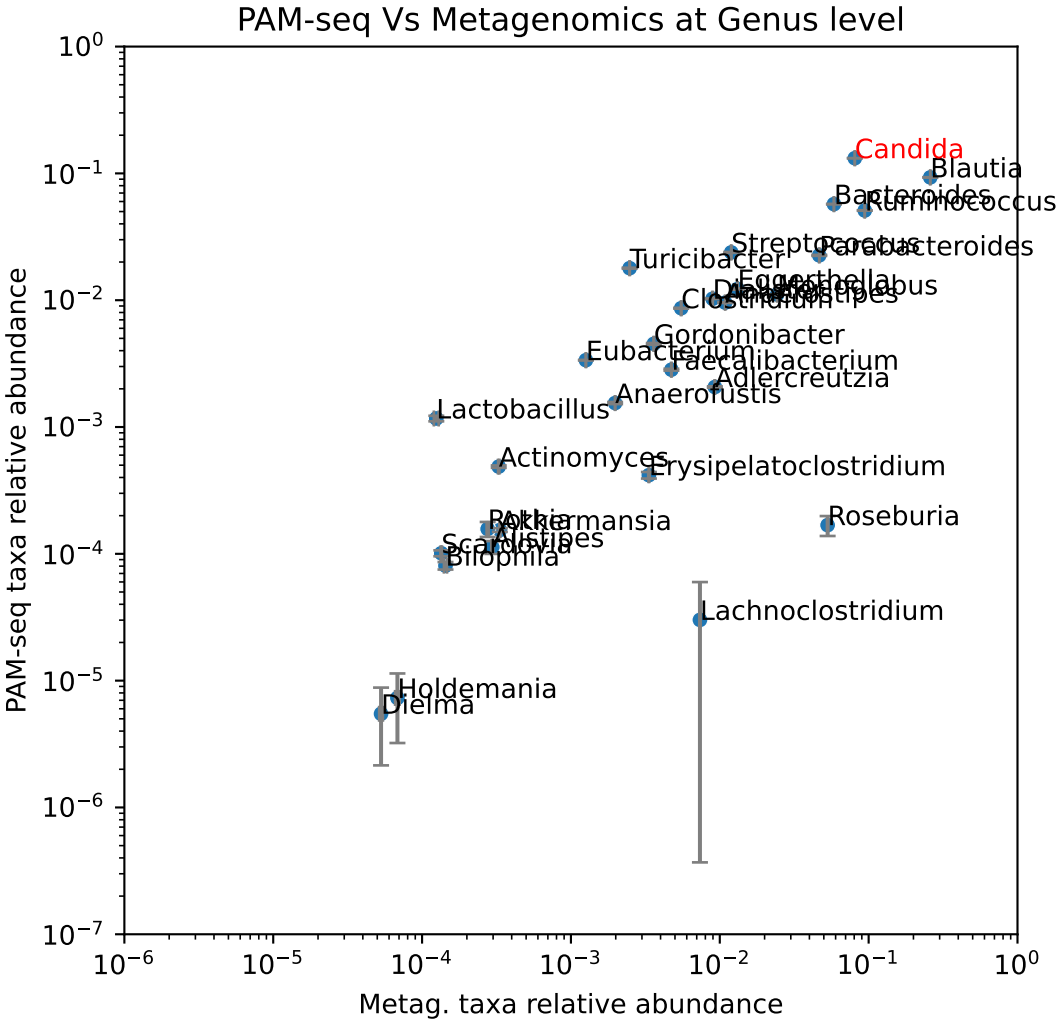
PAM-seq Vs Metagenomics at Genus level
PAM-seq taxa relative abundance
Metag. taxa relative abundance
Candida
Blautia
Bacteroides
Ruminococcus
Streptococcus
Parabacteroides
Turicibacter
Eggerthella
Clostridium
Gordonibacter
Eubacterium
Faecalibacterium
Adlercreutzia
Anaerofustis
Lactobacillus
Actinomyces
Erysipelatoclostridium
Roseburia
Akkermansia
Alistipes
Scardovia
Bifophila
Lachnoclostridium
Holdemania
Dielma