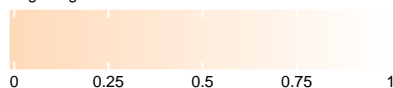
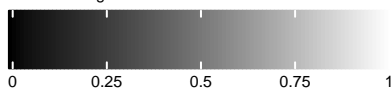


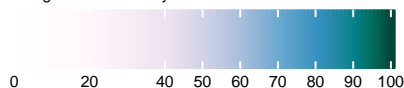
Aligned genome fraction



Genome length ratio



Intergenomic similarity



150000
1e+05
50000
0

Genome_length

	JN377895	MH179476	HM452125	MH179477	DQ529280	JN377894	JX306041	MF479730
JN377895	100.0	89.4	88.7	89.2	86.5	87.8	88.7	86.6
MH179476	0.9	100.0	91.5	91.3	89.8	86.7	89.8	88.3
HM452125	0.9	1.0	100.0	93.0	90.0	87.2	92.7	92.3
MH179477	0.9	1.0	1.0	100.0	88.2	86.5	91.1	90.1
DQ529280	0.9	1.0	1.0	0.9	100.0	86.6	89.8	87.1
JN377894	1.0	0.9	0.9	0.9	0.9	100.0	88.1	88.0
JX306041	0.9	1.0	1.0	1.0	1.0	0.9	100.0	91.2
MF479730	0.9	0.9	1.0	1.0	1.0	0.9	1.0	100.0