

Prediction of *Klebsiella* phage-host specificity at the strain level

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Supplementary Figure S1

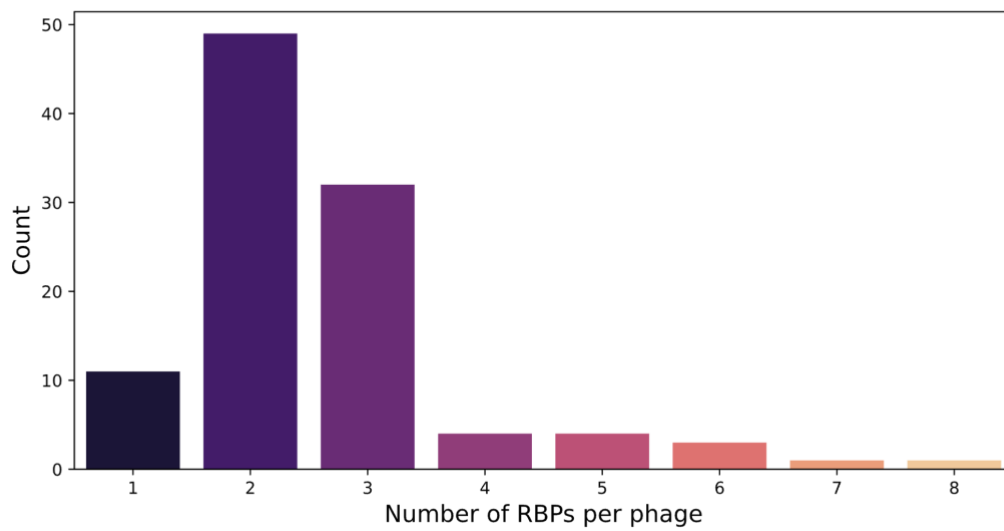


Figure S1: Count plot of the number of detected RBPs per phage across all the collected phage genomes (color scaled from low to high numbers).

Supplementary Figure S2

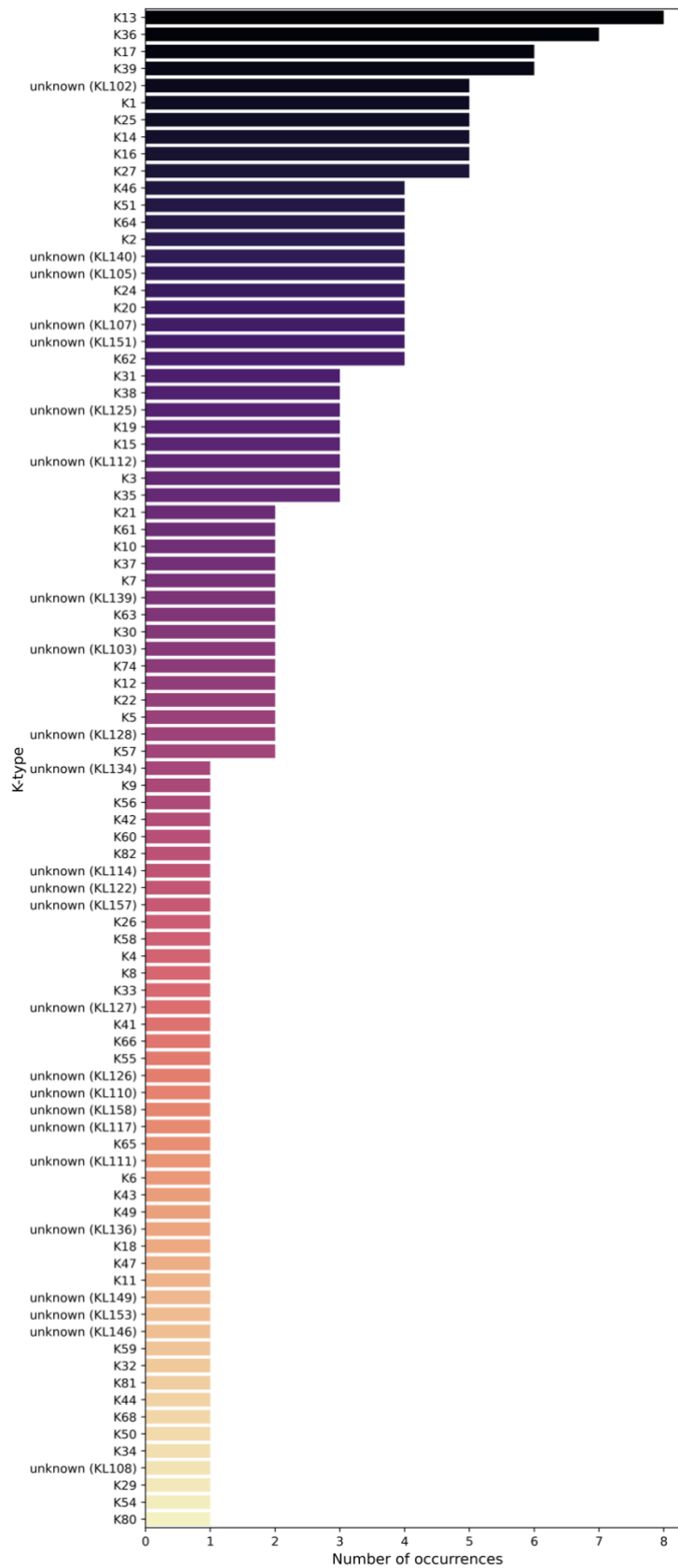


Figure S2: Count plot of the number of occurrences of each K-type across the 200 collected bacterial genomes (color scaled from high to low numbers).