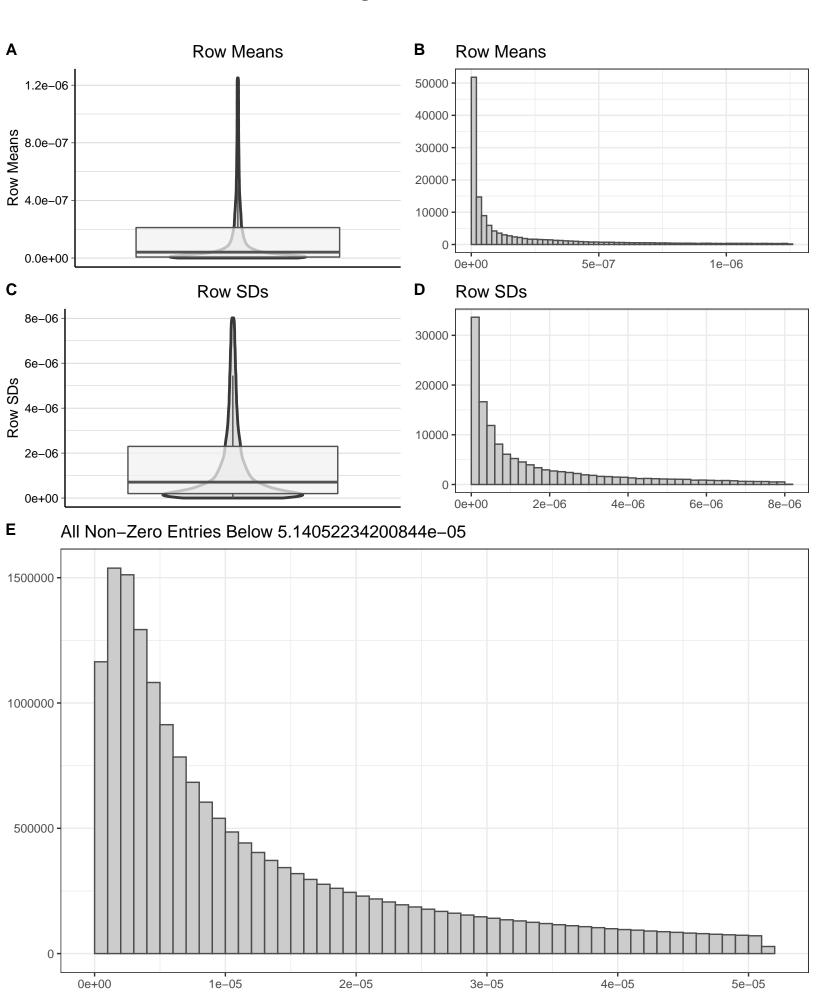
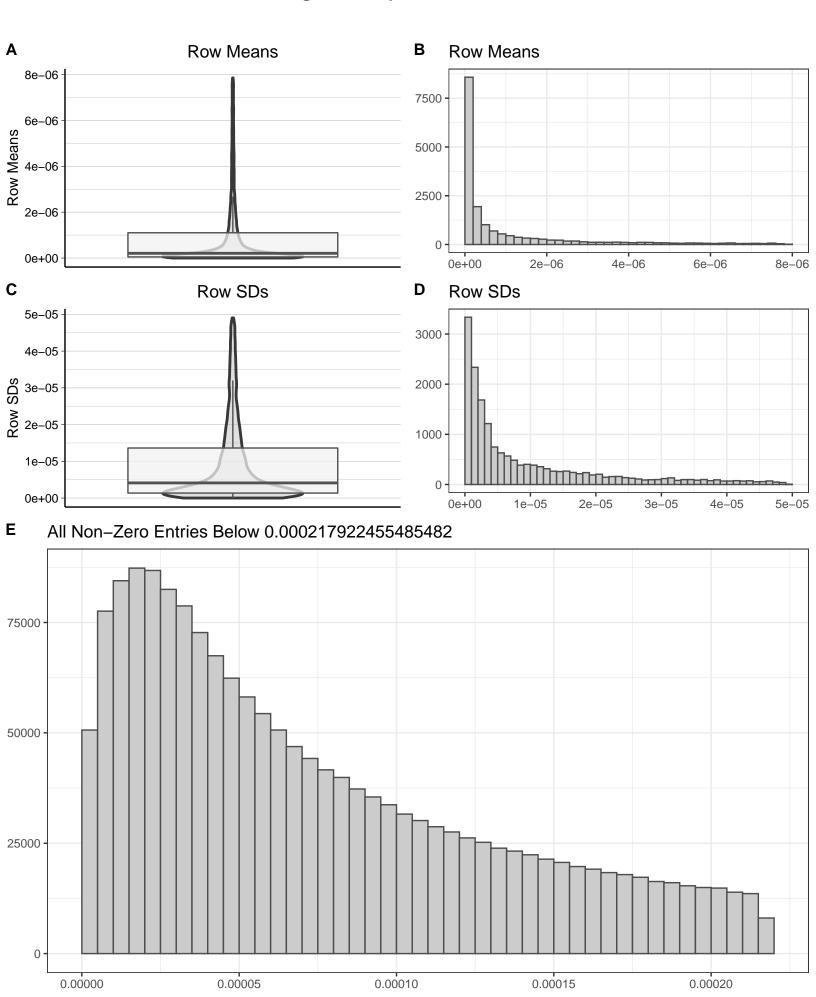


## Metagenomes ecs\_3



## Metagenomes pathabundances\_3



## Metagenomes taxonomic\_profiles\_3

