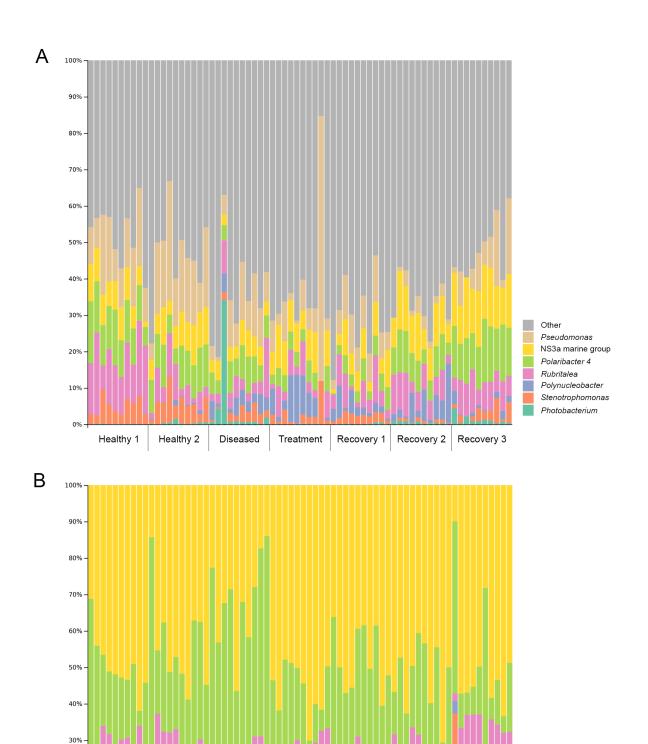
Supplementary information

Effects of disease, antibiotic treatment and recovery trajectory on the microbiome of farmed seabass (*Dicentrarchus labrax*)

Daniela Rosado, Raquel Xavier*, Ricardo Severino, Fernando Tavares, Jo Cable, Marcos Pérez-Losada

*Corresponding author: Raquel Xavier - raq.xavier@cibio.up.pt

Supplementary Figure



Supplementary Figure S1: Individual microbial variation of the most abundant (>5%) genera recovered from the gill (A) and skin (B) of the seabass. Distinctive bars represent relative abundance of each genus.

Recovery 1

Recovery 2

Recovery 3

10%

0%

Healthy 1

Healthy 2

Diseased

Treatment

Other

NS3a marine group
Polaribacter 4

Stenotrophomonas Pseudoalteromonas