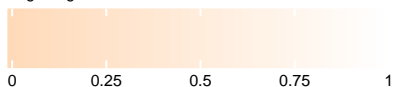
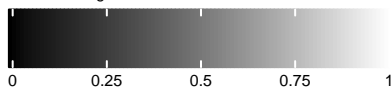


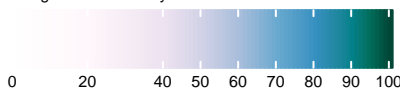
Aligned genome fraction



Genome length ratio



Intergenic similarity



2e+05
150000
1e+05
50000
0

Genome_length

	100.0	84.6	84.5	84.5	67.2	65.8	65.6	65.5	KJ019082
0.9		100.0	99.9	99.9	71.7	69.7	69.6	69.6	KJ019094
1.0		1.0							
0.9			100.0	99.9	71.7	69.7	69.6	69.6	KJ019054
1.0			1.0						
0.9				100.0	71.7	69.7	69.6	69.6	KJ019156
1.0				1.0					
0.9				1.0					
0.8	0.8	0.8	0.8	0.8	100.0	79.1	79.4	79.5	KJ019089
1.0	1.0	1.0	1.0	1.0					
0.8	0.8	0.8	0.8	0.8	0.8				
1.0	1.0	1.0	1.0	1.0	1.0	100.0	97.3	97.4	KU686202
0.8	0.8	0.8	0.8	0.8	0.9				
0.8	0.8	0.8	0.8	0.8	0.9	1.0			
1.0	1.0	1.0	1.0	1.0	1.0	1.0	100.0	99.4	KJ019069
0.8	0.8	0.8	0.8	0.8	0.9	1.0			
0.8	0.8	0.8	0.8	0.8	0.9	1.0	1.0		
1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	100.0	KU686200
0.8	0.8	0.8	0.8	0.8	0.9	1.0	1.0		
1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	1.0	KU686201
0.8	0.8	0.8	0.8	0.8	0.9	1.0	1.0	1.0	