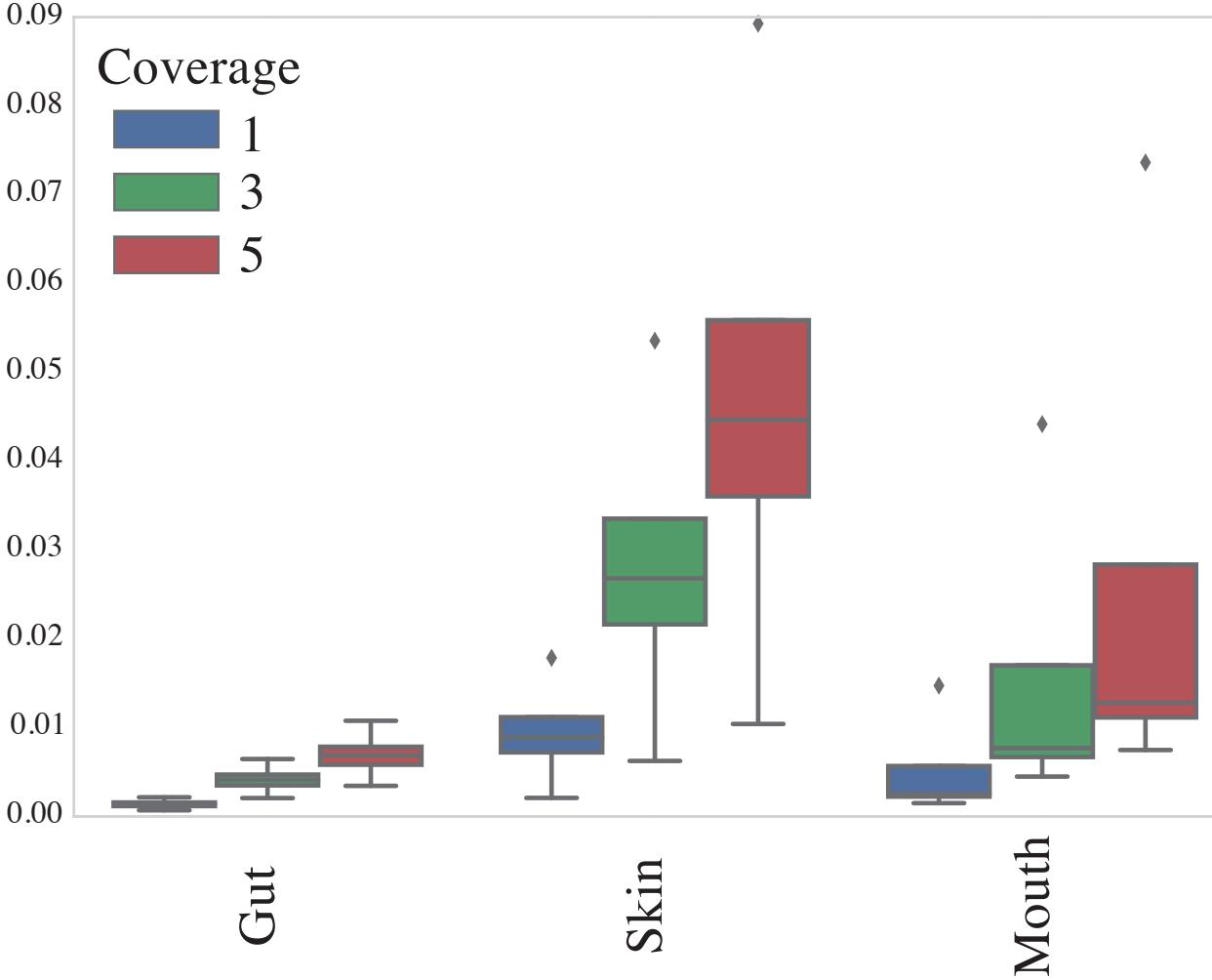


**A**

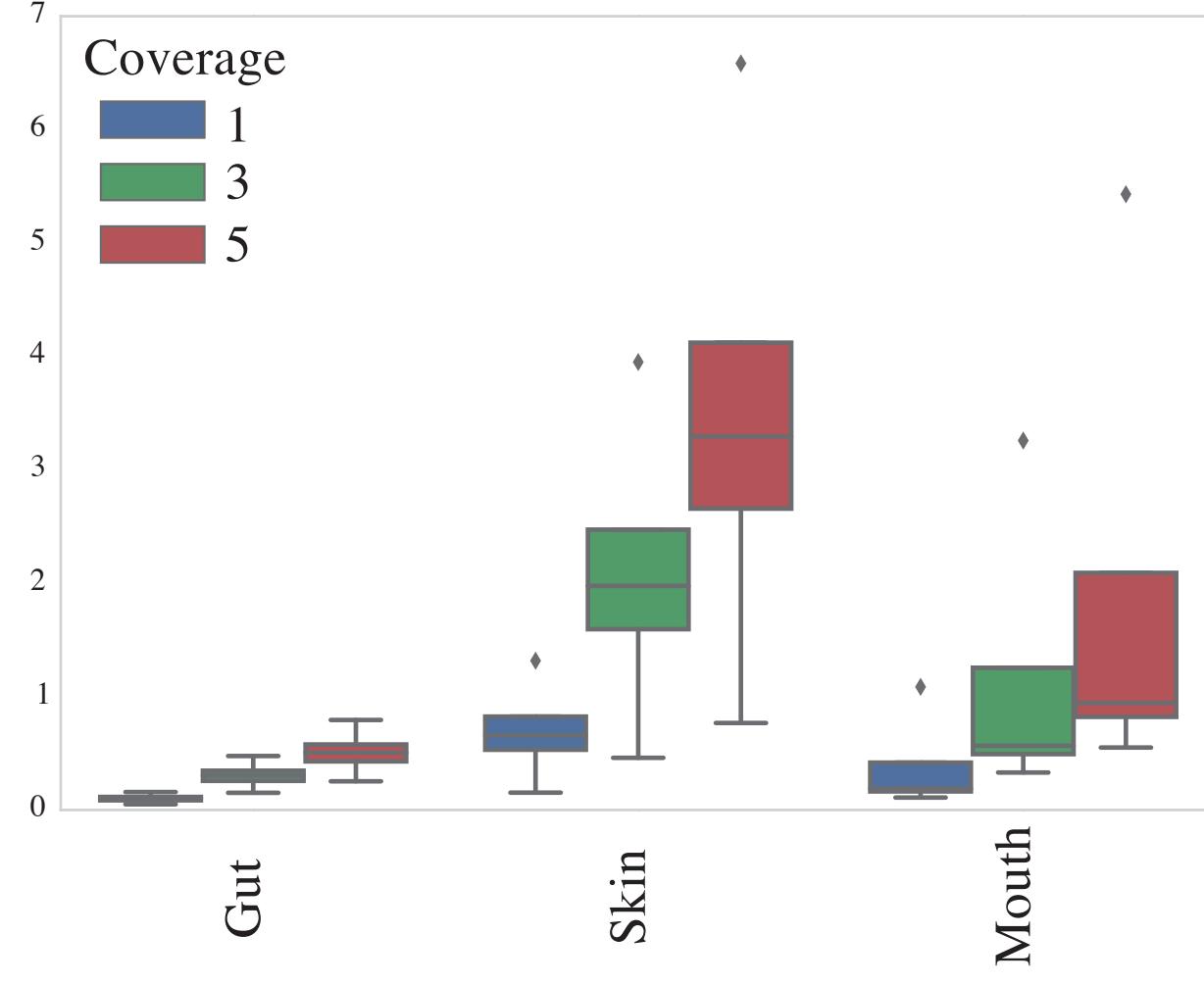
### Phage / plasmid genome

Percent of Project

**B**

### Bacterial Genome

Percent of Project



**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.