Genome-scale community modelling reveals conserved metabolic cross-feedings in epipelagic bacterioplankton communities

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Description of Additional Supplementary Files

Supplementary data legends:

File name: Supplementary Data 1

Description: List of all genomes (WGS, MAG, SAG) included in the study.

File name: Supplementary Data 2

Description: Genome summary statistics.

File name: Supplementary Data 3

Description: Scaling law-based comparative statistics for genome types.

File name: Supplementary Data 4

Description: Tara Oceans samples contextual parameters (metadata).

File name: Supplementary Data 5

Description: Principal Coordinates Analyses correlative statistics.

File name: Supplementary Data 6

Description: Scaling law-based comparative statistics for (co-)active genomes.

File name: Supplementary Data 7

Description: List of predicted metabolic exchanges in each co-active community.

File name: Supplementary Data

Description: Statistics of predicted metabolic exchanges in each co-active community type.