

V МЕЖДУНАРОДНАЯ КОНФЕРЕНЦИЯ "ПОСТГЕНОМ'2018" В ПОИСКАХ МОДЕЛЕЙ ПЕРСОНАЛИЗИРОВАННОЙ МЕДИЦИНЫ

Казань 29 октября – 2 ноября 2018

УДК 612:616.34-008.87-098

РАЗРАБОТКА НОВЫХ ПОДХОДОВ ВОЗДЕЙСТВИЯ НА АППЕТИТ ЧЕРЕЗ КИШЕЧНУЮ МИКРОБИОТУ

DEVELOPMENT OF MICROBIOTA-BASED APPROACHES FOR TARGETING HOST APPETITE

Sergueï O. Fetissov

University of Rouen Normandy, Inserm UMR1239, France

To develop microbiota-based approaches for targeting host appetite an important task is to understand molecular mechanisms linking the gut bacteria with the host pathways regulating feeding behavior (1). Using a proteomic approach, a molecular target has been identified in *Enterobacteriaceae* by linking these common gut bacteria with the host melanocortin system, critically involved in the regulation of energy balance. Bacterial protein homologue of caseinolytic protease B (ClpB) has been identified as a conformational mimetic of α -melanocyte-stimulating hormone (α -MSH), an anorexigenic neuropeptide (2). *E.coli* ClpB is present in both gut lumen and plasma and is able to activate directly intestinal and central satiety pathways, respectively (3). These results served for TargEDys SA, a start-up company, as the scientific background for development of the probiotic aimed to reduce appetite and body weight. Recent update on the development of this probiotic will be presented.

References

- Fetissov S.O. Role of the gut microbiota in host appetite control: bacterial growth to animal feeding behaviour. Nat Rev Endocrinol 13:11-25, 2017.
- Tennoune N., et al. Bacterial ClpB heat-shock protein, an antigen-mimetic of the anorexigenic peptide [alpha]-MSH, at the origin of eating disorders. Transl Psychiatry 4:e458, 2014.
- 3. Breton J., et al. Gut commensal E. coli proteins activate host satiety pathways following nutrient-induced bacterial growth. *Cell Metab* 23:1-11, 2016.