subject id	% identity	alignment l	mismatche	gap opens	evalue
Aphidoletes_aphidimyza	100	19	0	0	0.036
Eristalis_dimiata	100	19	0	0	0.036
Eristalis_dimiata	100	20	0	0	0.009
Eristalis_dimiata	100	19	0	0	0.036
Harmonia_axyridis	100	19	0	0	0.036
Lasius_niger	100	19	0	0	0.036
Lasius_niger	100	19	0	0	0.036
Myzus_persicae_genome	100	19	0	0	0.036
Myzus_persicae_genome	100	21	0	0	0.002
Myzus_persicae_genome	100	19	0	0	0.036
Myzus_persicae_genome	100	21	0	0	0.002
Myzus_persicae_genome	100	19	0	0	0.036
Myzus_persicae_genome	100	19	0	0	0.036
Myzus_persicae_genome	100	19	0	0	0.036
Myzus_persicae_genome	100	19	0	0	0.036
Myzus_persicae_genome	100	21	0	0	0.002
Myzus_persicae_genome	100	19	0	0	0.036
Myzus_persicae_genome	100	21	0	0	0.002
Myzus_persicae_genome	100	21	0	0	0.002
Myzus_persicae_genome	100	21	0	0	0.002
Myzus_persicae_genome	100	19	0	0	0.036
Myzus_persicae_transcriptome	100	21	0	0	0.002
Myzus_persicae_transcriptome	100	20	0	0	0.009
Myzus_persicae_transcriptome	100	21	0	0	0.002
Myzus_persicae_transcriptome	100	19	0	0	0.036
Myzus_persicae_transcriptome	100	19	0	0	0.036
Myzus_persicae_transcriptome	100	21	0	0	0.002
Myzus_persicae_transcriptome	100	19	0	0	0.036
Myzus_persicae_transcriptome	100	21	0	0	0.002
Myzus_persicae_transcriptome	100	21	0	0	0.002
Myzus_persicae_transcriptome	100	21	0	0	0.002
Myzus_persicae_transcriptome	100	21	0	0	0.002
Myzus_persicae_transcriptome	100	21	0	0	0.002
Myzus_persicae_transcriptome	100	19	0	0	0.036
Apis_mellifera_removed	100	19	0	0	0.036
Apis_mellifera_removed	100	19	0	0	0.036
Apis_mellifera_removed	100	19	0	0	0.036