



Escherichia\_phage\_vB\_EcoS\_AVIO78A,\_complete\_genome

NC\_047786.1\_Salmonella\_phage\_vB\_SenS\_Sasha,\_complete\_genome

KY002061.1\_Salmonella\_phage\_vB\_SenS\_Sergei,\_complete\_genome

MH586730.1\_Salmonella\_phage\_Solent,\_complete\_genome

100.0	73.8	75.5	75.4	Escherichia_phage_vB_EcoS_AVIO78A,_complete_genome
0.8				
0.9	100.0	93.6	93.7	NC_047786.1_Salmonella_phage_vB_SenS_Sasha,_complete_genome
0.8				
0.8	1.1			
1.0	1.0	100.0	99.9	KY002061.1_Salmonella_phage_vB_SenS_Sergei,_complete_genome
0.8	1.2			
0.8	0.9	1.0		
1.0	1.0	1.0	100.0	MH586730.1_Salmonella_phage_Solent,_complete_genome
0.8	1.0	1.0		