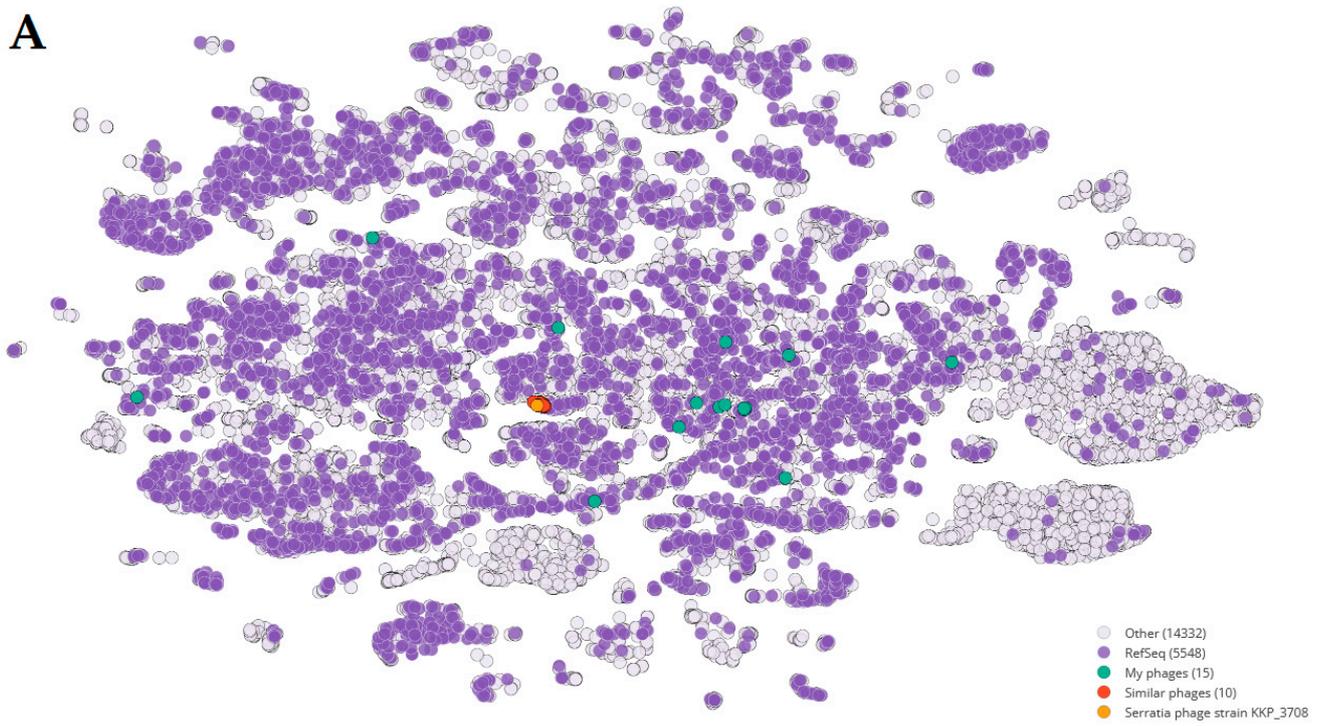
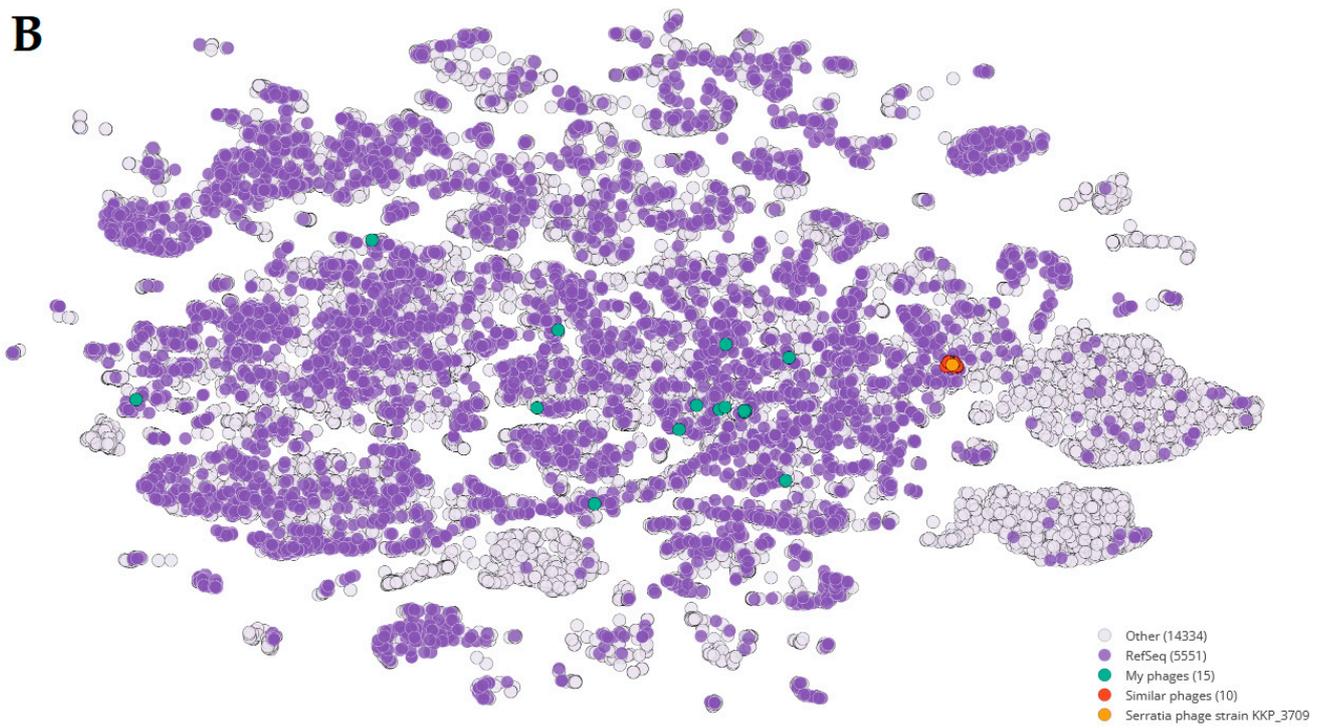
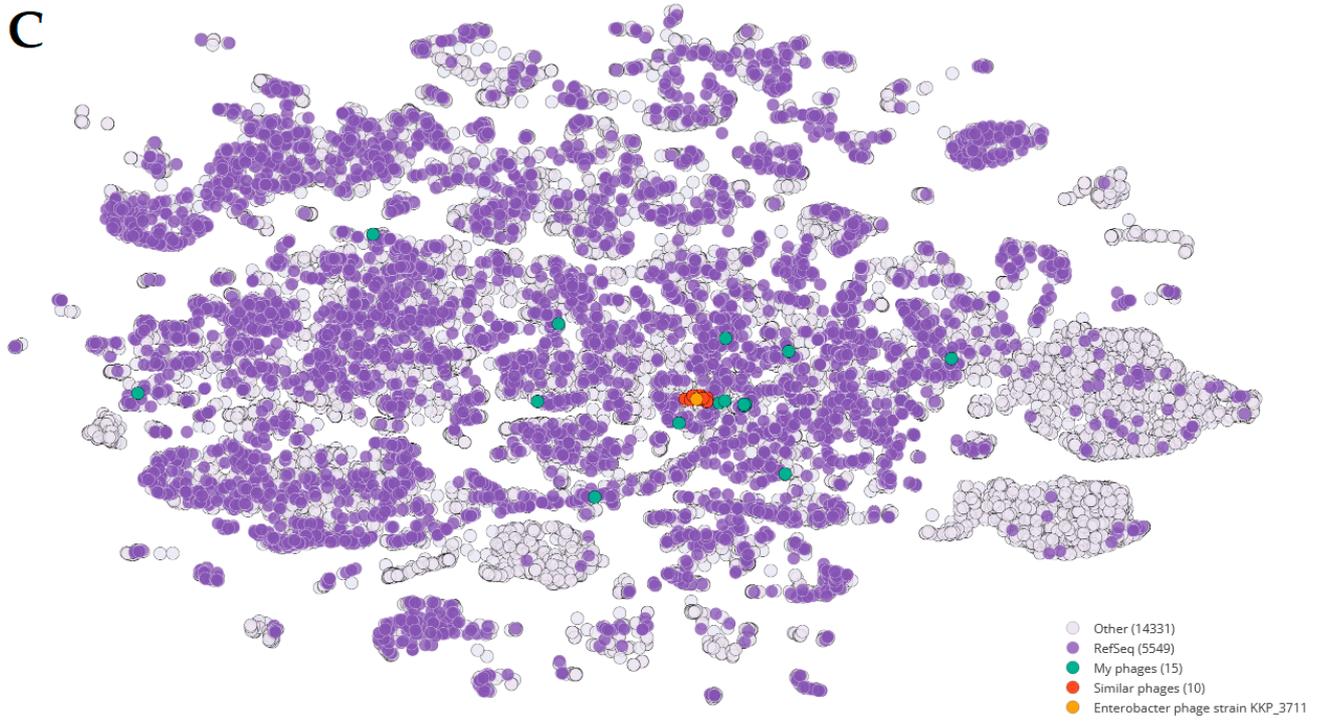


**A****B**



**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.