VEP: Variant Consequences Consequences (all), 6 samples transcript_ablation splice_acceptor_variant HBRRep1.haplotypecaller.filtered_VEP.ann splice_donor_variant stop_gained frameshift_variant inframe_insertion missense_variant HBRRep2.haplotypecaller.filtered_VEP.ann splice_donor_5th_base_variant splice_region_variant splice_donor_region_variant splice_polypyrimidine_tract_variant stop_retained_variant HBRRep3.haplotypecaller.filtered_VEP.ann synonymous_variant coding_sequence_variant mature_miRNA_variant 5_prime_UTR_variant UHRRep1.haplotypecaller.filtered_VEP.ann 3_prime_UTR_variant non_coding_transcript_exon_variant intron_variant NMD_transcript_variant non_coding_transcript_variant UHRRep2.haplotypecaller.filtered_VEP.ann upstream_gene_variant downstream_gene_variant regulatory_region_ablation regulatory_region_variant intergenic_variant UHRRep3.haplotypecaller.filtered_VEP.ann protein_altering_variant 2k 6k 4k 8k 10k Consequences (most severe)