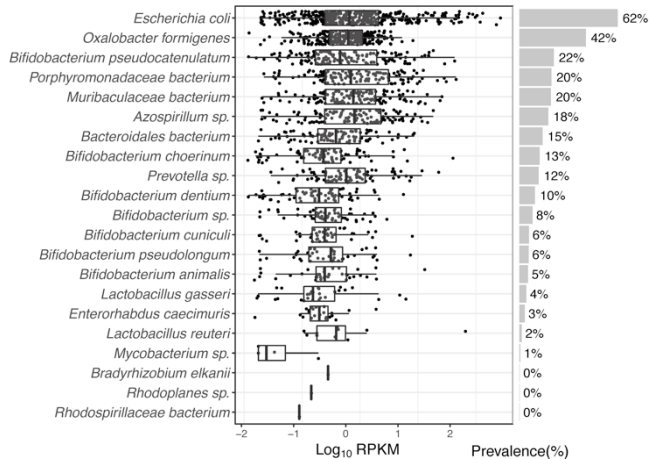


Metagenome($n=660$ subjects)Metatranscriptome($n=165$ subjects)