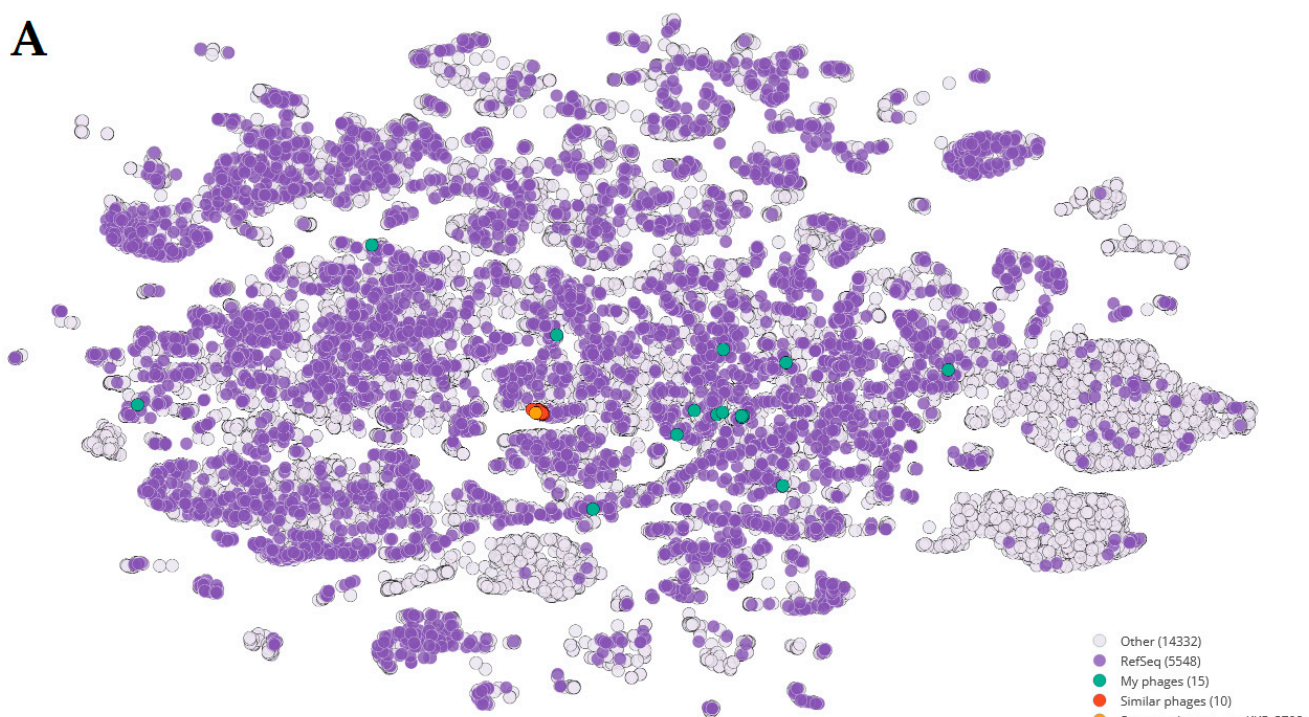
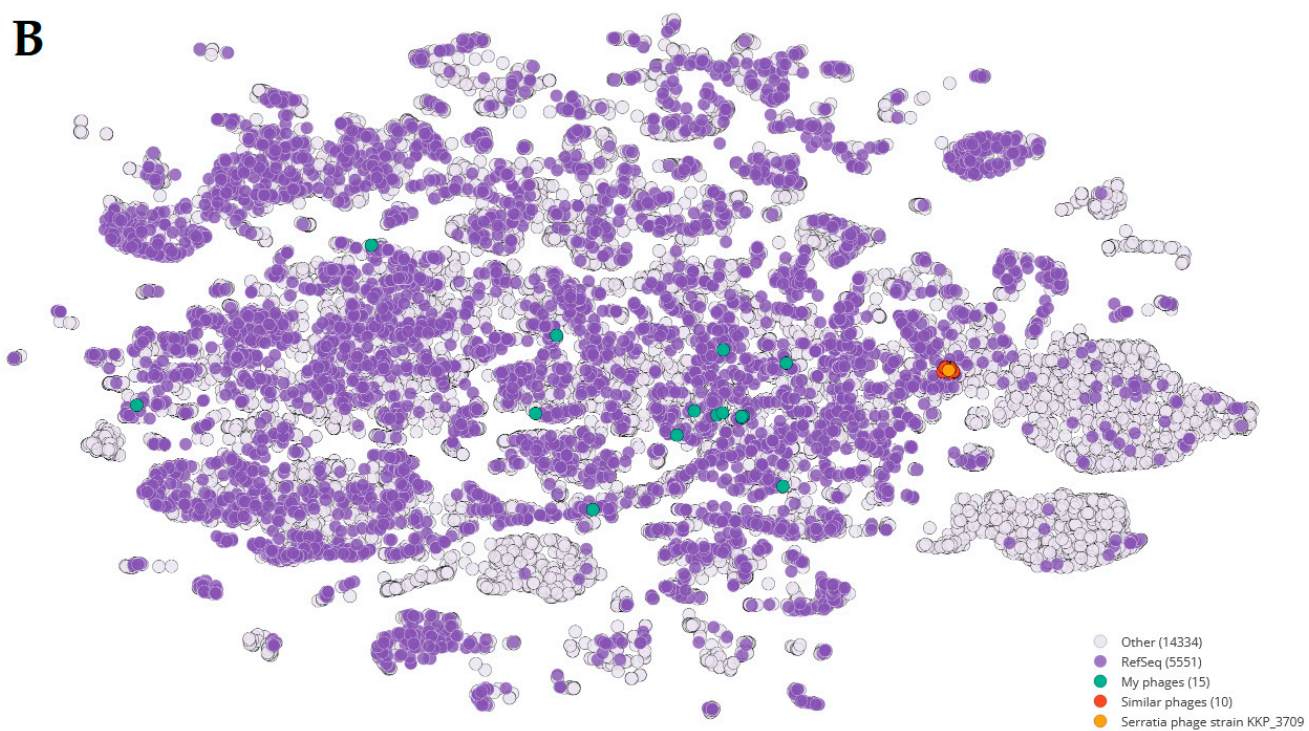


A**B**

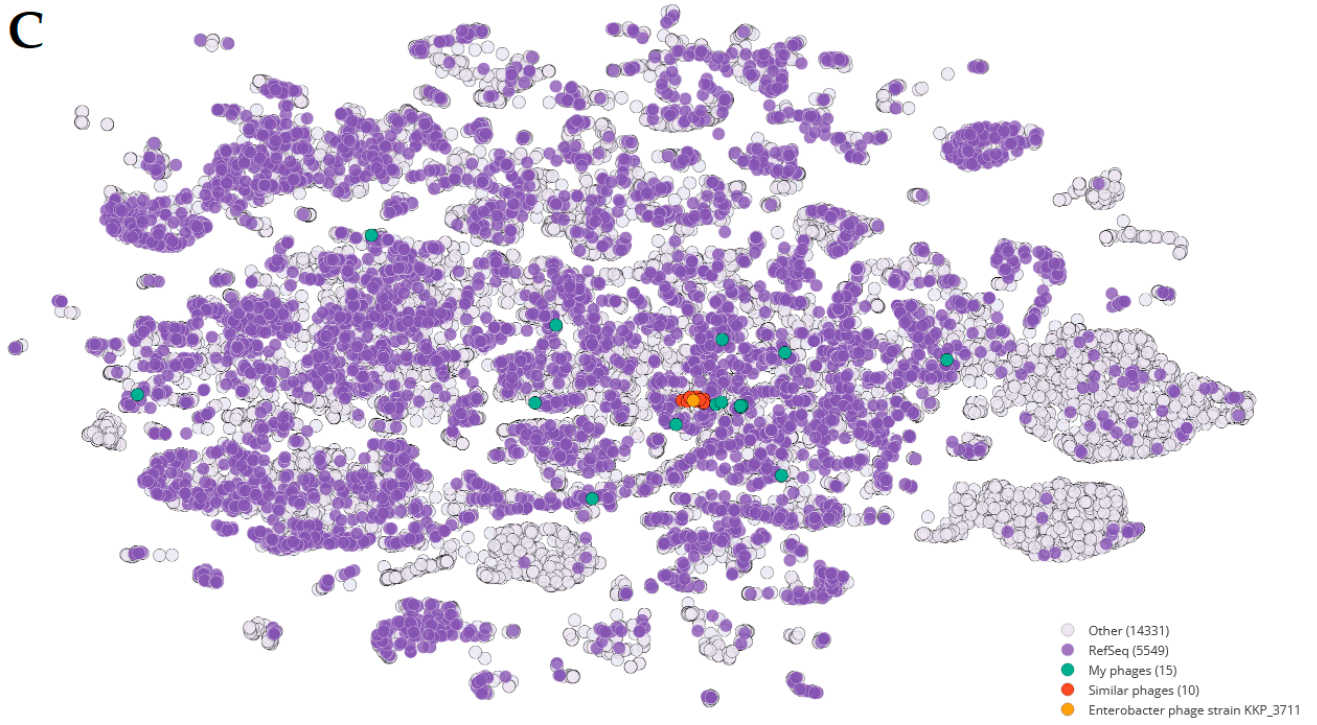


Figure S1. Phage similarity on 2D scatter plot computationally predicted and rendered through PhageAI software [89]: A – *Serratia* phage KKP_3708; B – *Serratia* phage KKP_3709; C – *Enterobacter* phage KKP_3711.