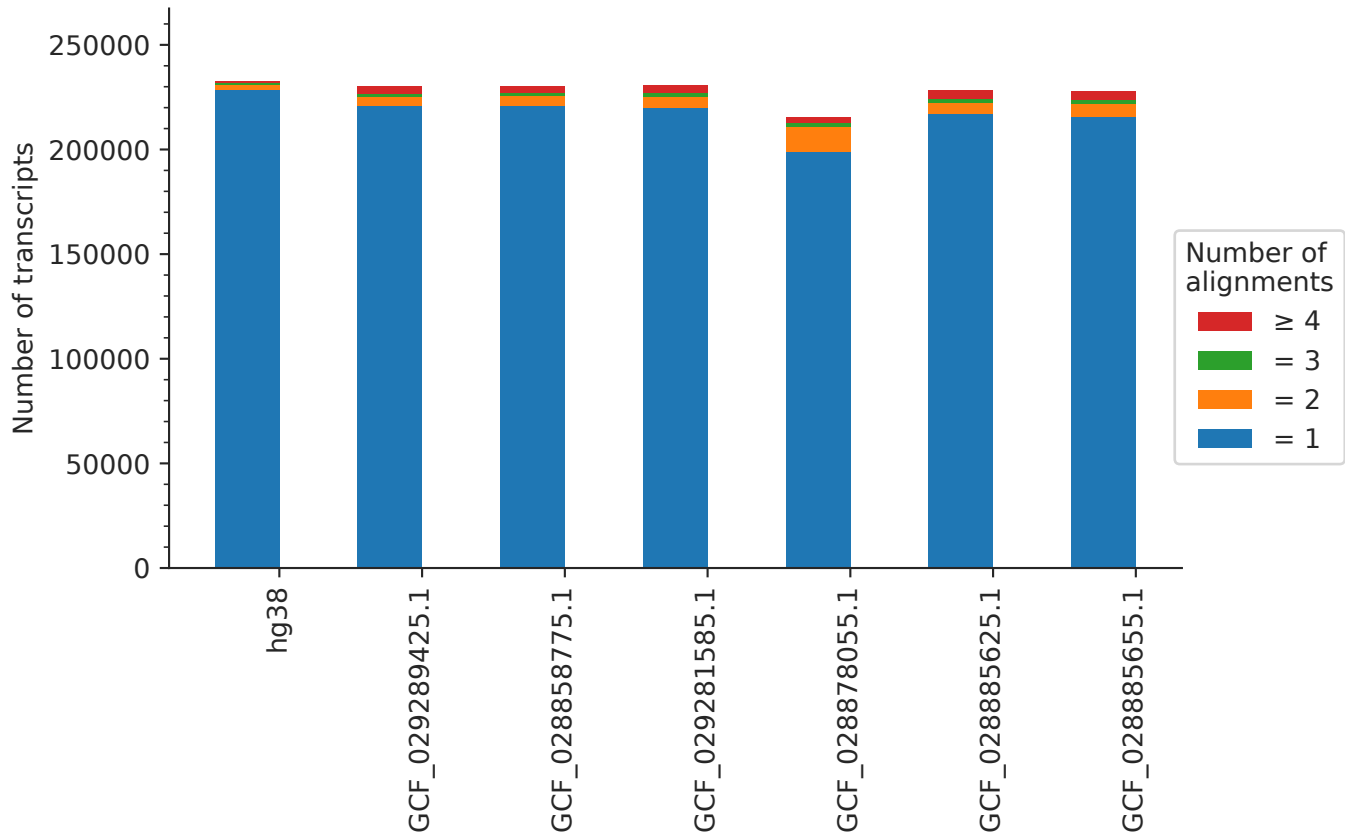
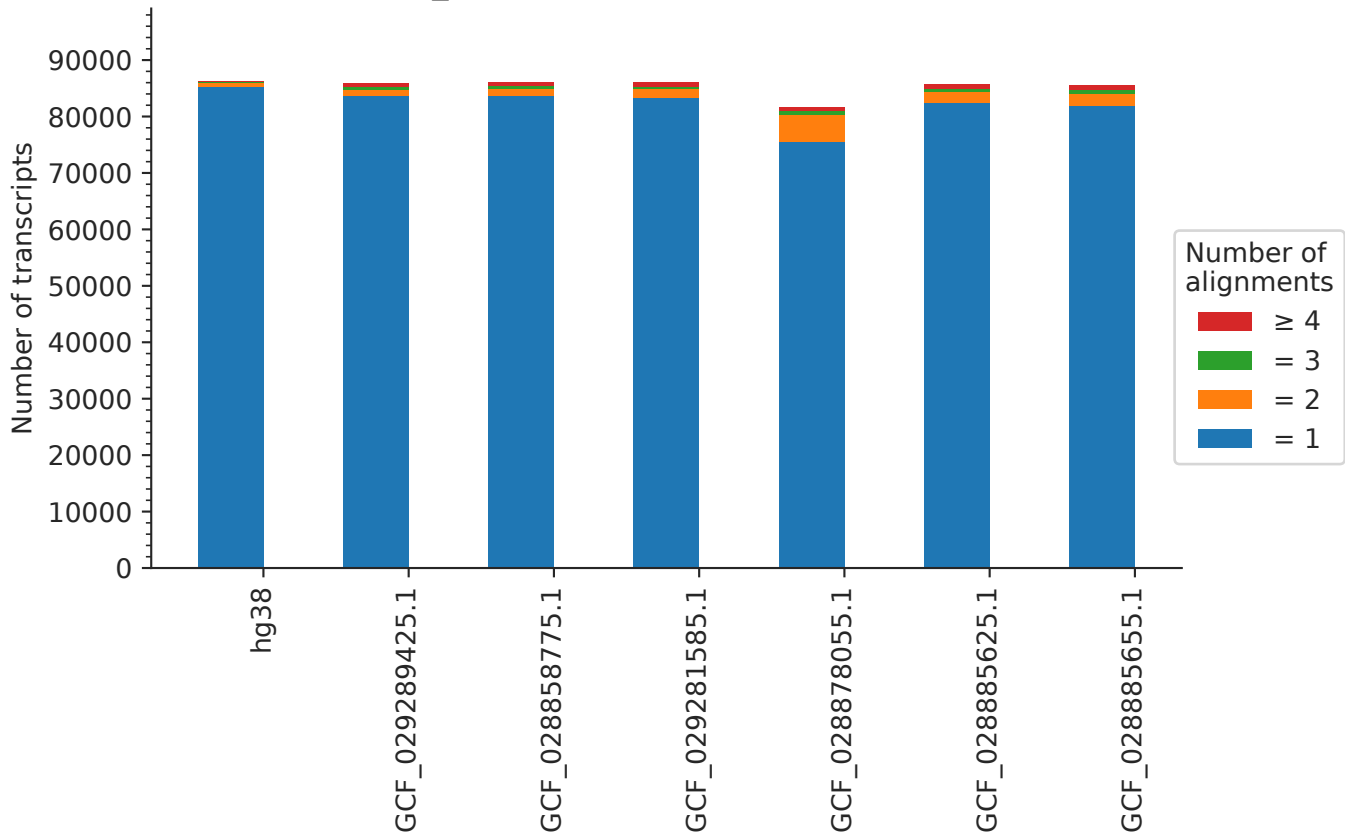


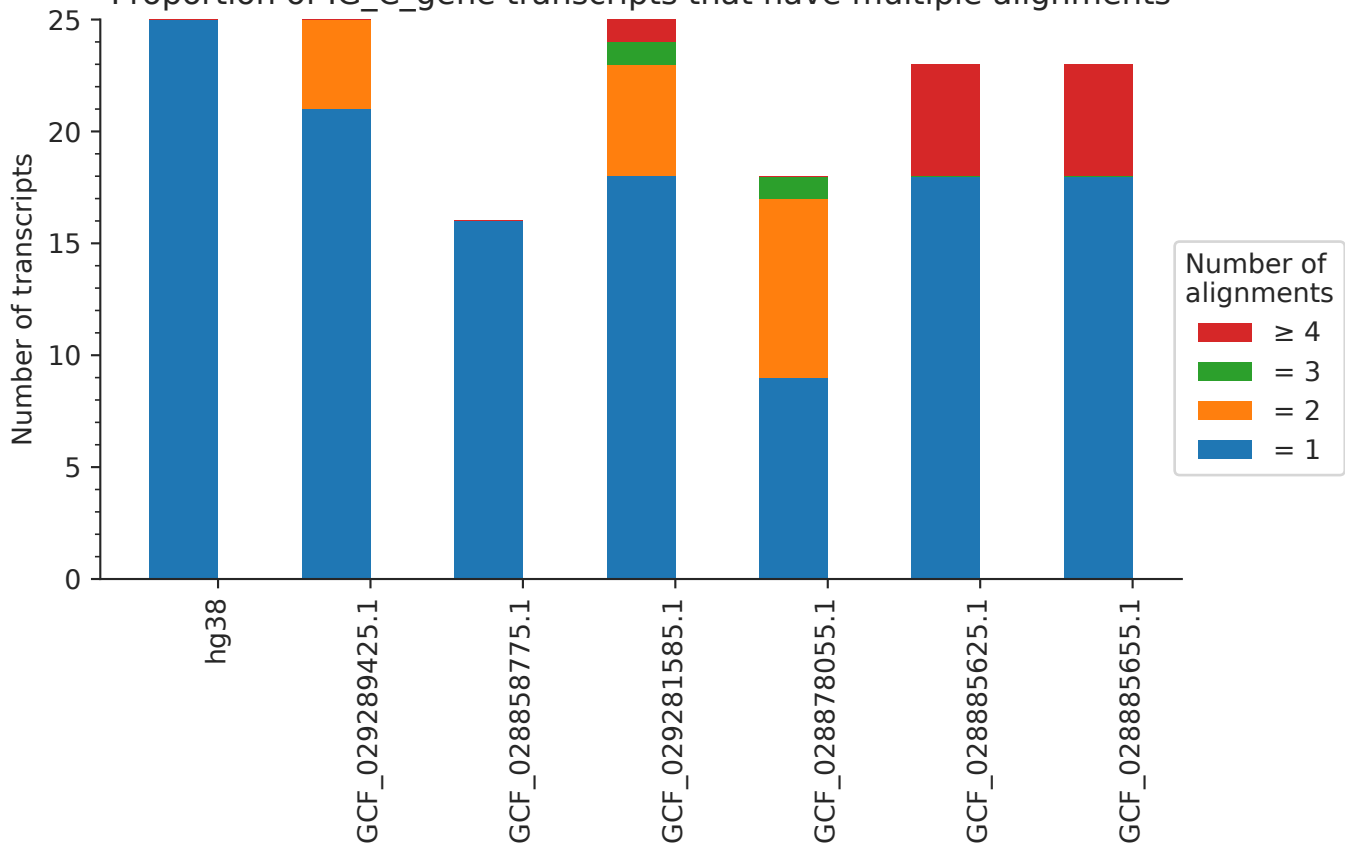
Proportion of transcripts that have multiple alignments



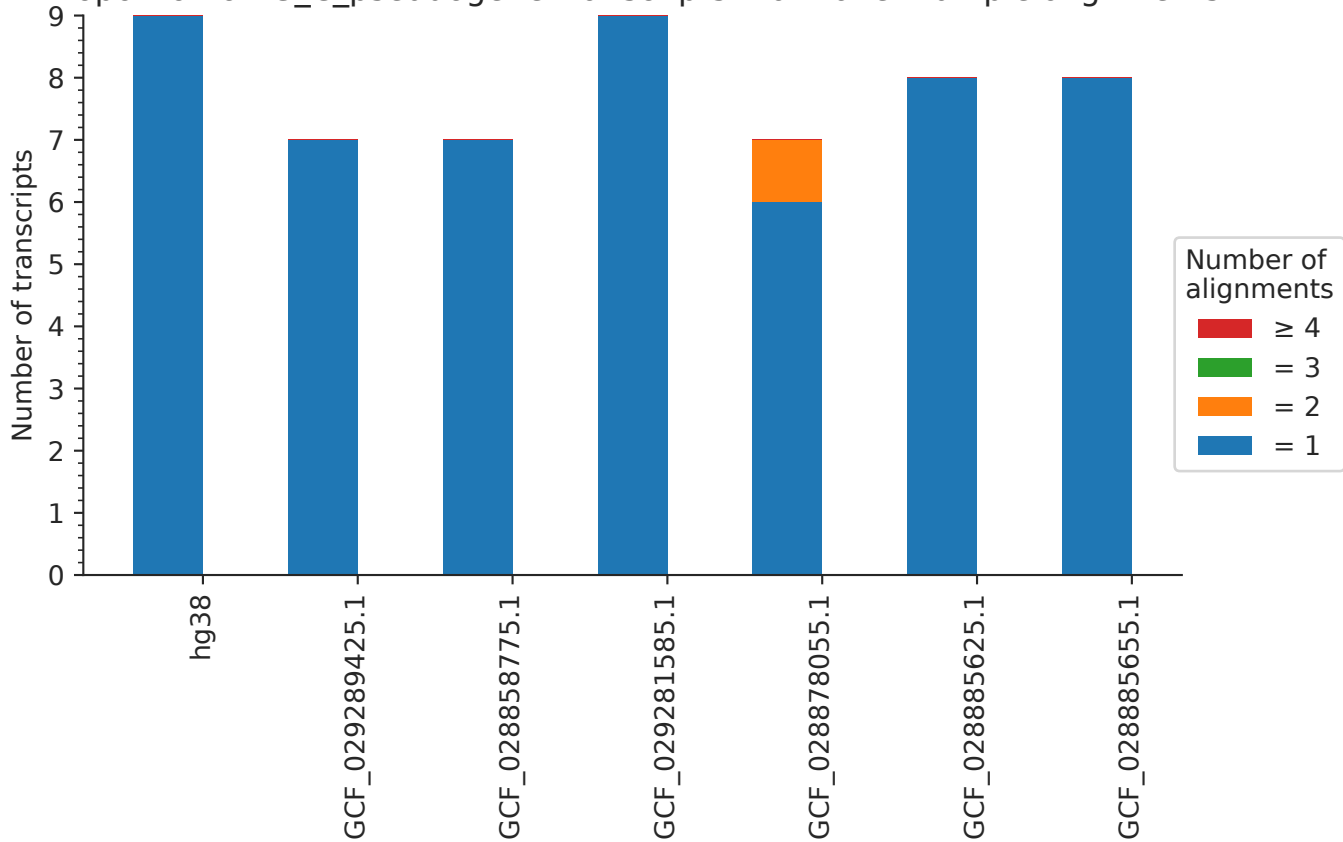
Proportion of protein_coding transcripts that have multiple alignments



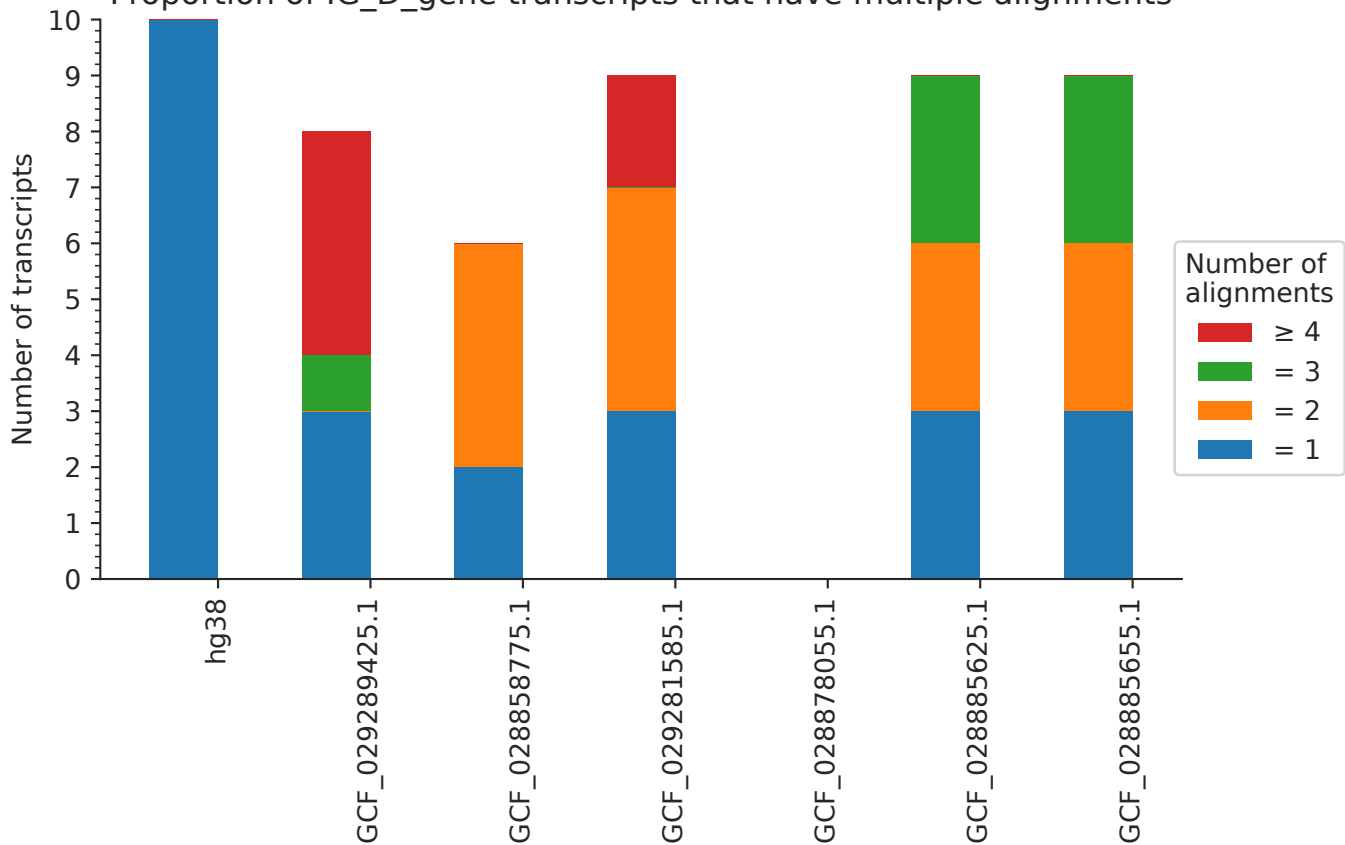
Proportion of IG_C_gene transcripts that have multiple alignments



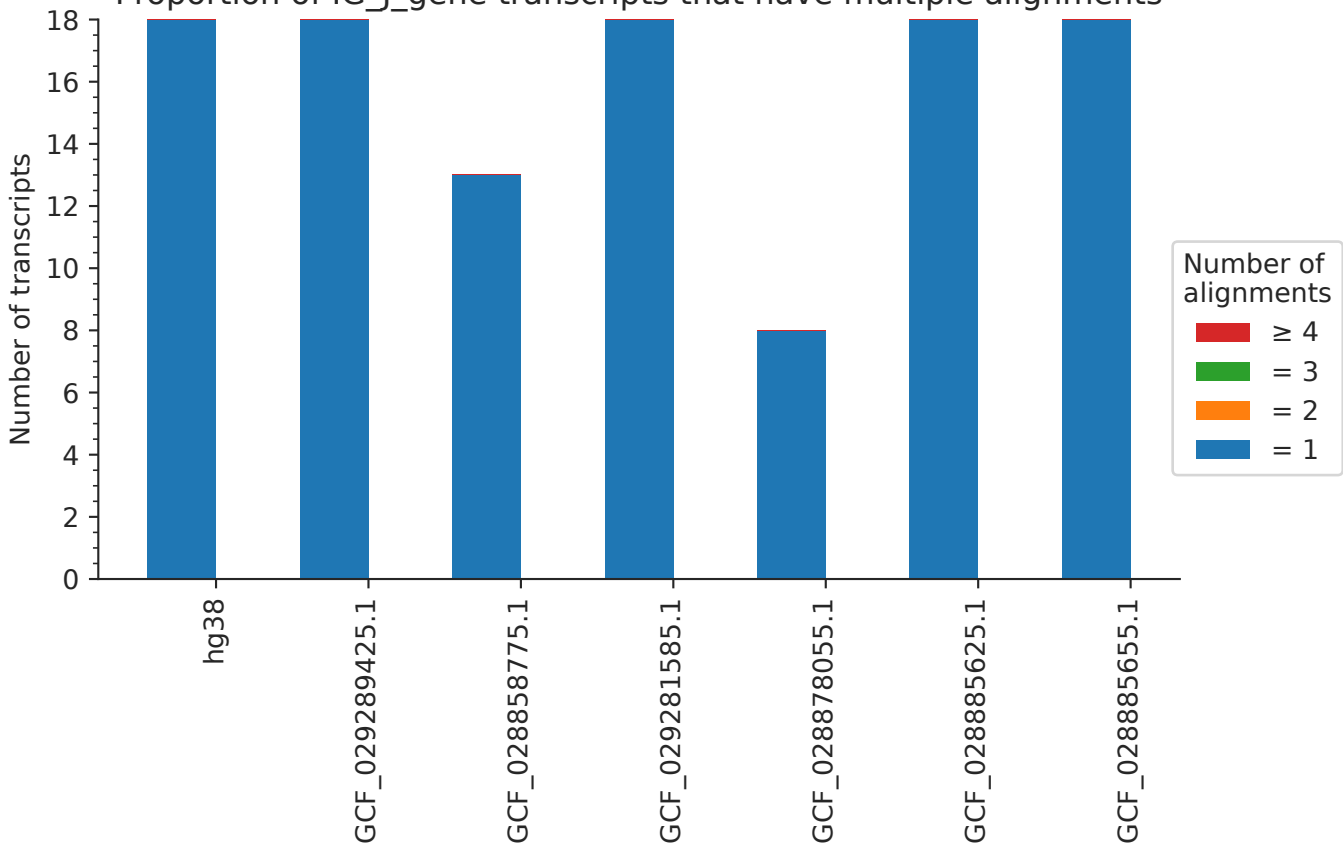
Proportion of IG_C_pseudogene transcripts that have multiple alignments



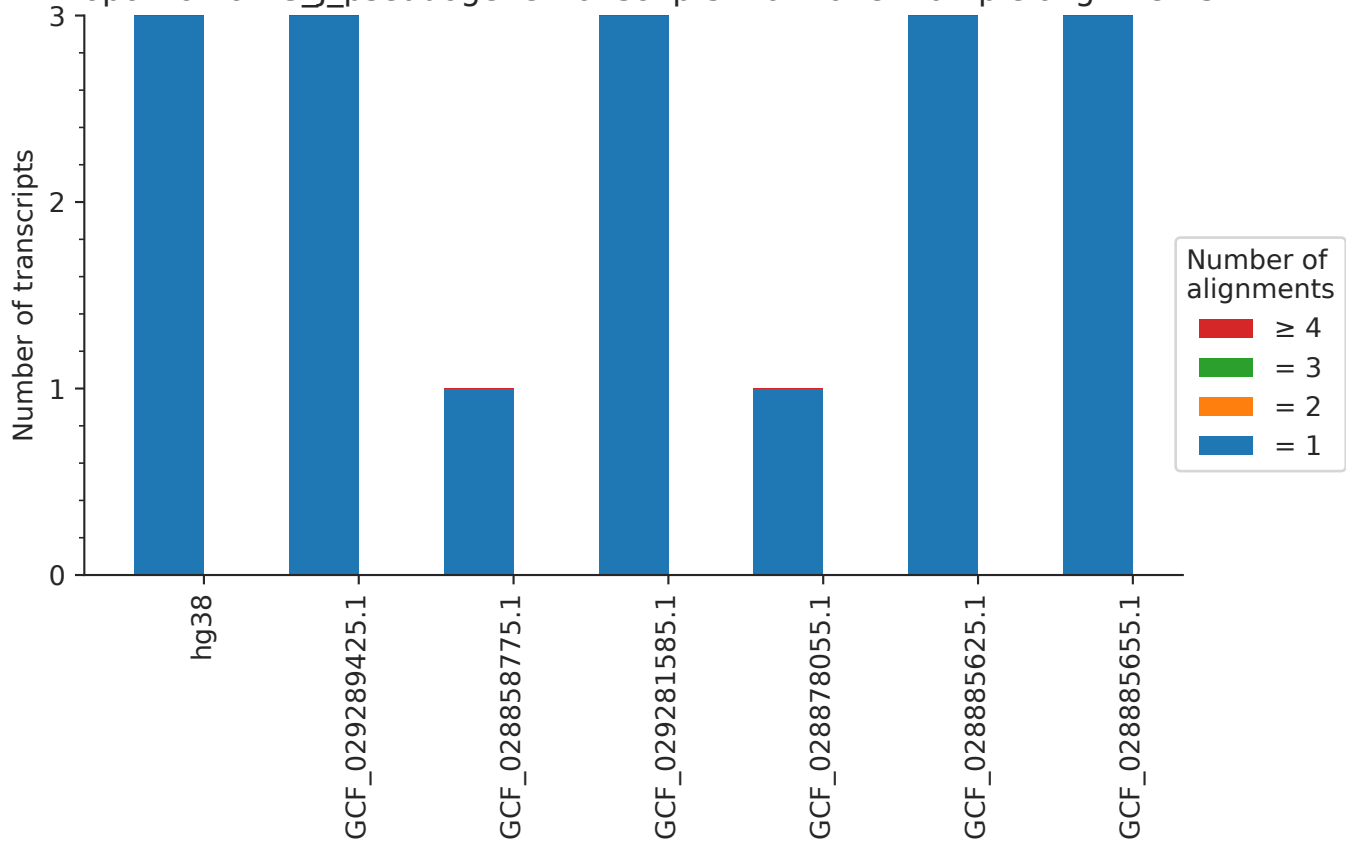
Proportion of IG_D_gene transcripts that have multiple alignments



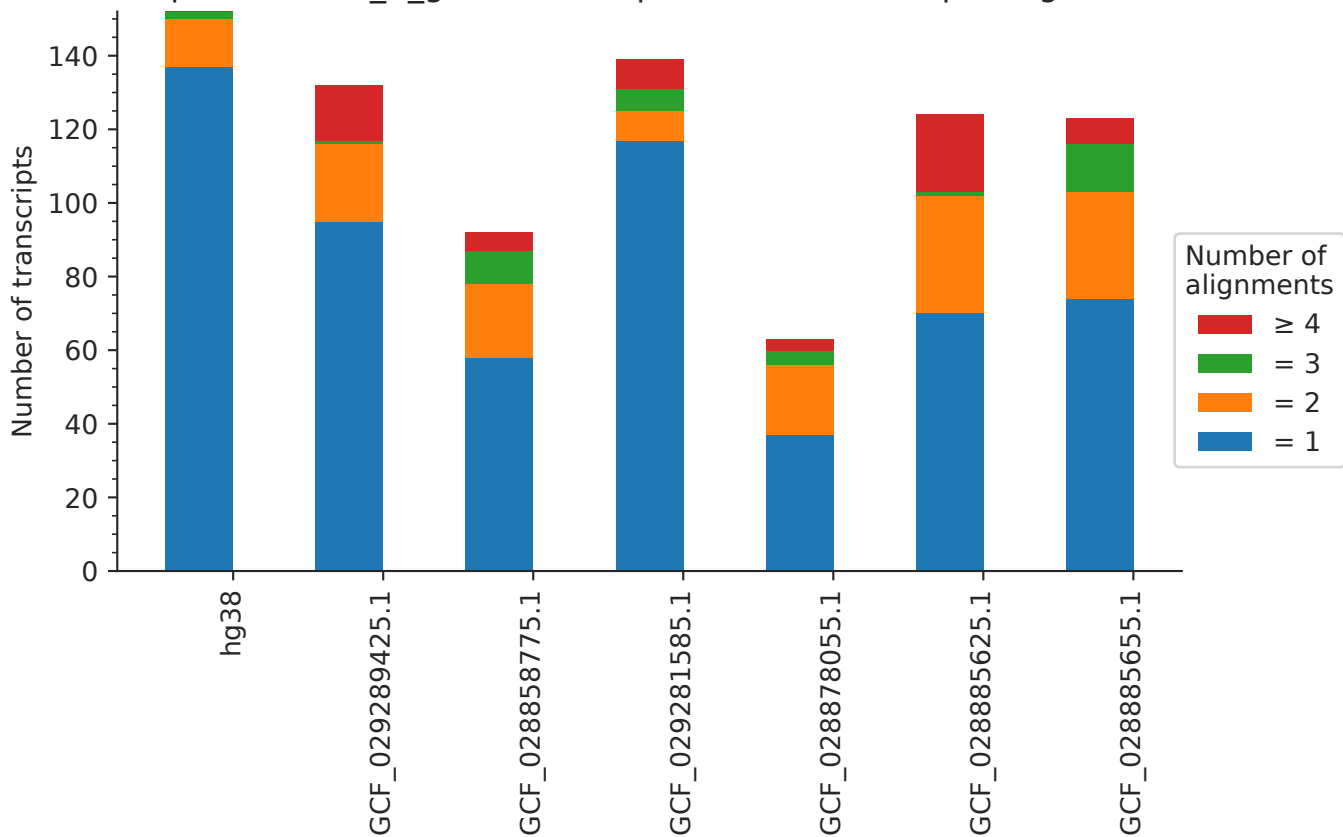
Proportion of IG_J_gene transcripts that have multiple alignments



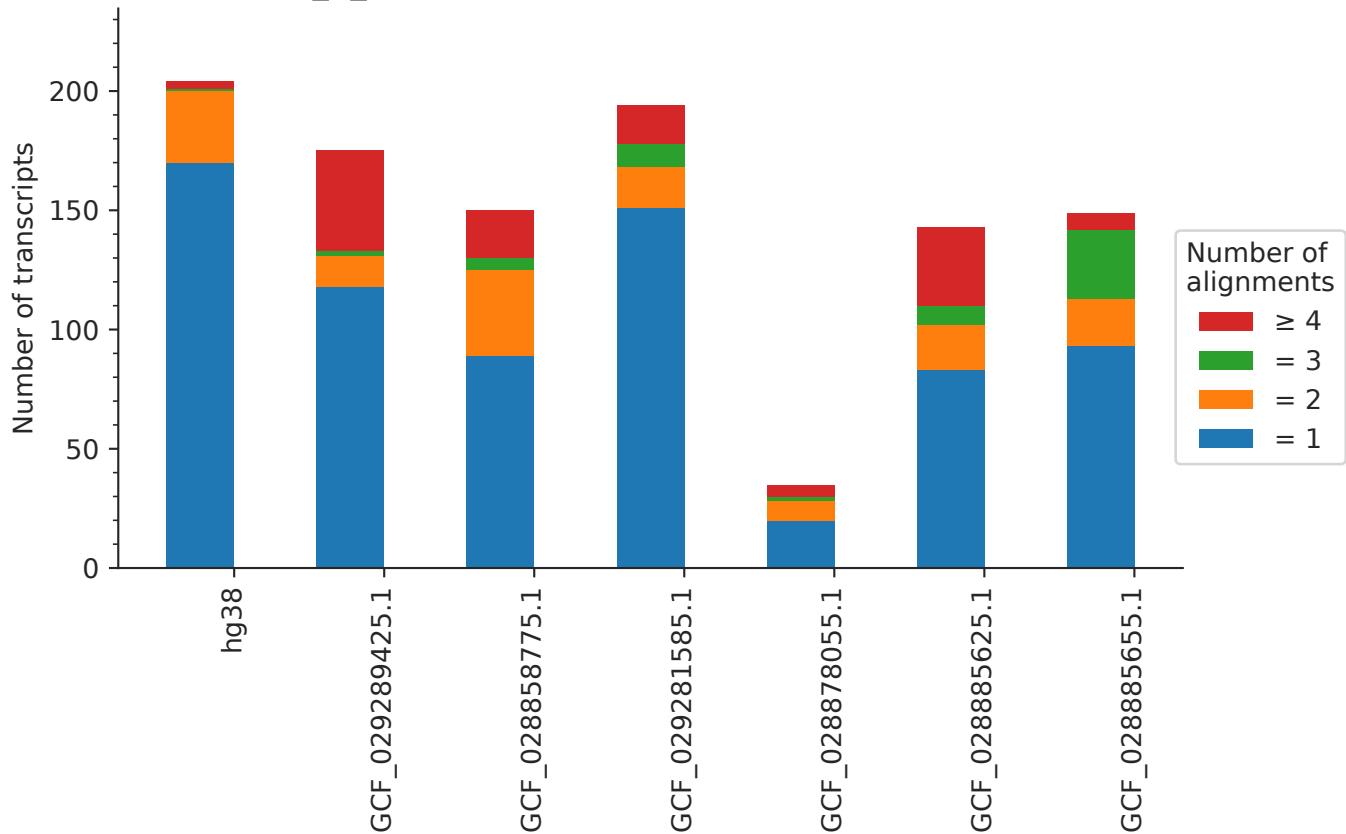
Proportion of IG_J_pseudogene transcripts that have multiple alignments



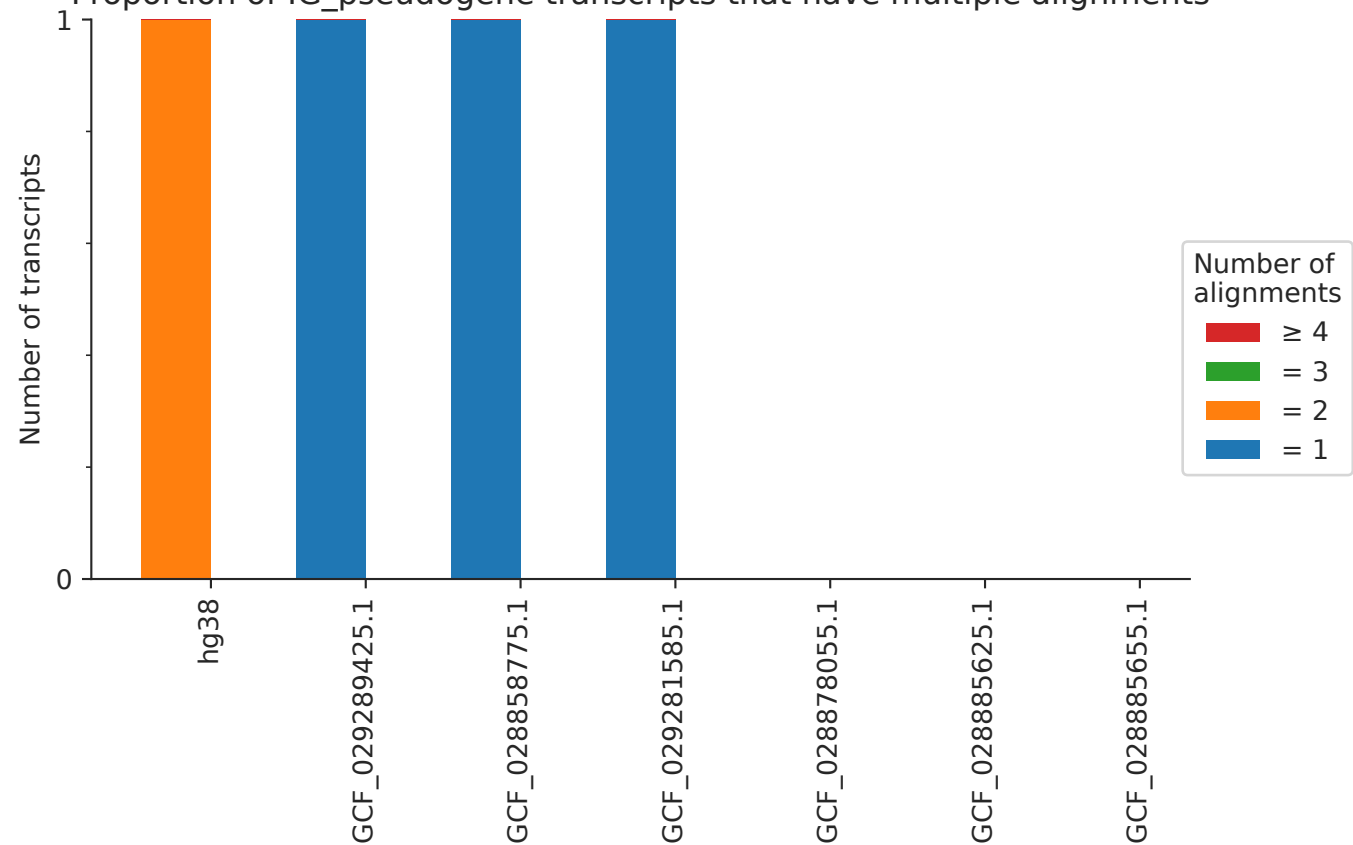
Proportion of IG_V_gene transcripts that have multiple alignments



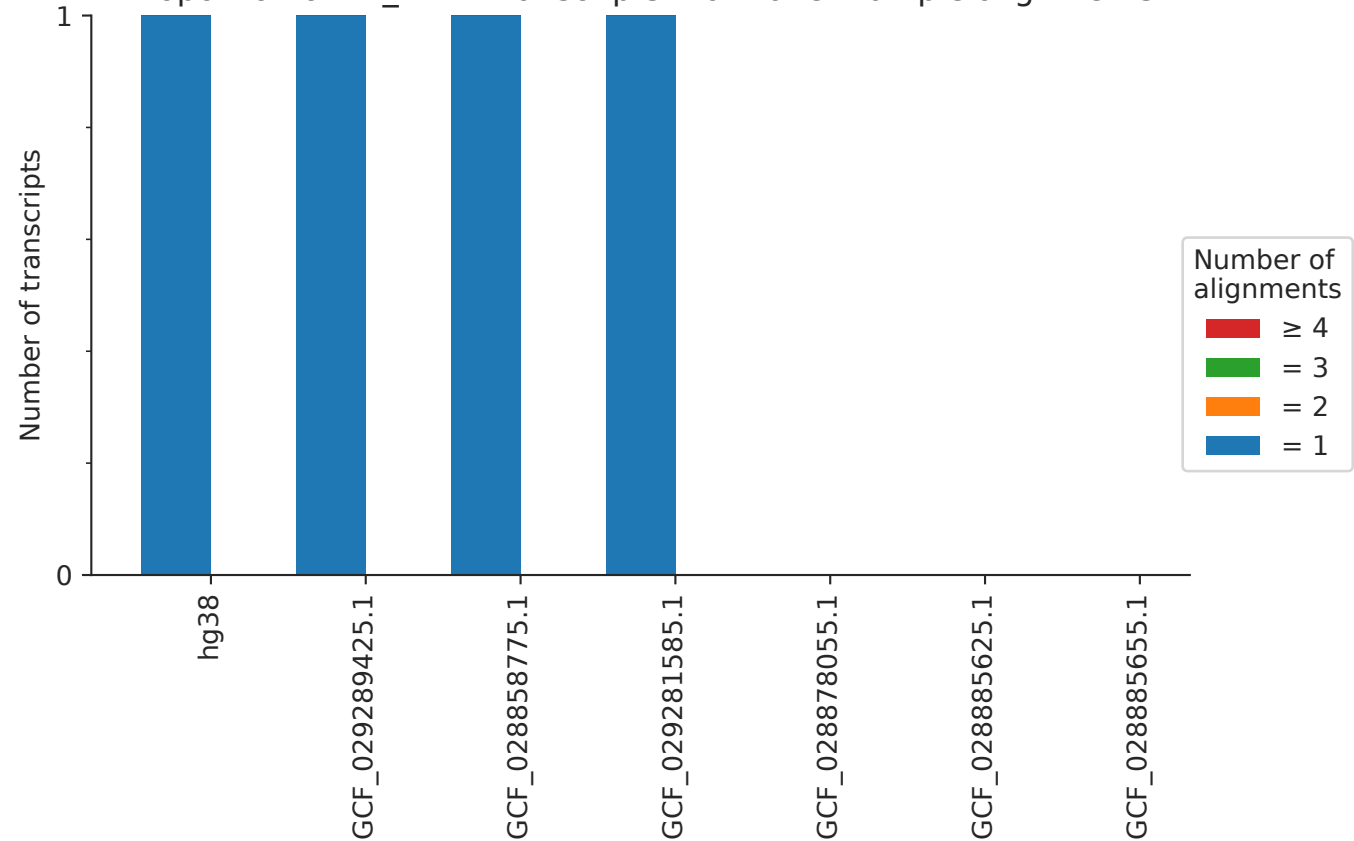
Proportion of IG_V_pseudogene transcripts that have multiple alignments



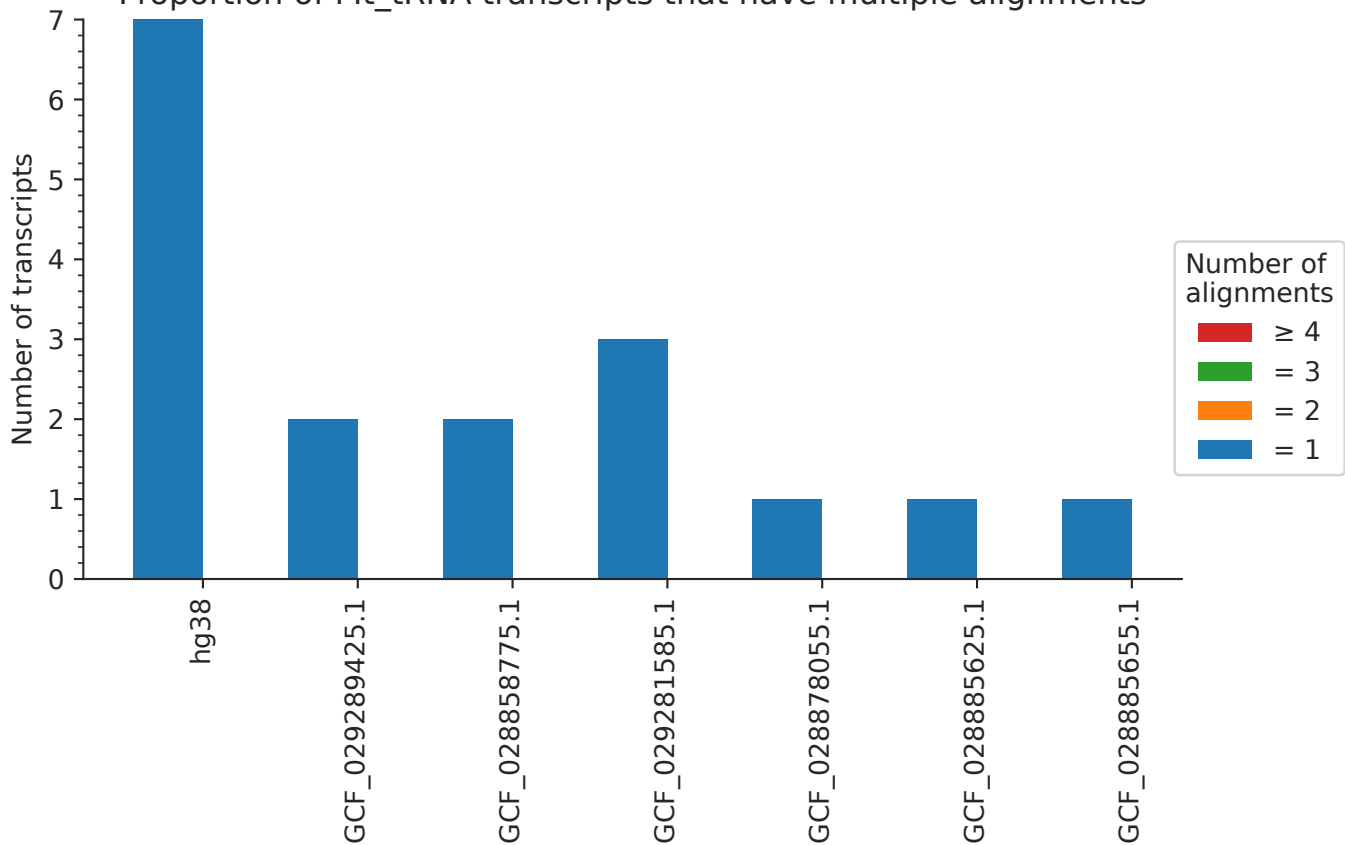
Proportion of IG_pseudogene transcripts that have multiple alignments



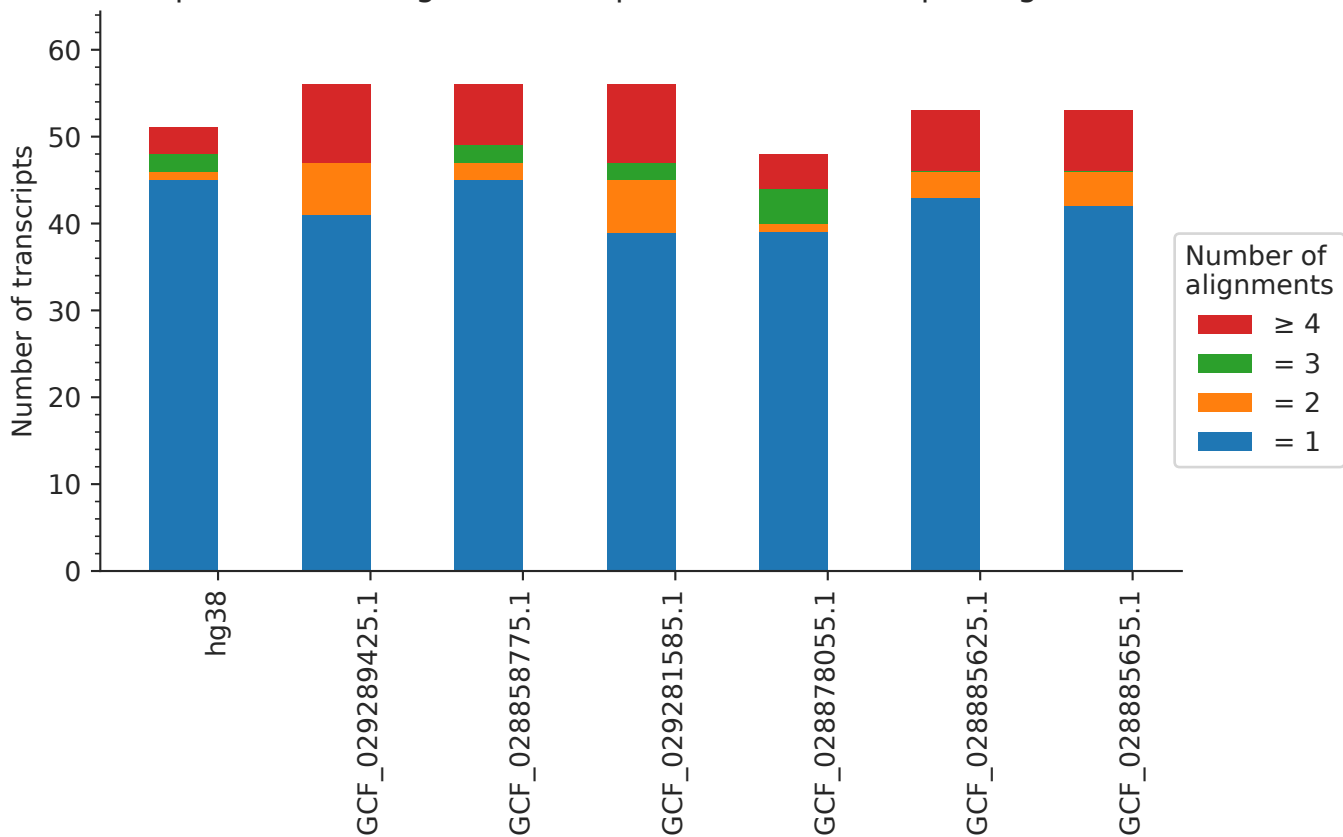
Proportion of Mt_rRNA transcripts that have multiple alignments



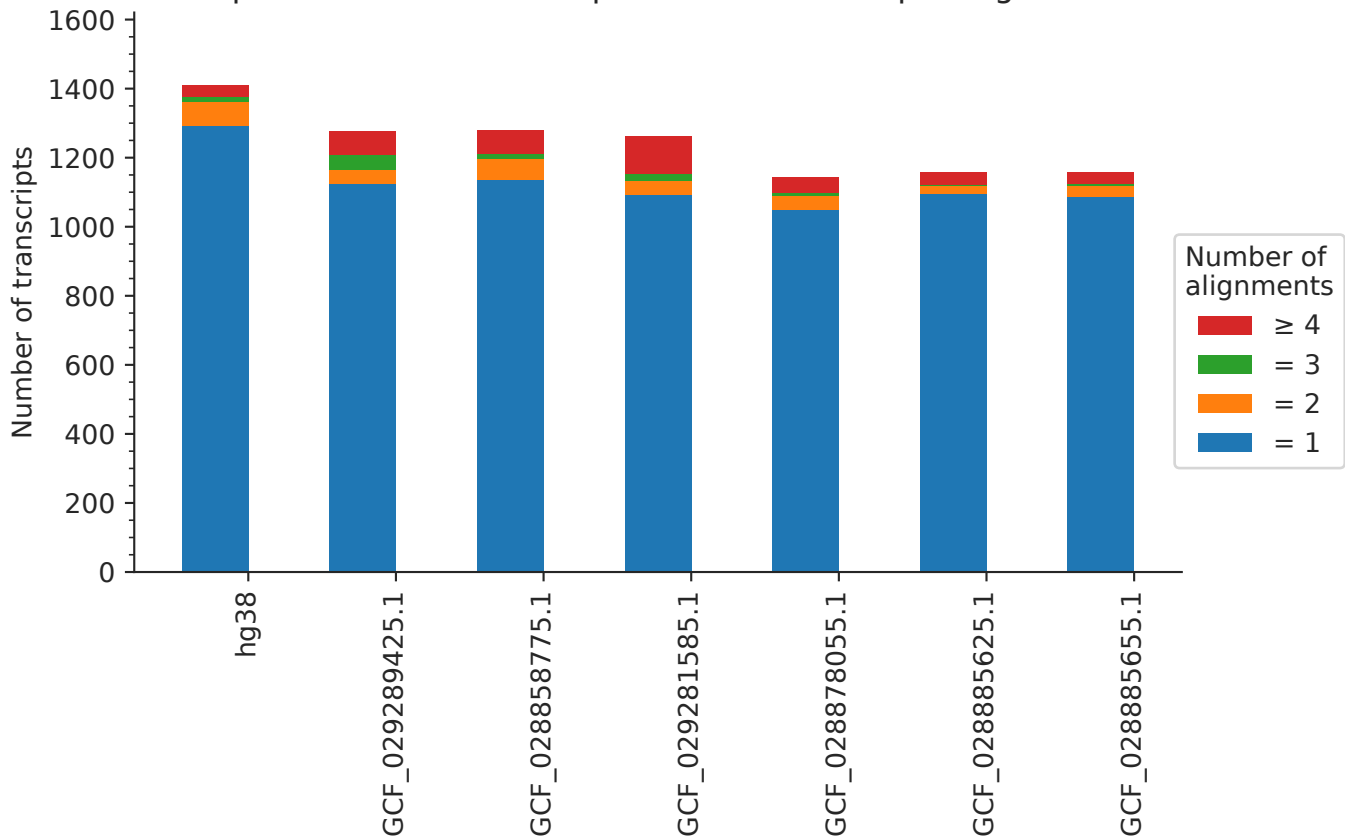
Proportion of Mt_tRNA transcripts that have multiple alignments



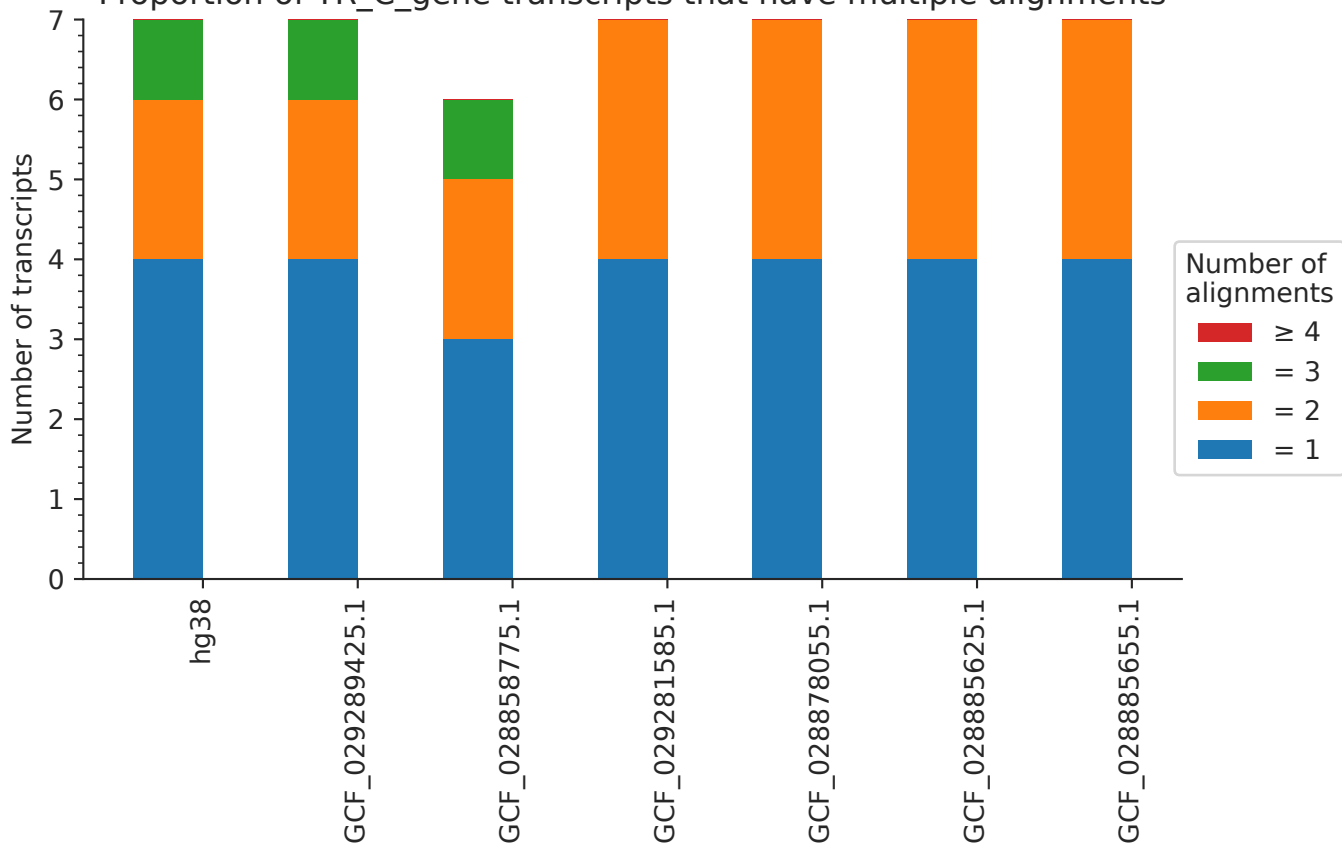
Proportion of StringTie transcripts that have multiple alignments



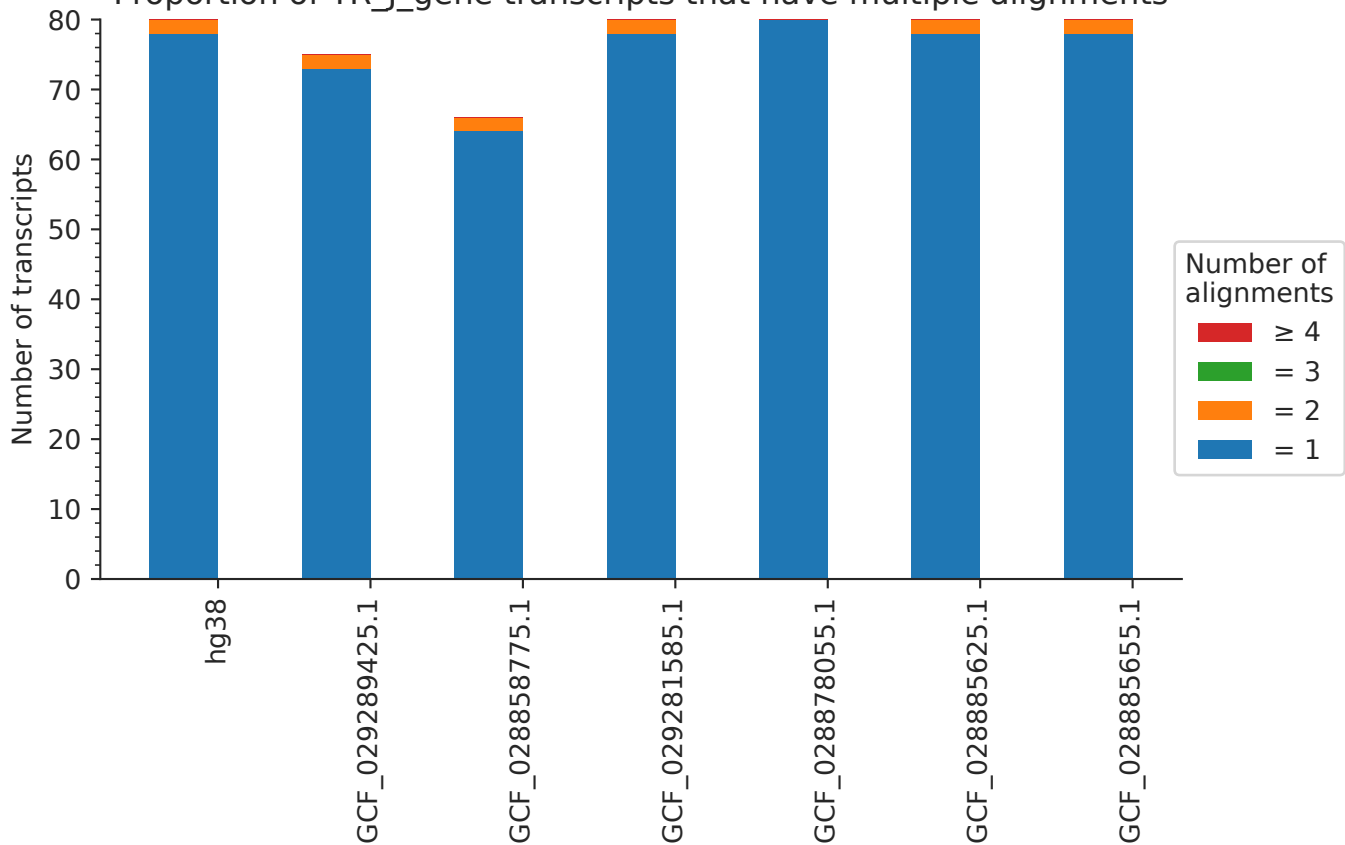
Proportion of TEC transcripts that have multiple alignments



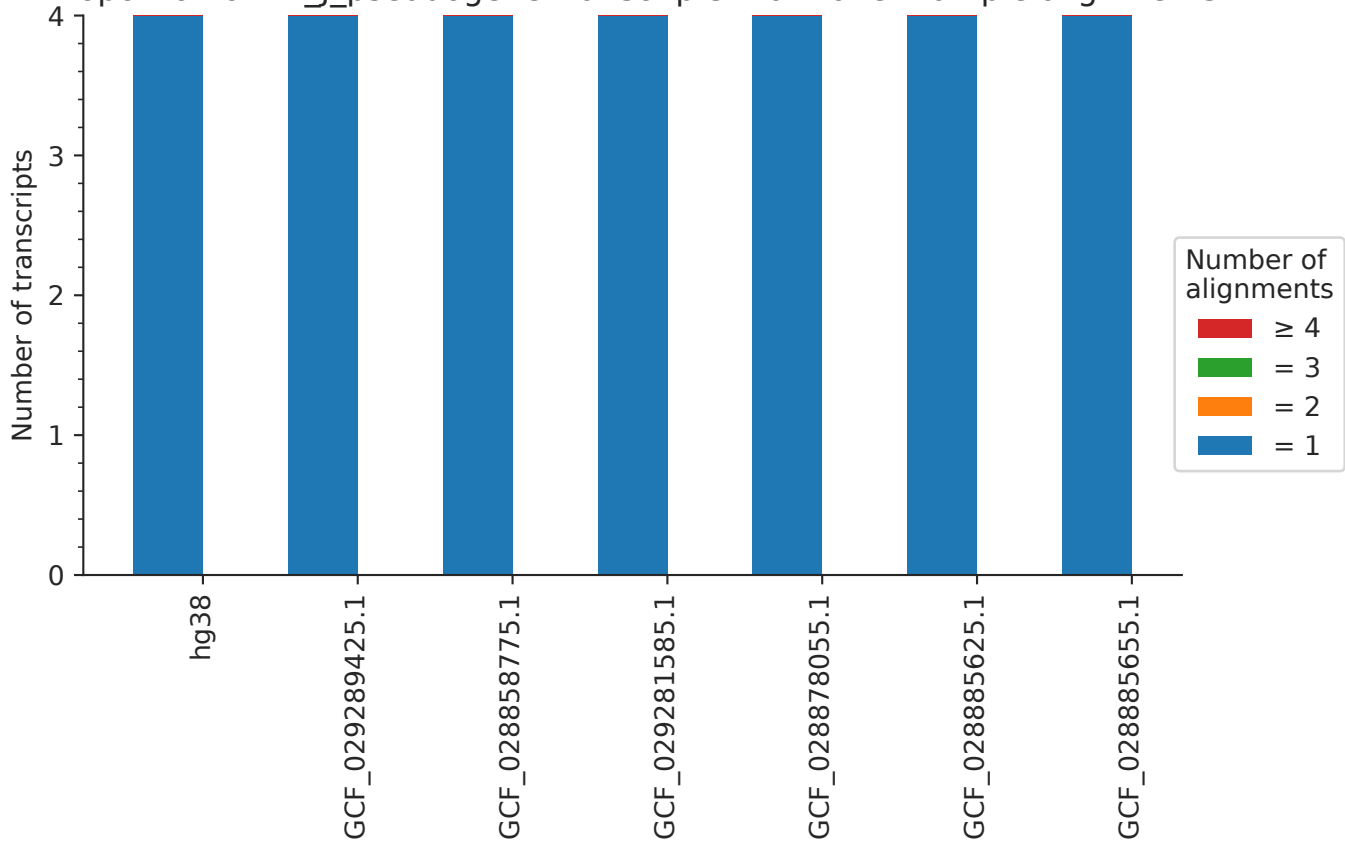
Proportion of TR_C_gene transcripts that have multiple alignments



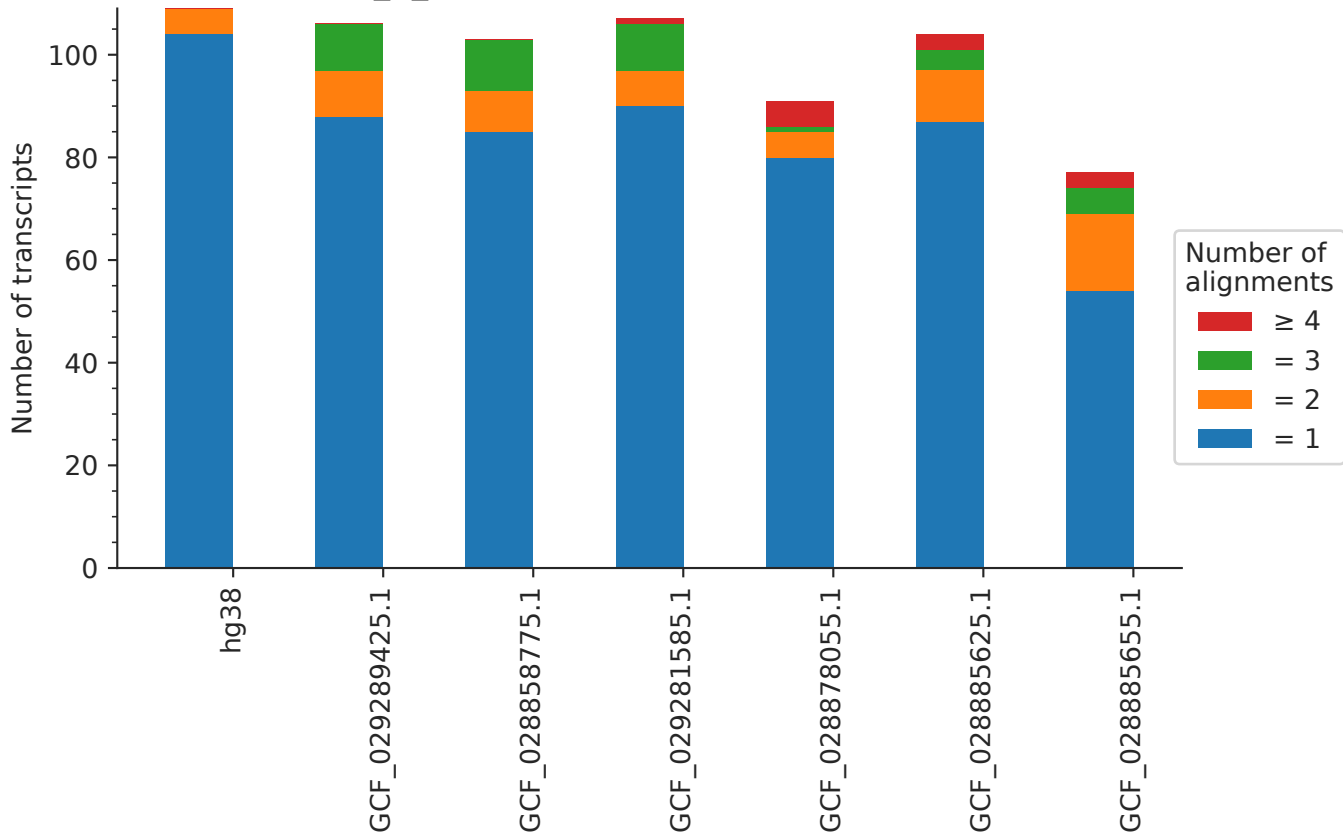
Proportion of TR_J_gene transcripts that have multiple alignments



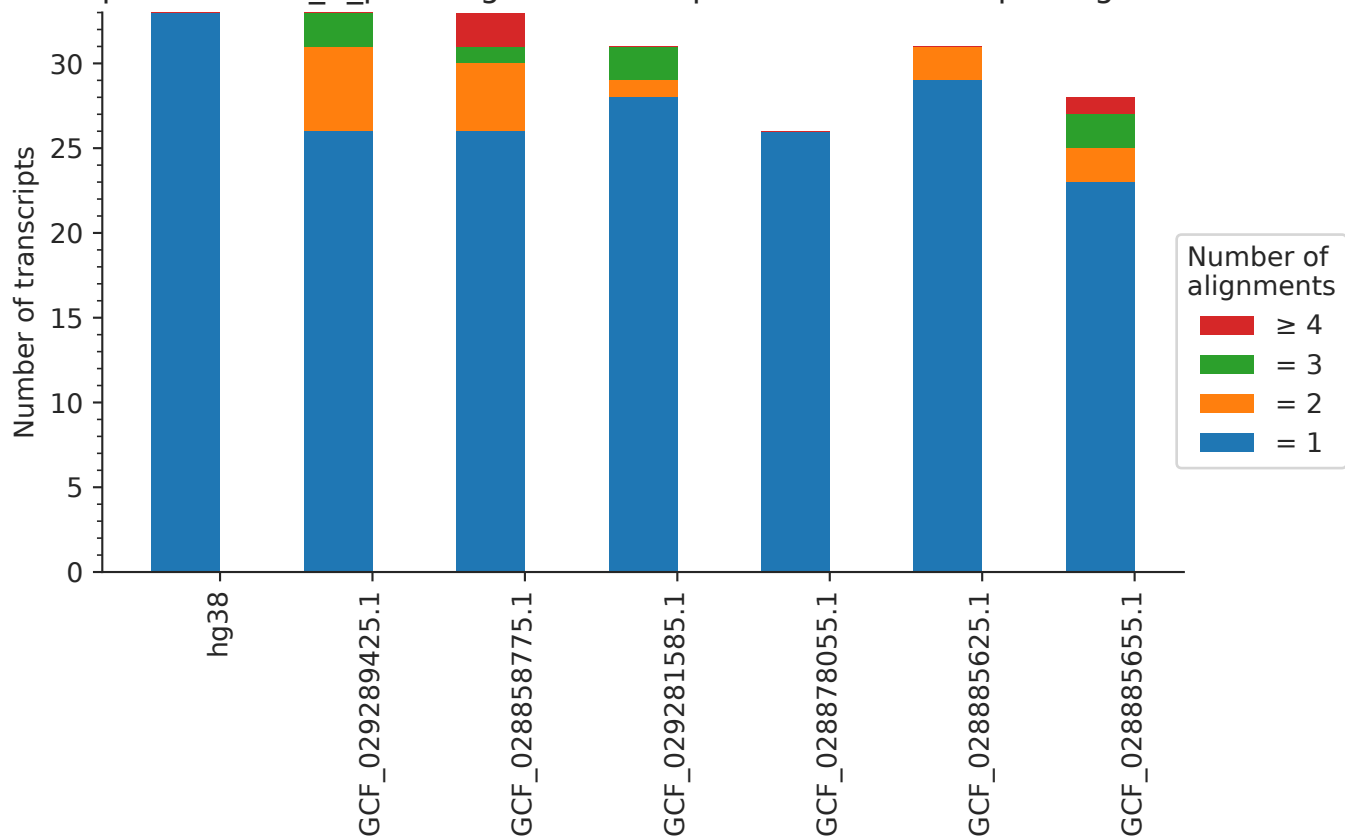
Proportion of TR_J_pseudogene transcripts that have multiple alignments



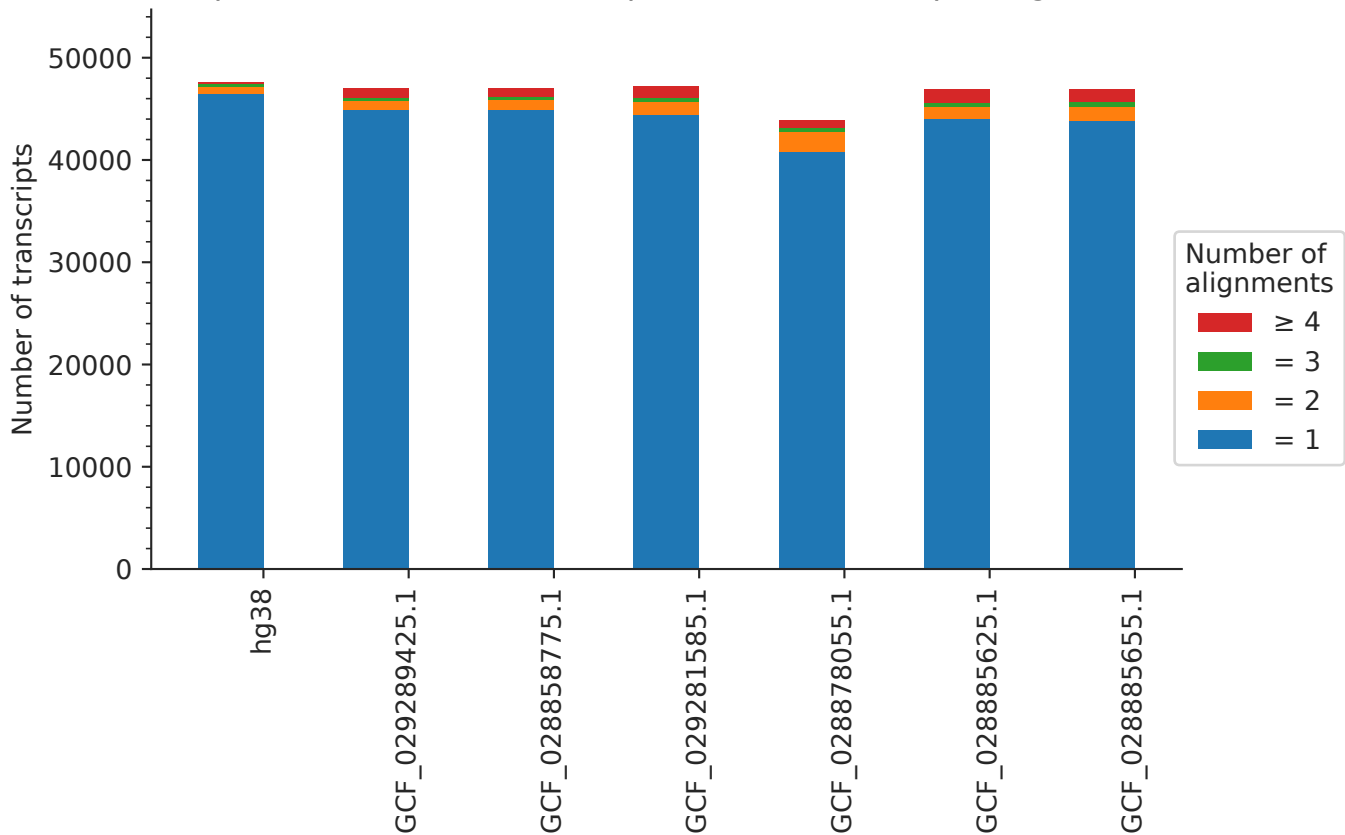
Proportion of TR_V_gene transcripts that have multiple alignments



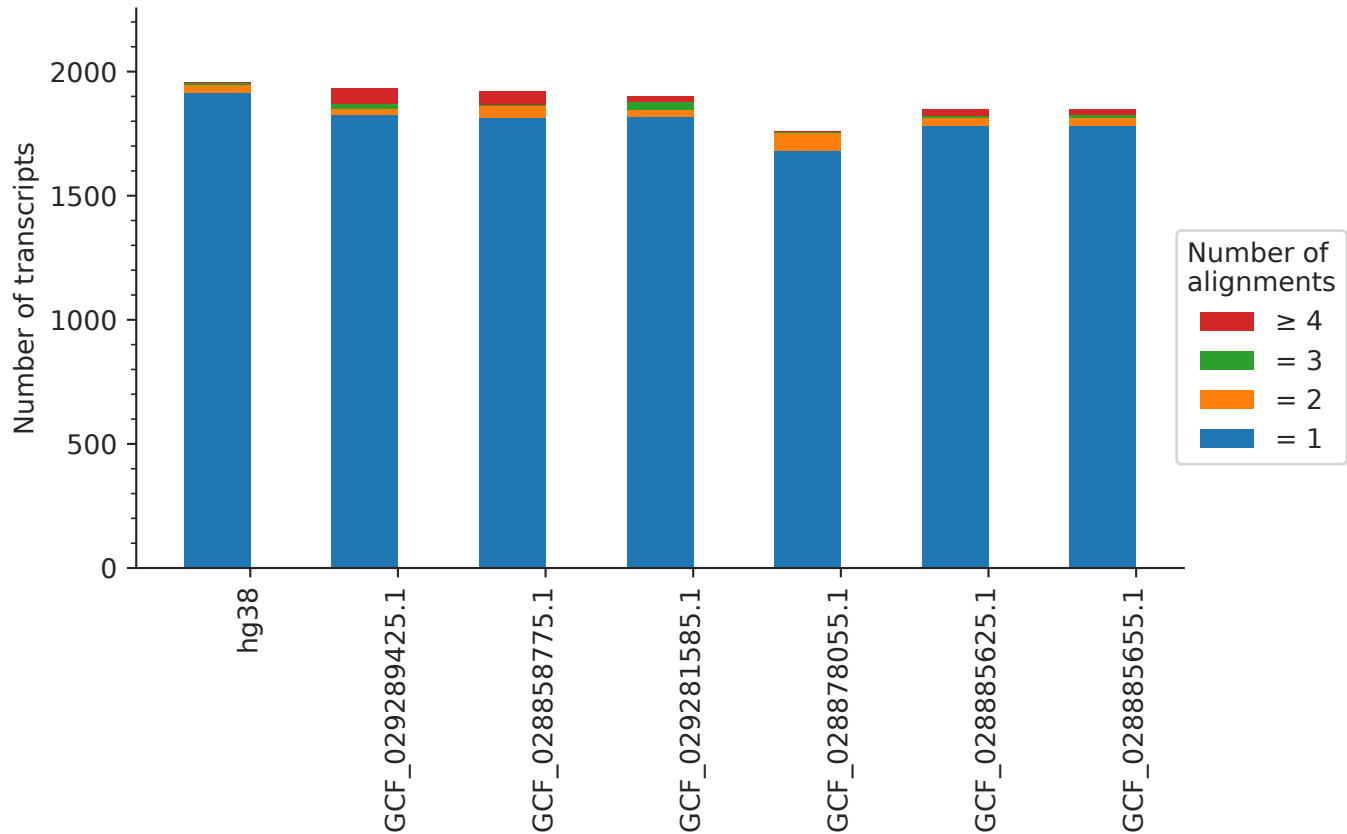
Proportion of TR_V_pseudogene transcripts that have multiple alignments



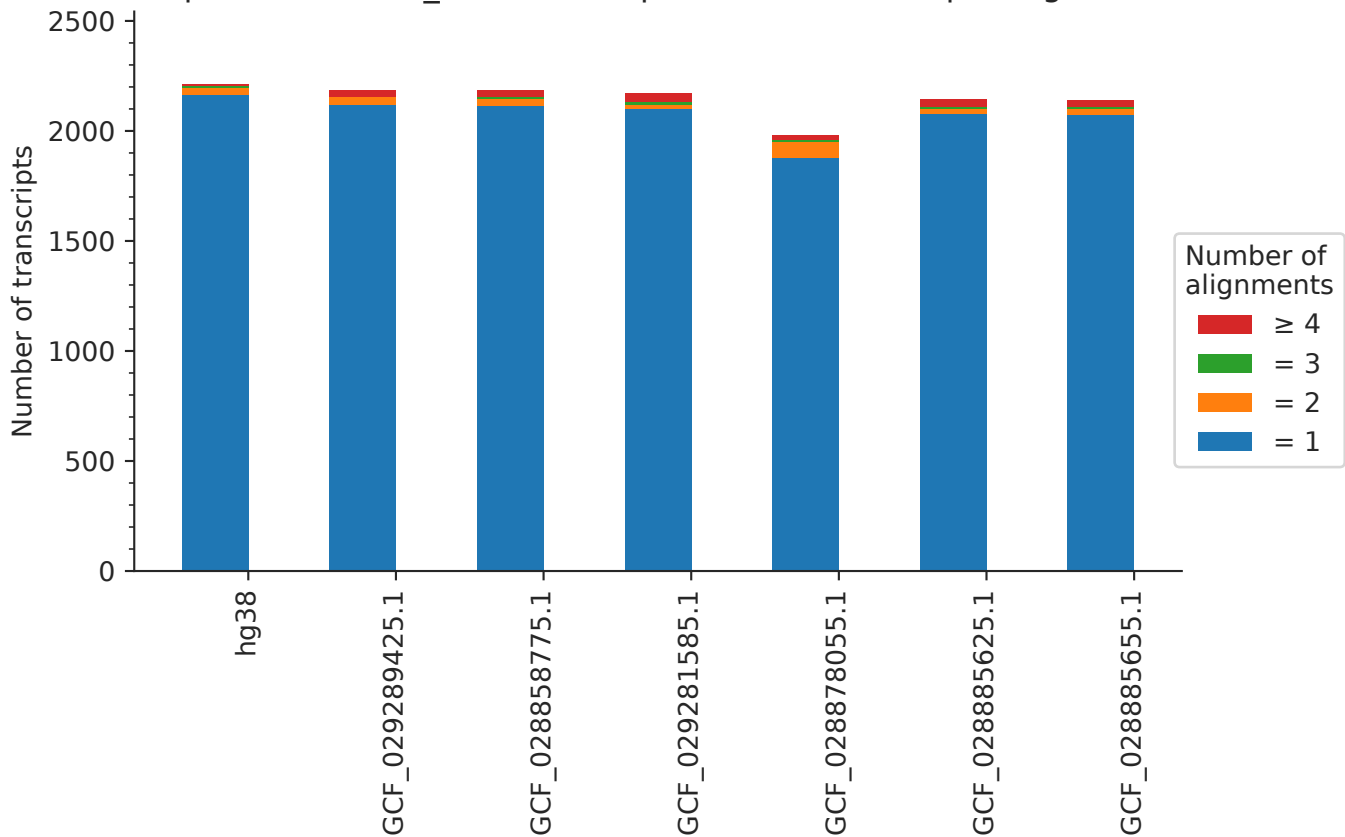
Proportion of lncRNA transcripts that have multiple alignments



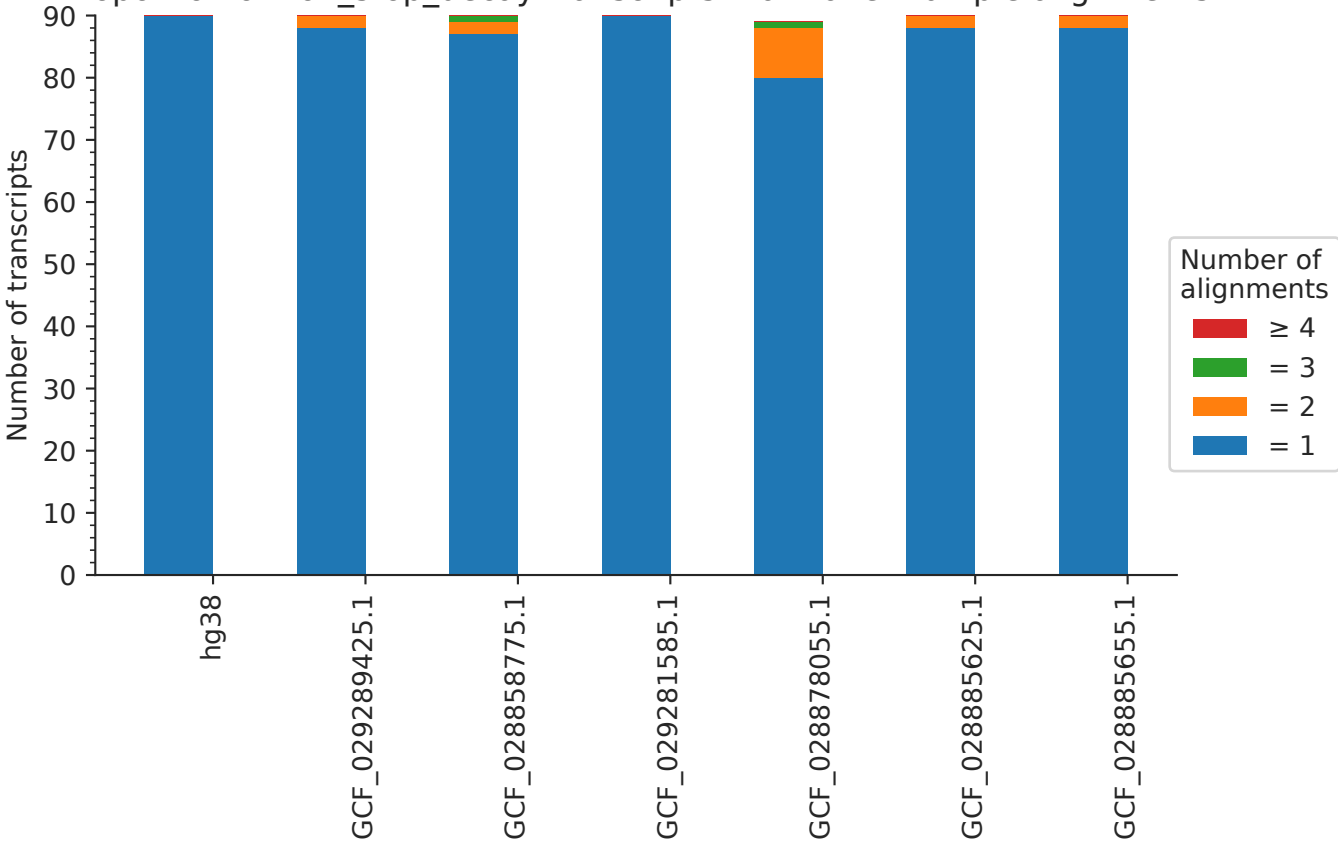
Proportion of miRNA transcripts that have multiple alignments



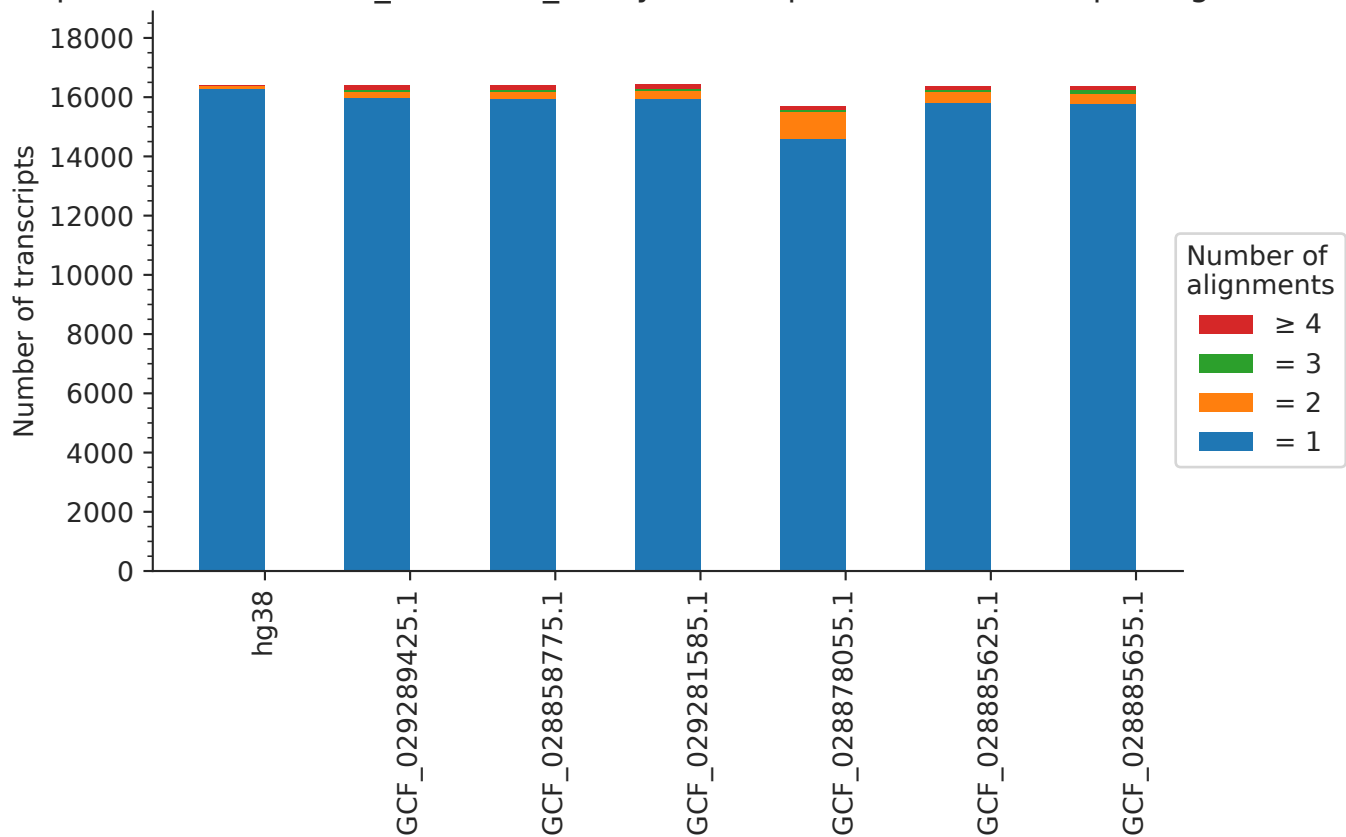
Proportion of misc_RNA transcripts that have multiple alignments



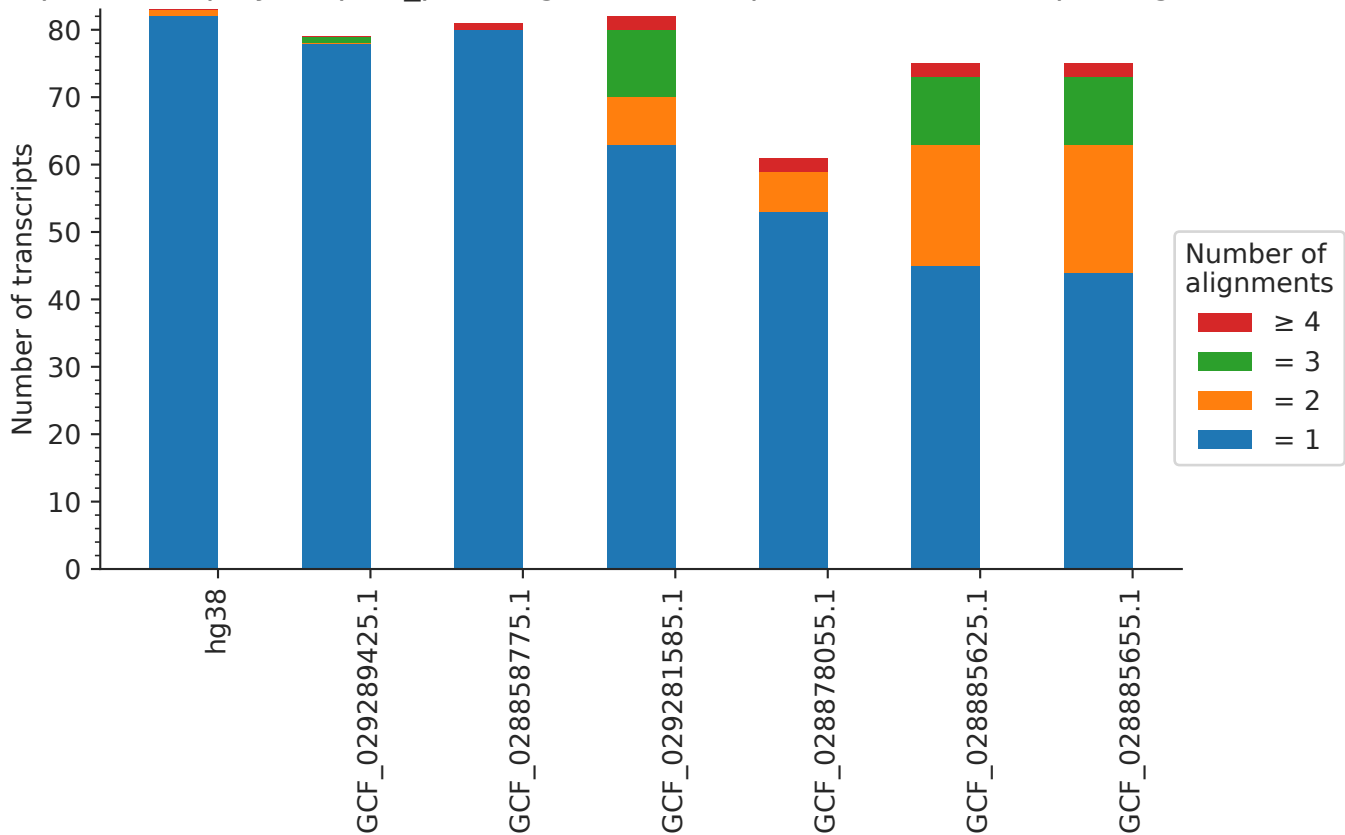
Proportion of non_stop_decay transcripts that have multiple alignments



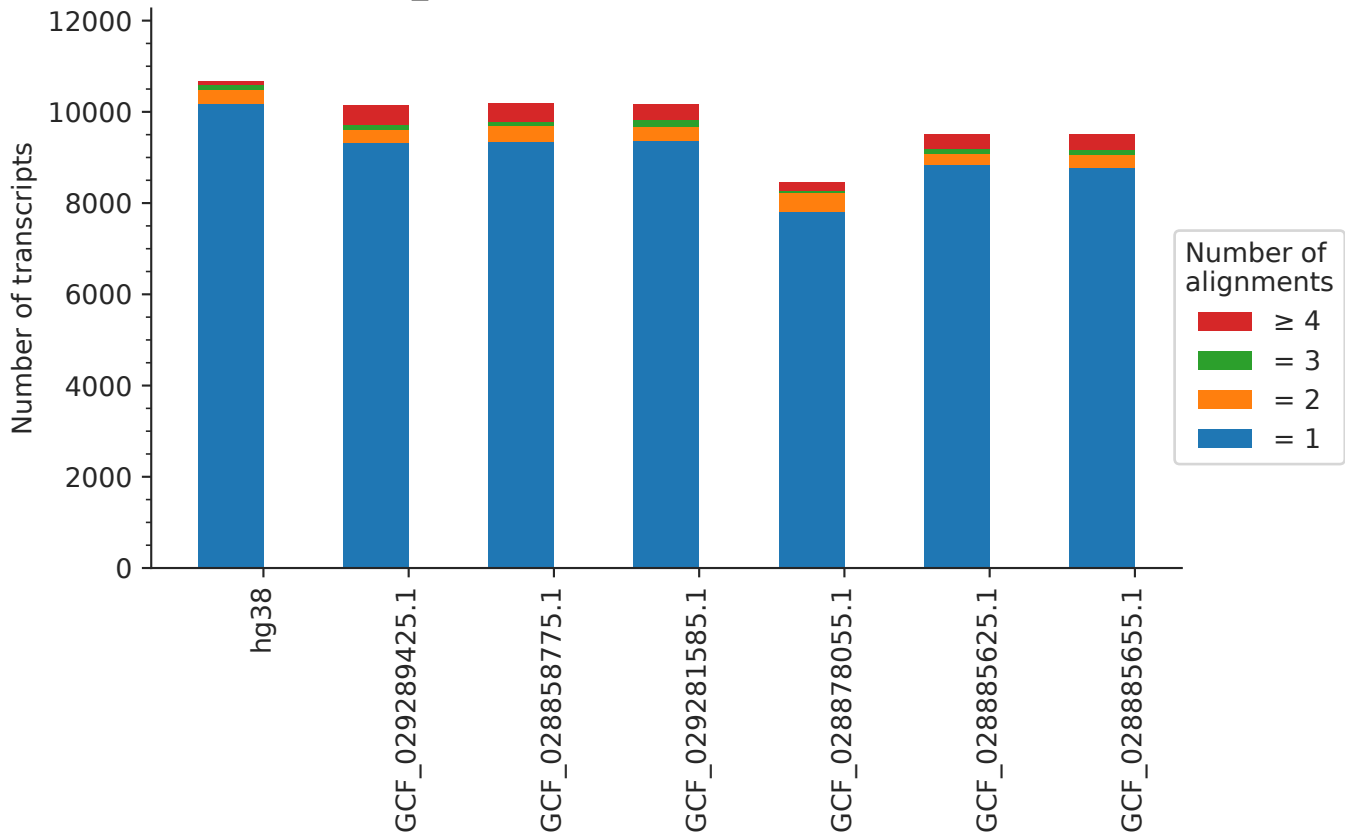
Proportion of nonsense_mediated_decay transcripts that have multiple alignments



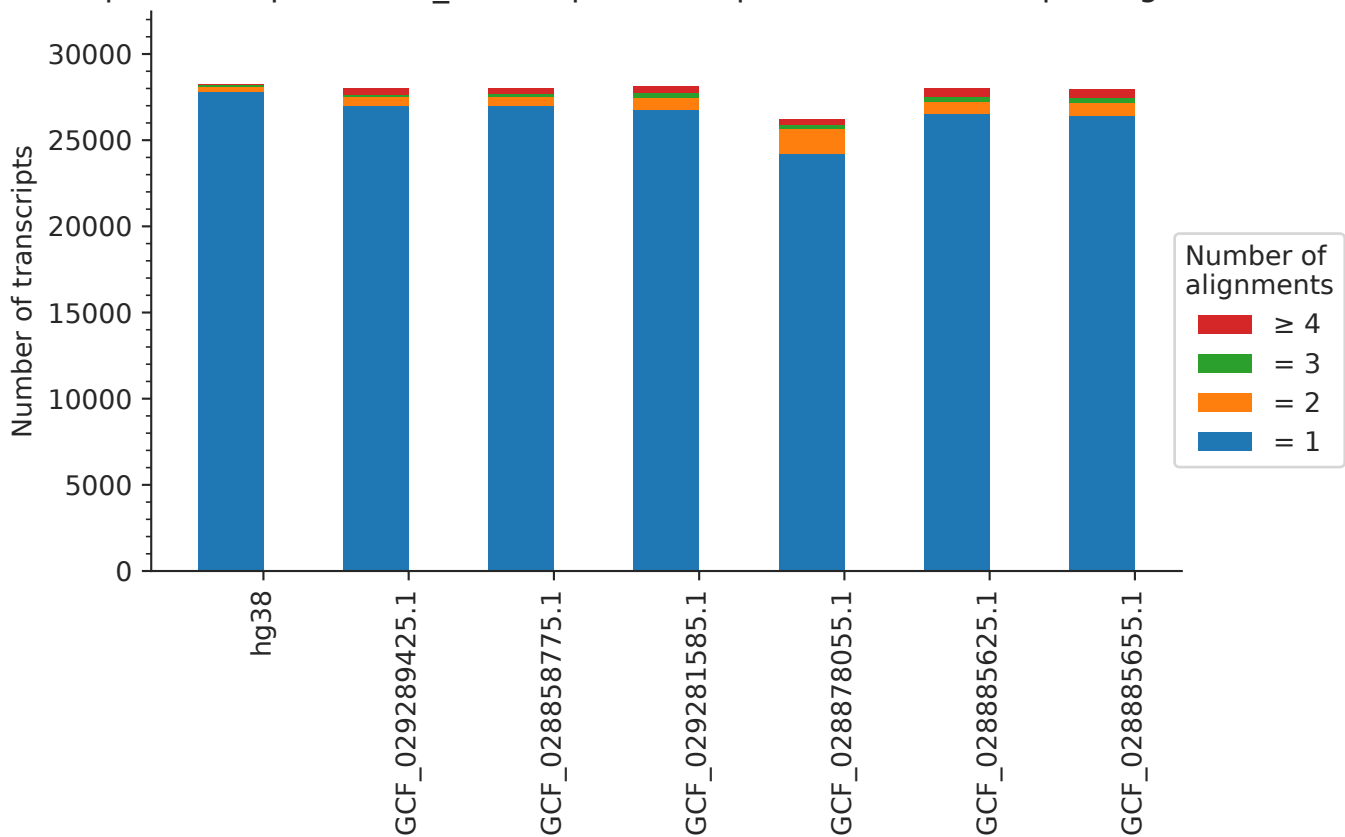
Proportion of polymorphic_pseudogene transcripts that have multiple alignments



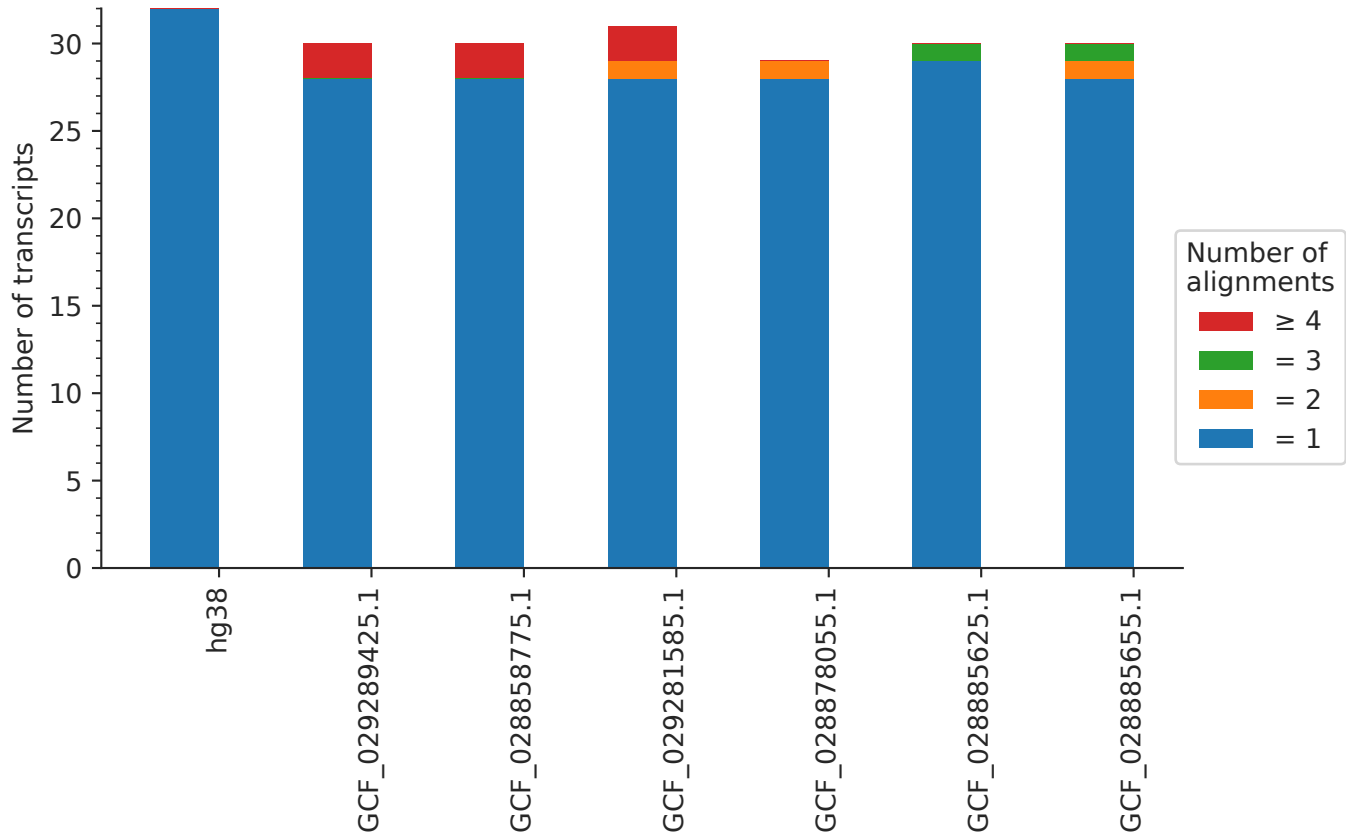
Proportion of processed_pseudogene transcripts that have multiple alignments



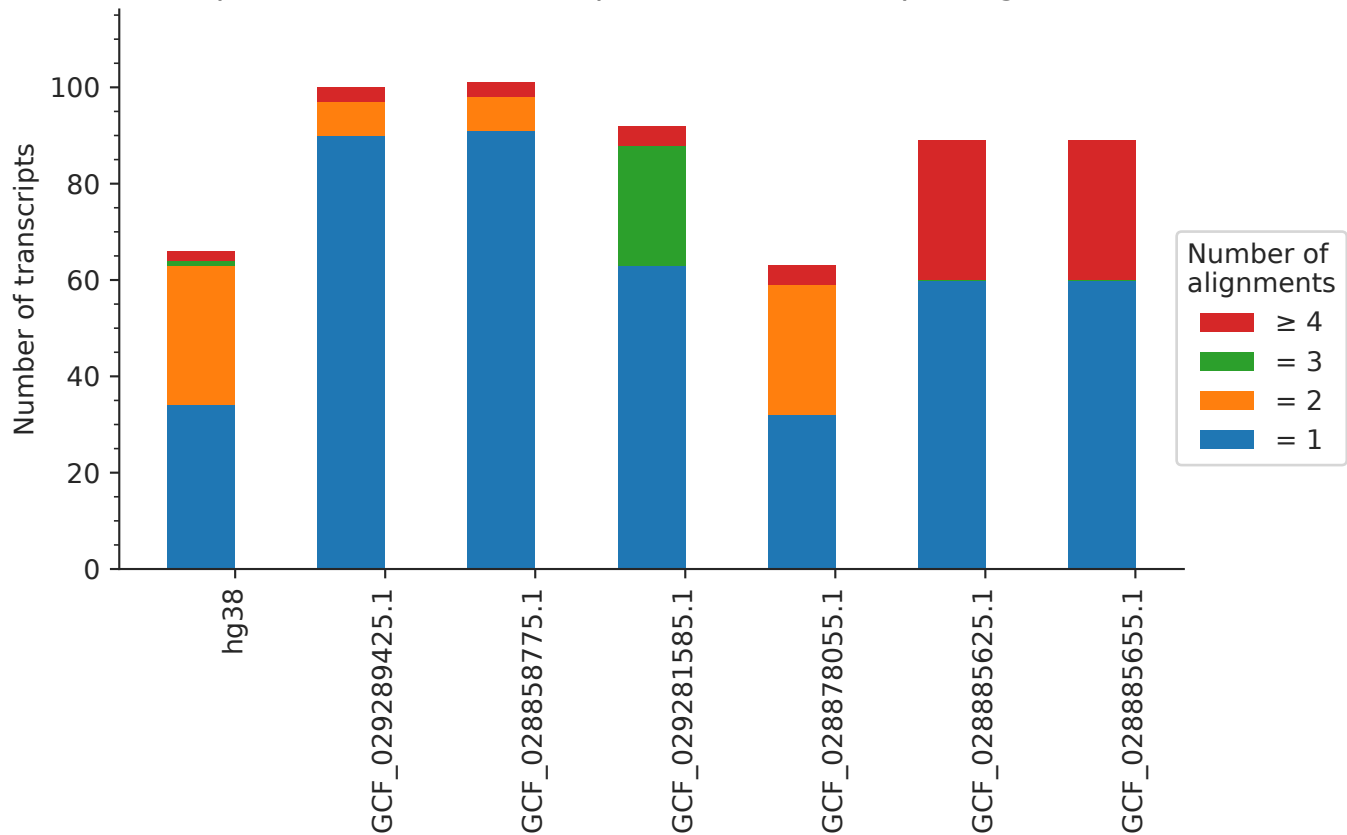
Proportion of processed_transcript transcripts that have multiple alignments



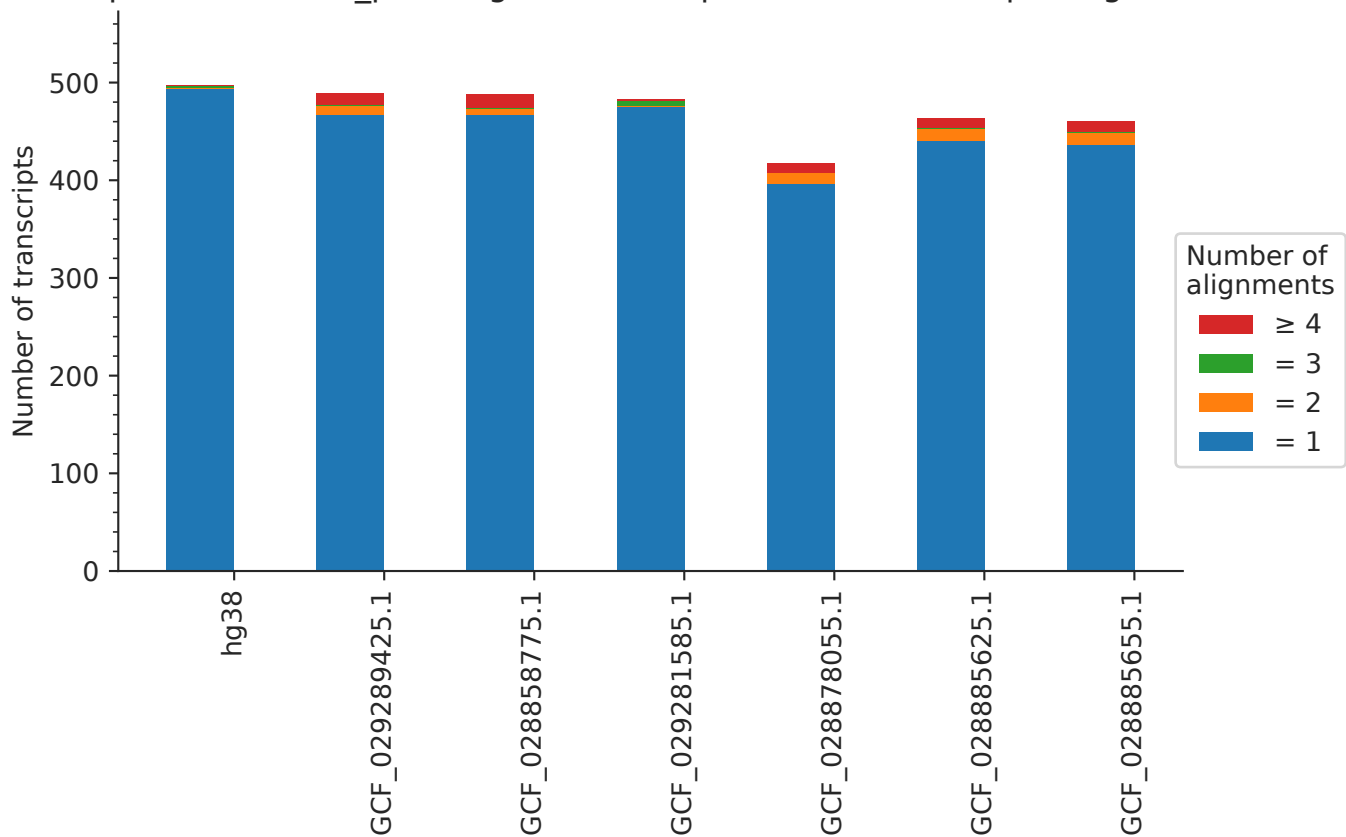
Proportion of pseudogene transcripts that have multiple alignments



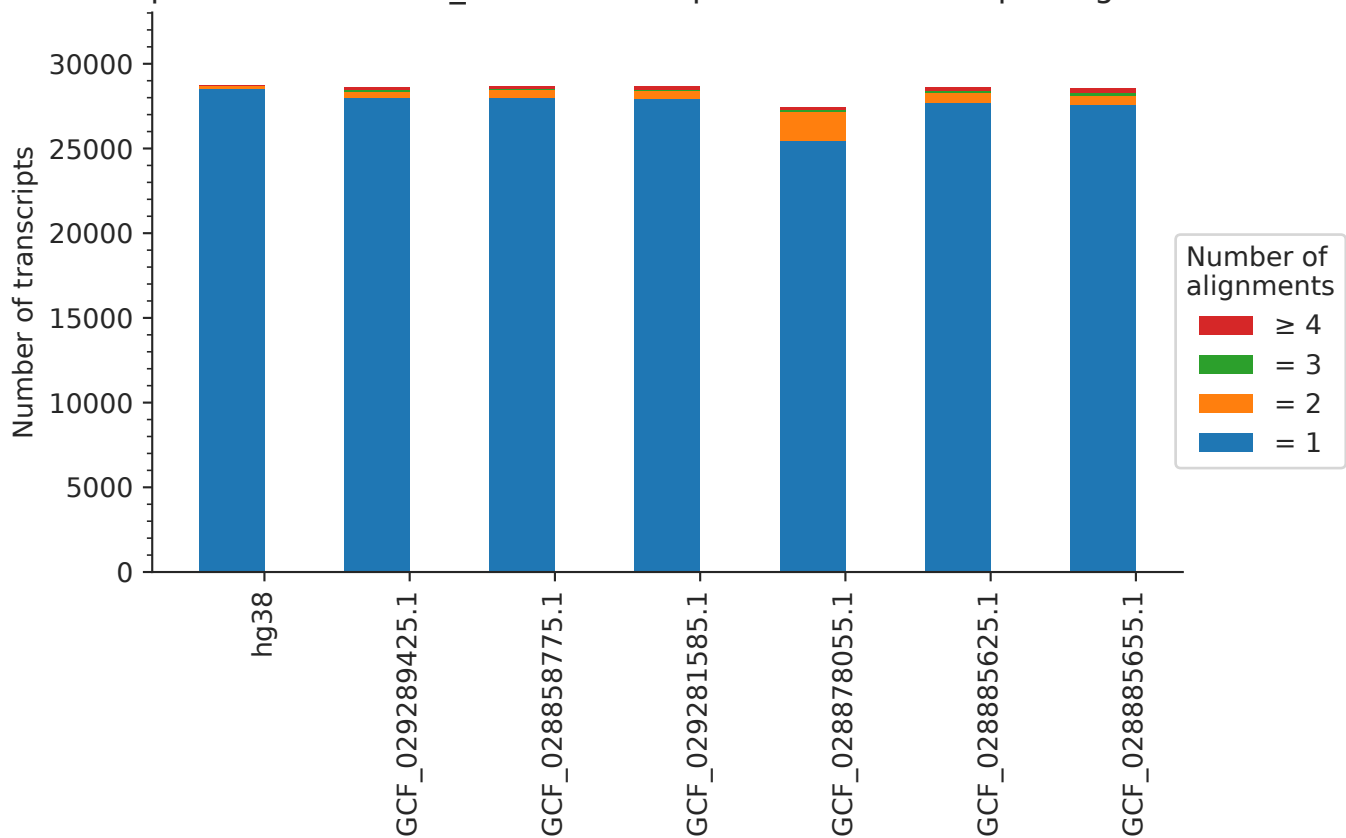
Proportion of rRNA transcripts that have multiple alignments



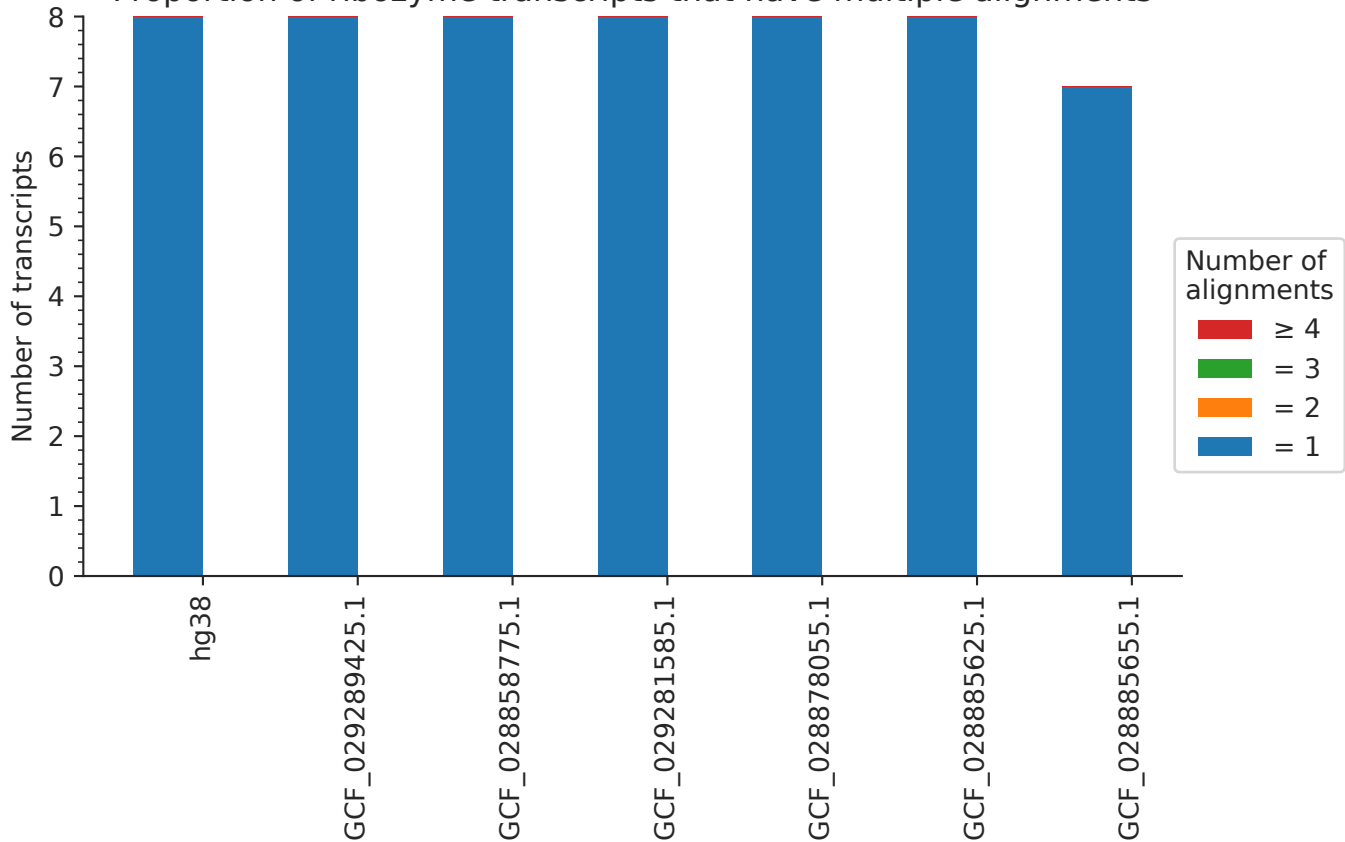
Proportion of rRNA_pseudogene transcripts that have multiple alignments



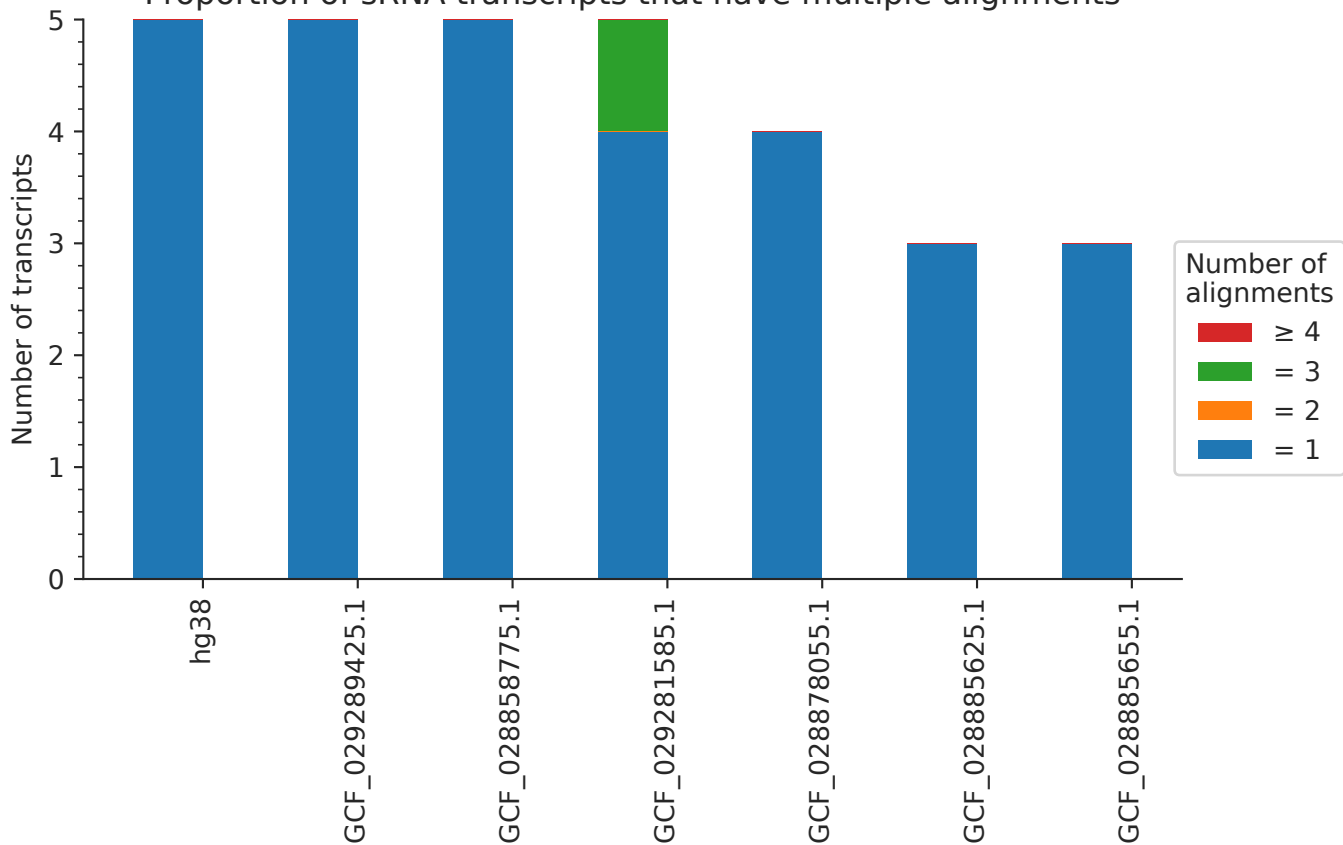
Proportion of retained_intron transcripts that have multiple alignments



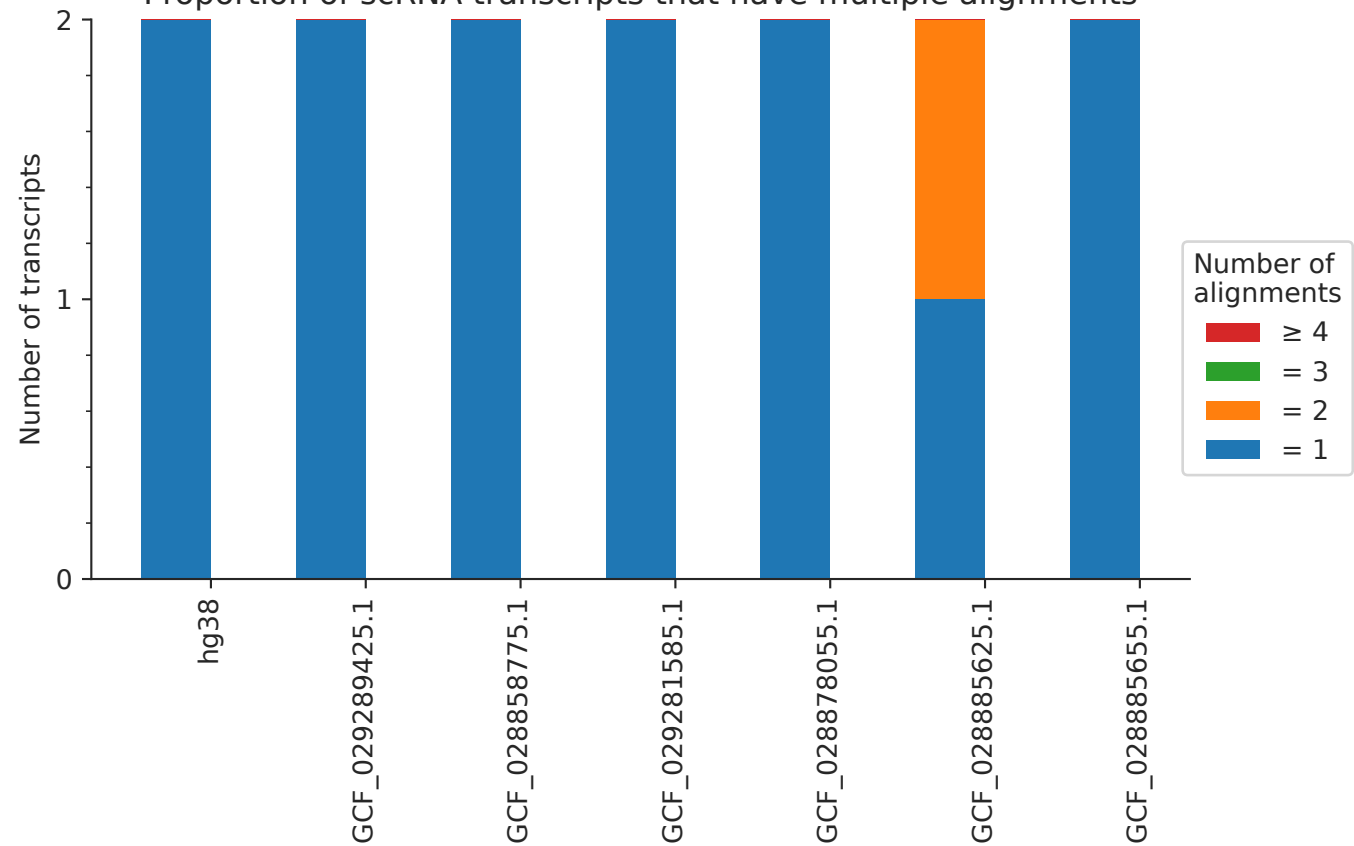
Proportion of ribozyme transcripts that have multiple alignments



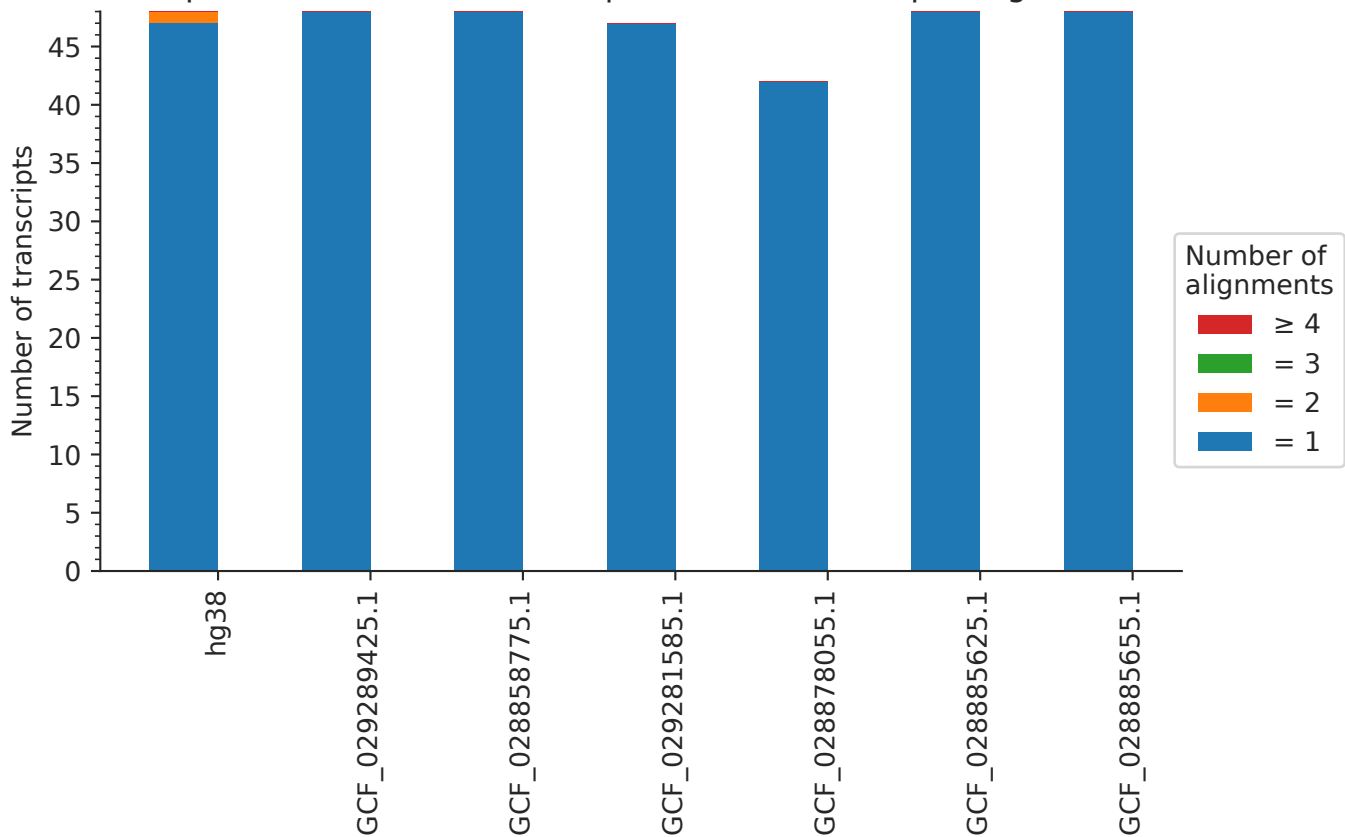
Proportion of sRNA transcripts that have multiple alignments



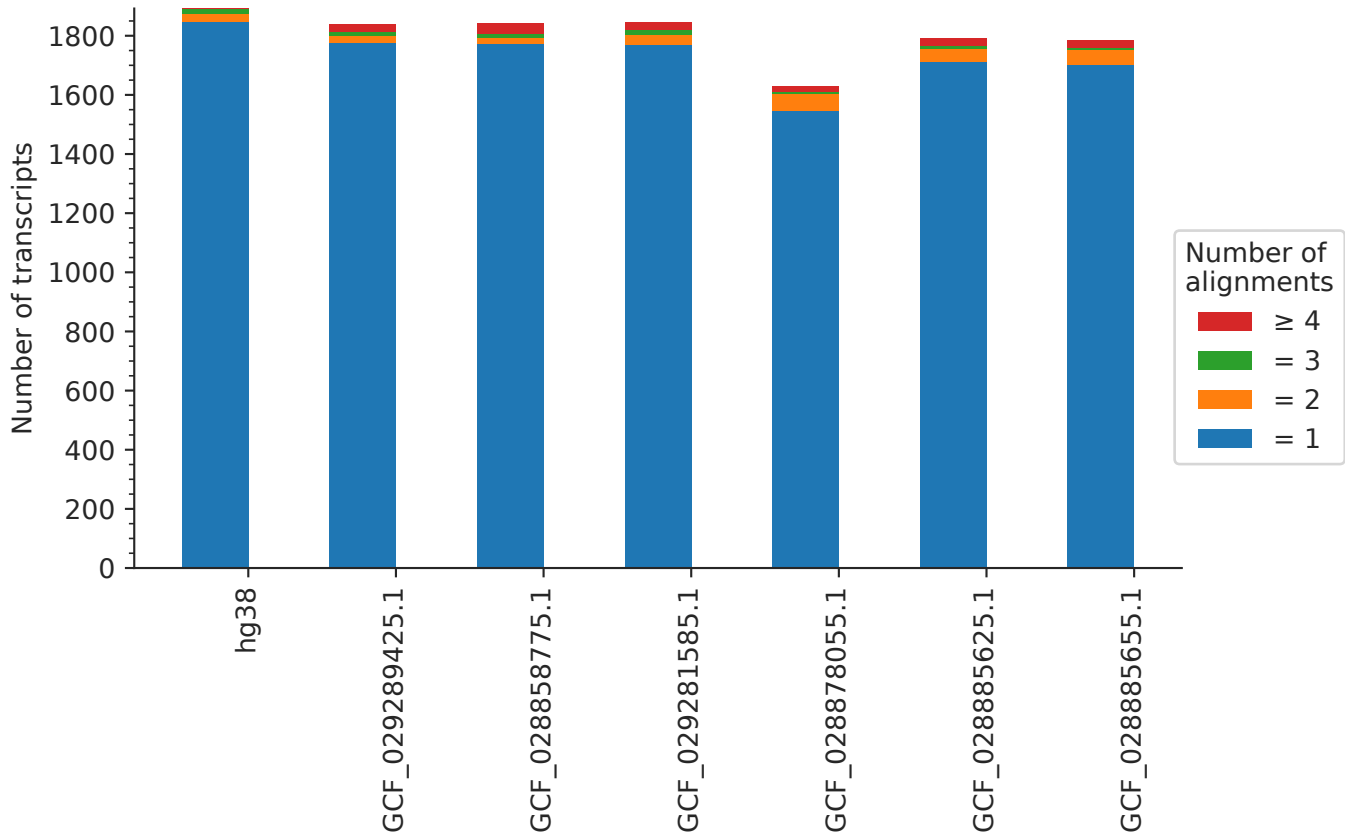
Proportion of scRNA transcripts that have multiple alignments



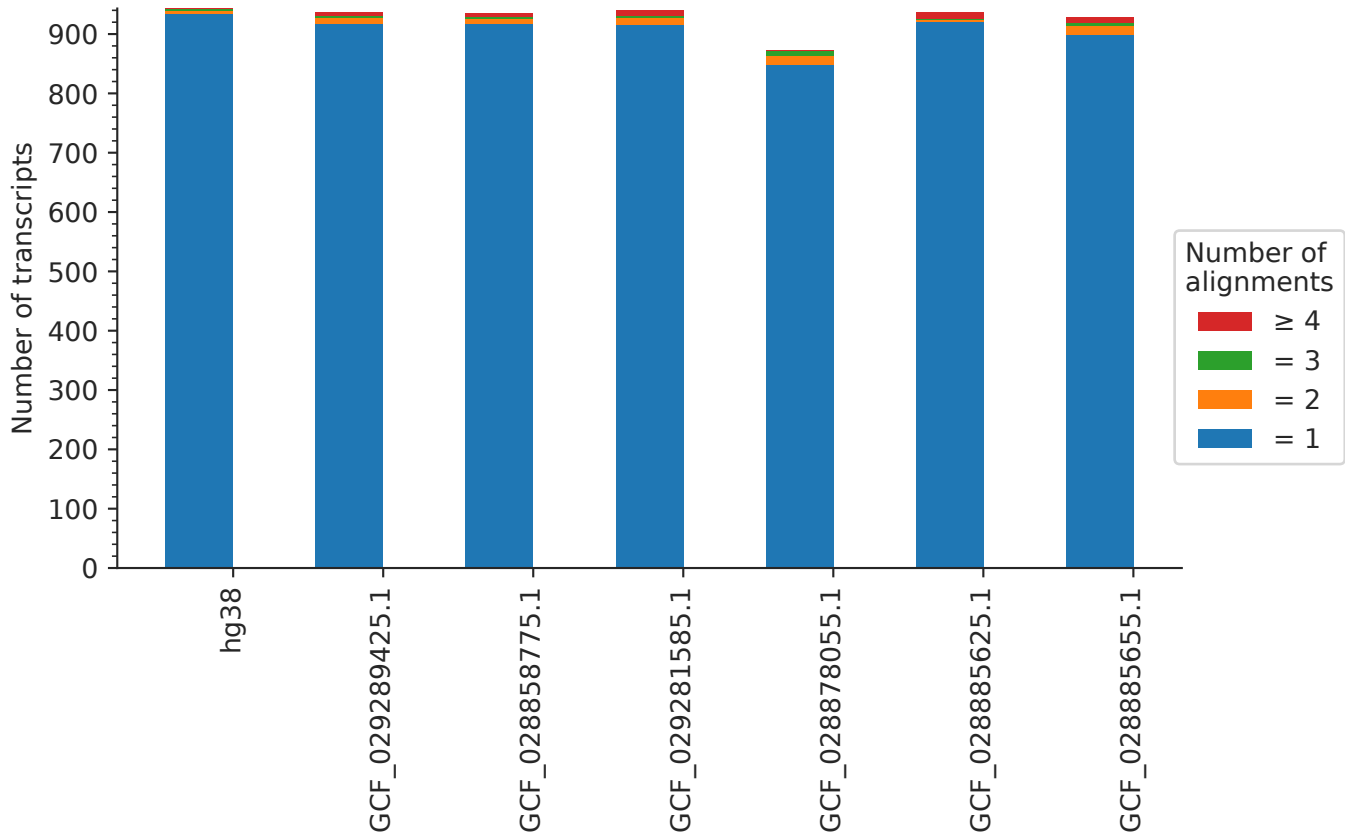
Proportion of scaRNA transcripts that have multiple alignments



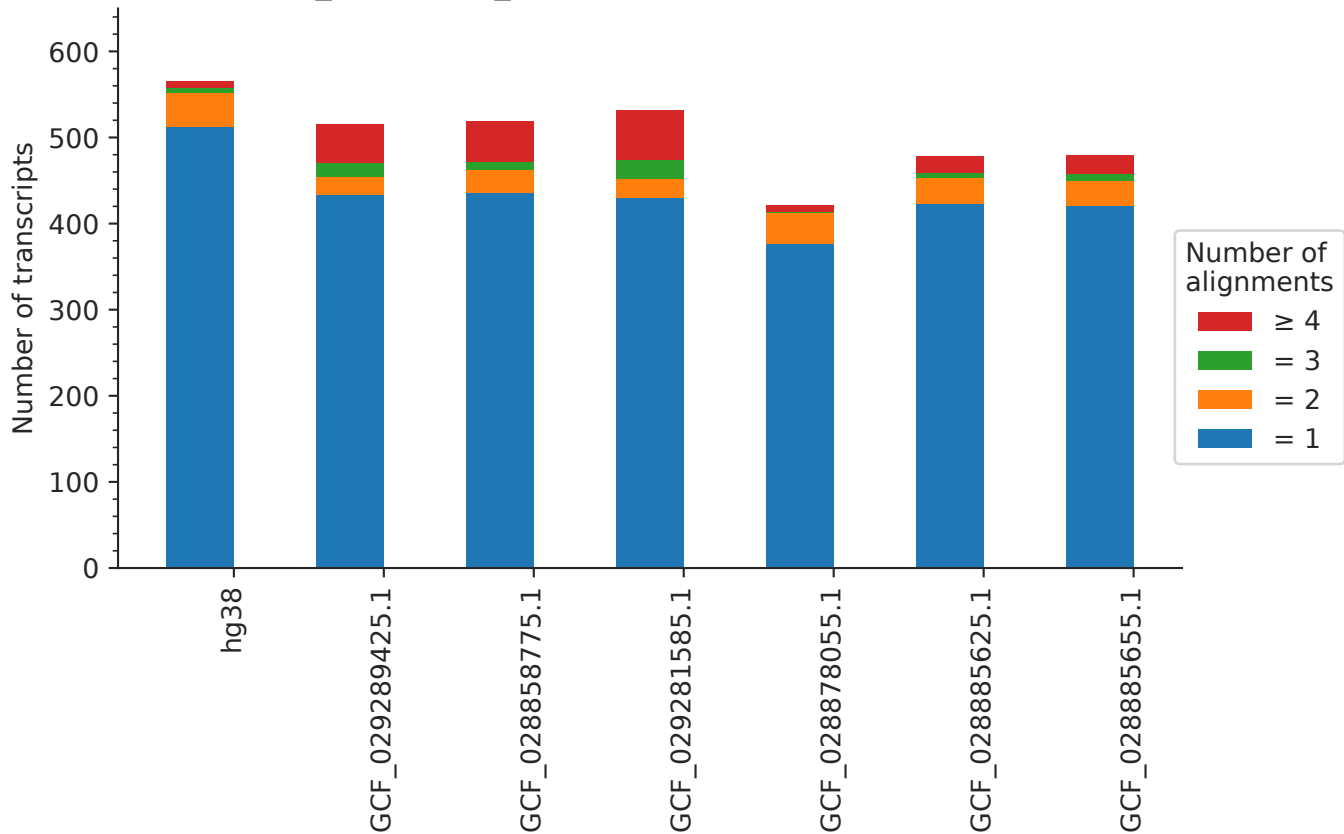
Proportion of snRNA transcripts that have multiple alignments



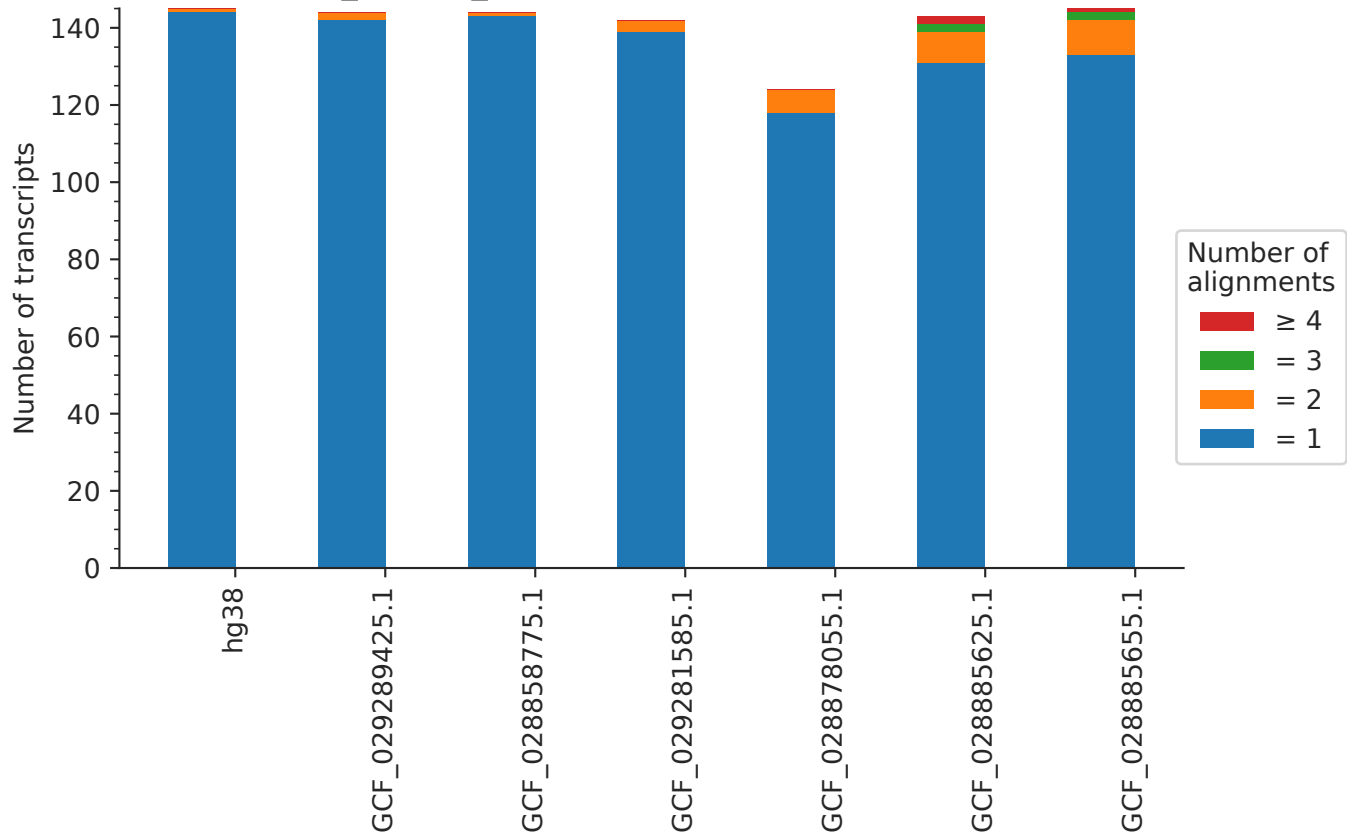
Proportion of snoRNA transcripts that have multiple alignments



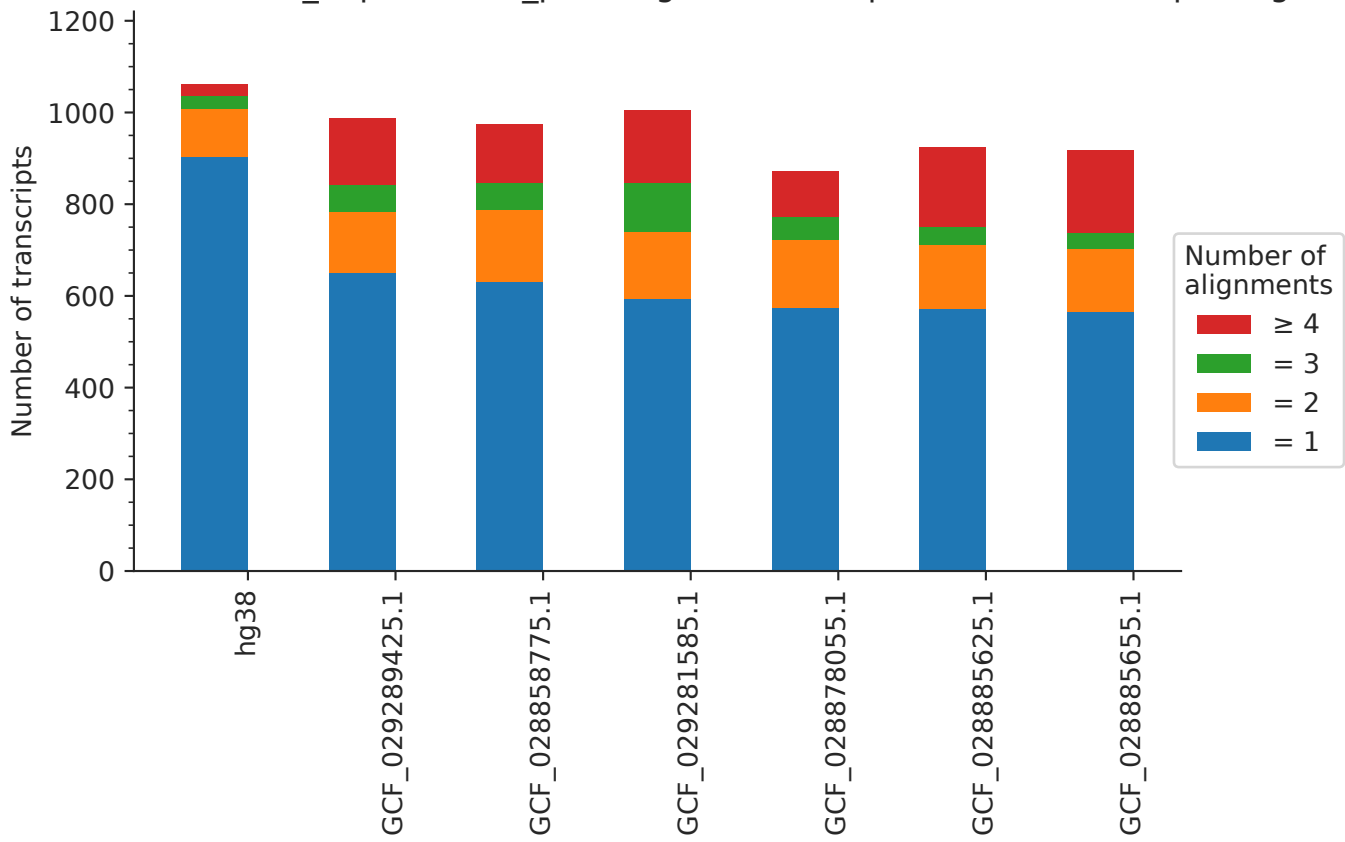
Proportion of transcribed_processed_pseudogene transcripts that have multiple alignments



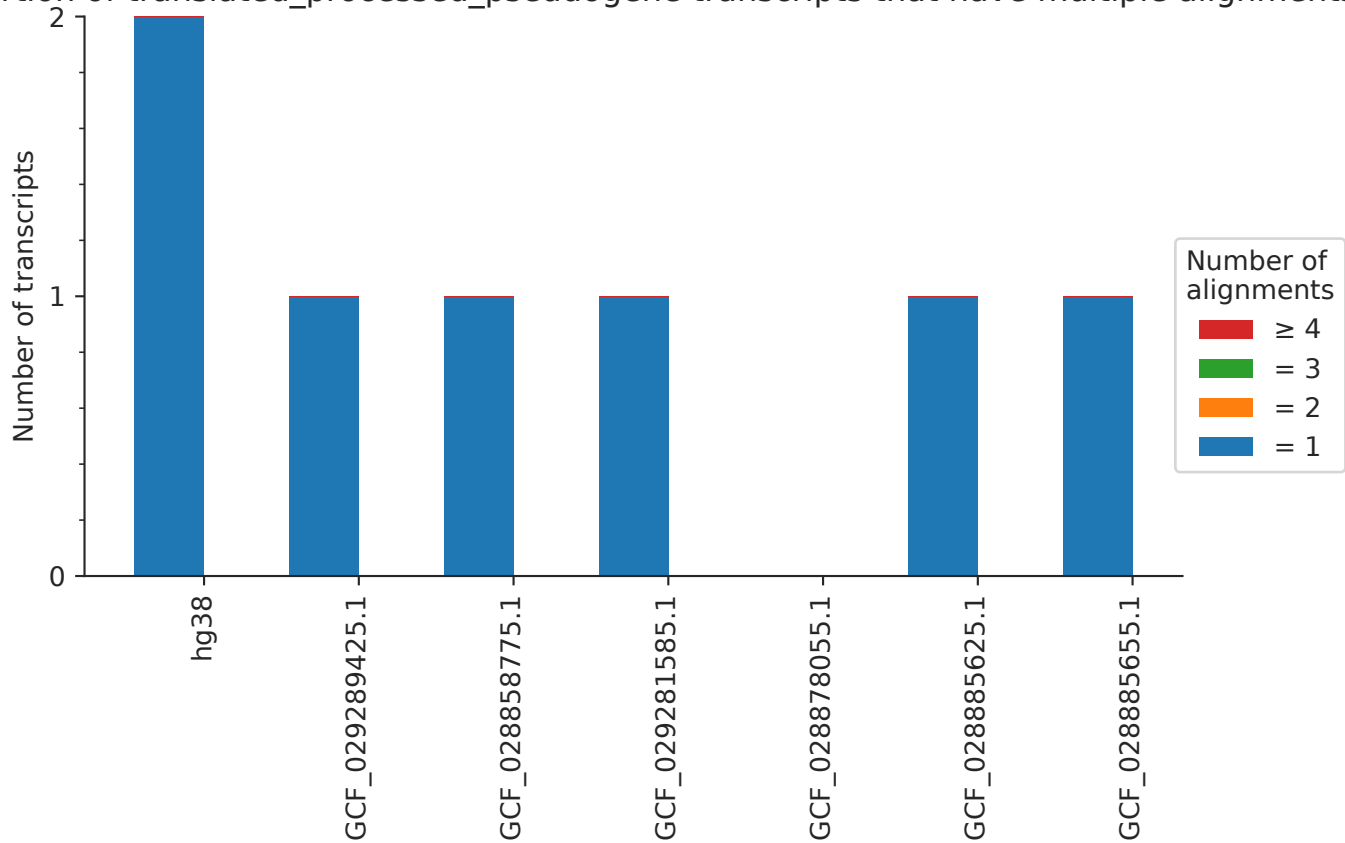
Proportion of transcribed_unitary_pseudogene transcripts that have multiple alignments



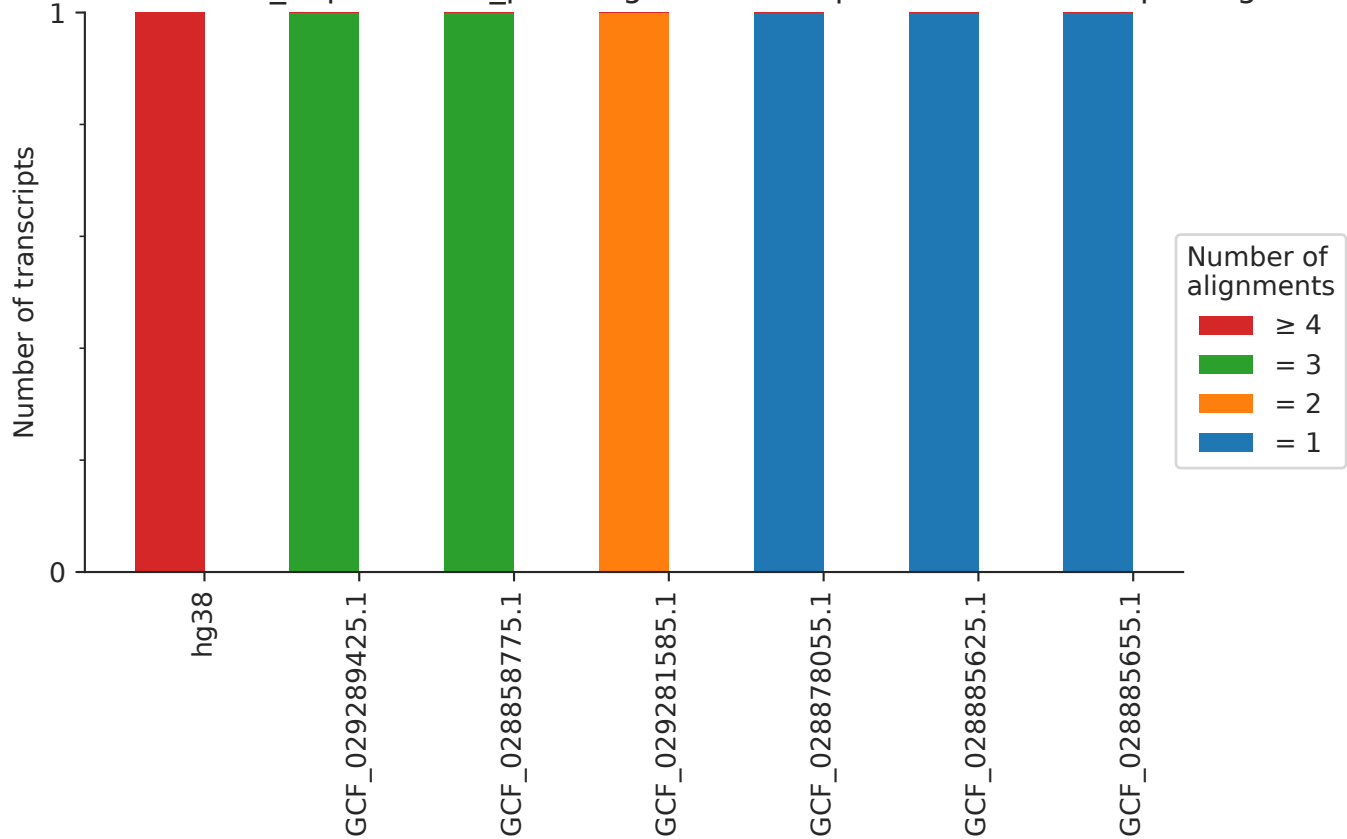
Proportion of transcribed_unprocessed_pseudogene transcripts that have multiple alignments



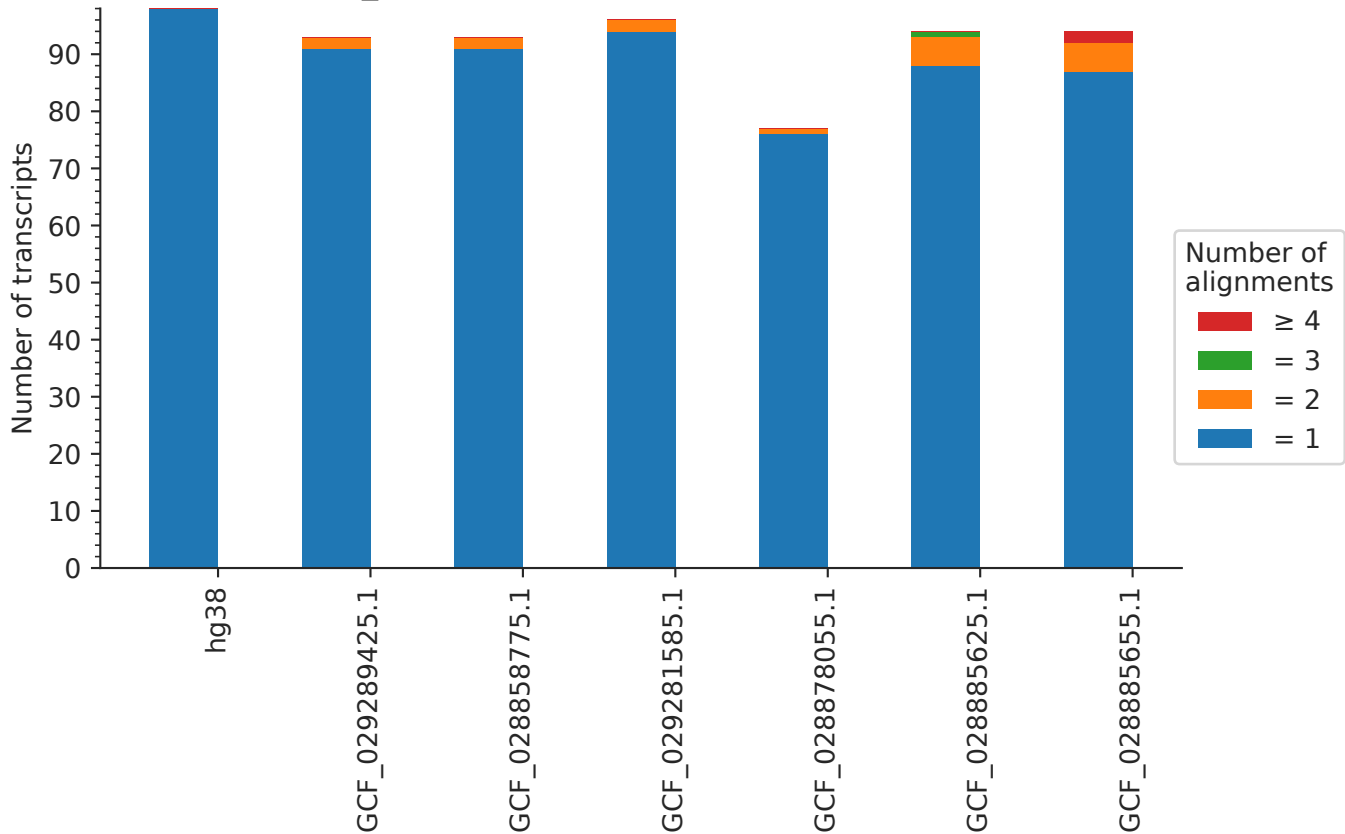
Proportion of translated_processed_pseudogene transcripts that have multiple alignments



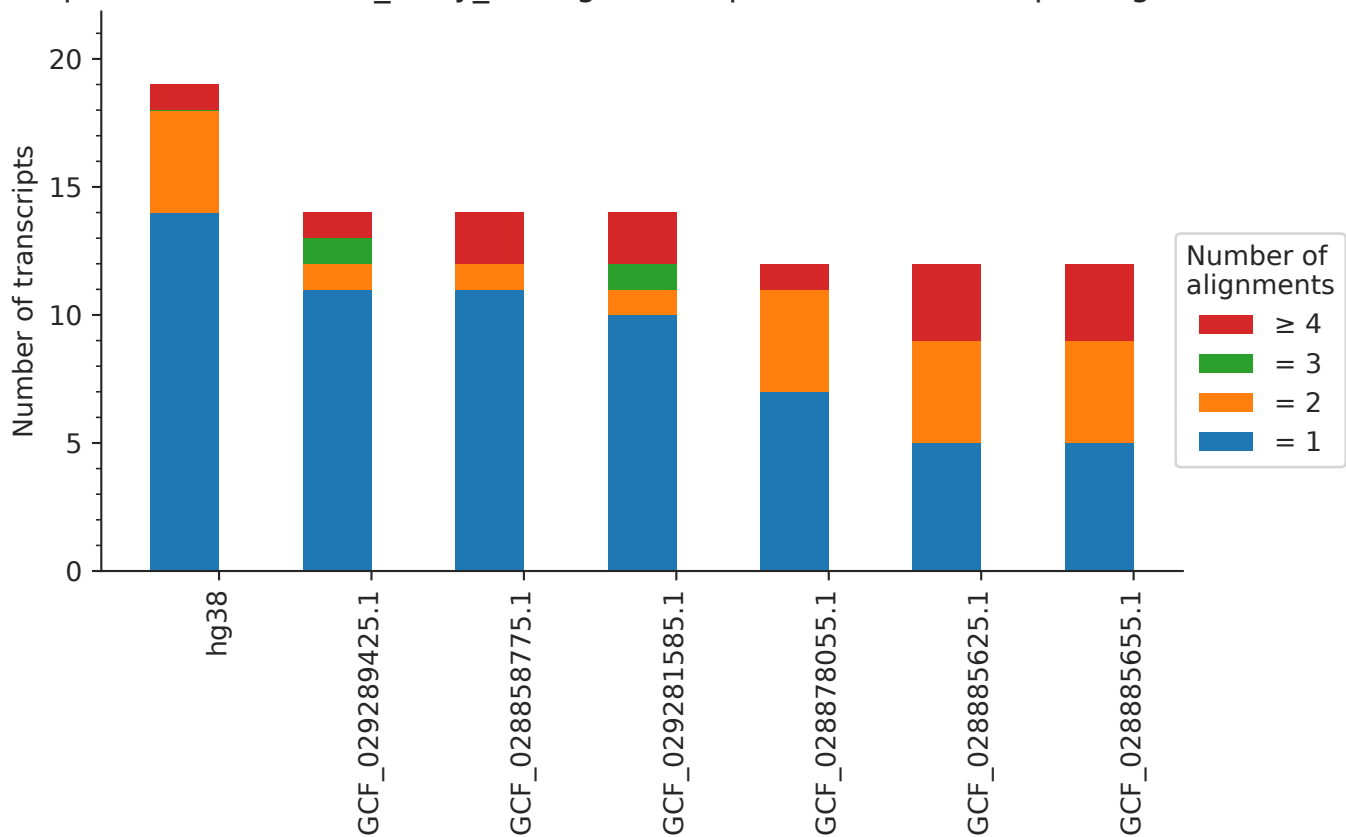
Proportion of translated_unprocessed_pseudogene transcripts that have multiple alignments



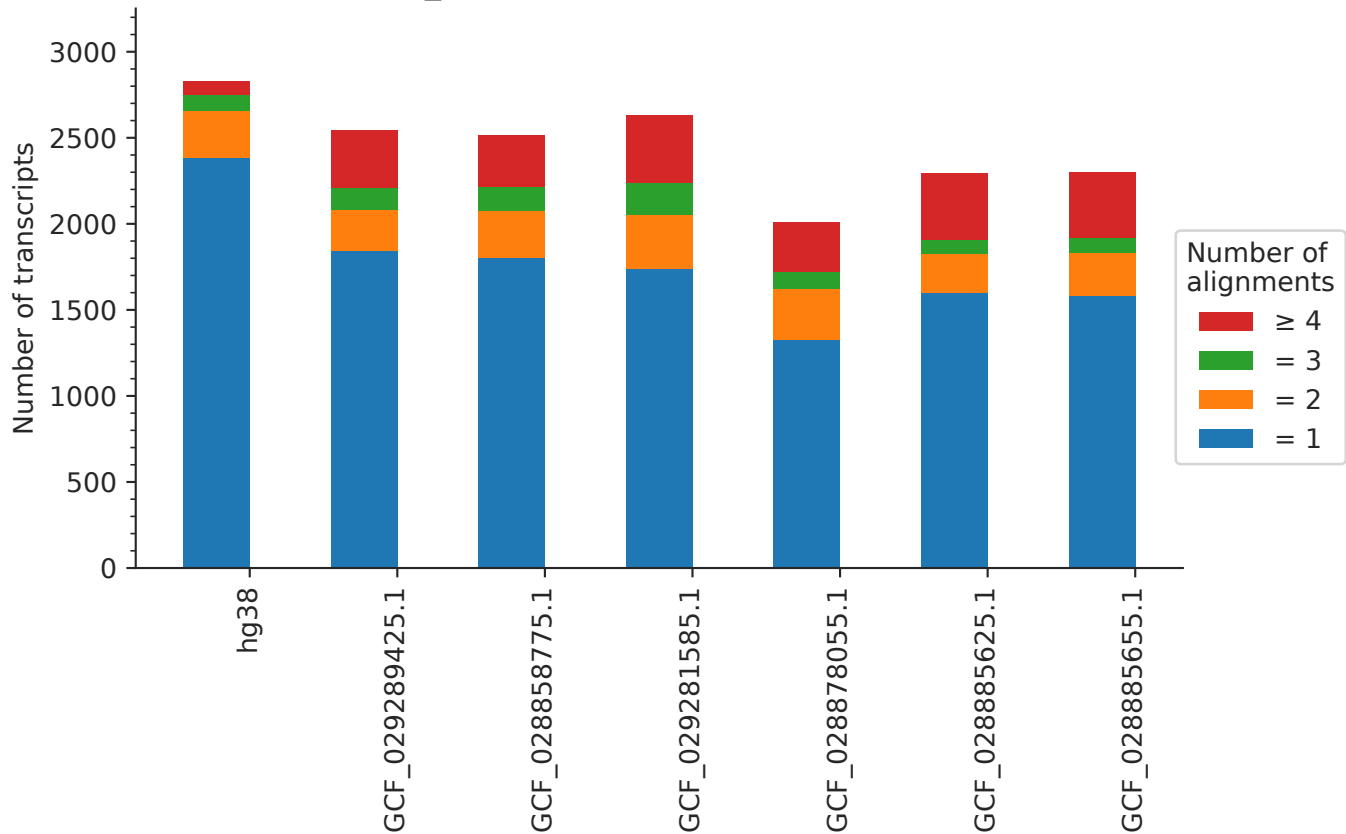
Proportion of unitary_pseudogene transcripts that have multiple alignments



Proportion of unknown_likely_coding transcripts that have multiple alignments



Proportion of unprocessed_pseudogene transcripts that have multiple alignments



Proportion of vault_RNA transcripts that have multiple alignments

