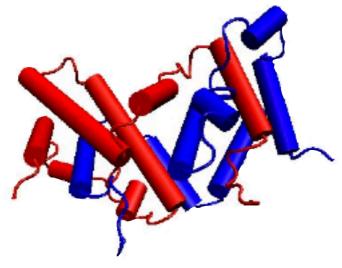


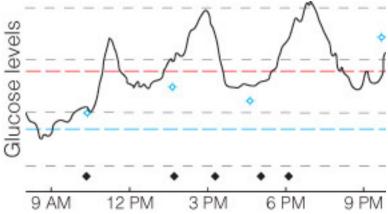


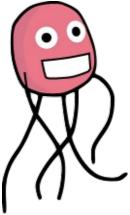
Observar e criar hipótese

- IFN-gamma atrapalha o metabolismo de glicose.
- Diabetes tipo 2 é um desequilíbrio no metabolismo.
- Akkermansia muciniphila associada à diabetes.
- Inflamação associada à diabetes.
 "Chronic inflammation impairs metabolic homeostasis
- and is intimately correlated with the pathogenesis
- of type 2 diabetes. The pro-inflammatory
- cytokine IFN-gamma is an integral part of the metabolic
- inflammation circuit and contributes significantly
- to metabolic dysfunction. The underlying mechanism,
- however, remains largely unknown."













Interferon gamma (IFN-γ) disrupts energy expenditure and metabolic homeostasis by suppressing SIRT1 transcription

Ping Li^{1,6}, Yuhao Zhao^{1,6}, Xiaoyan Wu^{2,6}, Minjie Xia¹, Mingming Fang^{1,4}, Yasumasa Iwasaki⁵, Jiahao Sha¹, Qi Chen¹, Yong Xu^{1,*} and Aiguo Shen^{3,*}

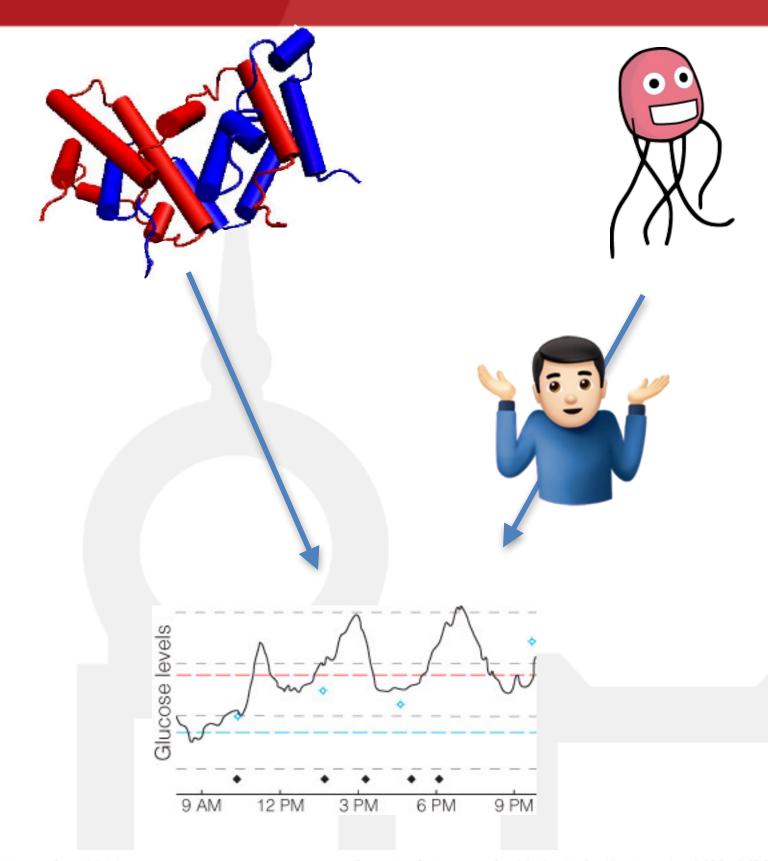
¹State Key Laboratory of Reproductive Medicine and Department of Pathophysiology, Key Laboratory of Cardiovascular Disease, ²Laboratory Center for Basic Medical Sciences, ³Institute of Gerontology, The Second Affiliated Hospital, Nanjing Medical University, Nanjing, China, ⁴Jiangsu Jiankang Vocational Institute, Nanjing, China and ⁵Health Care Center, Kochi University, Kochi, Japan

Observar e criar hipótese

- IFN-gamma atrapalha o metabolismo de glicose.
- Diabetes tipo 2 é um desequilíbrio no metabolismo.
- Akkermansia muciniphila associada à diabetes.
- Inflamação associada à diabetes.

"Chronic inflammation impairs metabolic homeostasis and is intimately correlated with the pathogenesis of type 2 diabetes. The pro-inflammatory cytokine IFN-gamma is an integral part of the metabolic inflammation circuit and contributes significantly to metabolic dysfunction. The underlying mechanism, however, remains largely unknown."





Published online 7 November 2011

Nucleic Acids Research, 2012, Vol. 40, No. 4 1609–1620 doi:10.1093/nar/gkr984

Interferon gamma (IFN-γ) disrupts energy expenditure and metabolic homeostasis by suppressing SIRT1 transcription

Ping Li^{1,6}, Yuhao Zhao^{1,6}, Xiaoyan Wu^{2,6}, Minjie Xia¹, Mingming Fang^{1,4}, Yasumasa Iwasaki⁵, Jiahao Sha¹, Qi Chen¹, Yong Xu^{1,*} and Aiguo Shen^{3,*}

¹State Key Laboratory of Reproductive Medicine and Department of Pathophysiology, Key Laboratory of Cardiovascular Disease, ²Laboratory Center for Basic Medical Sciences, ³Institute of Gerontology, The Second Affiliated Hospital, Nanjing Medical University, Nanjing, China, ⁴Jiangsu Jiankang Vocational Institute, Nanjing, China and ⁵Health Care Center, Kochi University, Kochi, Japan

Testar!



