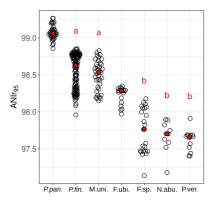
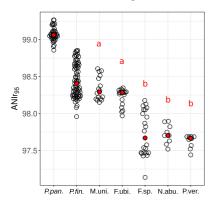
Including home habitats of reference genomes



Excluding home habitats of reference genomes



Suppl. Fig. S3: ANIr $_{95}$ values for the seven species when metagenomes sampled from the habitat where the reference genome was obtained from are included (left) and excluded (right). Each dot refers to one metagenome. Red dots show the medians. Within each plot, pairs that are not significantly different (p >0.05) according to Wilcoxon-Mann-Whitmey Rank Sum test are marked with a common letter.