VEP: Variant Consequences

Coding consequences, 6 samples

Legend:
- stop_gained
- frameshift_variant
- inframe_insertion
- missense_variant
- stop_retained_variant
- synonymous_variant
- coding_sequence_variant
- protein_altering_variant

Samples (top to bottom):
- HBRRep1.haplotypecaller.filtered_VEP.ann
- HBRRep2.haplotypecaller.filtered_VEP.ann
- HBRRep3.haplotypecaller.filtered_VEP.ann
- UHRRep1.haplotypecaller.filtered_VEP.ann
- UHRRep2.haplotypecaller.filtered_VEP.ann
- UHRRep3.haplotypecaller.filtered_VEP.ann

X-axis: Consequences (most severe) — 0%, 20%, 40%, 60%, 80%, 100%

Created with MultiQC