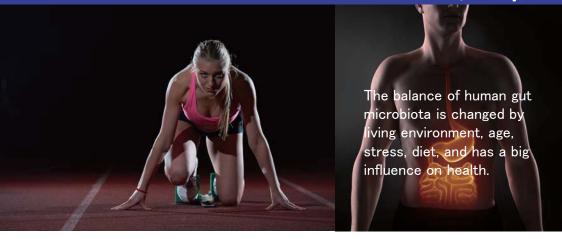
Evaluation system of human healthy by its microbiome

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ORIGINAL ARTICLE

Exercise and associated dietary extremes impact on gut microbial diversity

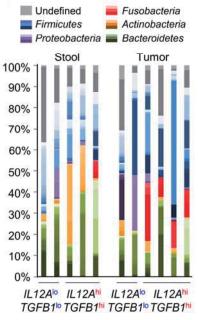
Clarke et al, Gut, 2014

Athletes and controls differed significantly with respect to plasma creatine kinase (a marker of extreme exercise), and inflammatory and metabolic markers. More importantly, athletes had a higher diversity of gut micro-organisms, representing 22 distinct phyla, which in turn positively correlated with protein consumption and creatine kinase.



Gut microbiota

Dysbiosis of gut microbiota on various disease

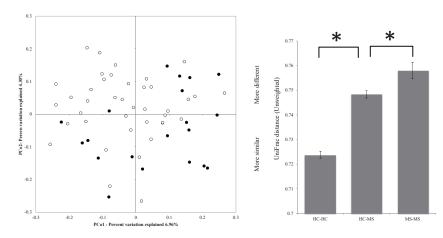


Two FOXP3+CD4+ T-cell subpopulations distinctly control the prognosis of colorectal Sequencing of bacterial 16S ribosomal DNA detected Fusobacterium nucleatum in IL12AhiTGFB1hi, but not in IL12AloTGFB1lo CRC tissues or the stool of either type. Saito T, Nishikawa H, Morita H, Honda K and Sakaguchi S, et al., Nature Medicine, in press

Higher diversity of gut microbiota of top athletes

PC2 (4.8%) Healthy group thlete group

Dysbiosis of gut microbiota of multiple sclerosis (MS)



Unweighted UniFrac Principal Coordinate (PCoA) and UniFrac distance analysis for $HC40(\bigcirc)$ and $MS20(\bigcirc)$. Miyake S, Morita H, Hattori M, Yamamura T, et al., PLoS One, 2015

Our health is kept to good condition by various metabolites, which were produced by gut microorganisms, derived from proteins, vitamins, minerals and sugars.