Underlying mechanisms & causal links: high fat feeding & circadian rhythms

Fondation Merieux Conference, April 6-8, 2016 Better Foods for Better Health Microbiota & Health: The Challenges of a Promising Approach





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The obesity and diabetes epidemic





High fat diets affect the murine enteric microbiome differently Suzanne Devkota





16S rRNA gene sequences determined by Sanger-based clone library sequencing

Western-diet induced changes in humanized mouse gut microbiota promote adipose tissue

* 3.0 * 2.5 weight pad Microbiota from 2.0 Epidydymal fat Human subjects on of total body 1.5 different diets transferred to GF mice 1.0 0.5 LF/PP – Low fat, plant polysaccharide-rich Western – High fat, high simple sugar 0.0 Western LF/PP LF/PP Western **Generation 1 Generation 2**



Turnbaugh, et al. Sci. Transl. Med. 2010

FMT from lean donors reduces insulin resistance in obese subjects



Hepatic genes upregulated in GF versus conventionalized mice

p<0.05 Androgen and Estrogen Metabolism **PXR/RXR** Activation Methionine Metabolism LXR/RXR Activation Fatty Acid Metabolism Linoleic Acid Metabolism Circadian Rhythm Signaling Xenobiotic Metabolism Signaling Metabolism of Xenobiotics by Cytochrome P450 **Biosynthesis of Steroids** LPS/IL-1 Mediated Inhibition of RXR Function 0 3 6 9



Vanessa Leone

CHICAGC MEDICINE

Circadian clock networks regulate daily metabolic functions



Bass, J. Nature 2012



Germ free (GF) mice are resistant to obesogenic diets



n = 17 or 18 age-matched individually housed male mice/trt group

Germ-free mice have altered circadian mediobasal hypothalamic and liver circadian clock gene expression profiles



High fat dietary intake shifts cecal microbial diversity and taxonomic structure PCoA – PC1 vs. PC2 6.5₁ *** 0.15 Shannon Diversity Index -0. -2. -0. -0. -2. -0. -0. -0. **Highfat** _owfat 0.10 PC2 - 13.95% 0.05 0.00 3.5 Highfat Lowfat -0.05100% 90% -0.10 80% -0.15-0.10-0.05 0.00 0.05 0.10 0.15 0.20 70% PC1 - 56.20%

V4-V5 16s rRNA amplicon sequencing performed using Illumina Miseq

Leone, et al. Cell Host & Microbe (2015)

Verrucomicrobia Spirochaetes Cyanobacteria 60% Proteobacteria 50% Actinobacteria 40% Tenericutes 30% Bacteroidetes 20% 10% Firmicutes 0% Bacteria;Other Highfat Lowfat

Shifts in microbial diversity exhibit diurnal oscillations that are influenced by diet



Enteral feeding is not the only driver of diurnal variations in gut microbes





Bacteria;[Thermi]

- Verrucomicrobia
- Tenericutes
- Proteobacteria
- Gemmatimonadetes
- Firmicutes
- Cyanobacteria
- Chloroflexi
- Bacteroidetes
- Actinobacteria
 - Bacteria;Other

Oscillatory cecal OTUs found in mice fed regular, low fat chow (RC) diets



High fat diet-induced gut microbes exhibit altered diurnal oscillations of known microbial metabolites

Effects of SCFAs and H₂S (NaHS) on hepatocyte circadian clock function

MEDICINE

Effects of intraperitoneal administration of butyrate on CC networks of GF mice

Gut microbiota sense dietary cues and translating into an output that needed for Circadian networks

Impact of gut microbes on host metabolism

The challenge to treating obesity – changing energy balance

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