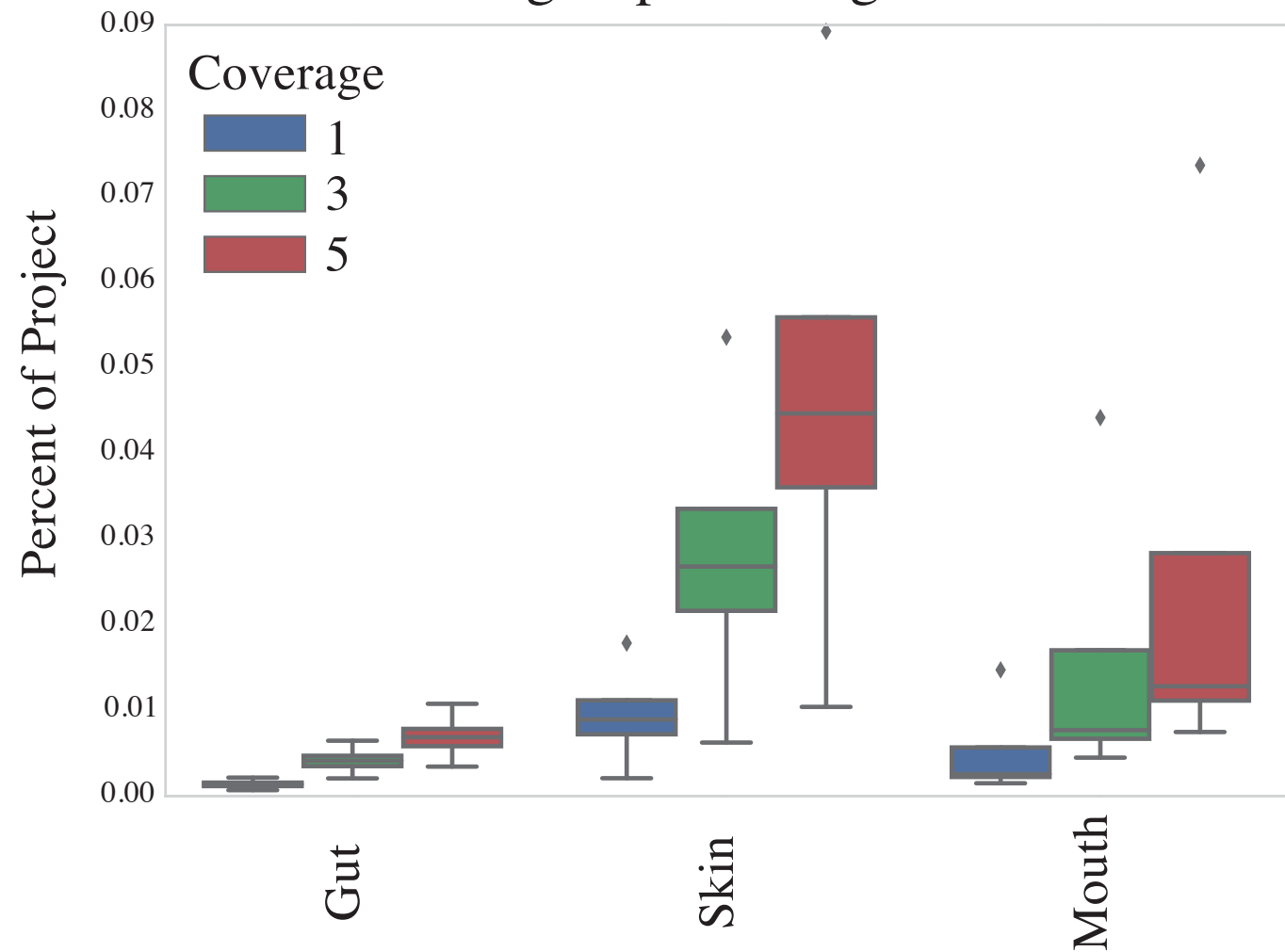


A

Phage / plasmid genome

**B**

Bacterial Genome

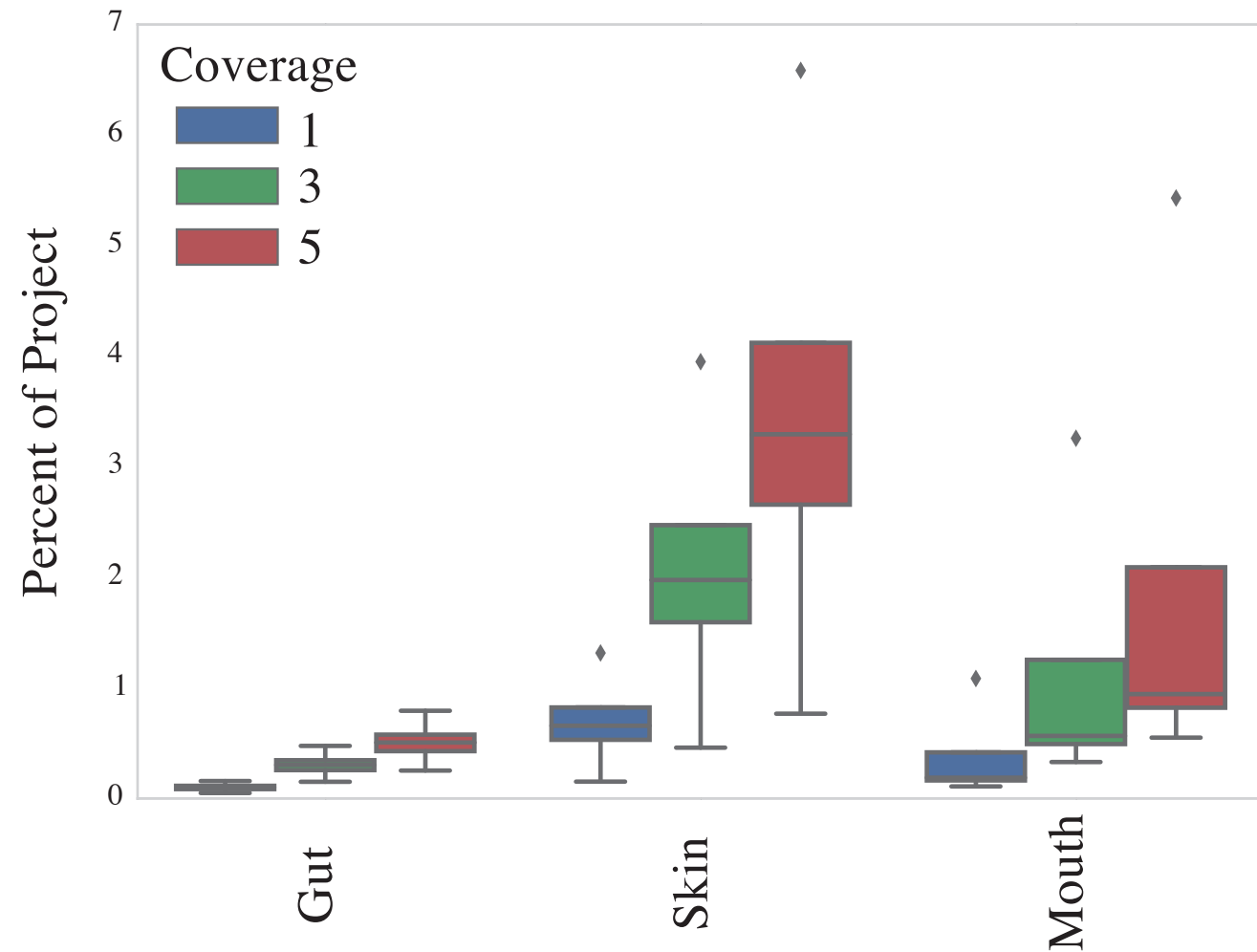


Figure S4 Body sites have different depths of sequencing, due largely to extensive human DNA contamination. For the average size phage/plasmid (a) (44 kb) and bacterial (b) (3.3 Mbp) genome, the percentage of project reads that correspond with different coverage levels is shown for all three body sites.