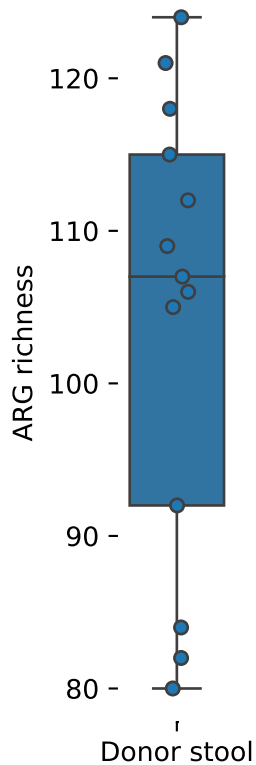
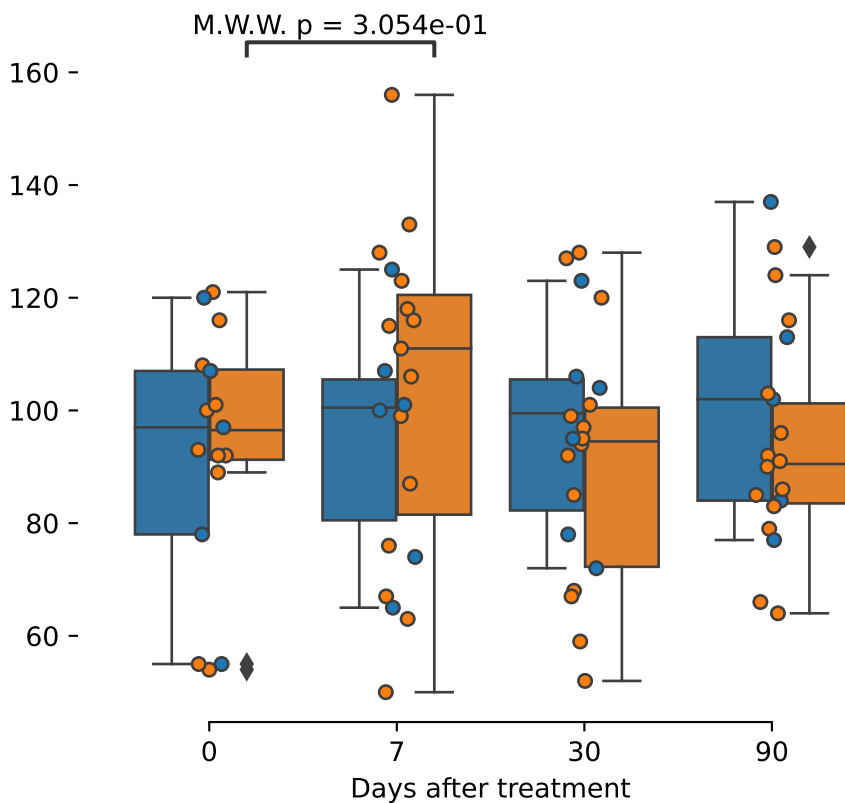


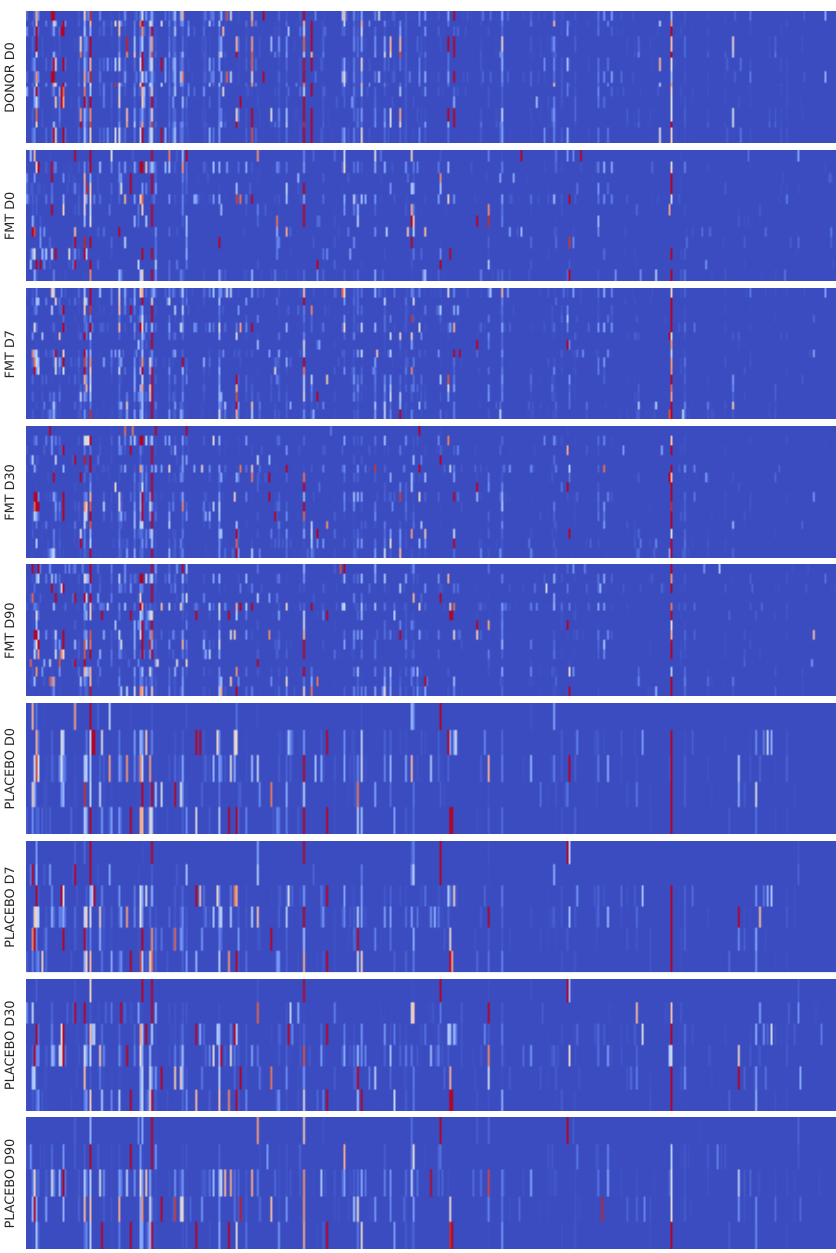
donor



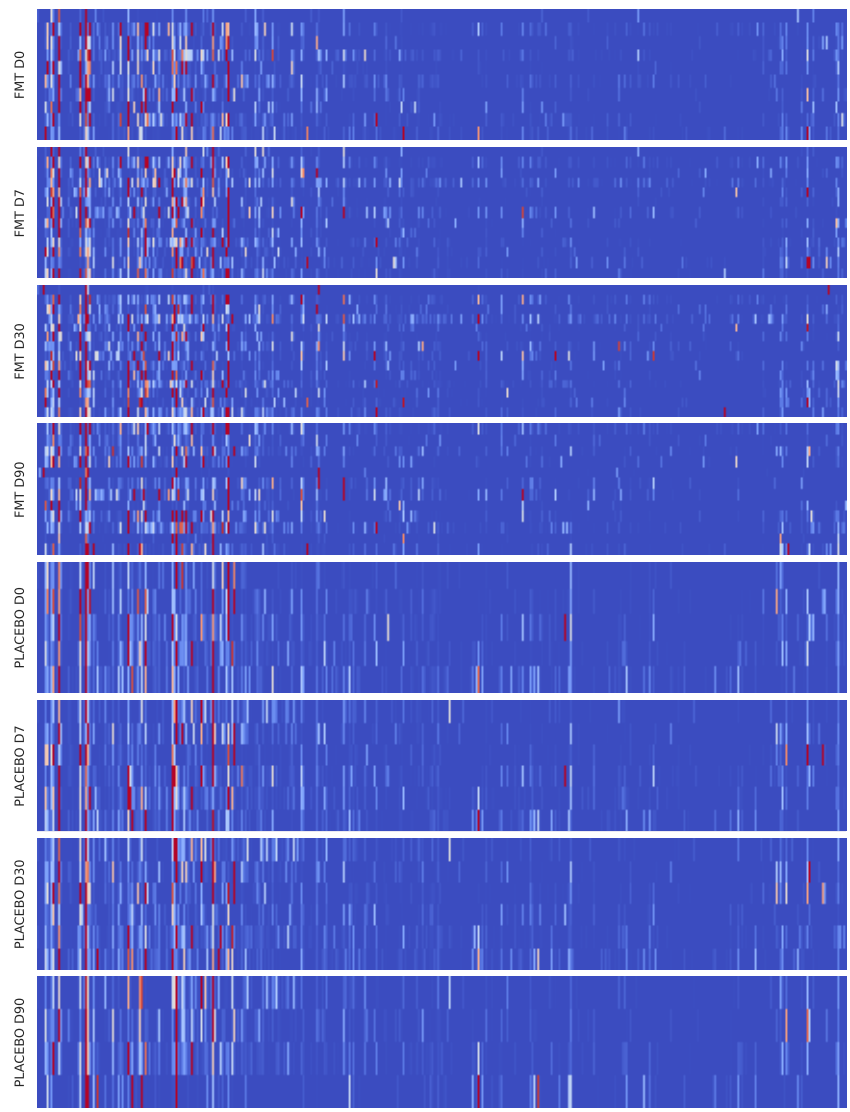
gut

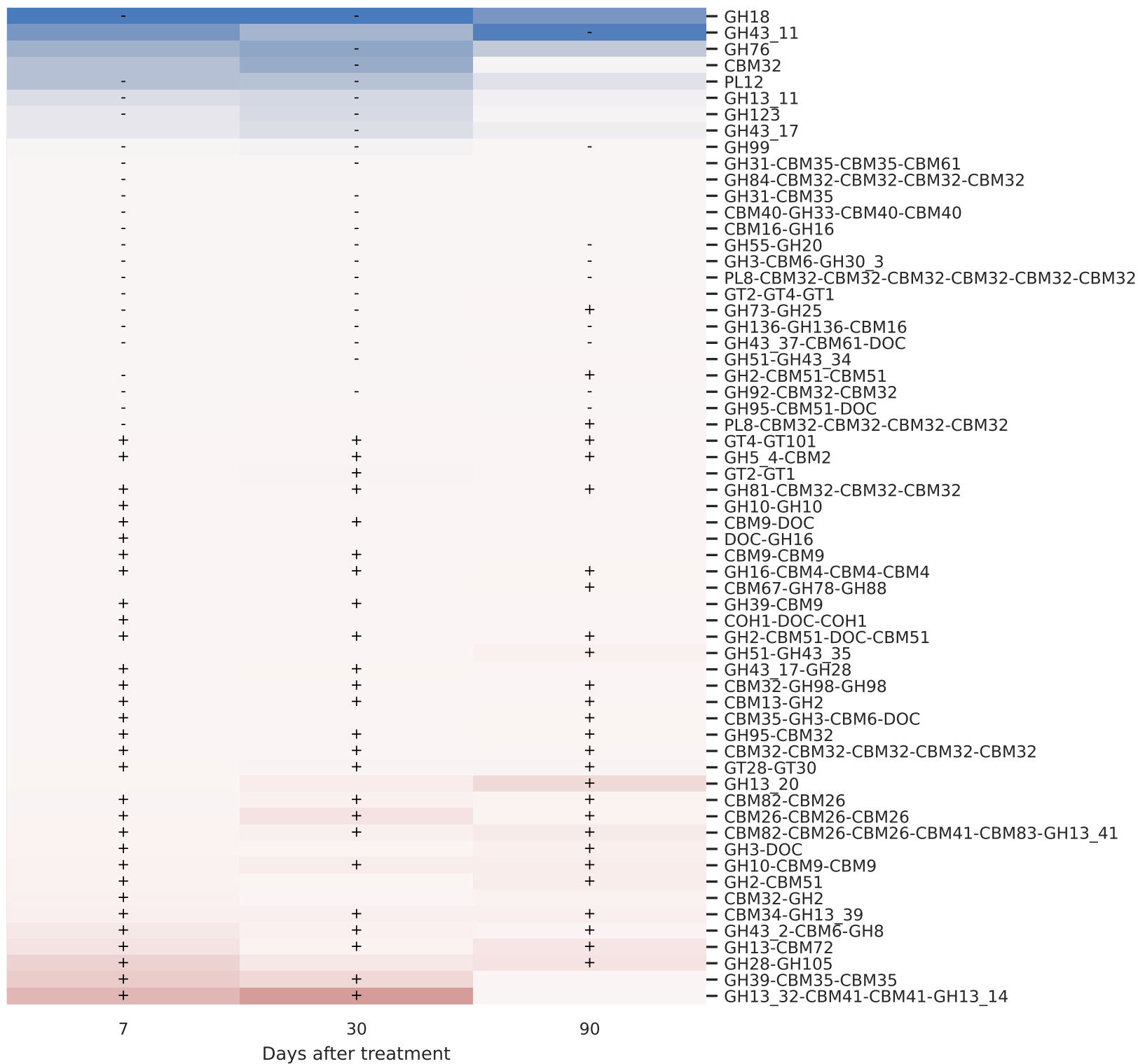
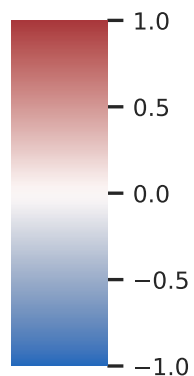


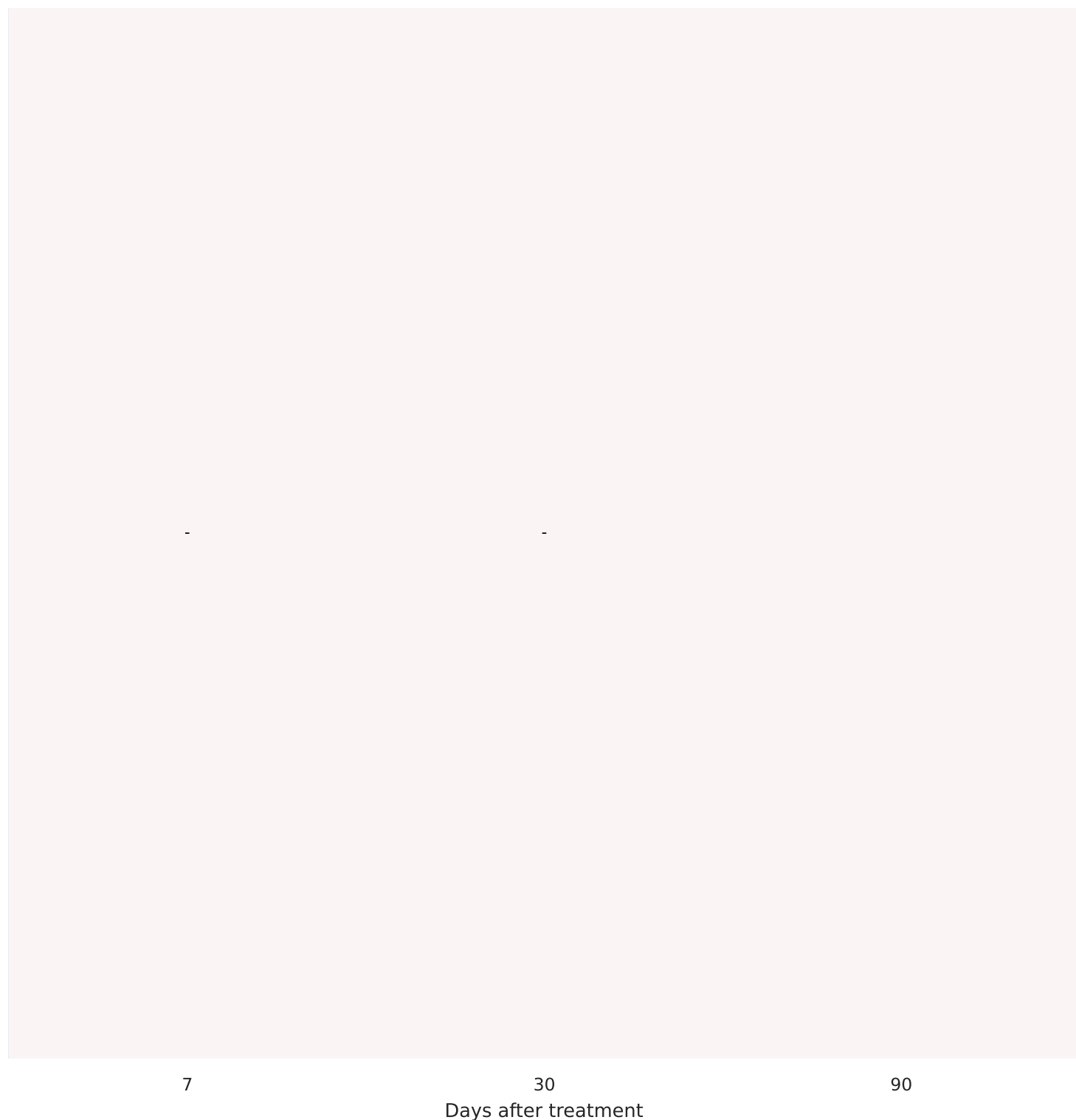
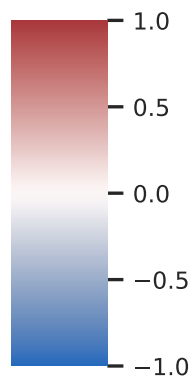
Stool

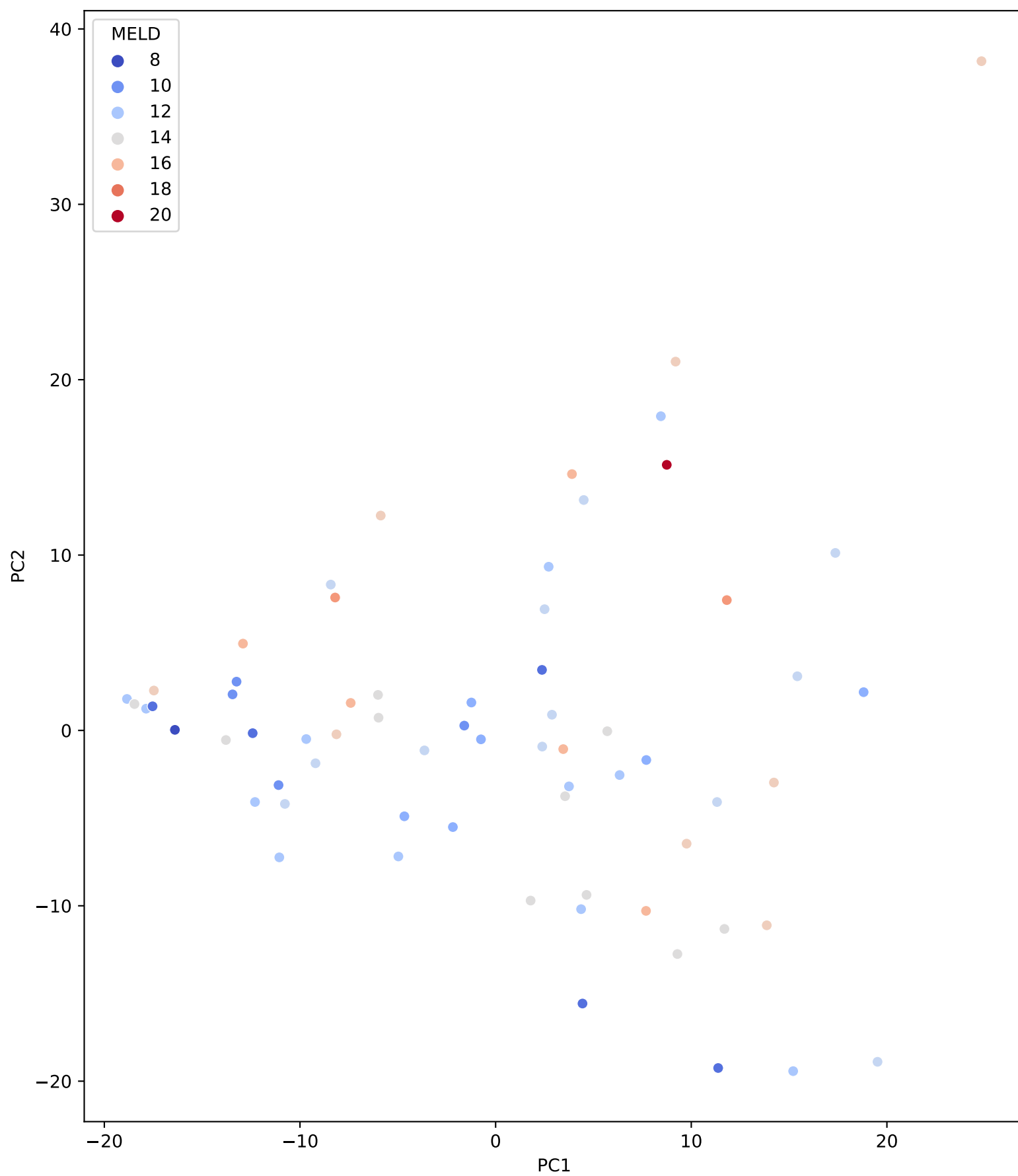


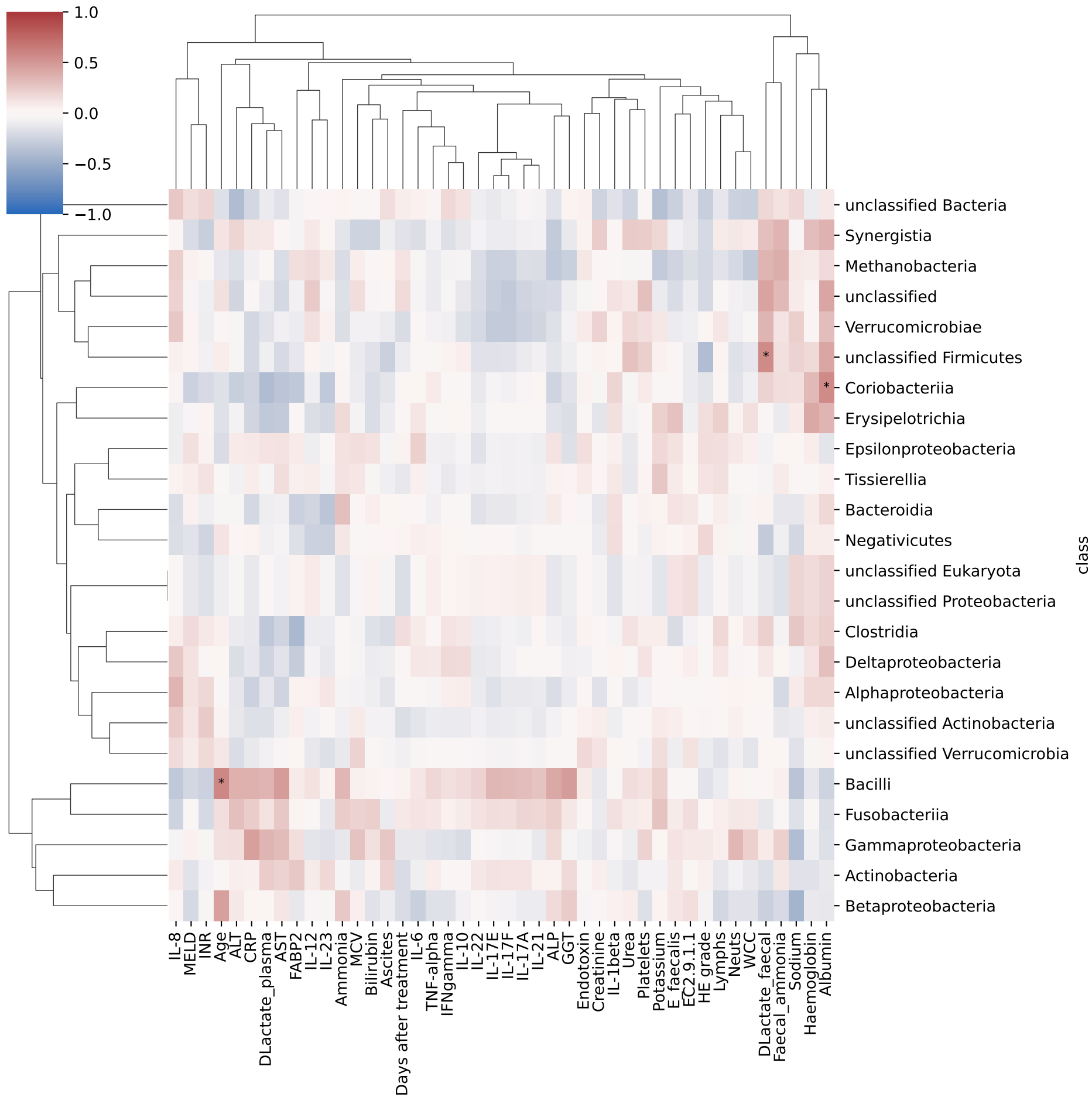
Saliva



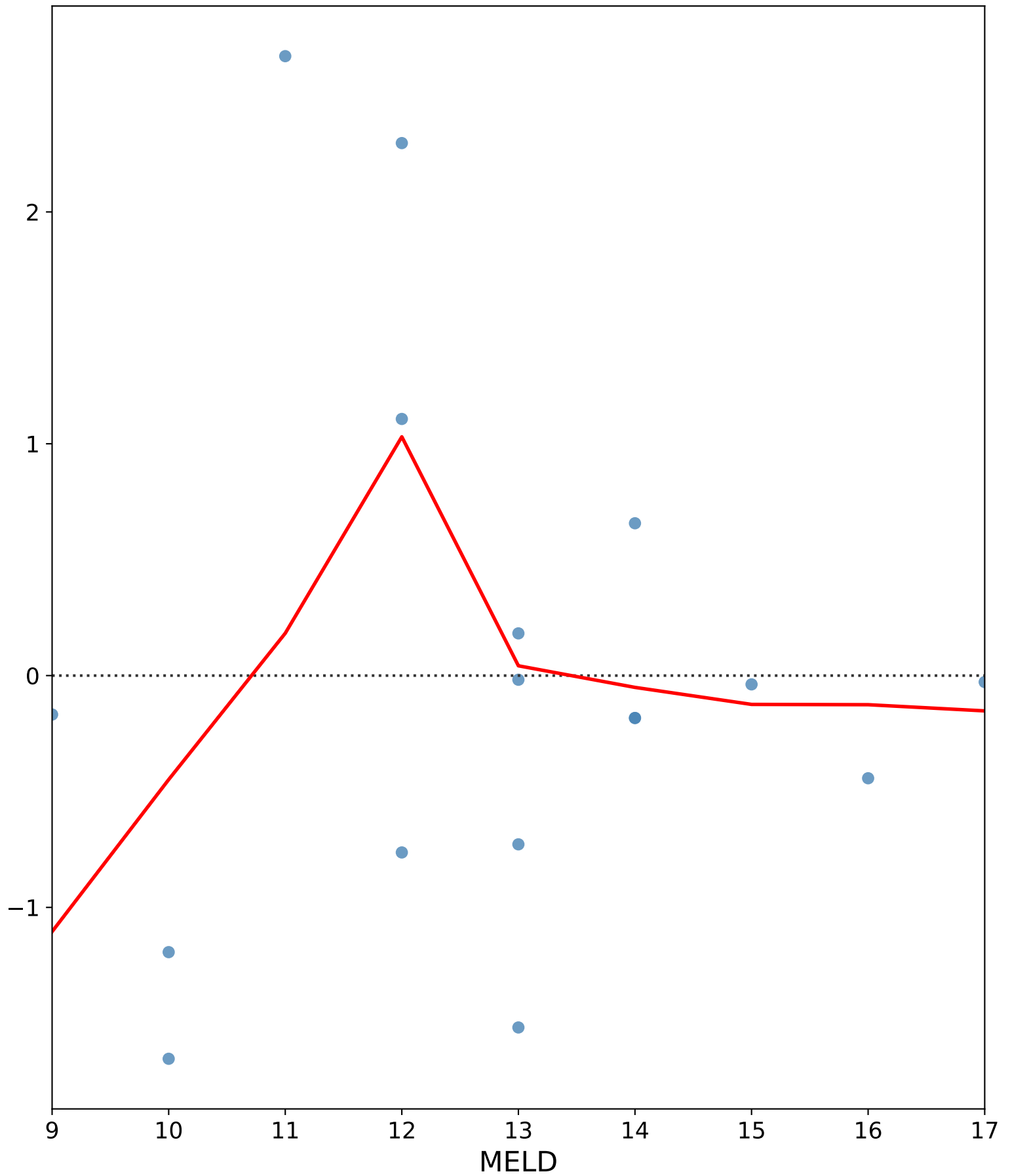


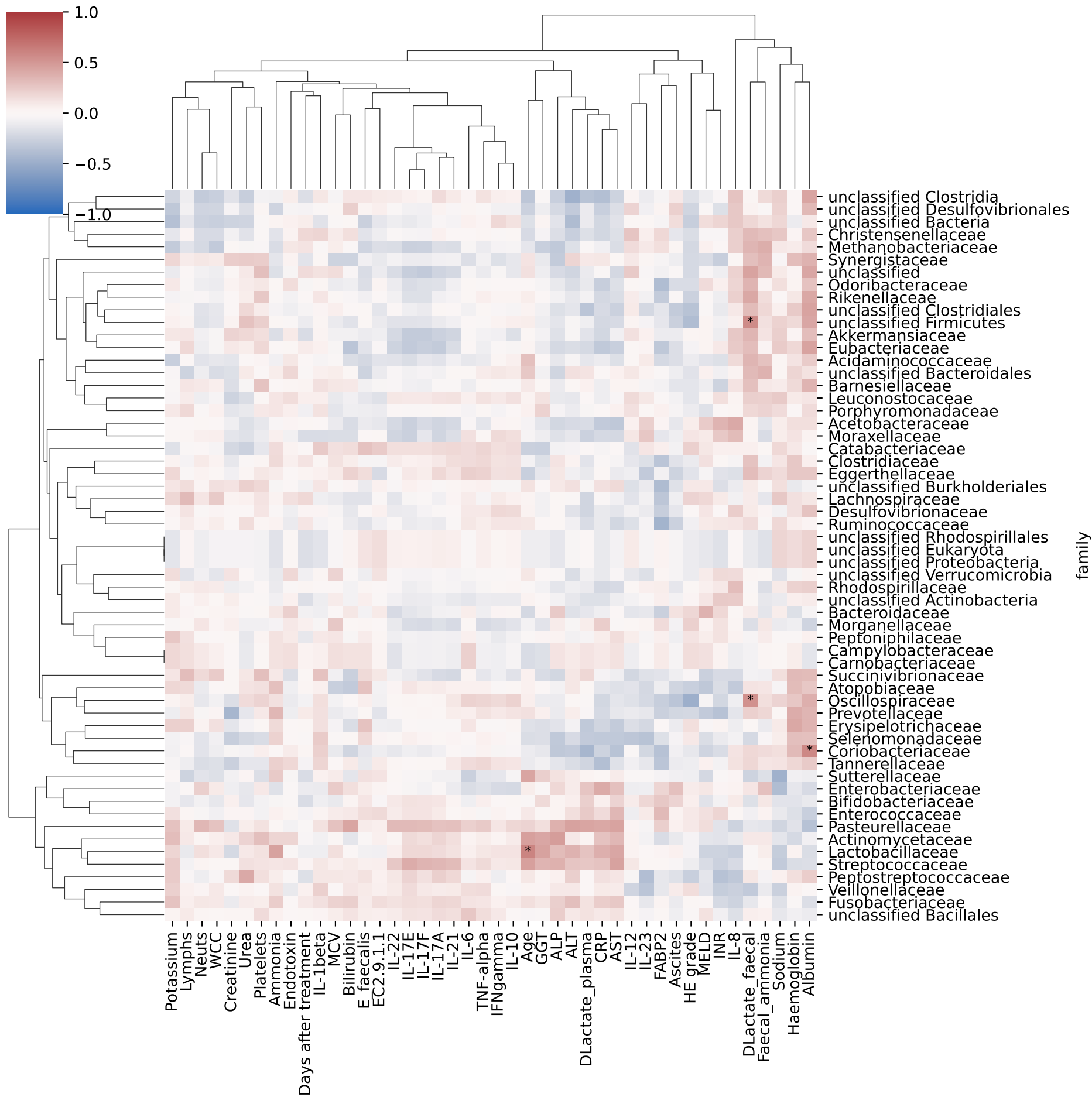


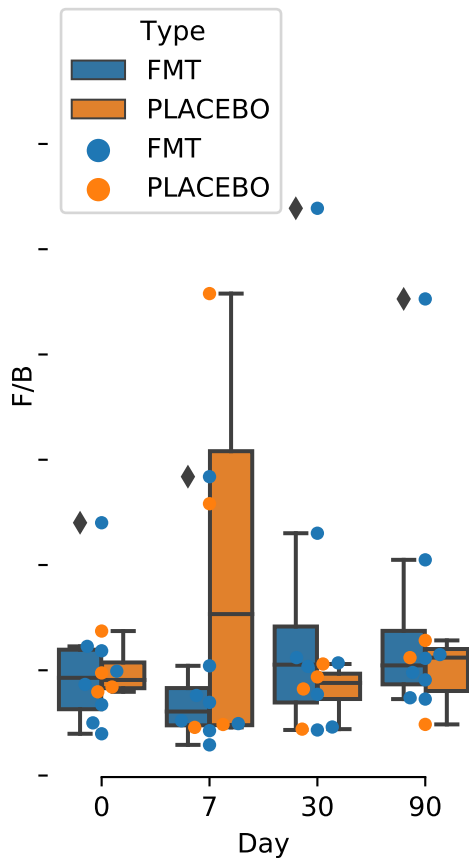
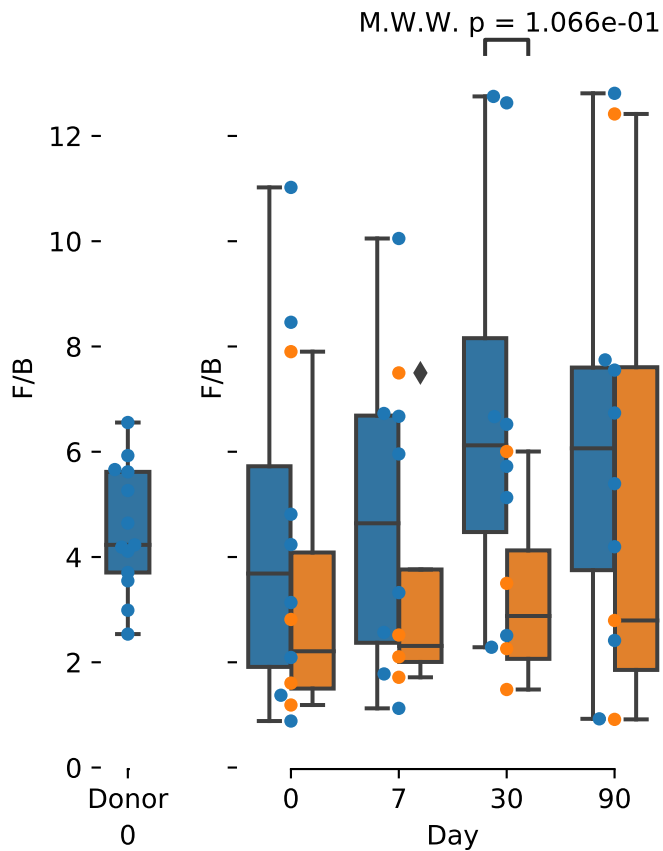


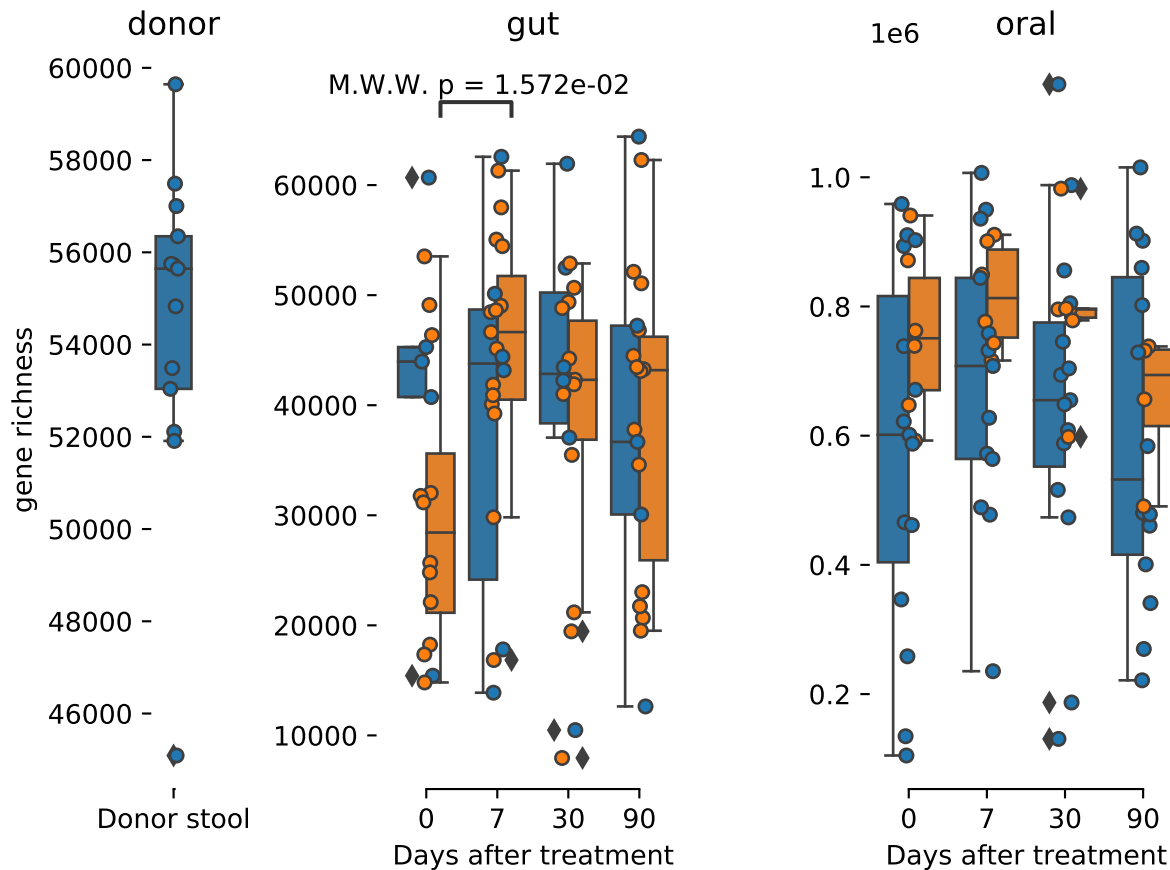


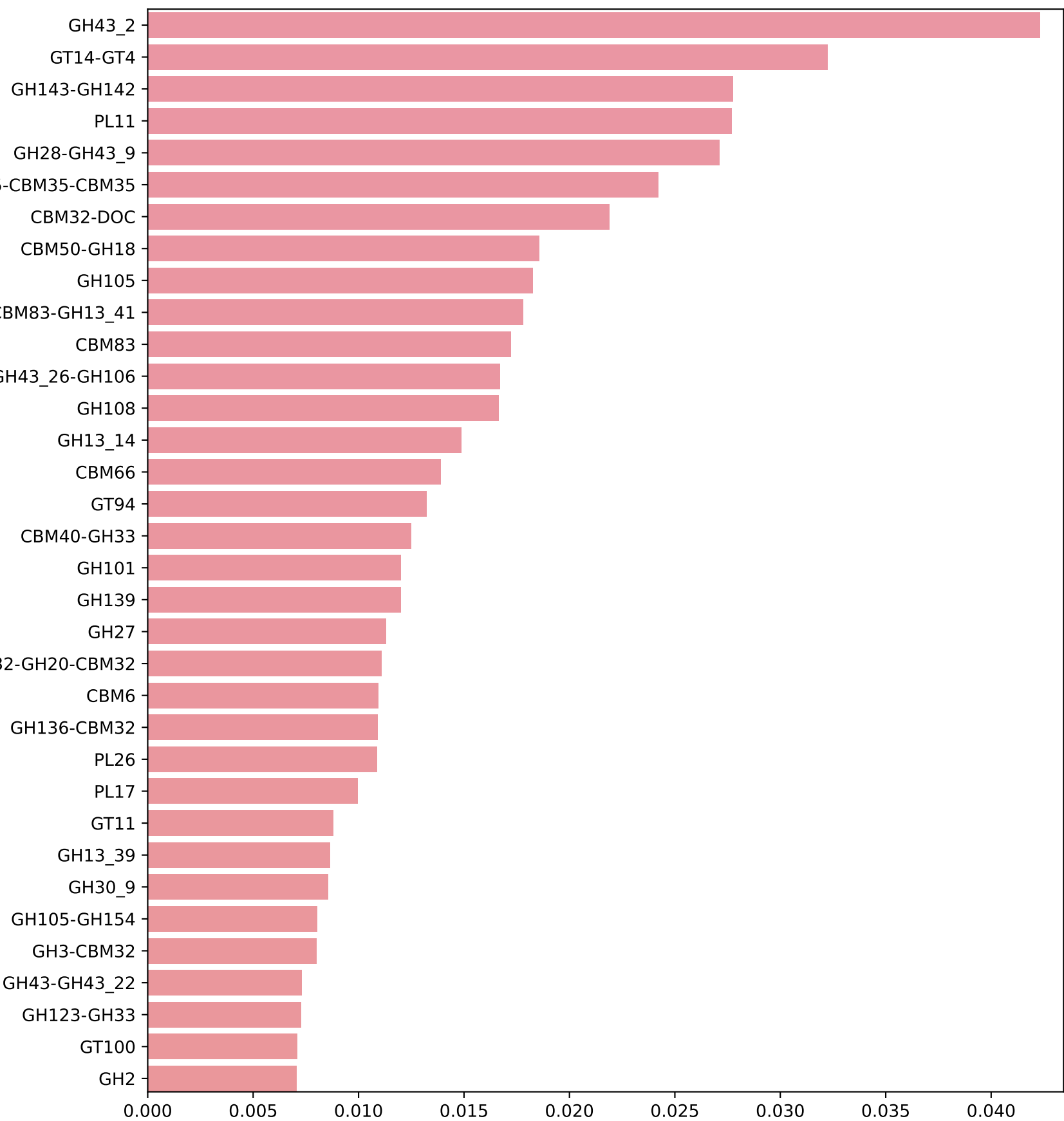
MELD vs Residuals for Histogram-based Extra Trees Regressor



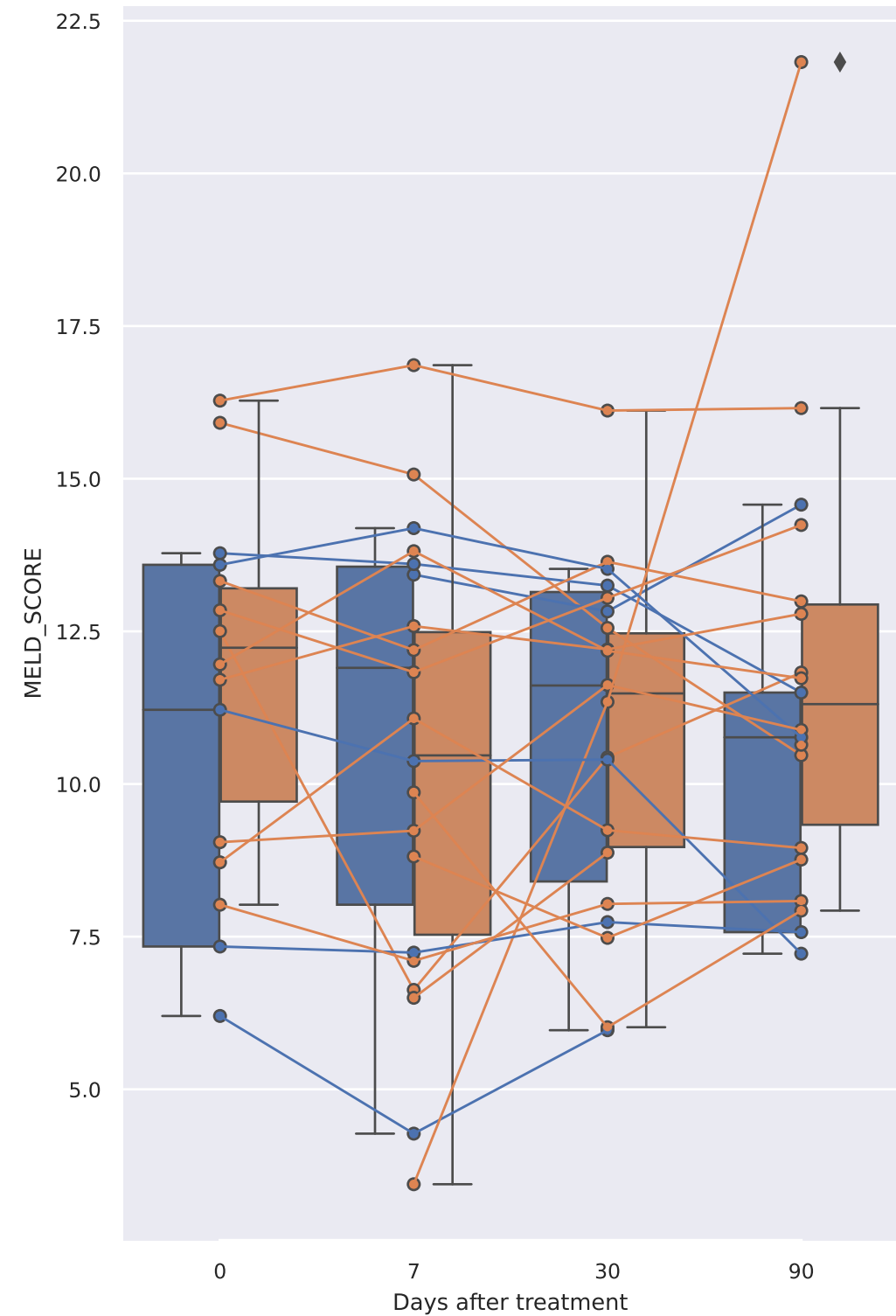




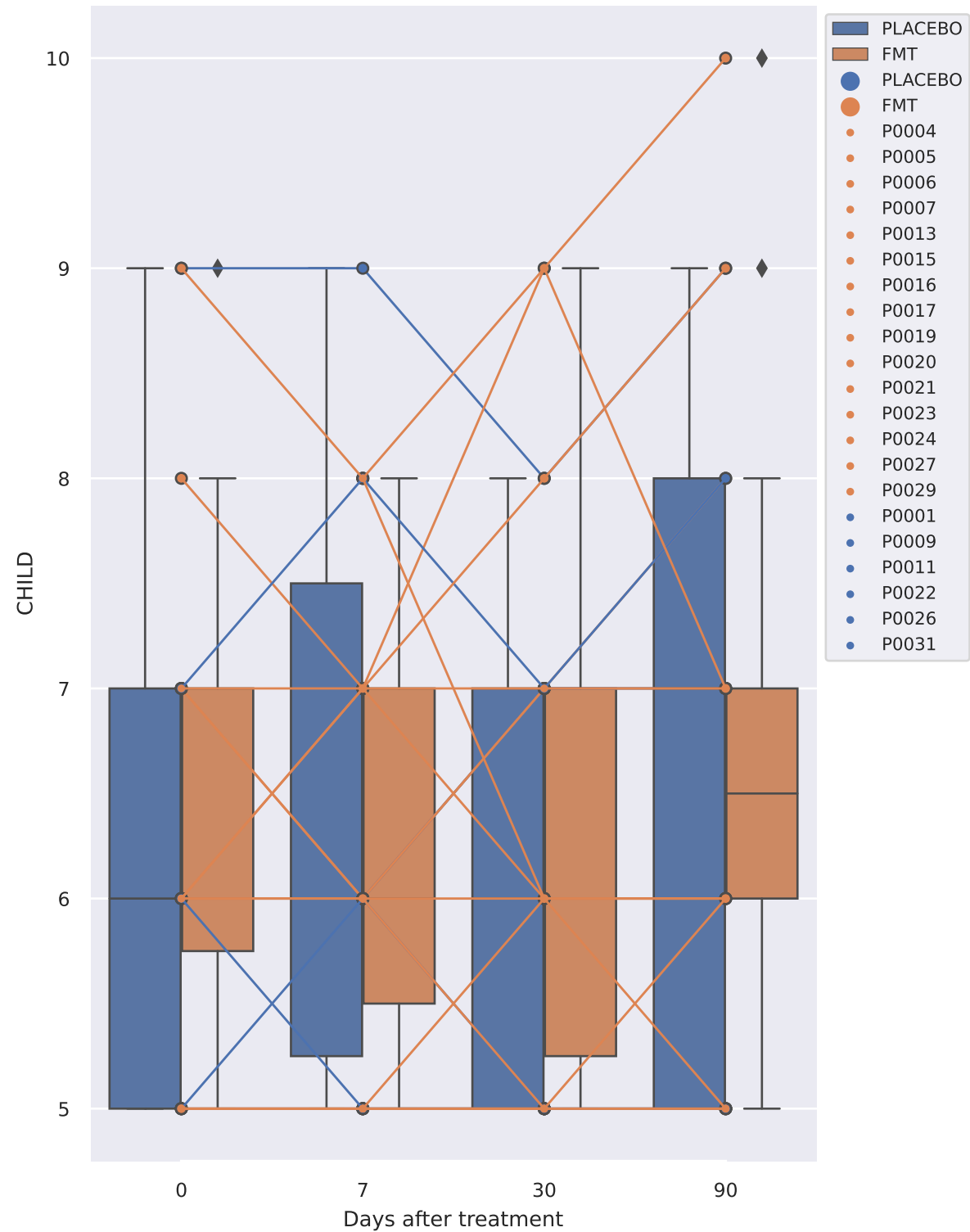


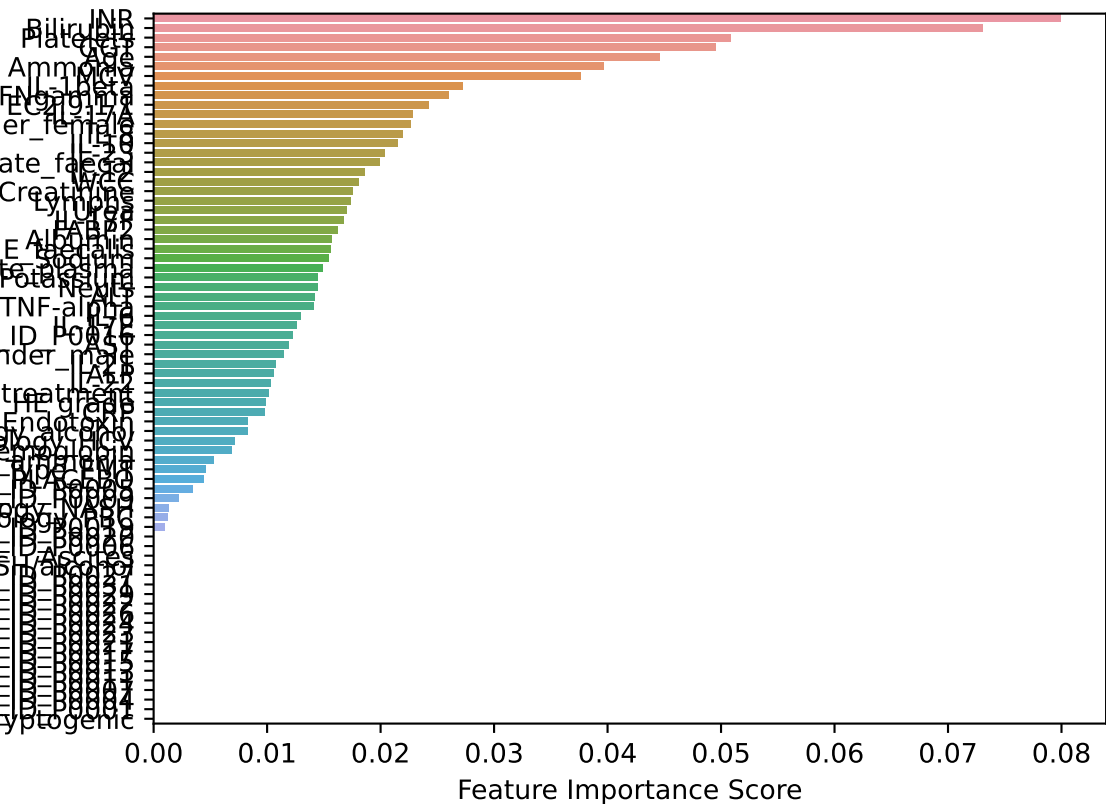


Patient stool

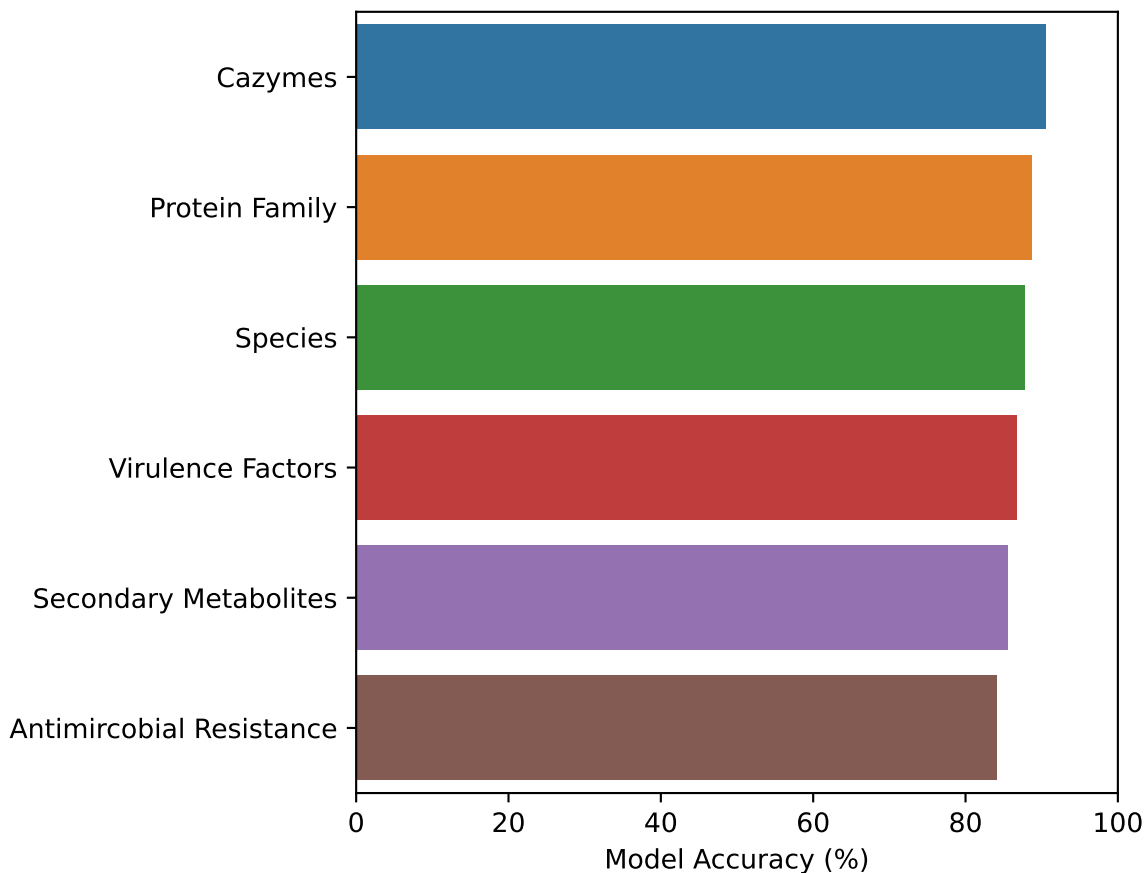


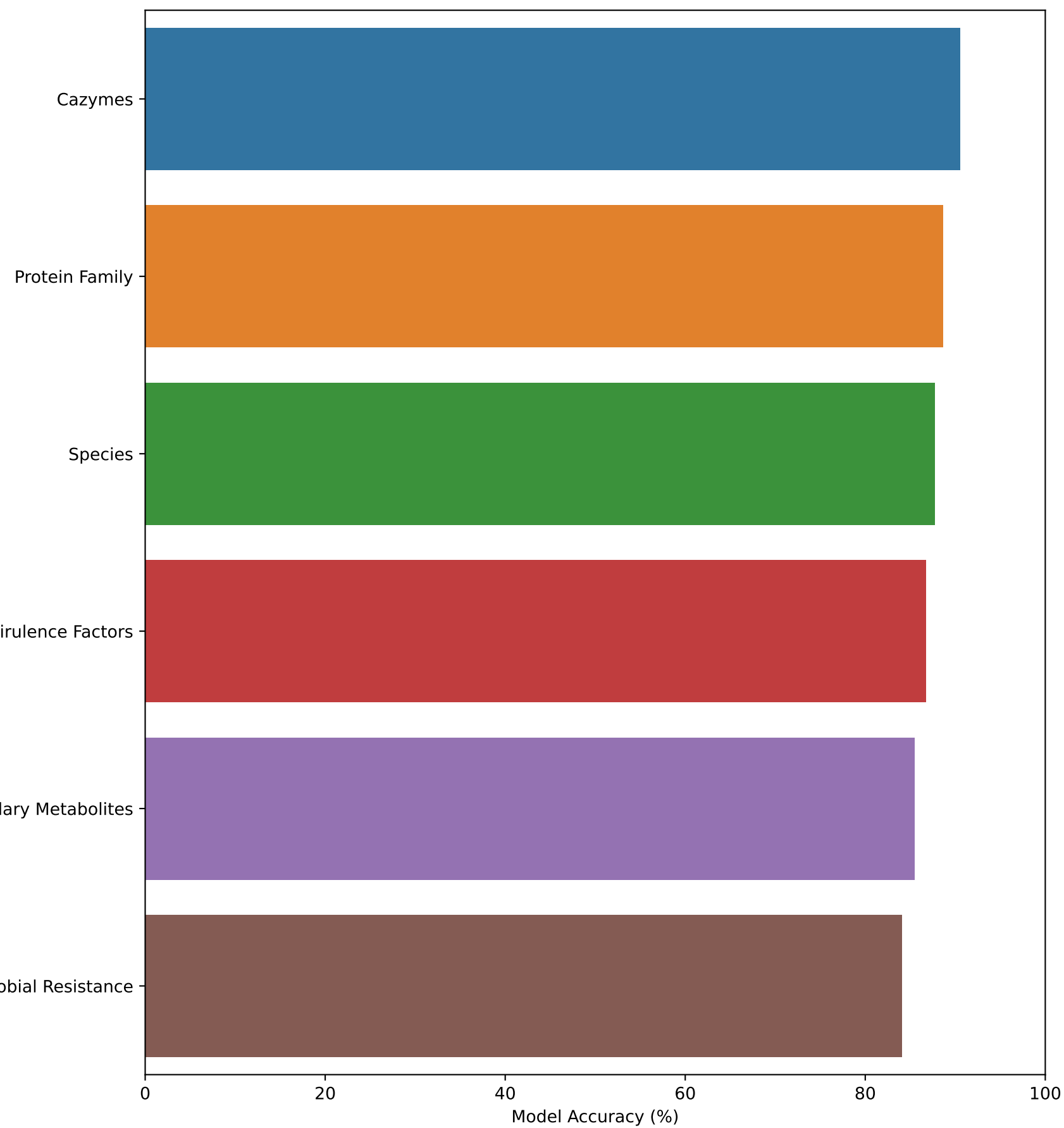
Patient stool



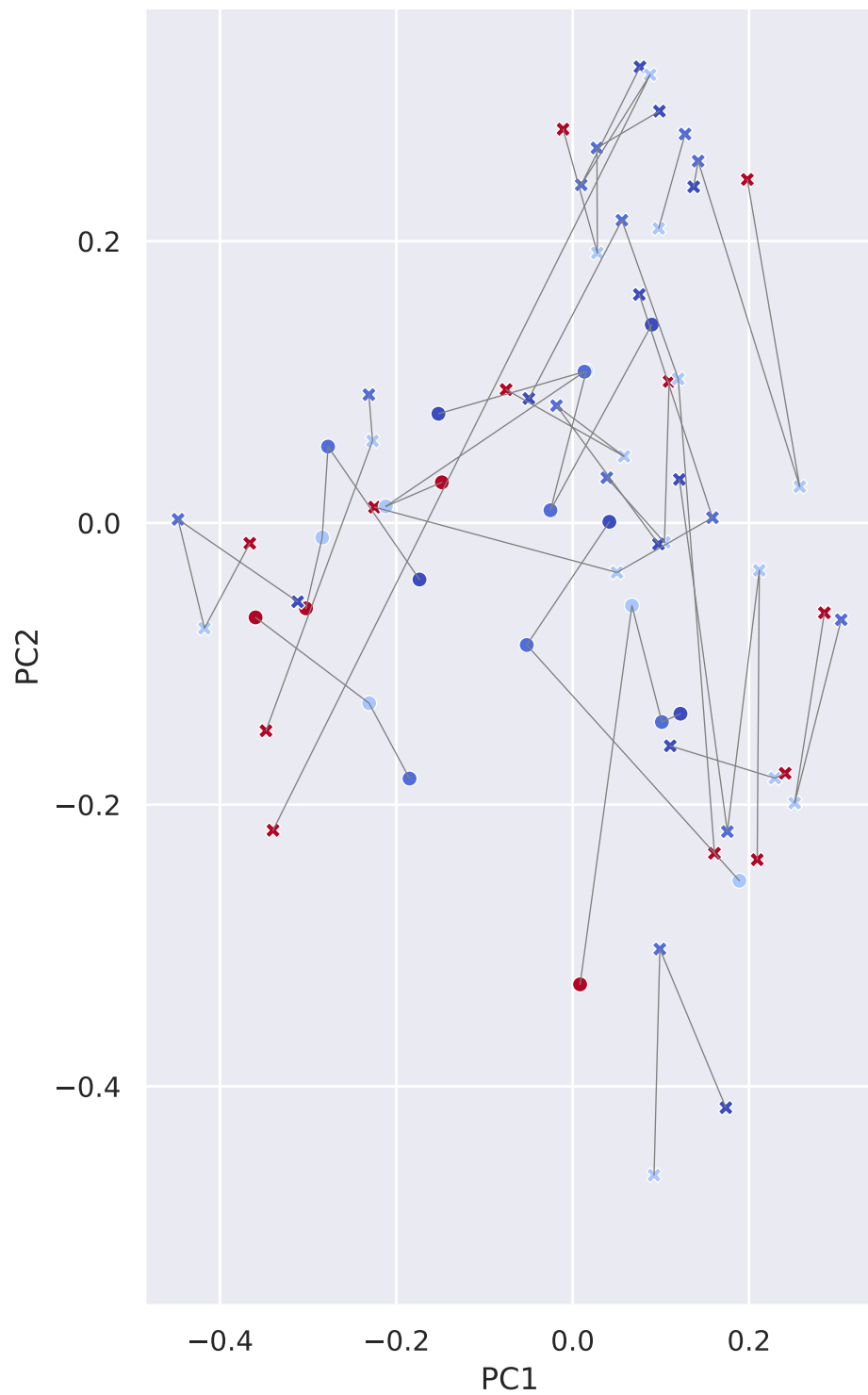


Dataset used for MELD prediction

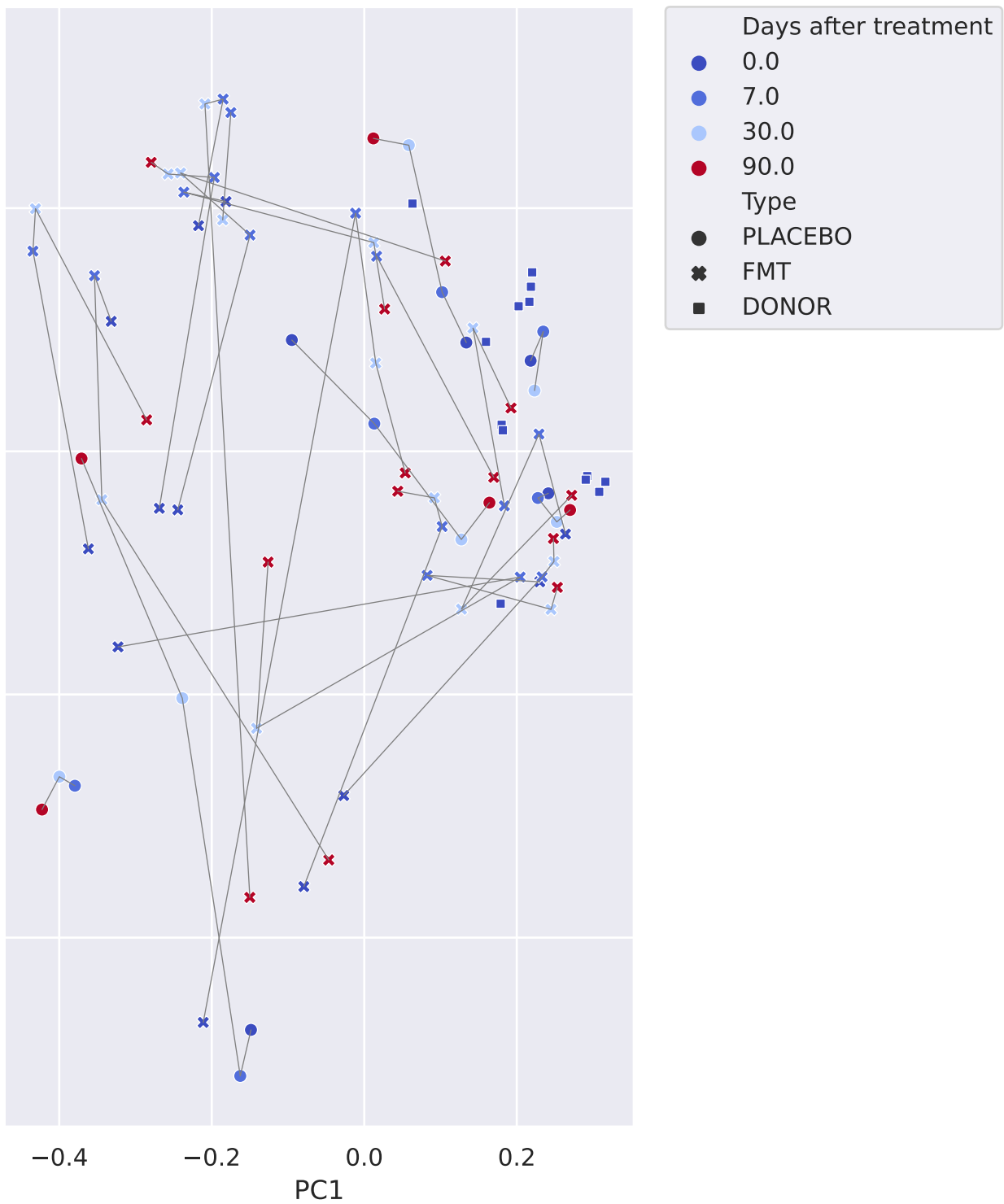


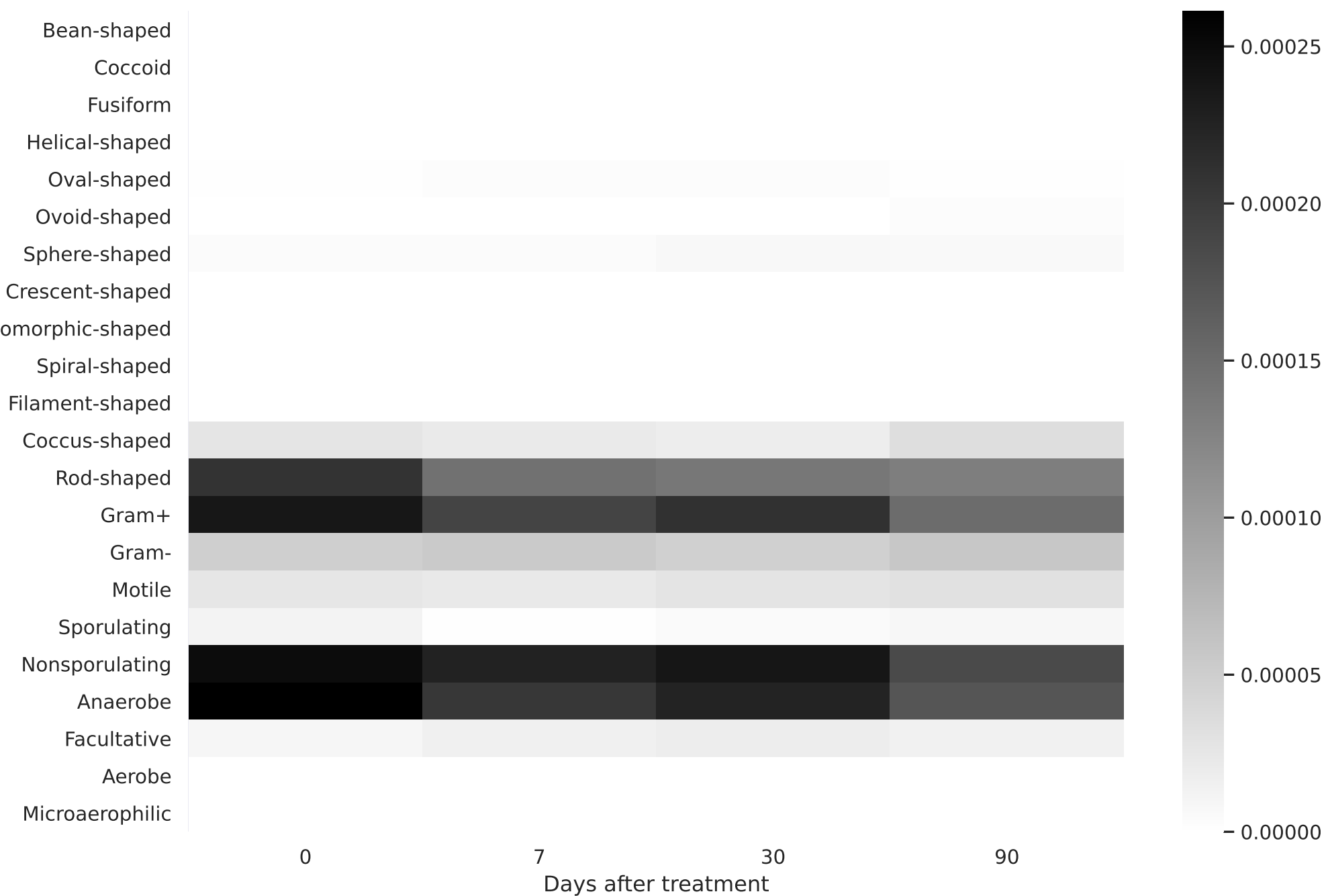


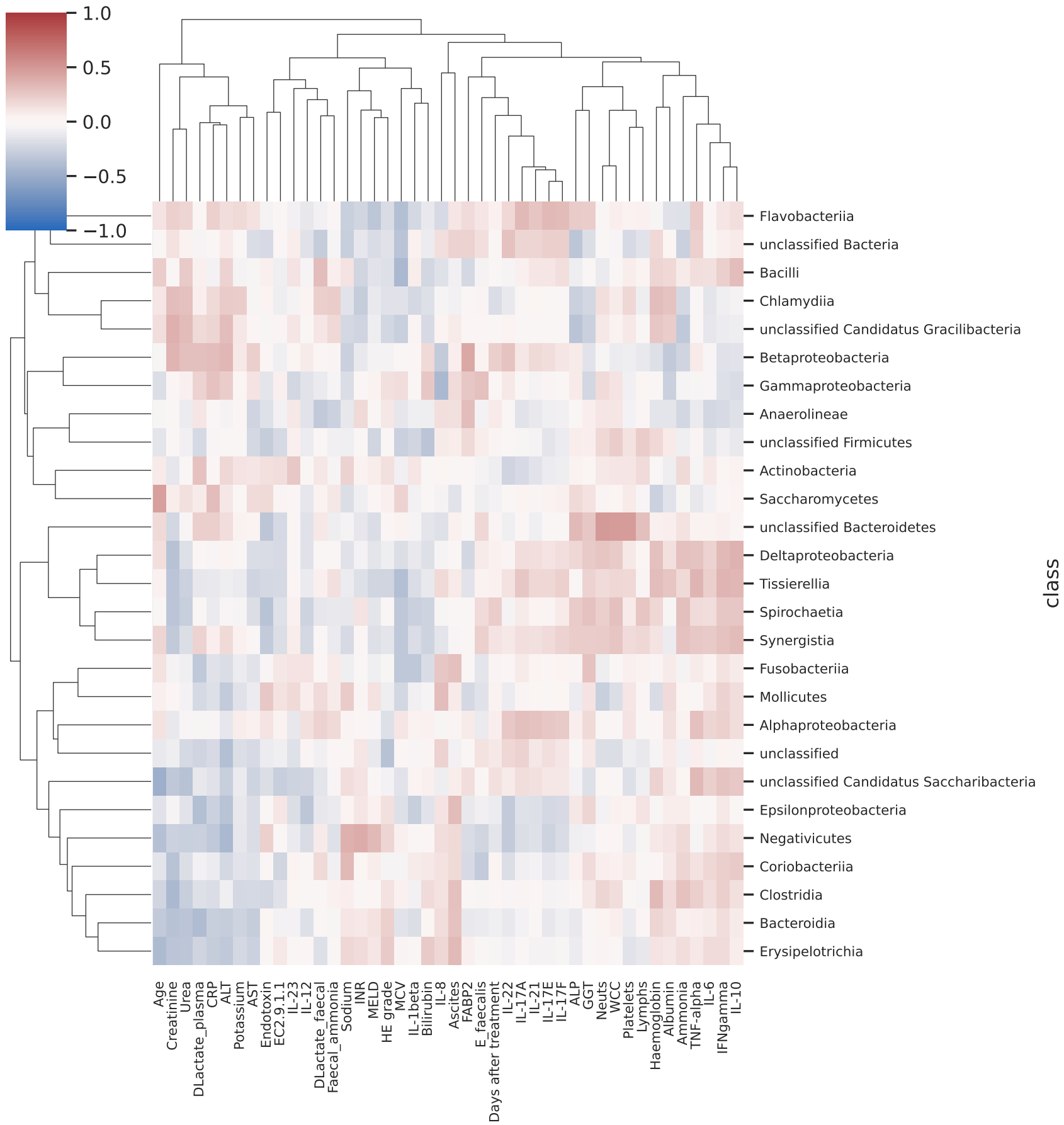
Oral

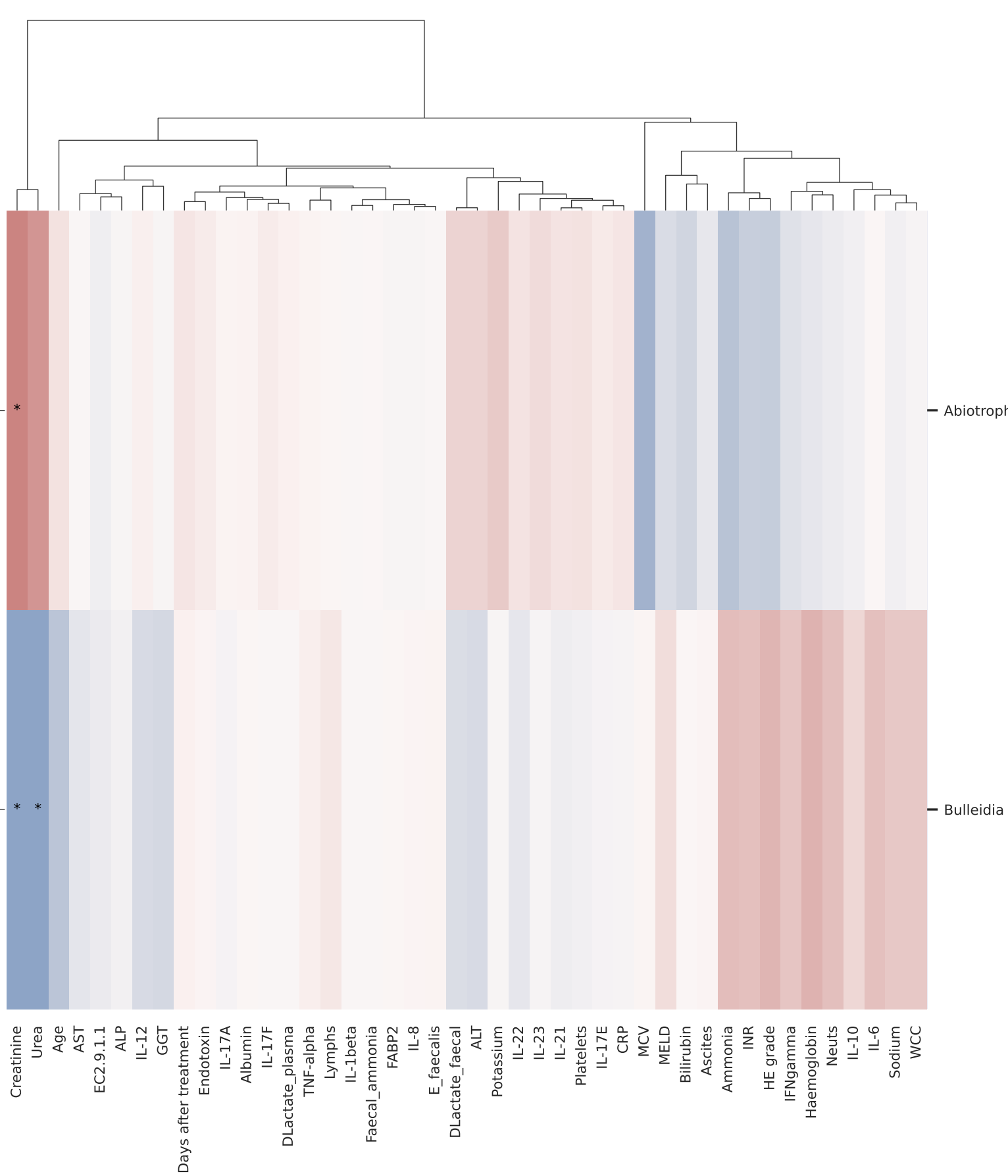
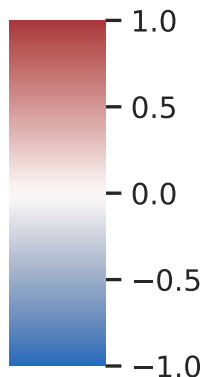


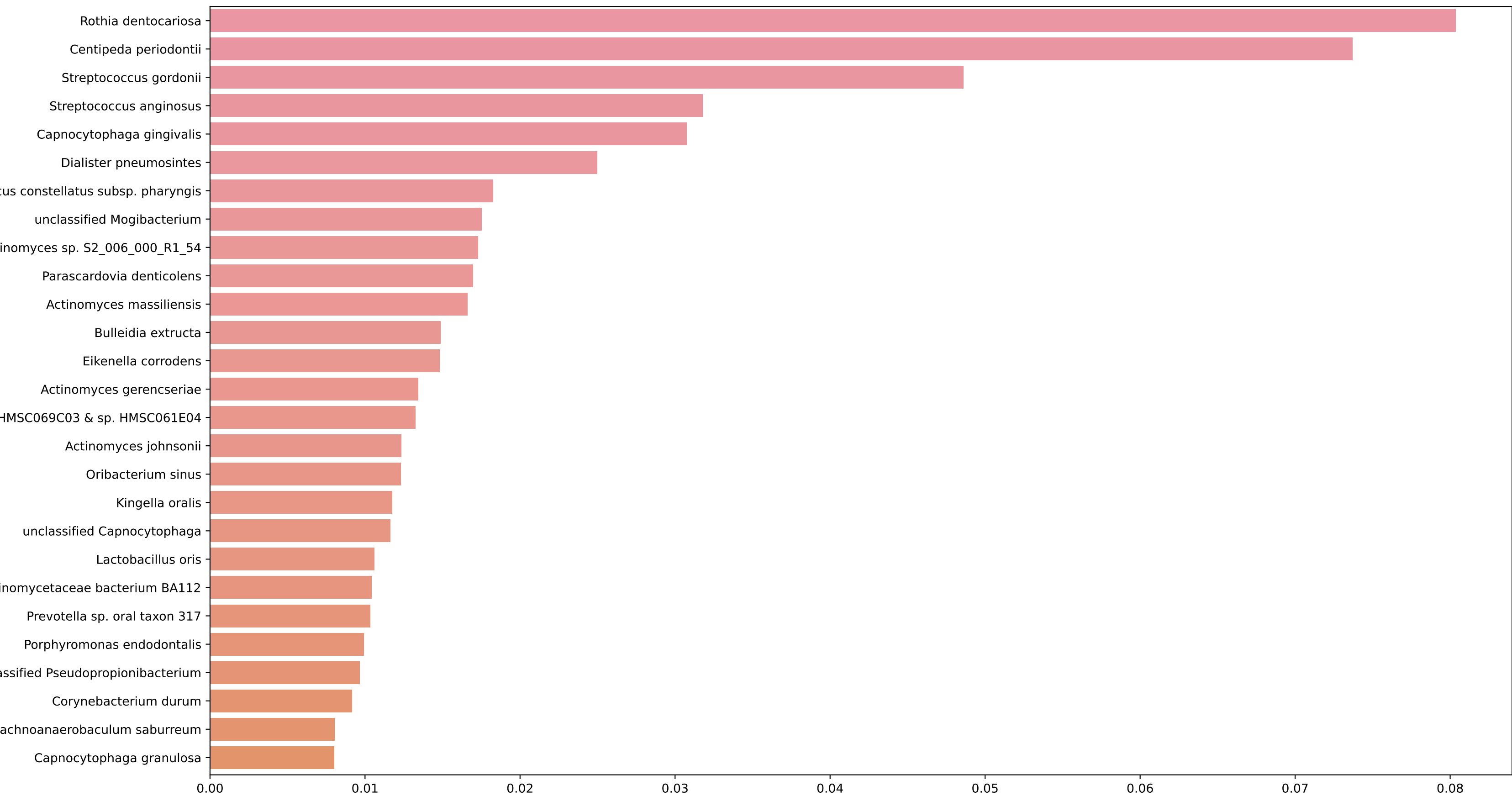
Gut

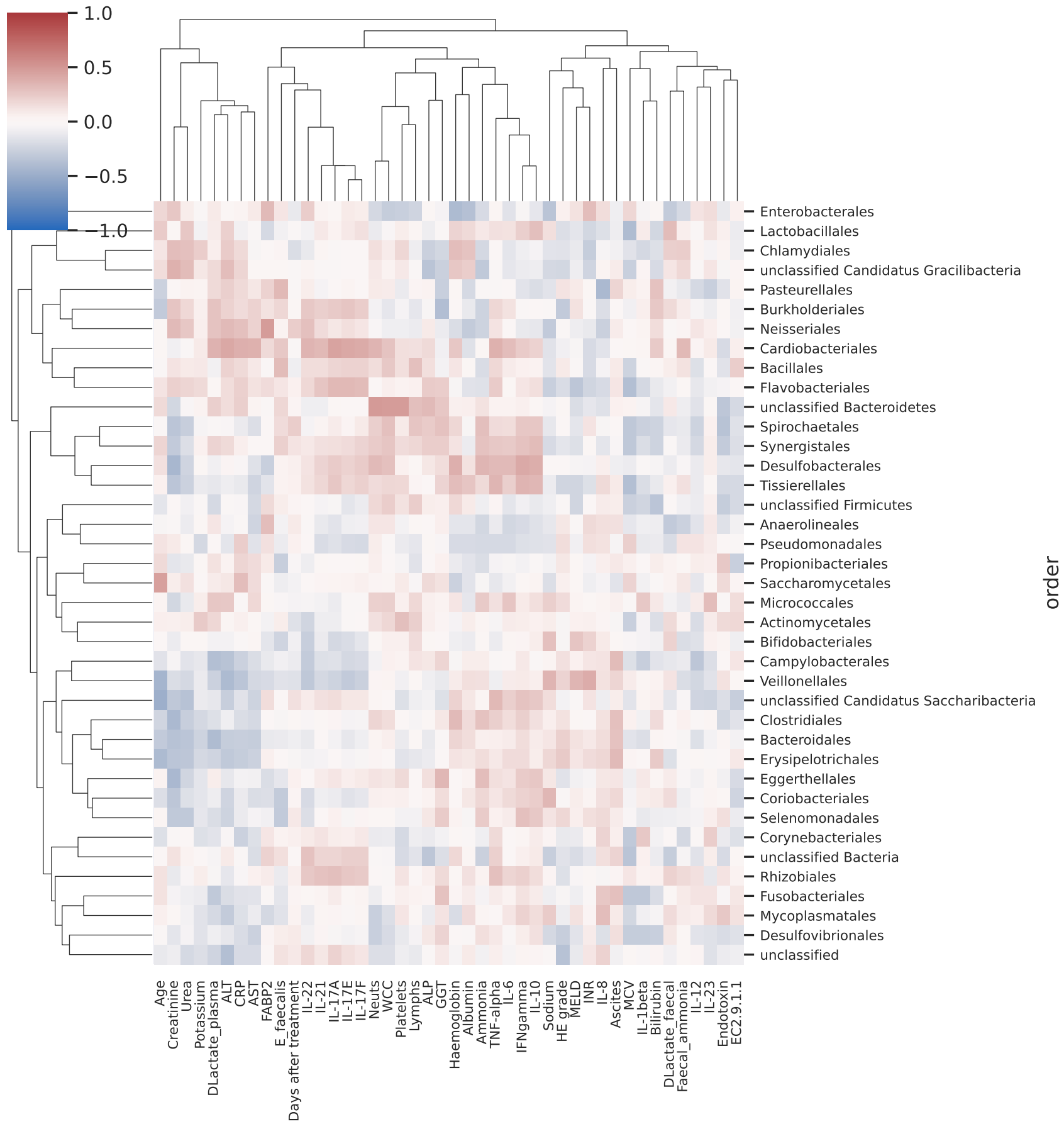


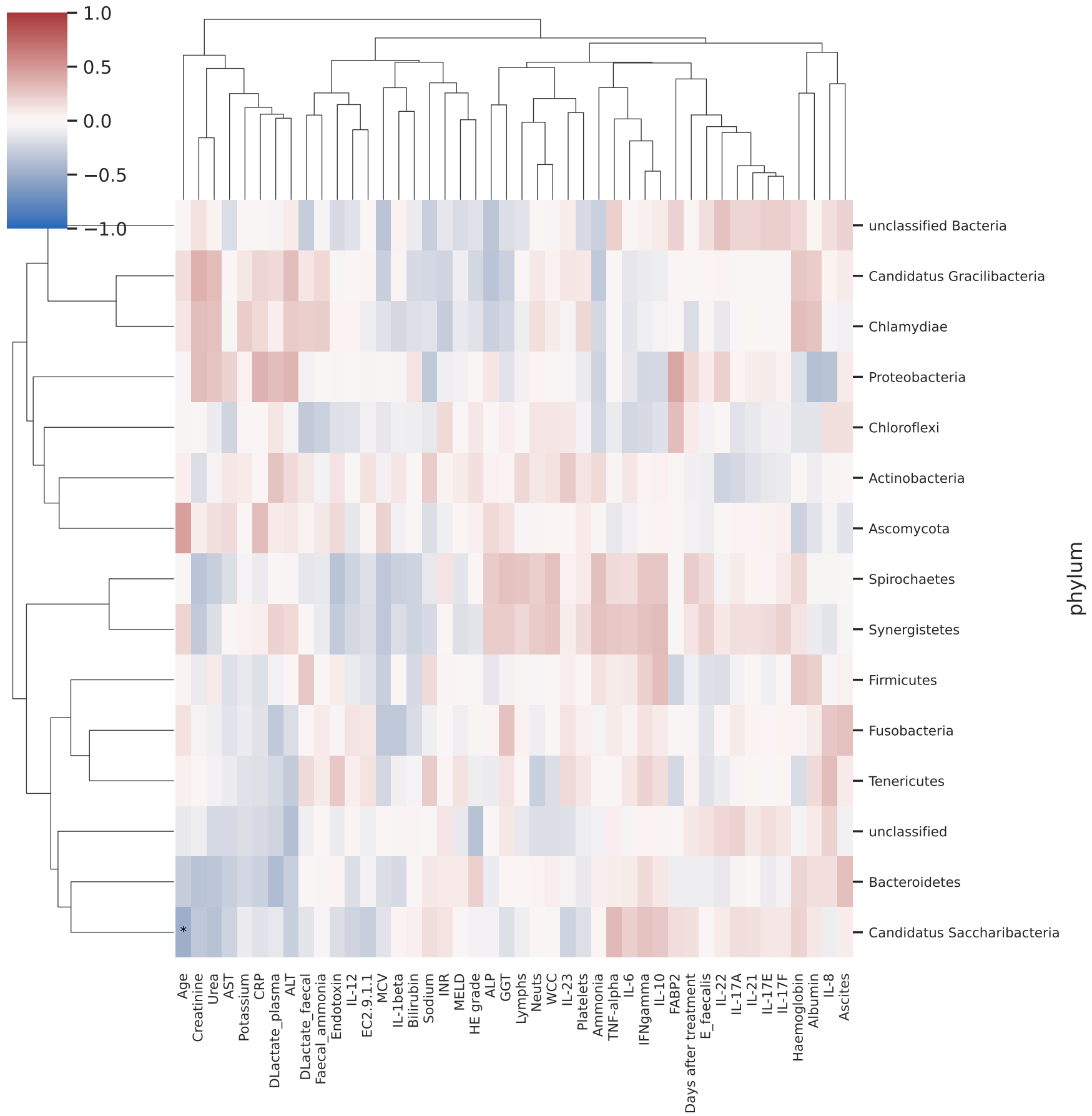


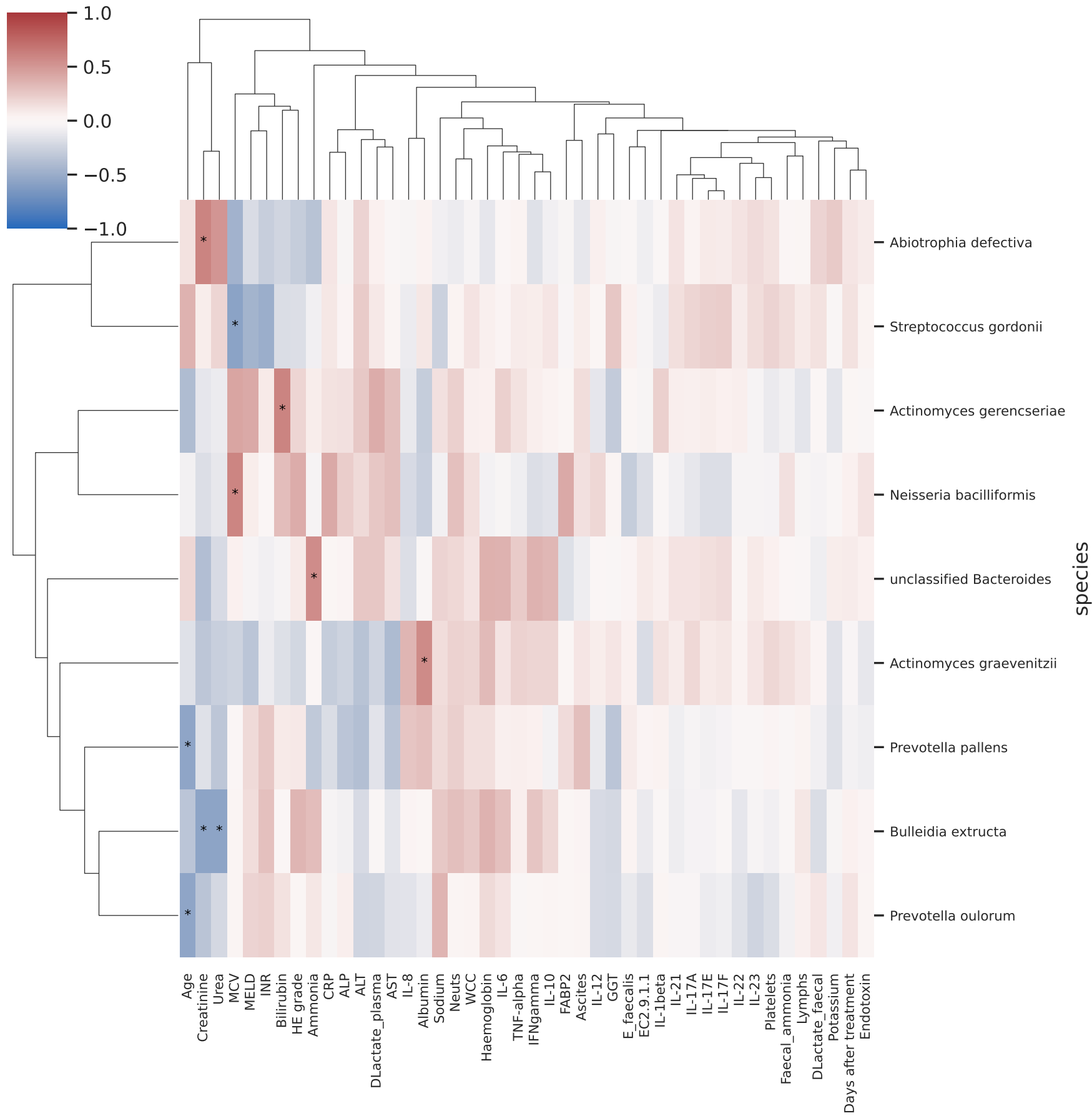


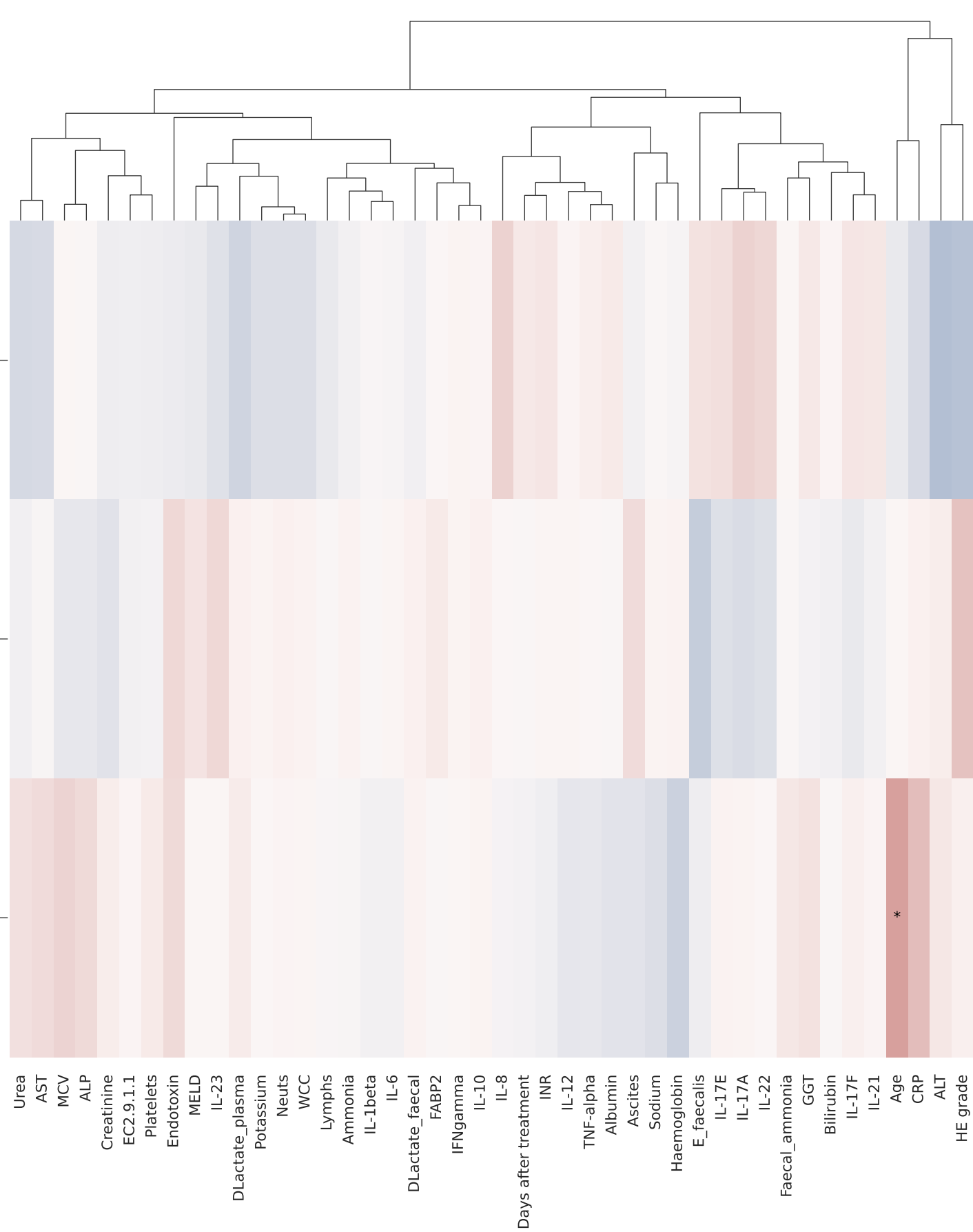
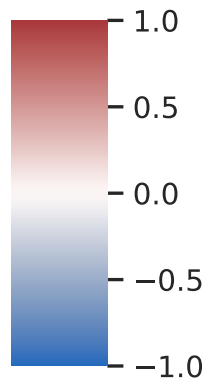










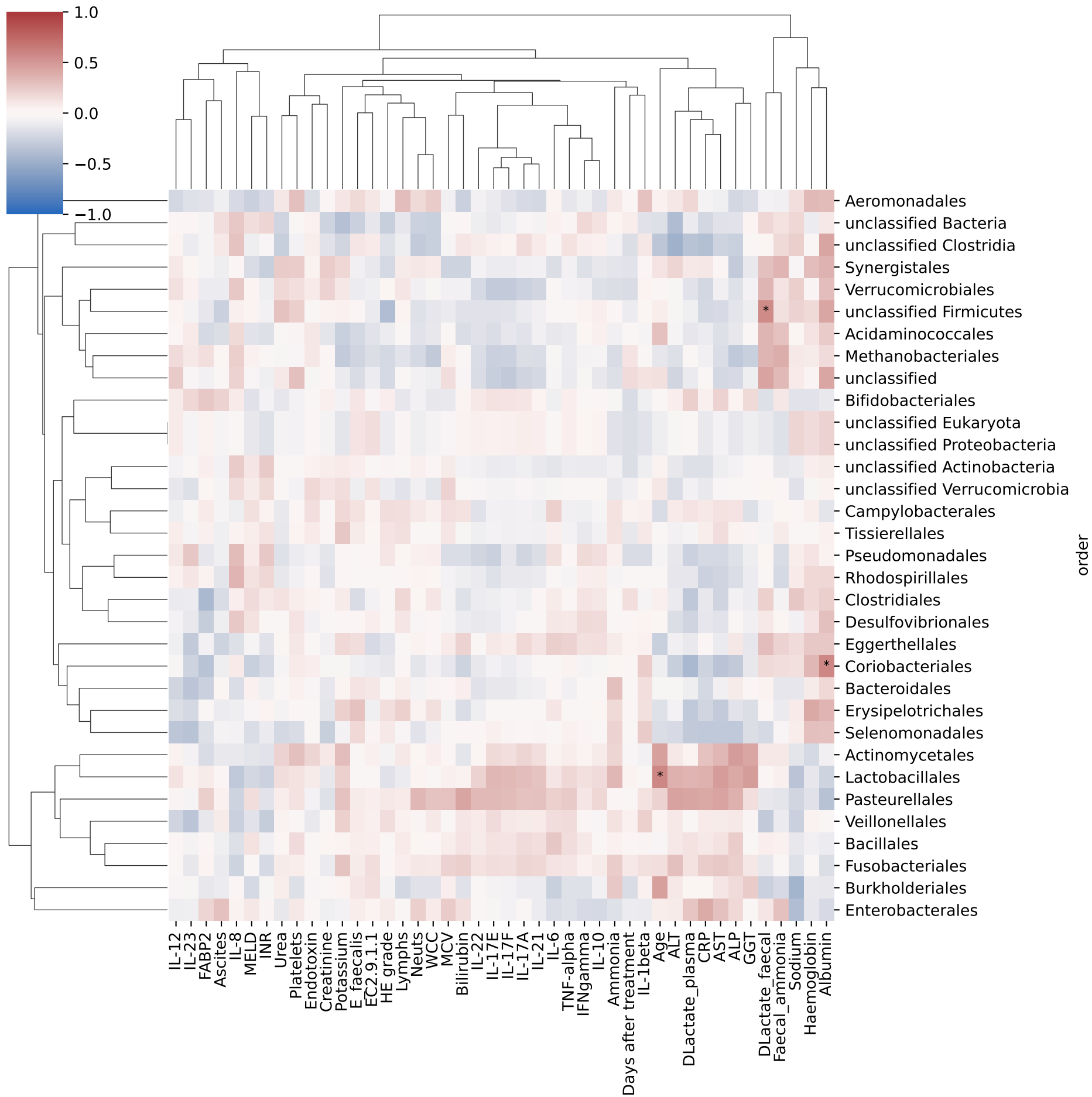


unclassified

Bacteria

Eukaryota

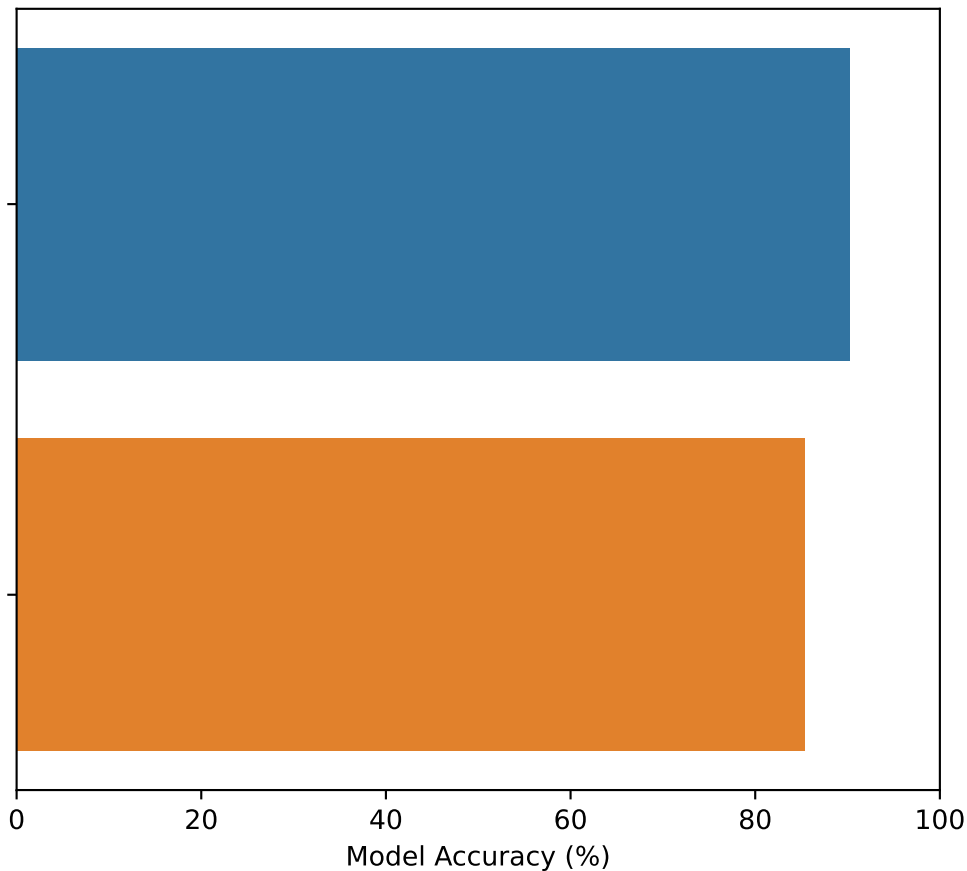
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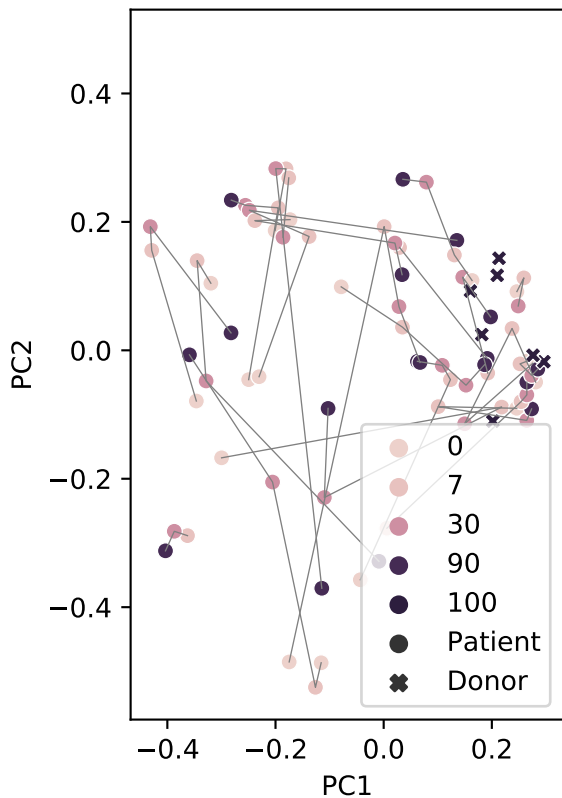
Dataset used for MELD prediction

Oral Species

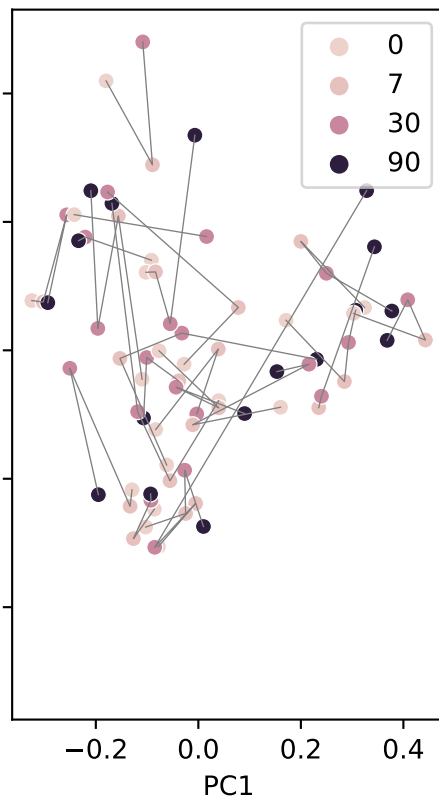
Gut Species

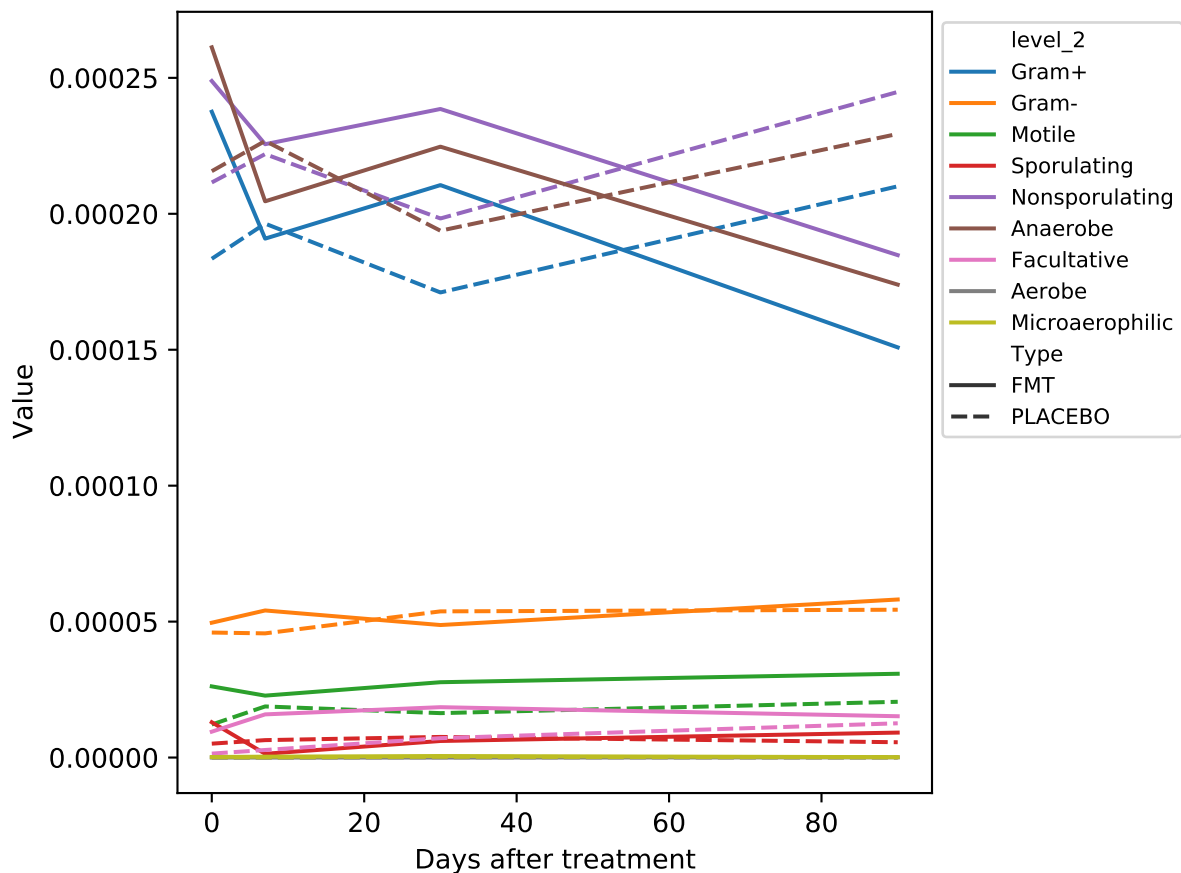


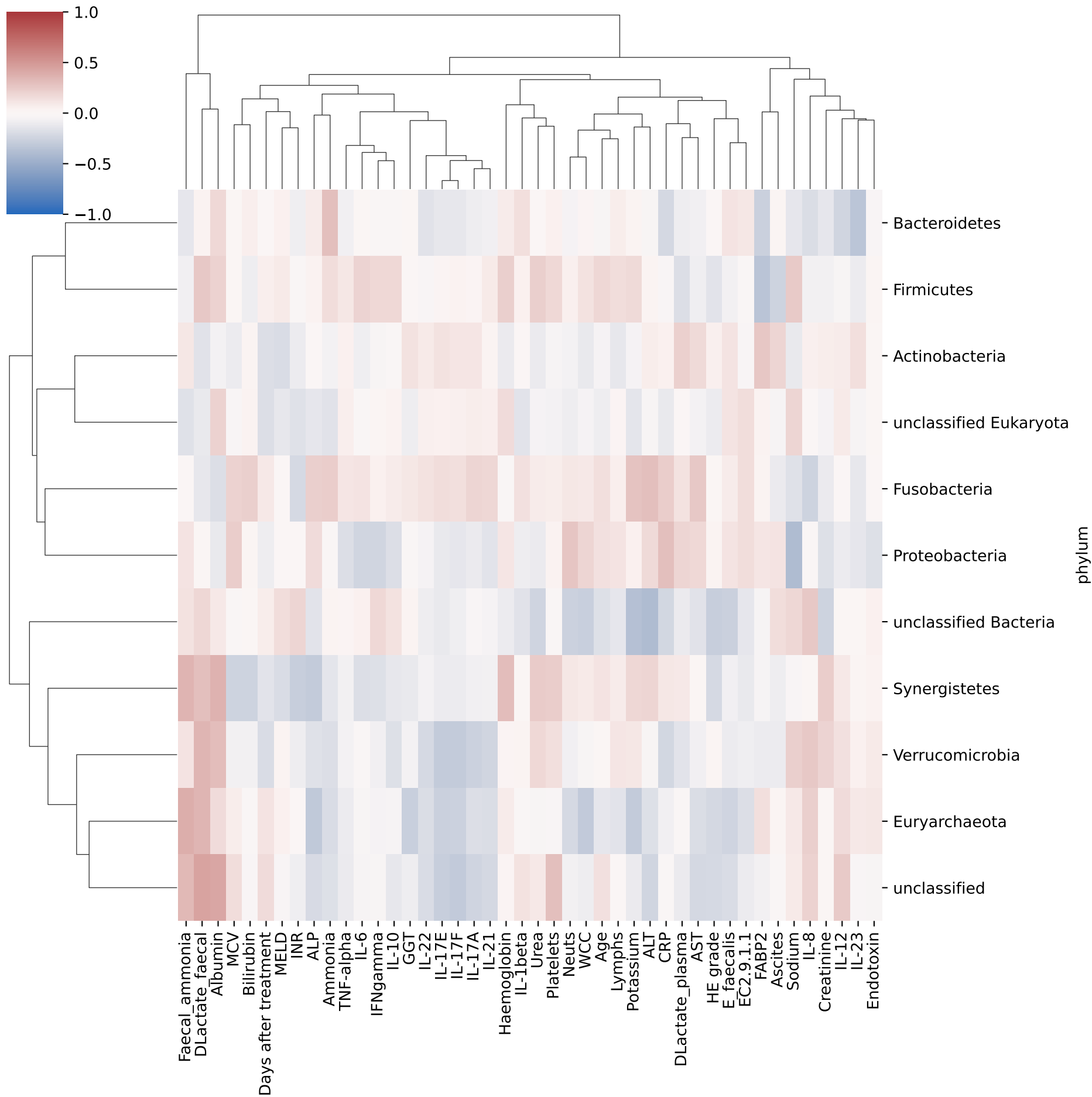
Gut

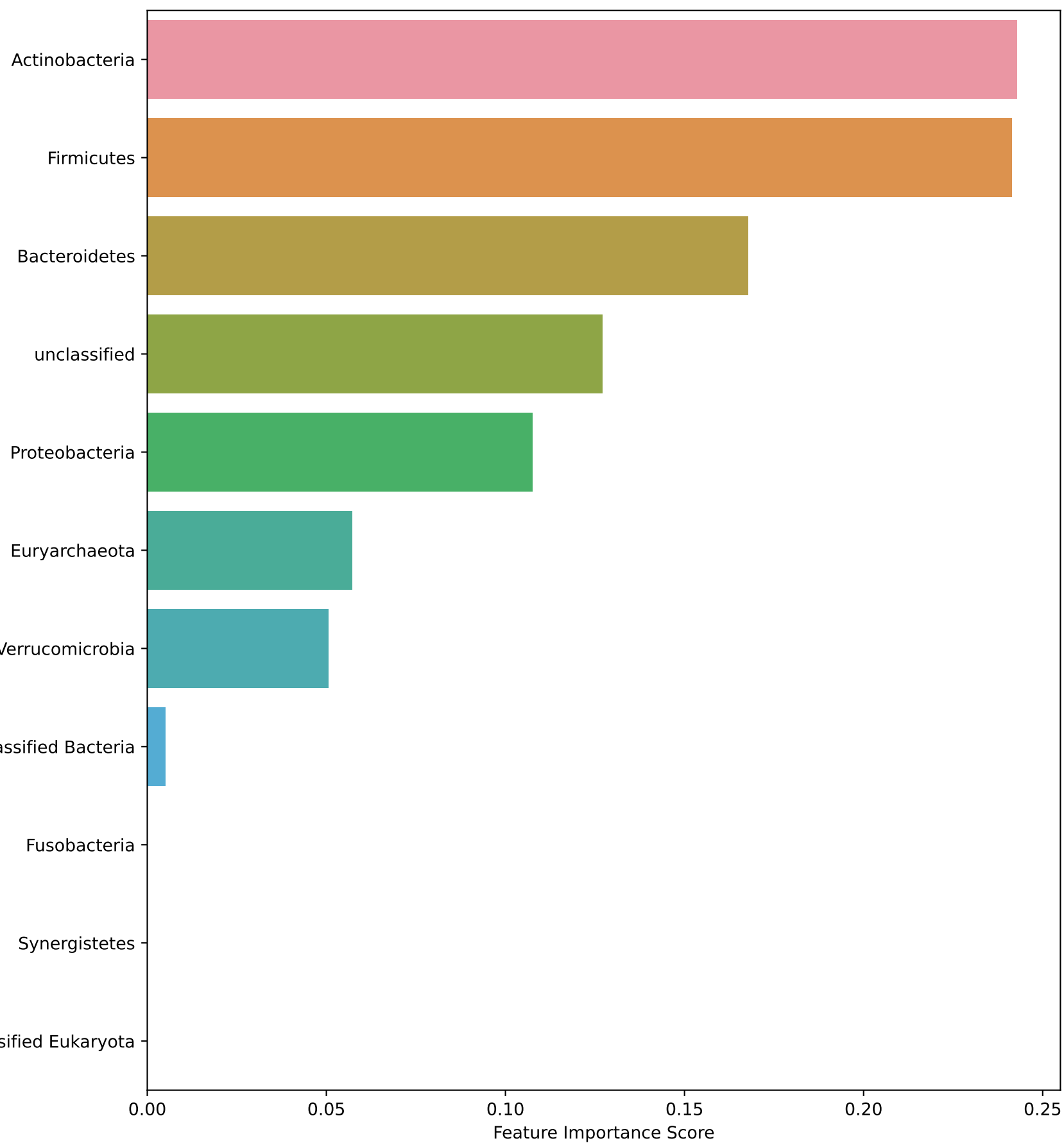


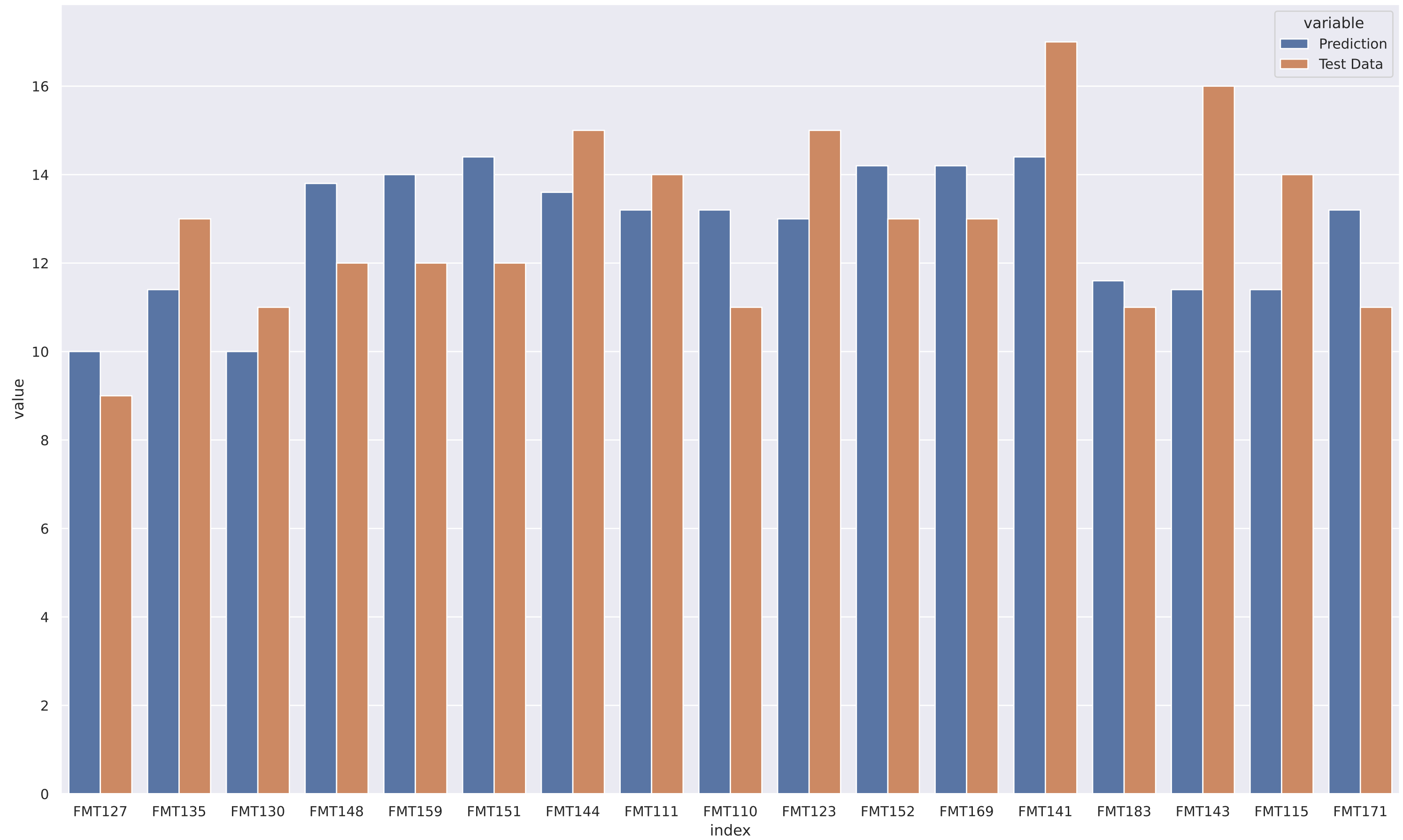
Oral

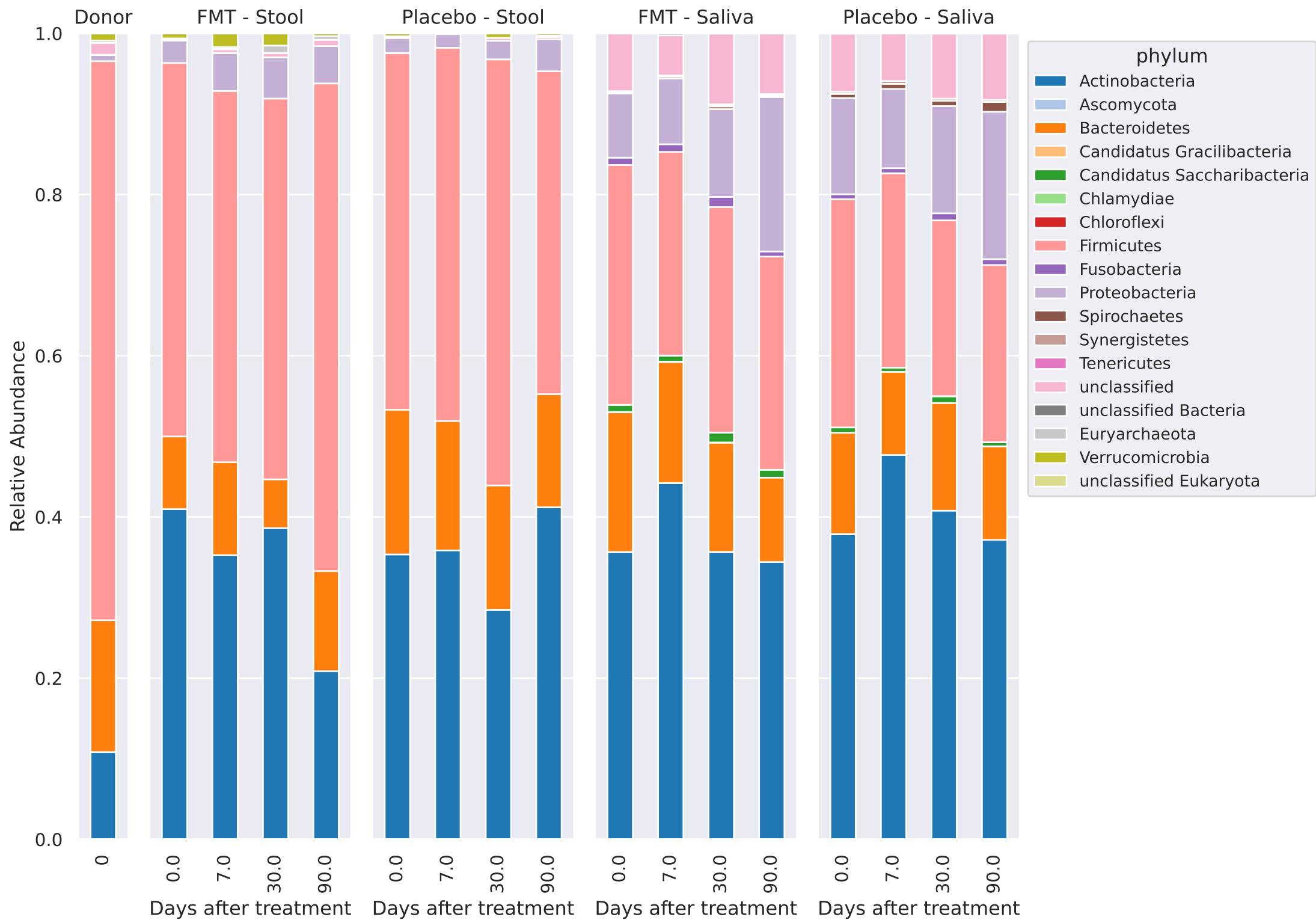




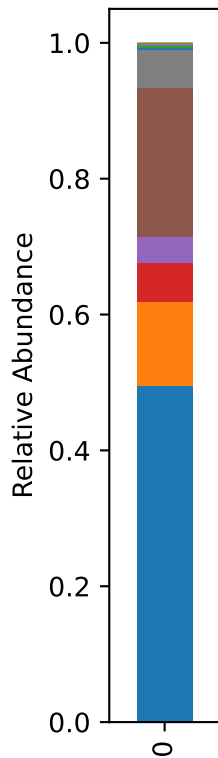




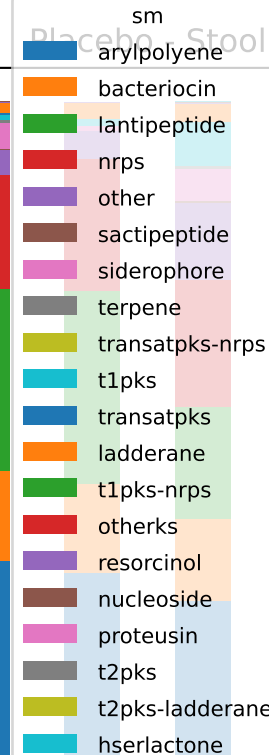
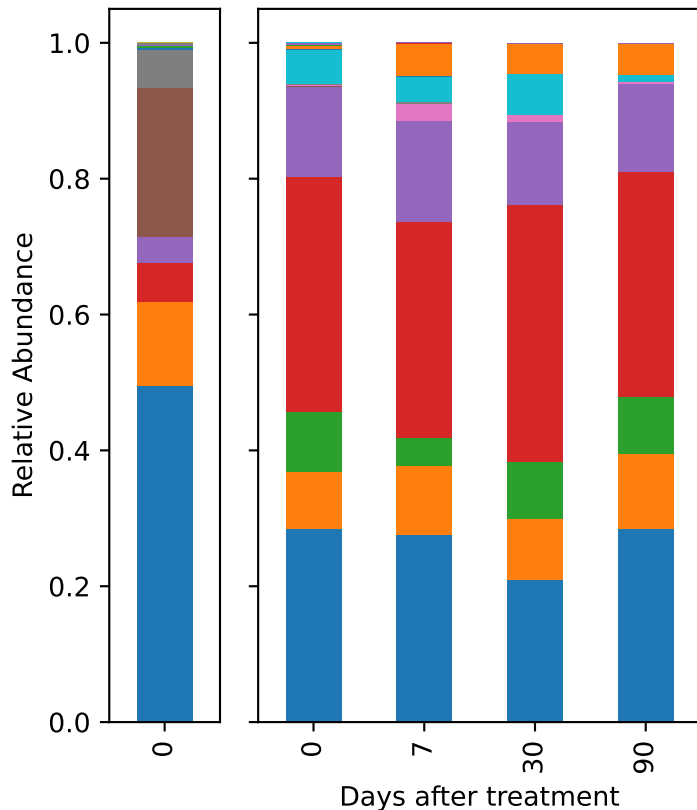




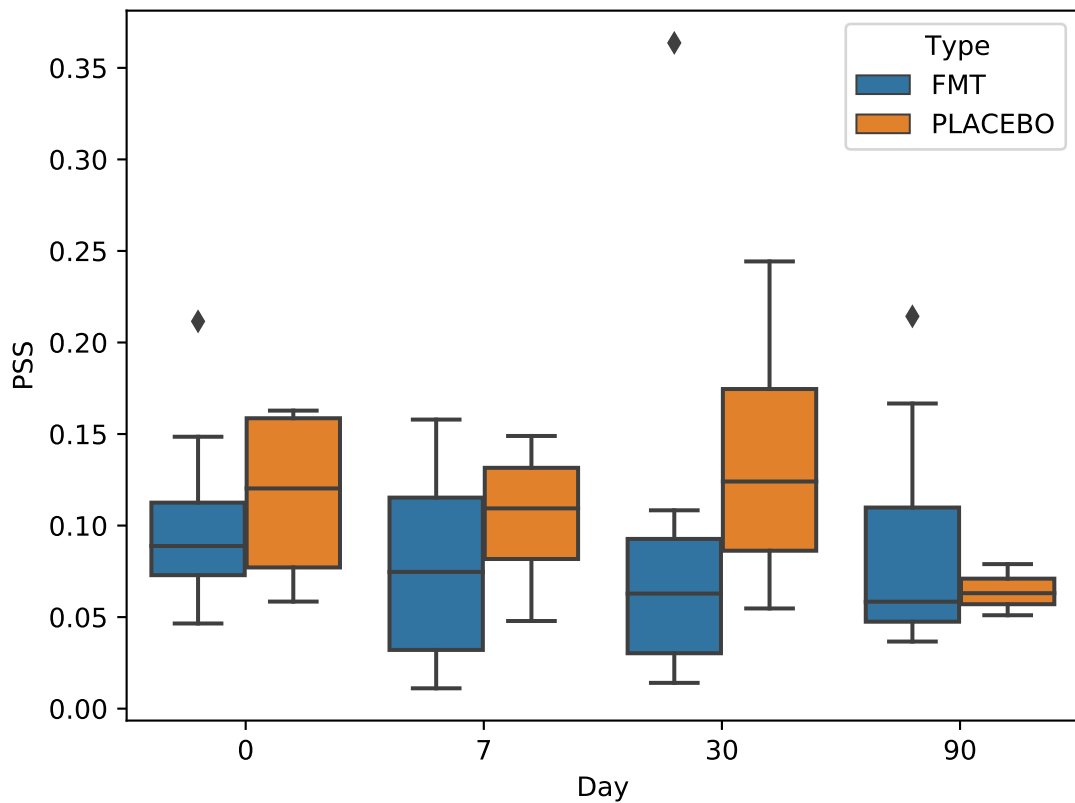
Donor



FMT - Stool



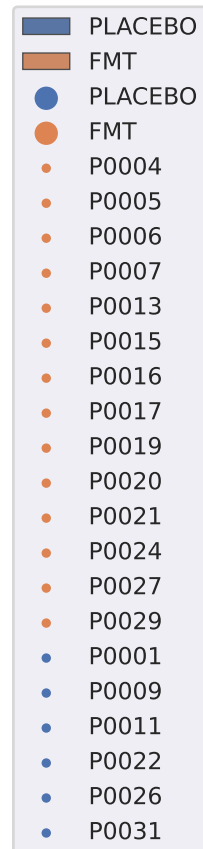
Days after treatment



Donor stool

Patient stool

Patient Saliva



Shannon Index

1

2

3

4

5

6

DONOR

0.0

7.0

30.0

90.0

Days after treatment

0.0

7.0

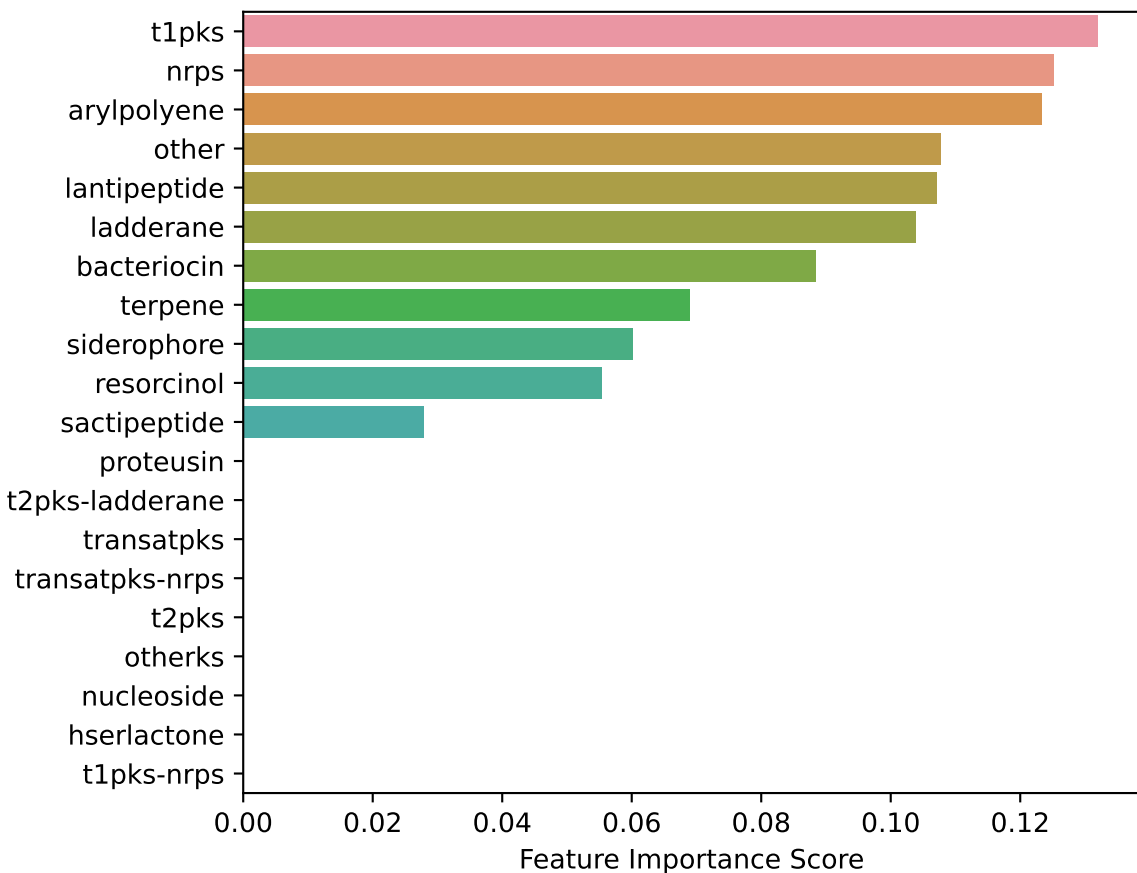
30.0

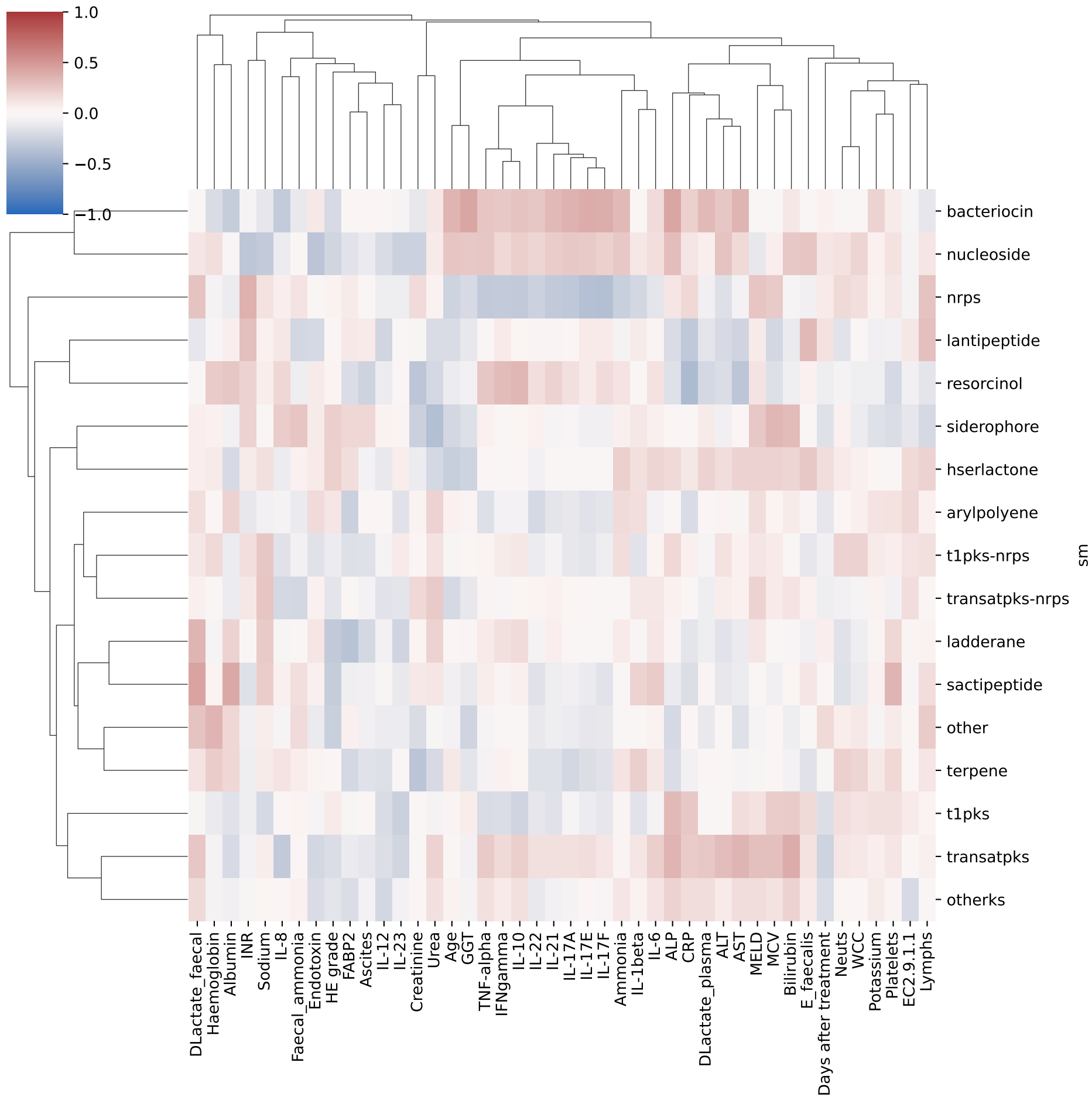
90.0

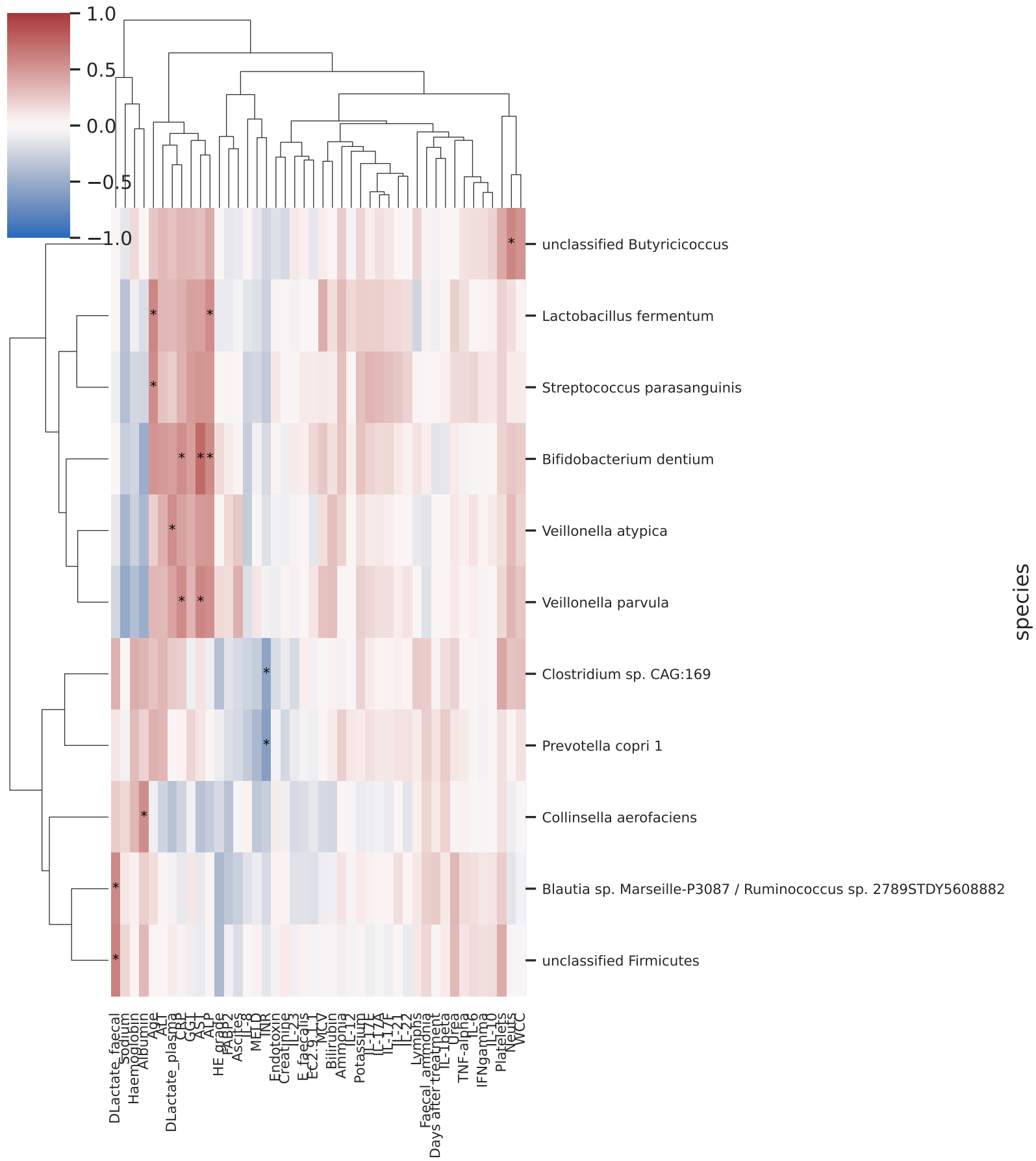
Days after treatment

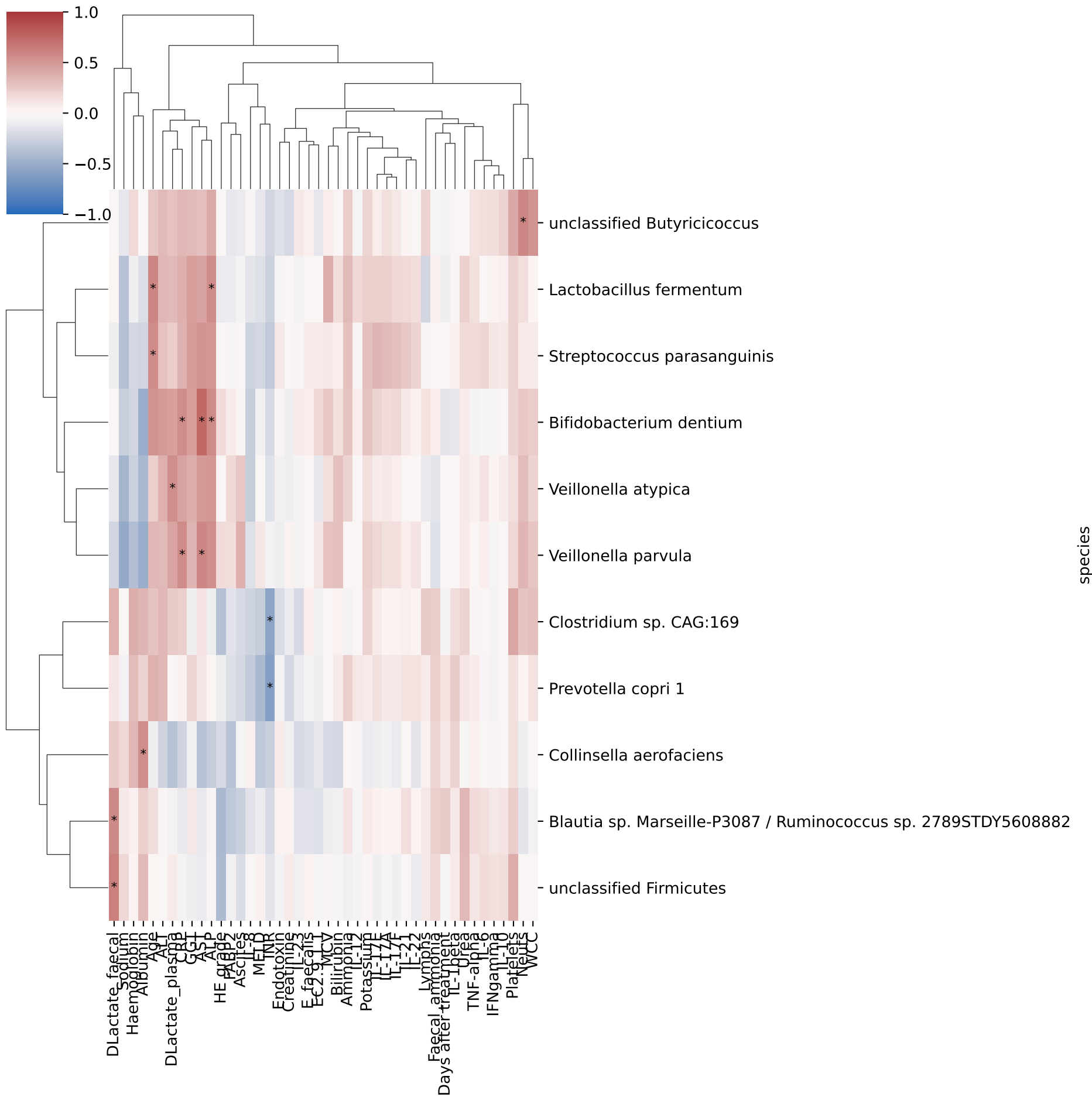
M.W.W. $p = 6.030e-02$

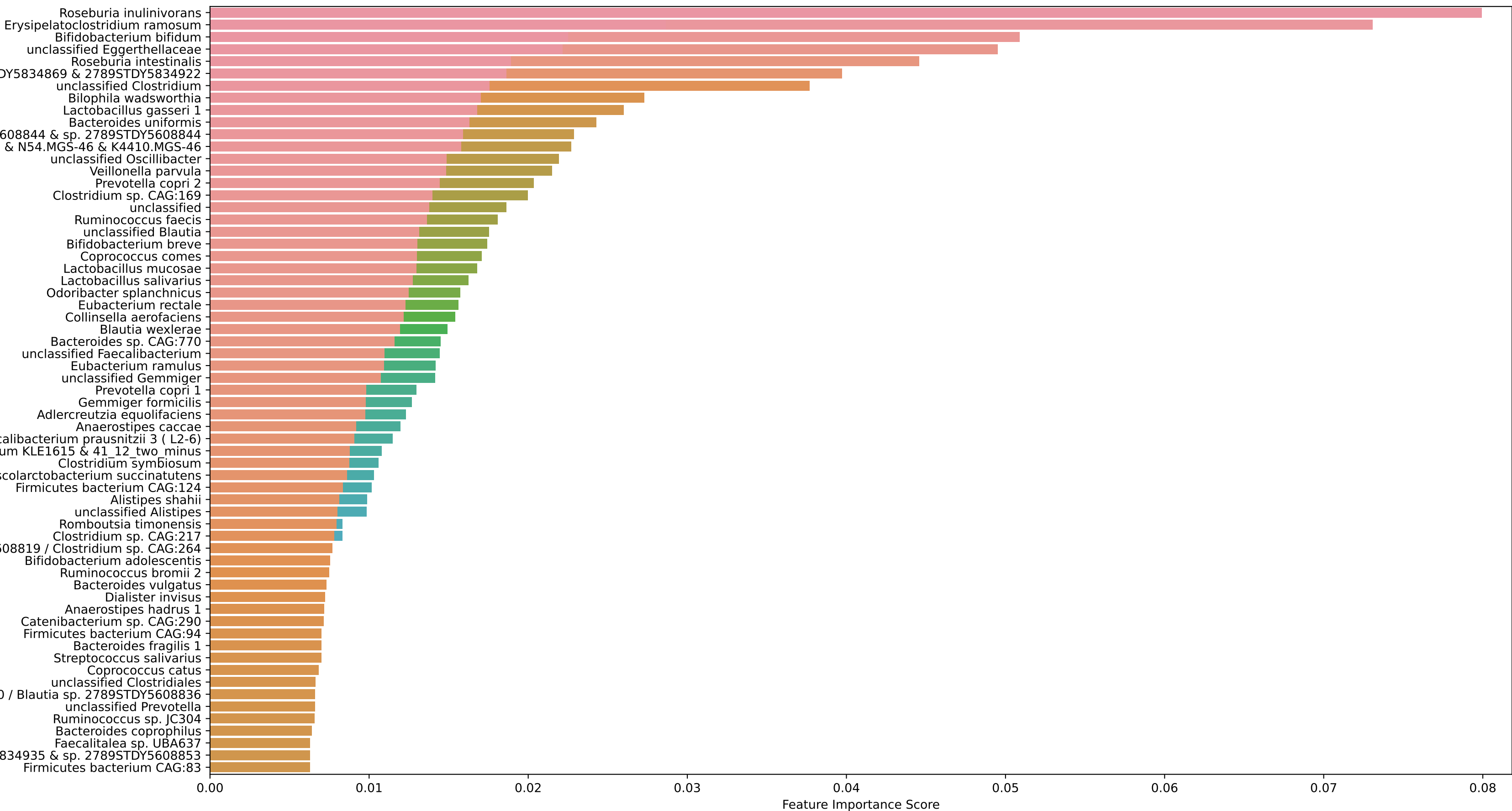
Secondary Metabolites











Donor stool

Patient stool

Patient Saliva

