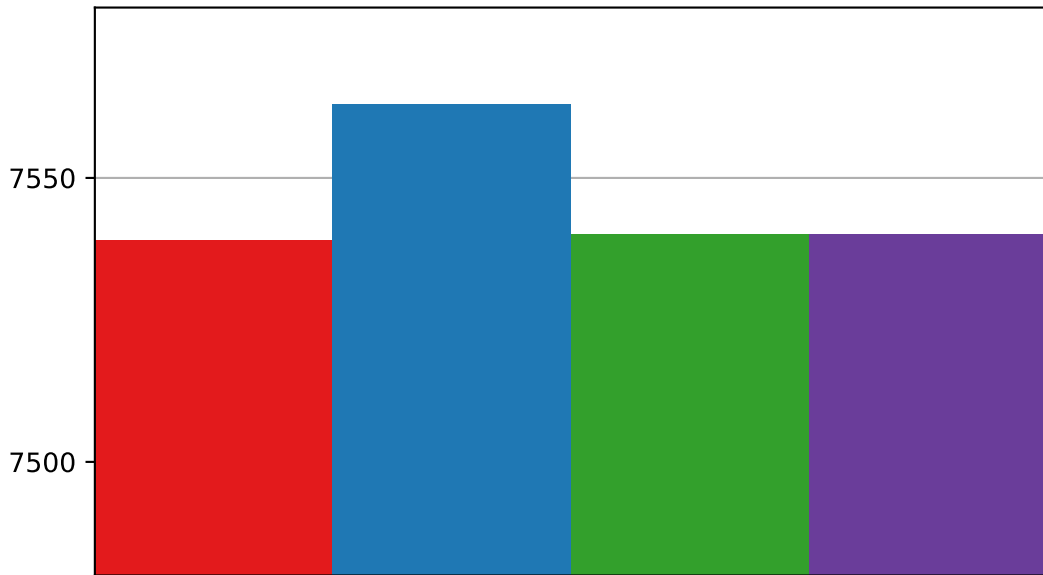


complete genomic features



spades_default

megahit_default

megahit_min_count_3

spades_kmers_careful