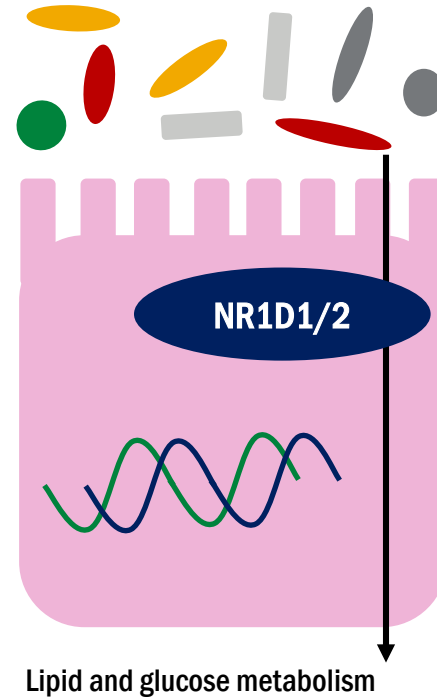


The gut microbiota regulates host metabolism through circadian clock genes *Nr1d1* and *Nr1d2*

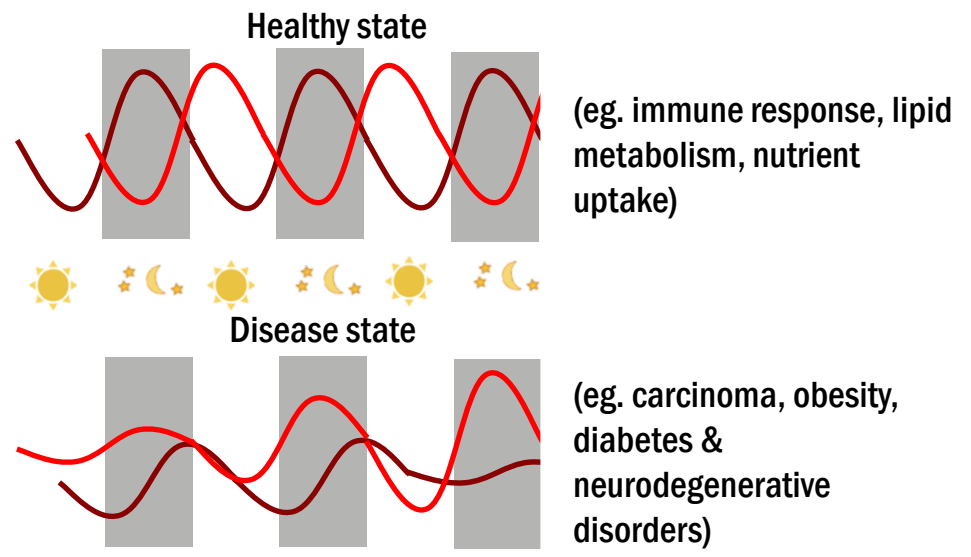
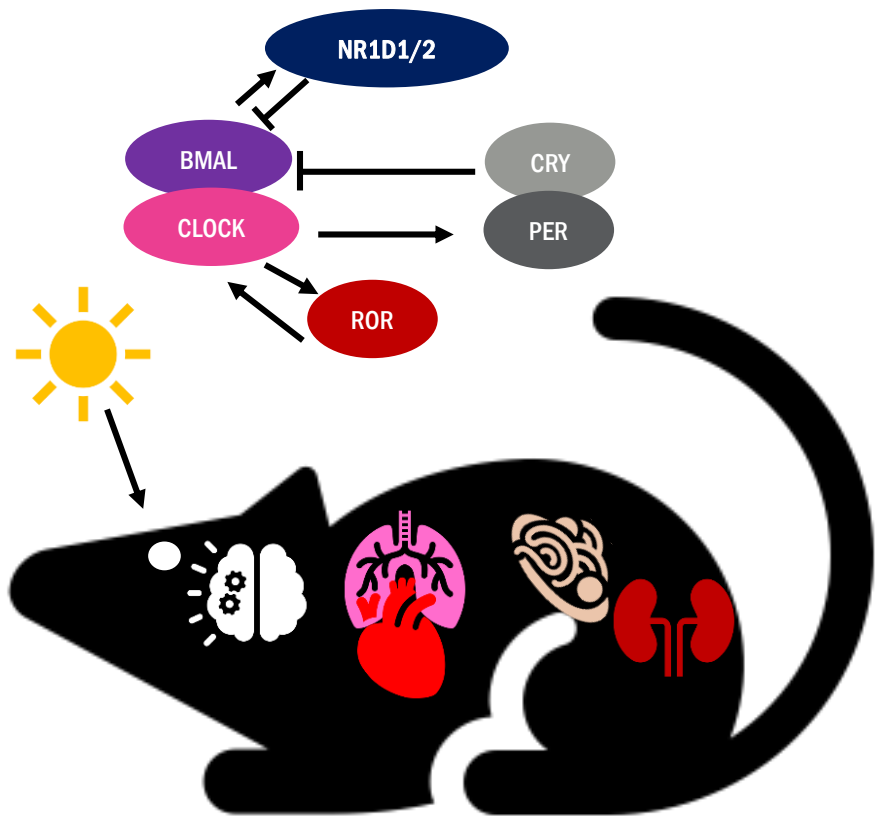
Samskrathi Sharma
Ph.D. student
Department of Biological Sciences

Advisor: Dr. Zheng Kuang

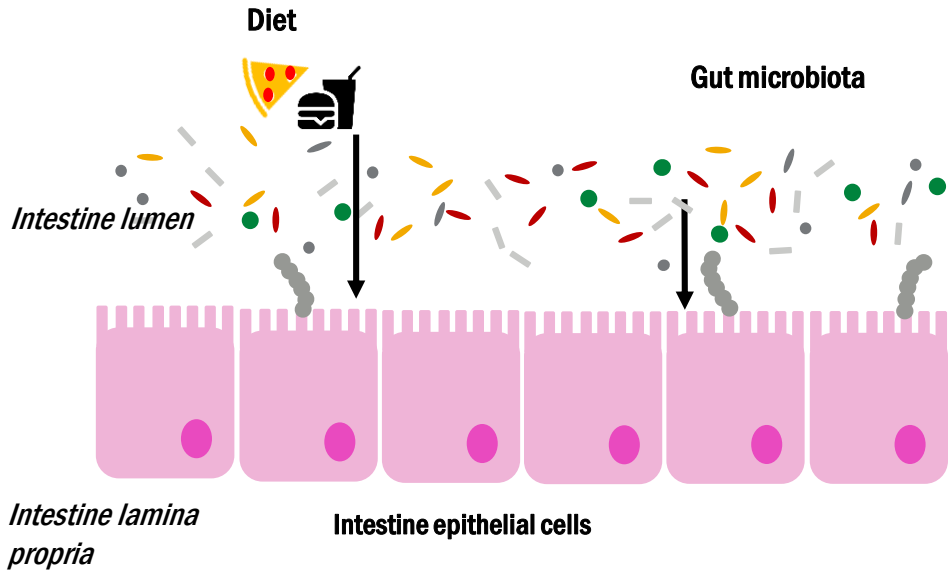
9th December 2023
6th International Conference on Microbiome Engineering
International House at UC Berkeley



Circadian rhythms are vital to health and are disrupted in disease



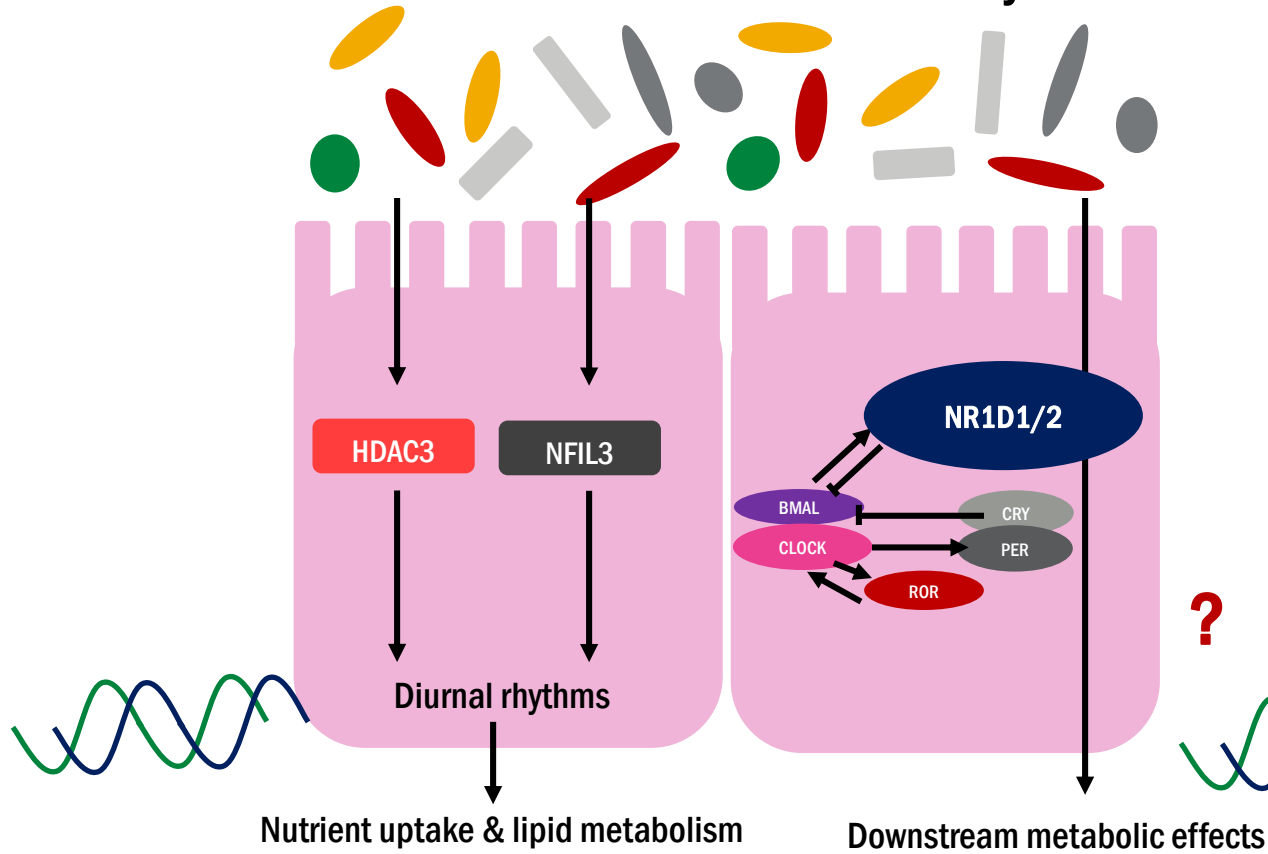
The intestine epithelial cells (IECs) are at the interface of host-gut microbiota interactions

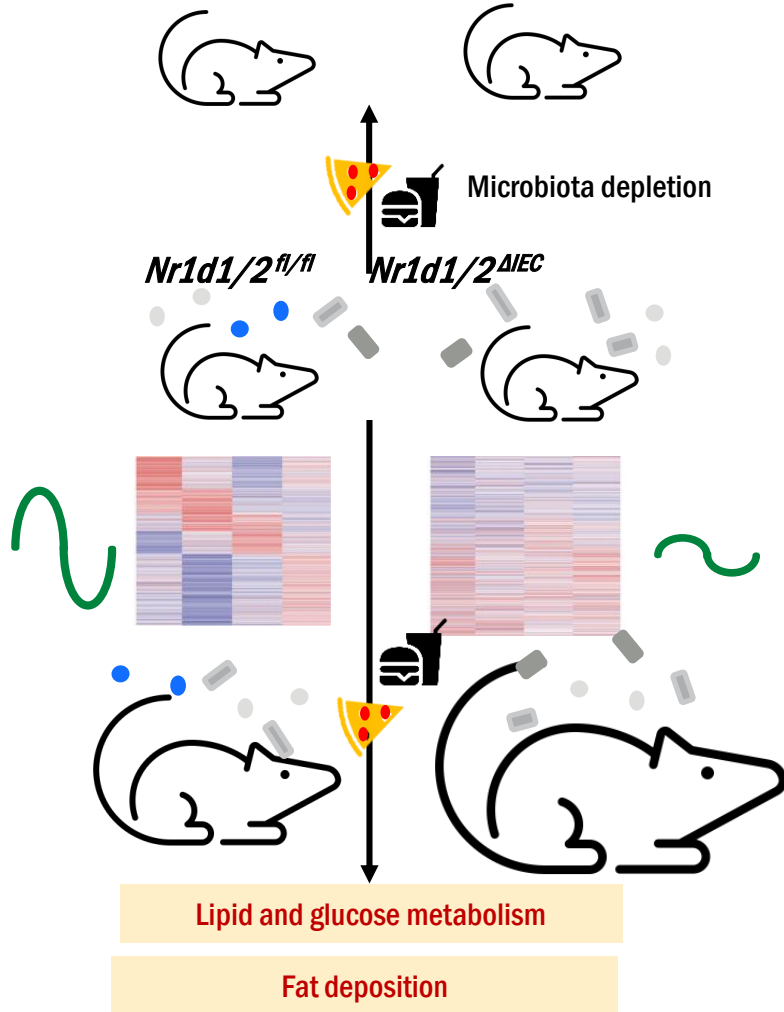


IECs are important for

- **nutrient digestion and absorption**
- **interacting with microbial components**
- **immunity**

How does the gut microbiota regulate host metabolism through core circadian clock machinery?





What did we find?

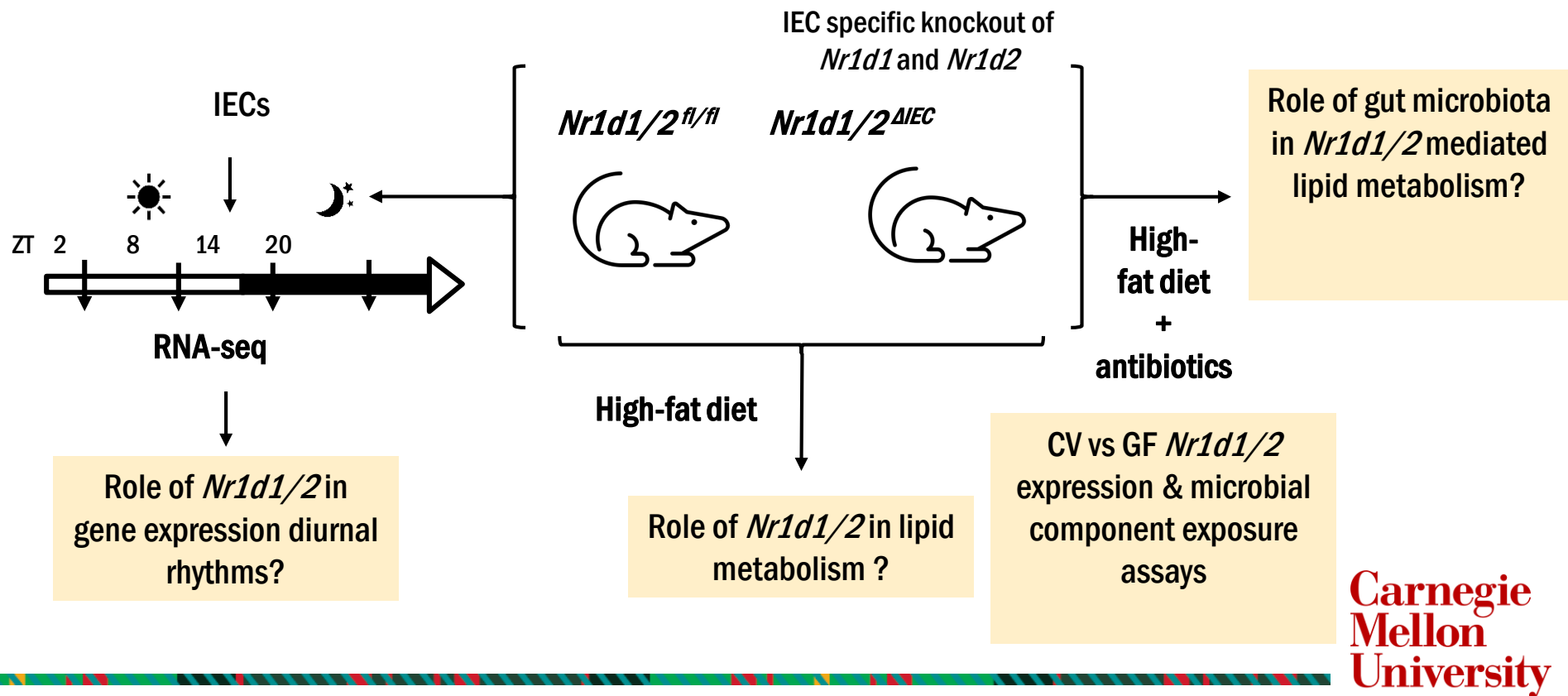
$Nr1d1/2^{\Delta IEC}$ mice have disrupted circadian rhythms of expression

$Nr1d1/2^{\Delta IEC}$ mice are more obese on a high-fat diet

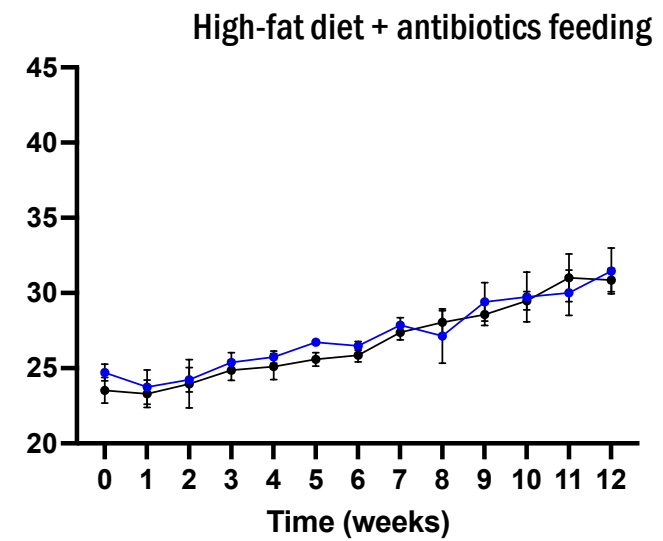
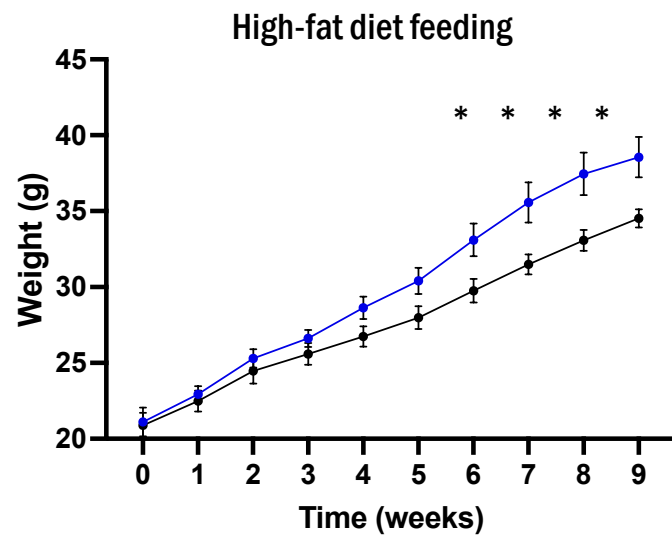
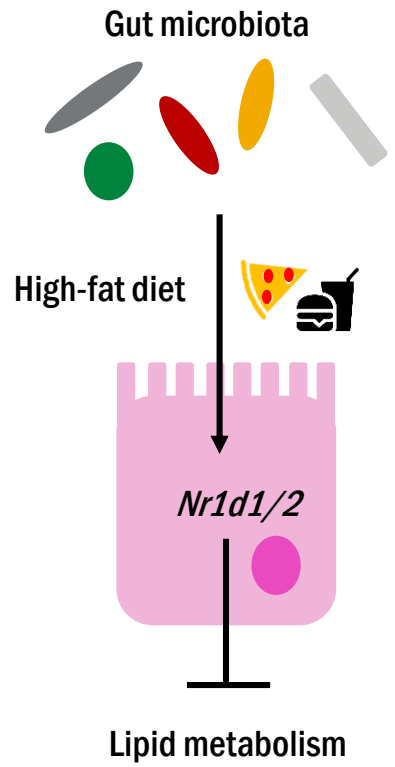
The gut microbiota is important for $Nr1d1/2^{\Delta IEC}$'s higher obesity

Microbial components LPS and flagellin modulate $Nr1d1/2$

What did we do?



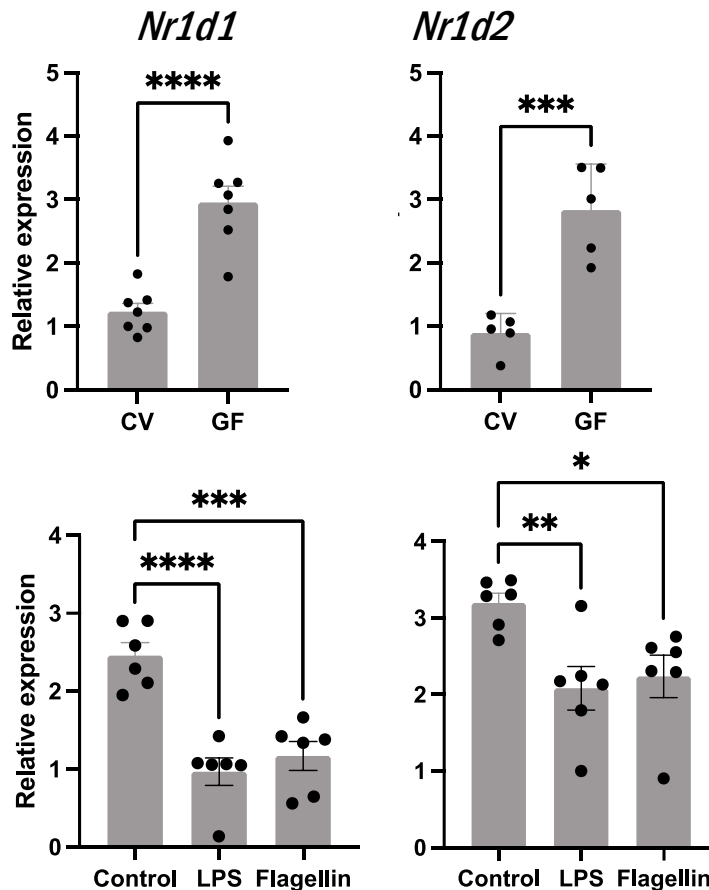
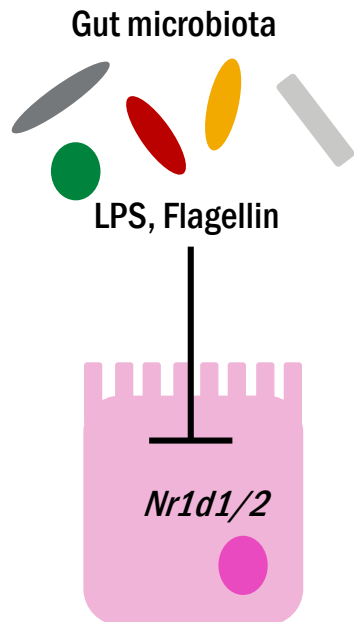
The gut microbiota is important for the higher obesity of *Nr1d1/2^{ΔIEC}* mice on a high-fat diet



● *Nr1d1/2^{fl/fl}*
 ● *Nr1d1/2^{ΔIEC}*

n=8 per group, Data is plotted as Mean±SEM

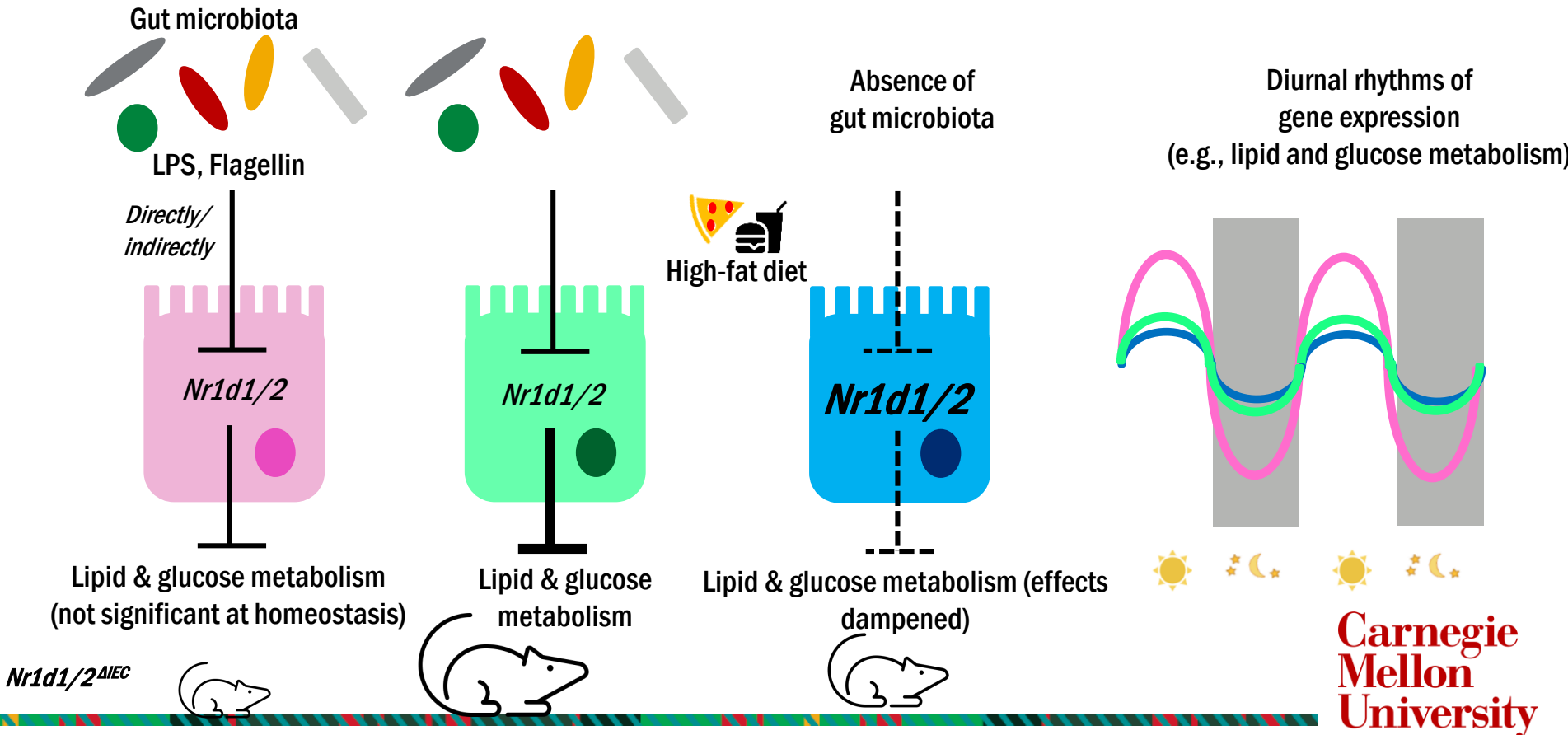
The gut microbiota represses *Nr1d1/2*



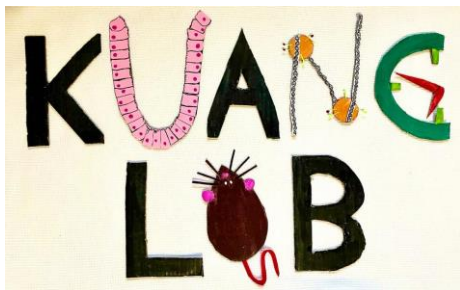
CV: Conventional mice
 GF: Germ-free mice
 LPS: Lipopolysaccharides

The gut microbiota regulates host metabolism through circadian clock

genes *Nr1d1* and *Nr1d2*



Acknowledgements



Dr. Zheng Kuang
Dr. Junjie Ma
Dr. Jianglin Zhang
Sarah Oladejo
Harihara Prakash
Iris Reed
Bhavana Kalidindi

Hooper Lab
 (UT Southwestern Medical Centre)
 CMU Animal Facility staff



R00 DK120897 and
DP2DK136278



AICHe Travel Grant



The Global Home of Chemical Engineers



U.S. DEPARTMENT OF
ENERGY

Office of Science



Margaret Carver Travel Grant
 Department of Biological Sciences

Carnegie
Mellon
University

Pronovost & Hsiao (2017)

