

The mechanistic link between health and gut microbiota diversity

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SUPPLEMENTARY INFORMATION

Simulations. All simulations were performed using R statistical software (version 3.4.0) within an RStudio environment (Version 1.0.143). Simulations were executed on a HP Z640 workstation containing an Intel(R) Xeon(R) CPU E5-2620 v4 @ 2.10GHz, with 64GB RAM. Simulations were partly performed in parallel (multicore).