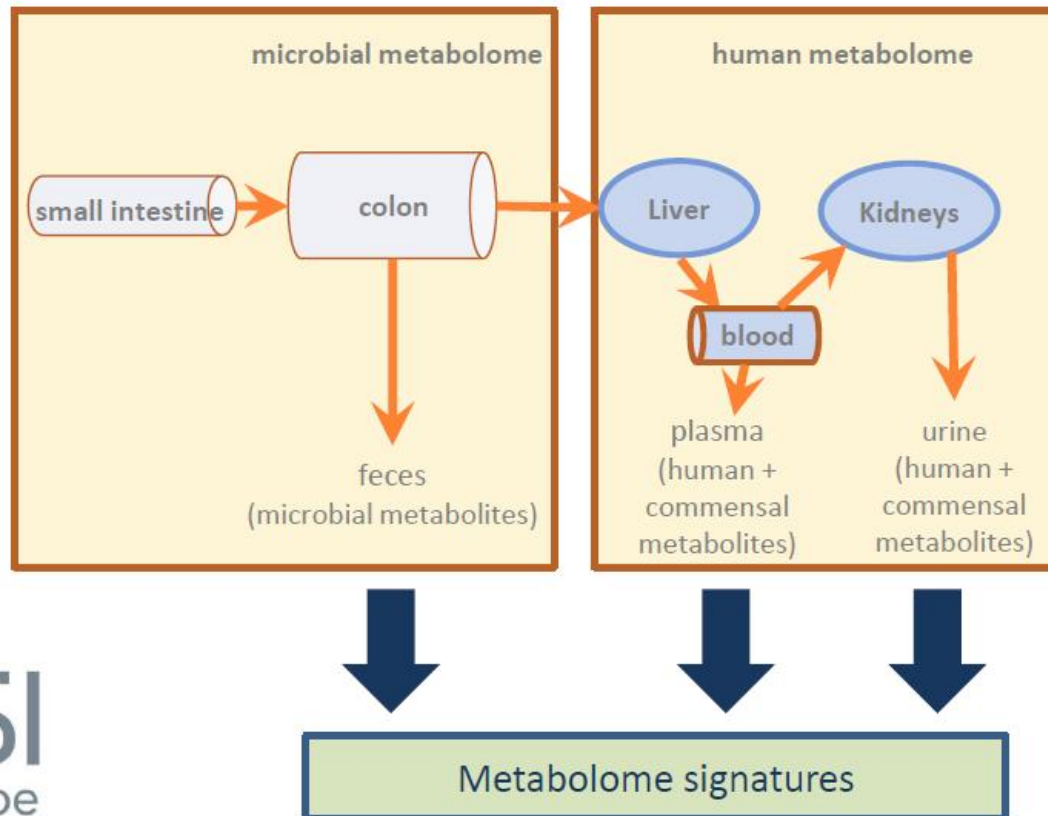
# Available assays

Endpoint	Assay
Genotoxicity	Bacterial mutagenicity
	Comet assay in mammalian cells
Cell toxicity	Cytotoxicity
	Barrier function
	Invasive potential
	Red cell lysis
Cell proliferation	Cell number
	Cell cycle analysis
Immune modulation	Expression of inflammatory markers
Gene expression	AP-1, COX-2,

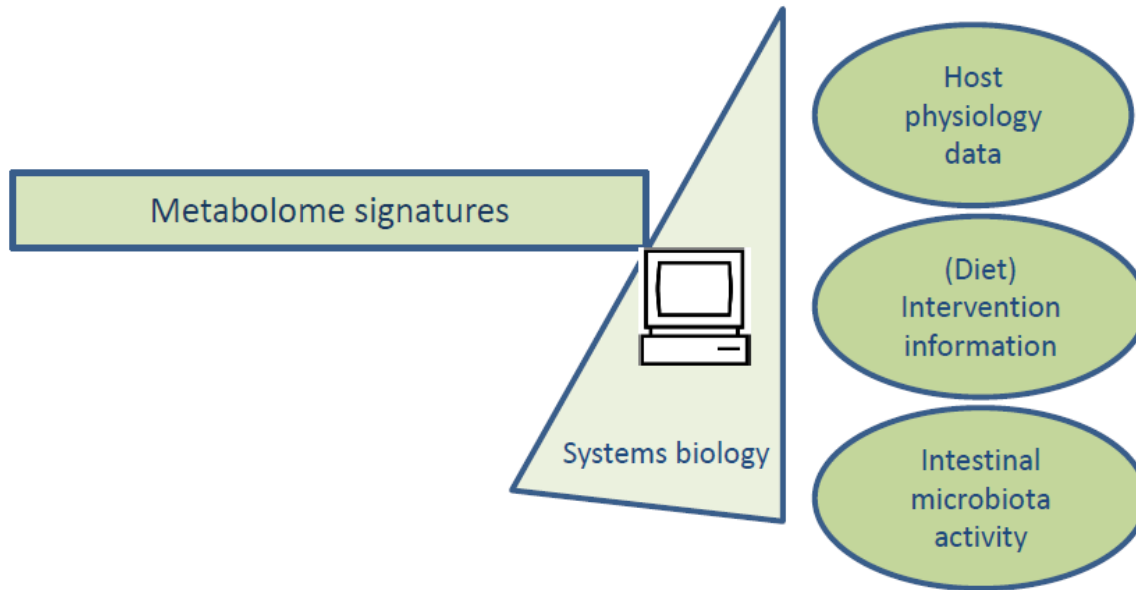
- Limitation: lack of standardisation in target cells and sample preparation
- Few studies applied the approach to evaluate prebiotic effects

# Metabolomics

- simultaneous monitoring of changes in a wide range of metabolites
  - ⇒ No need for an *a priori* hypothesis
- $^1\text{H-NMR}$ , LC-MS, GC-MS
- matrix: feces, urine, plasma, tissue homogenates

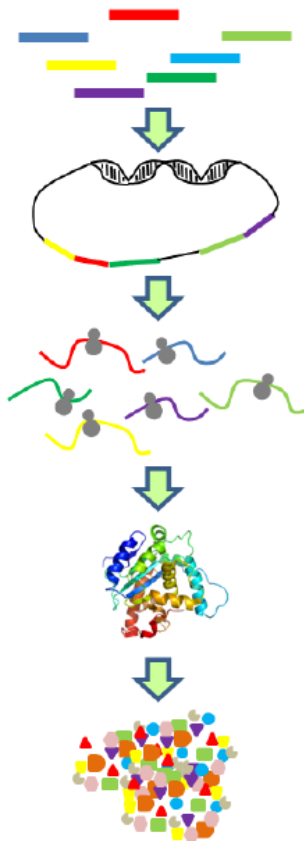


# Applicability of systems biology



- ⇒ Metabolome signatures are analysed in the context of overall biochemistry of the tissue or sample
- ⇒ “Biochemical connectivity”
- ⇒ Enables to analyse the overall impact of the microbiota on host-biochemistry and -function

# Tools for stools!



Quantified microbiota species (OTU) composition (16S rRNA based)

Availability of methods



Quantified microbiota function profile (metagenome)



Quantified microbiota gene expression profile (metatranscriptome)

emerging

Quantified microbiota protein expression profile (metaproteome)

emerging

Quantified microbiota metabolite profiles (metametabolome)



# Conclusions

- No formal systematic reviews evaluating the physiological or toxicological properties of bacterial fermentation metabolites were found.
- End products of saccharolytic fermentation, SCFA, may have varying effects on colonic health, host physiology, lipoprotein metabolism and appetite.
- Comprehensive reviews and experimental studies indicated that protein fermentation metabolites (phenol, p-cresol, indole, ammonia), typically considered as harmful metabolites, occur at concentration ranges in the colon such that no toxic effects are expected either locally or following systemic absorption.

# Conclusions

- There is insufficient data published to support the use of any individual bacterial metabolite as faecal biomarker of gut health.
- Way to go:
  - Profiling of metabolites in the context of overall tissue biochemistry
  - correlation of (multivariate) metabolome signatures with microbial, dietary and physiological data
    - ⇒ evaluation of the overall impact of the microbiota on host health and gut function
- Current limitation
  - the bioinformatics integration and interpretation of the data
  - the lack of studies measuring metabolite fluxes in different body compartments or biofluids to provide an accurate picture of colonic metabolite nutrkinetics.

# Thank you!



Nutrition Research Reviews

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*Nutrition Research Reviews* (2015), **28**, 42–66

doi:10.1017/S0954422415000037

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## Towards microbial fermentation metabolites as markers for health benefits of prebiotics

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on behalf of the ILSI Europe Prebiotics Task Force Expert Group 'Microbial metabolism and fermentation'



Acknowledgements  
**Prebiotic Task Force, ILSI Europe**