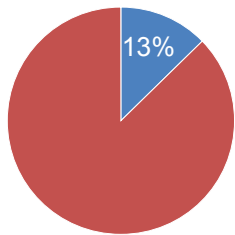
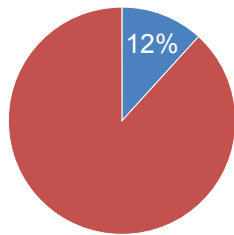


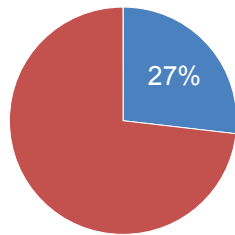
V. parahaemolyticus
genome (4458)



GC \geq 40 %
(4186)



GC < 40 %
(272)



Classification in WT *V. parahaemolyticus*

■ Underrepresented

■ Neutral

Supplementary Figure S1