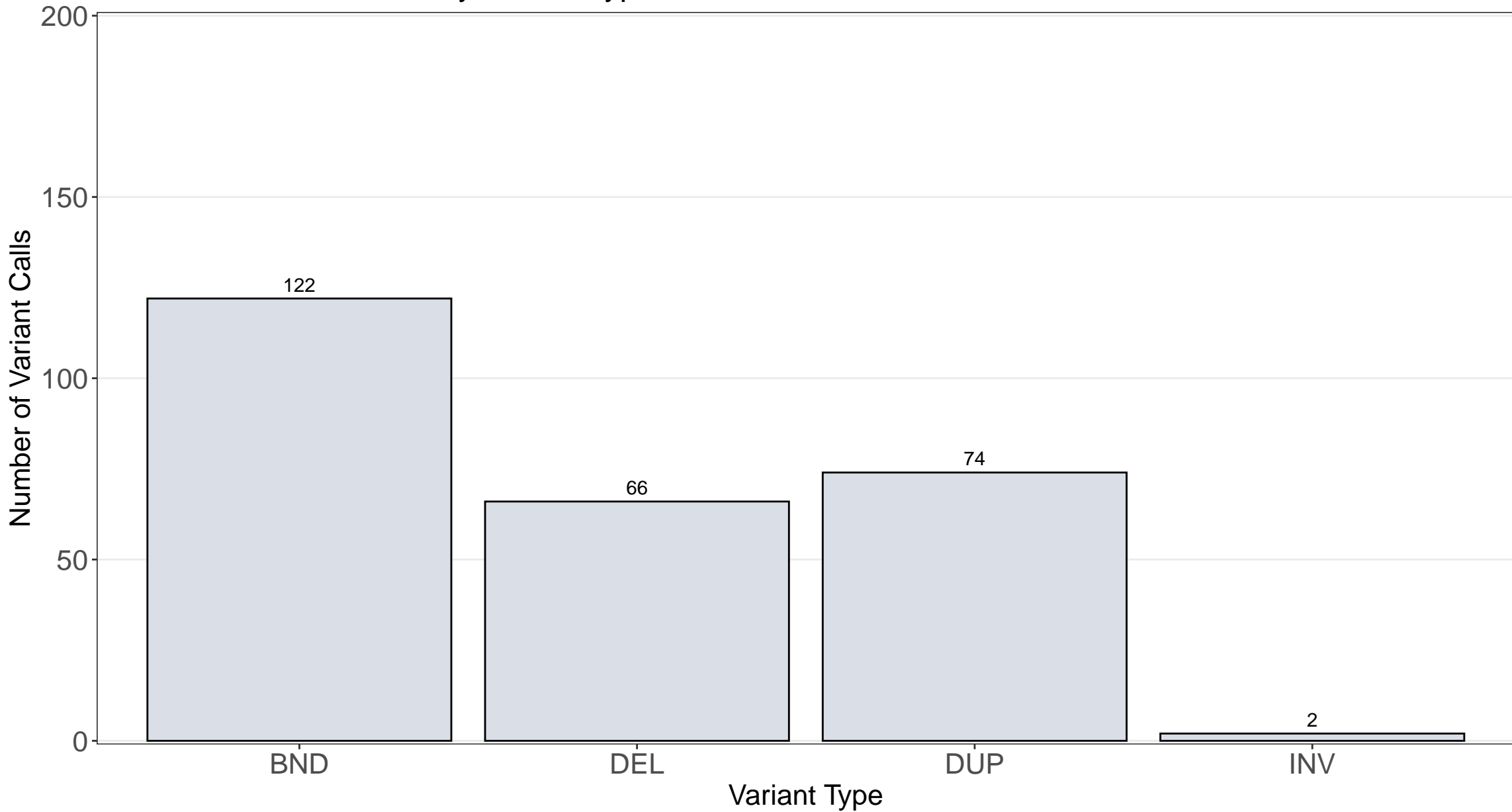
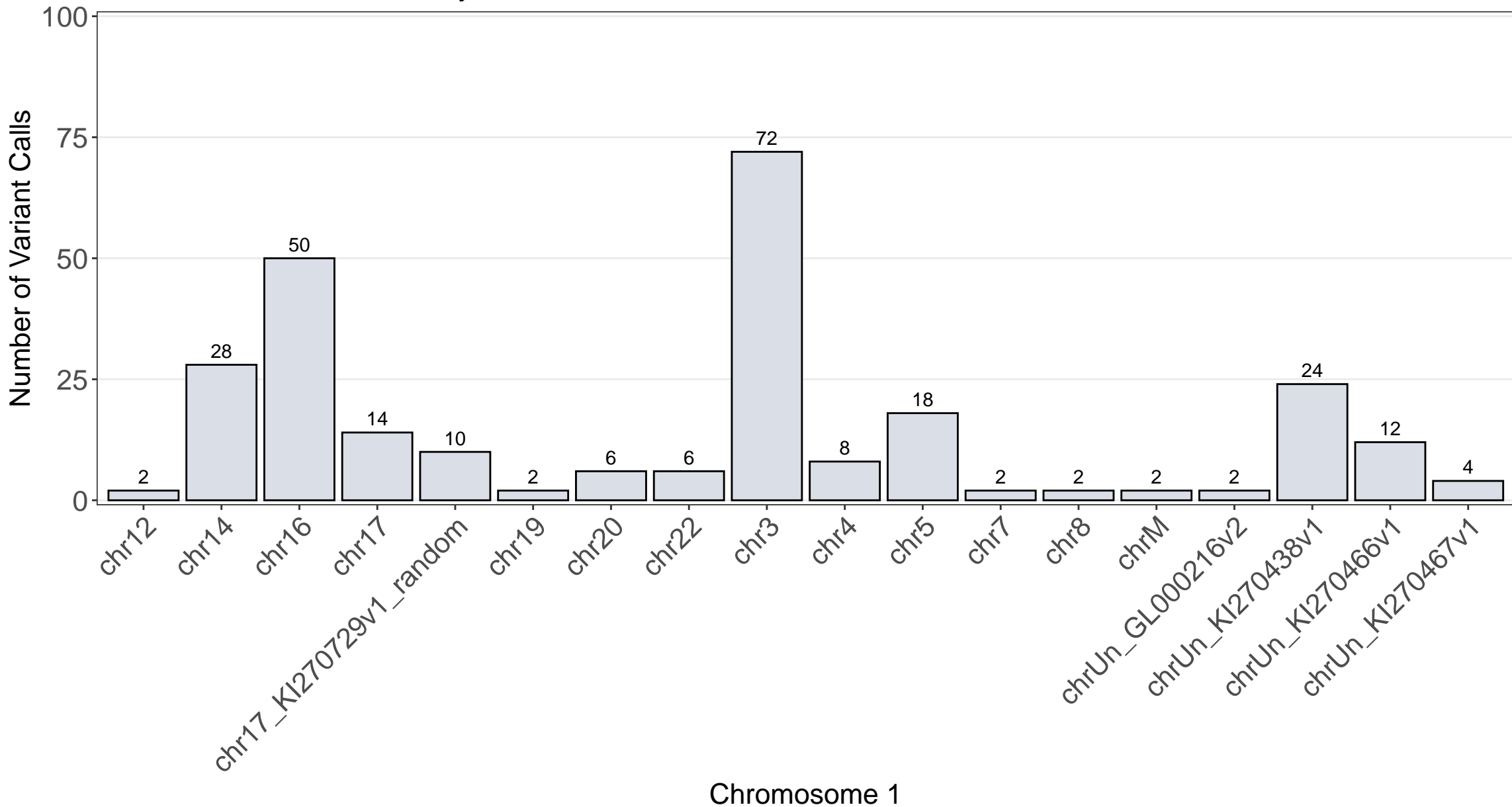


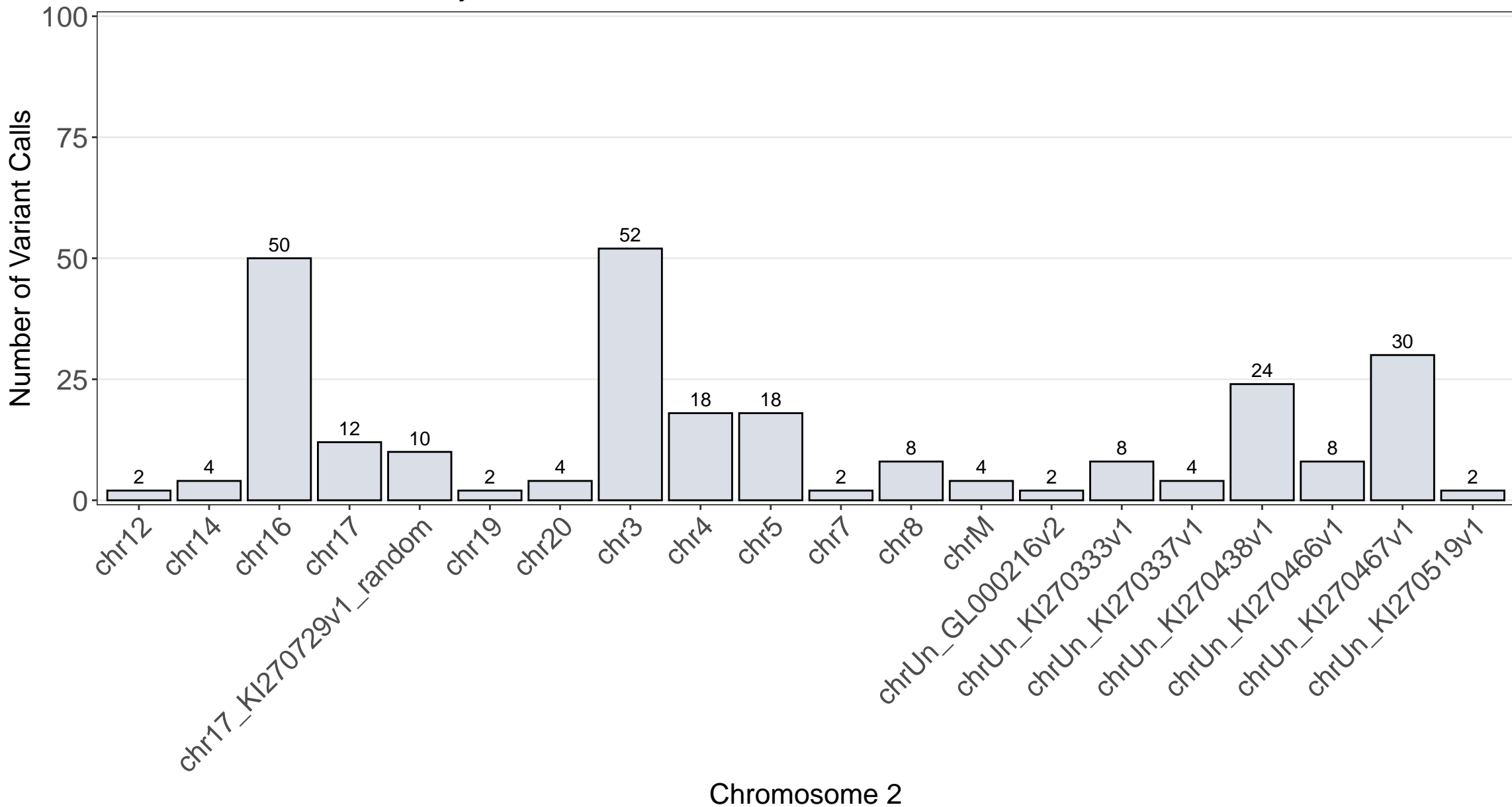
Number of Variant Calls by variant_type

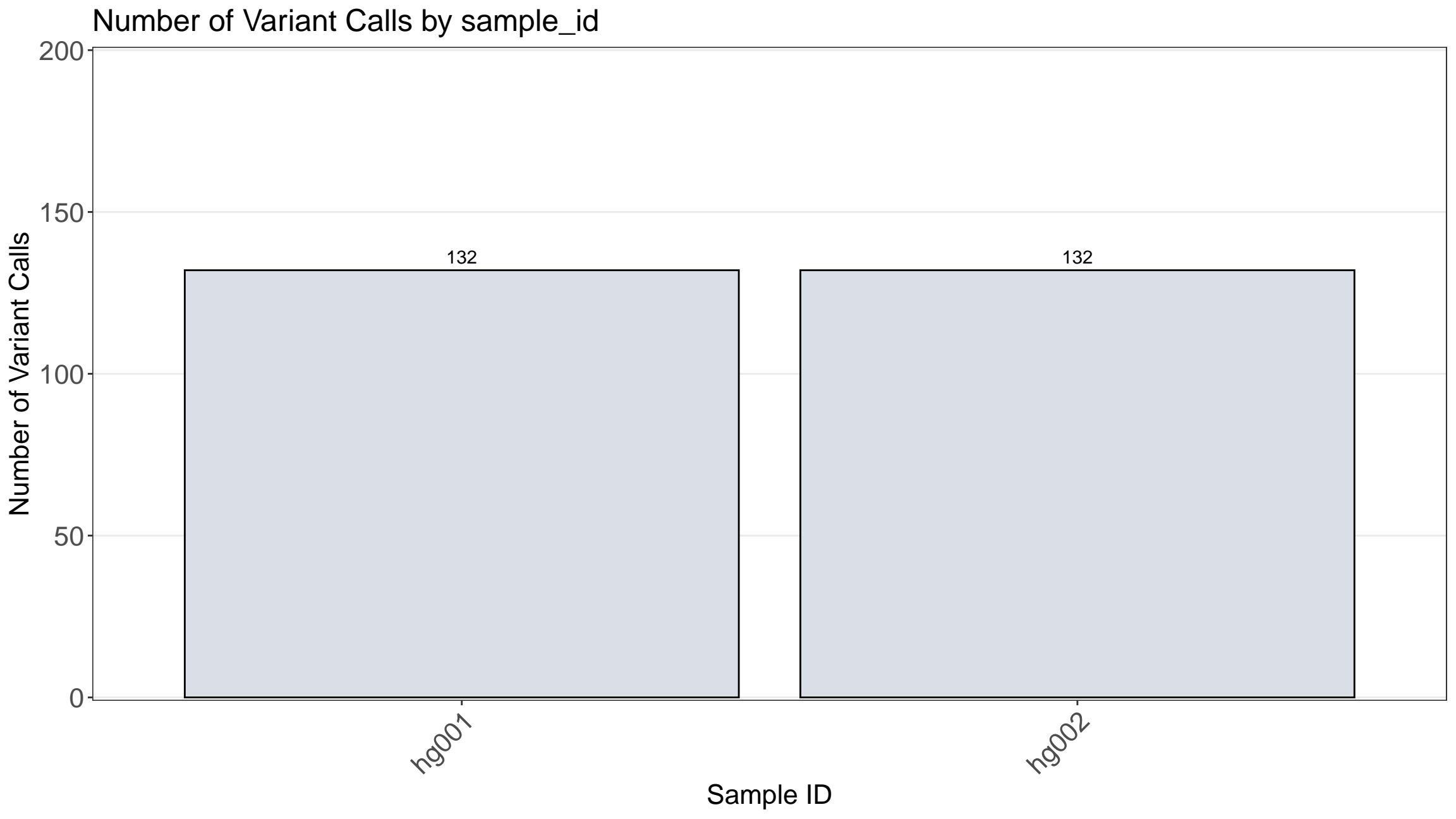


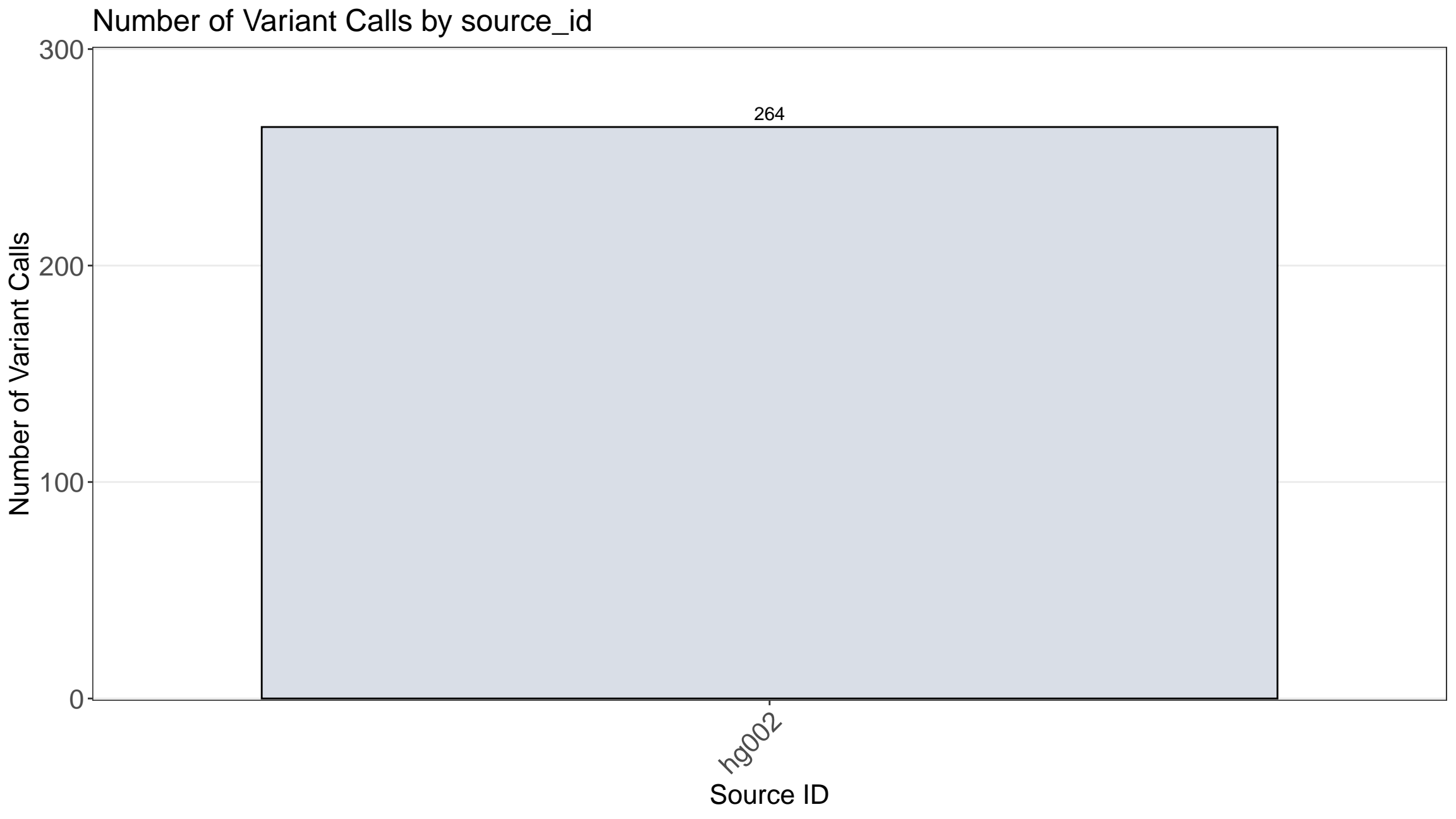
Number of Variant Calls by chromosome_1



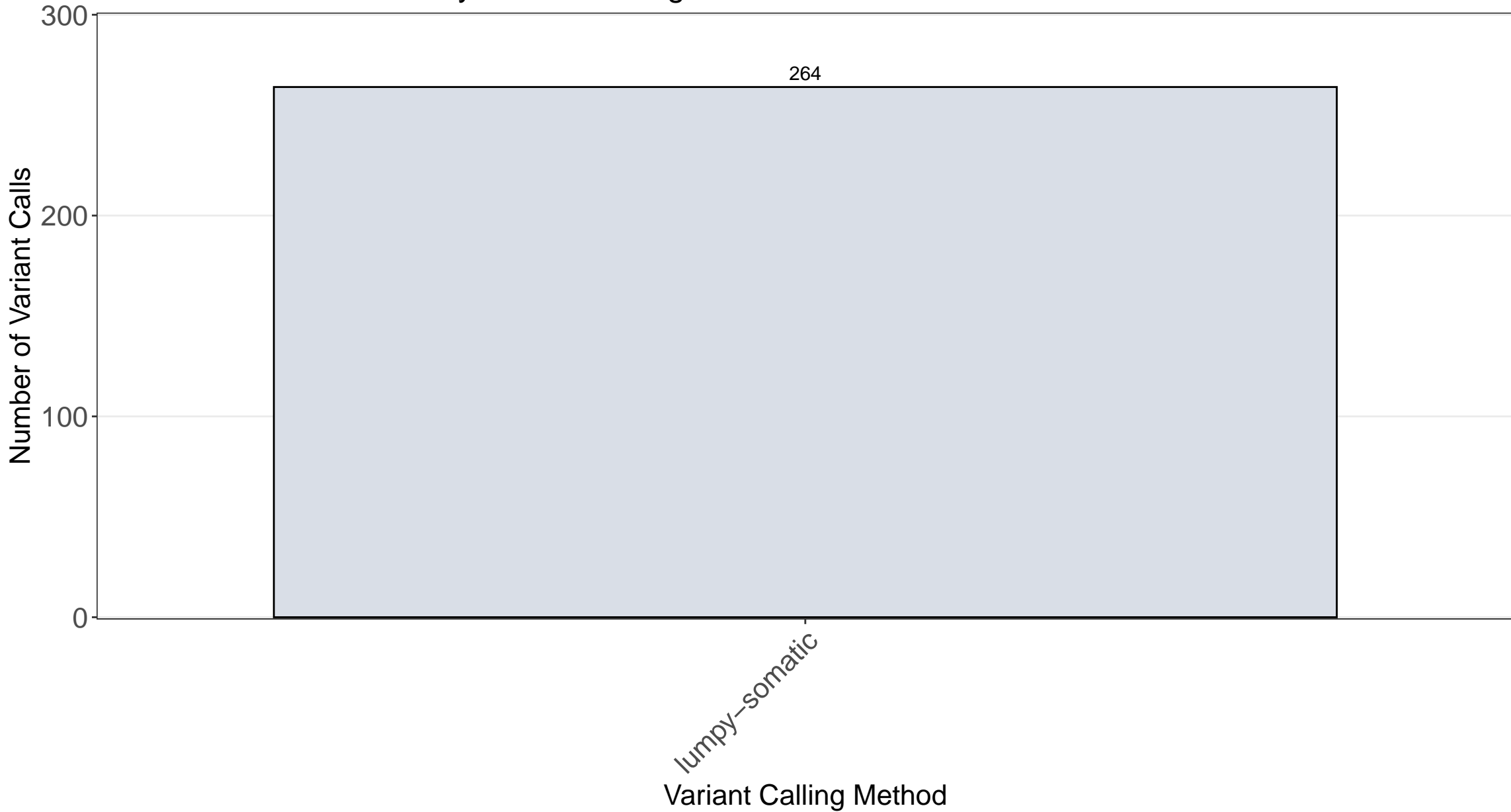
Number of Variant Calls by chromosome_2



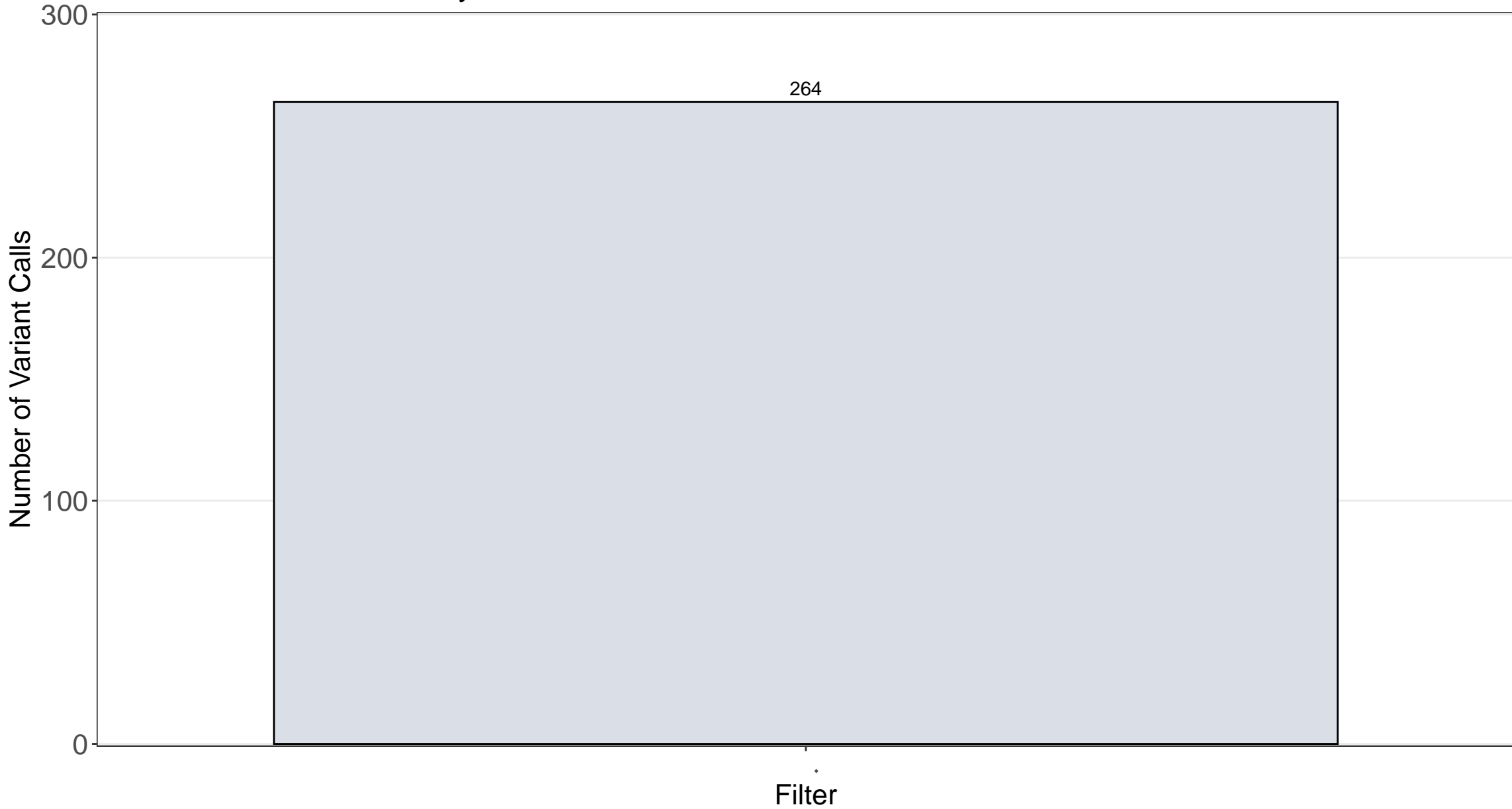




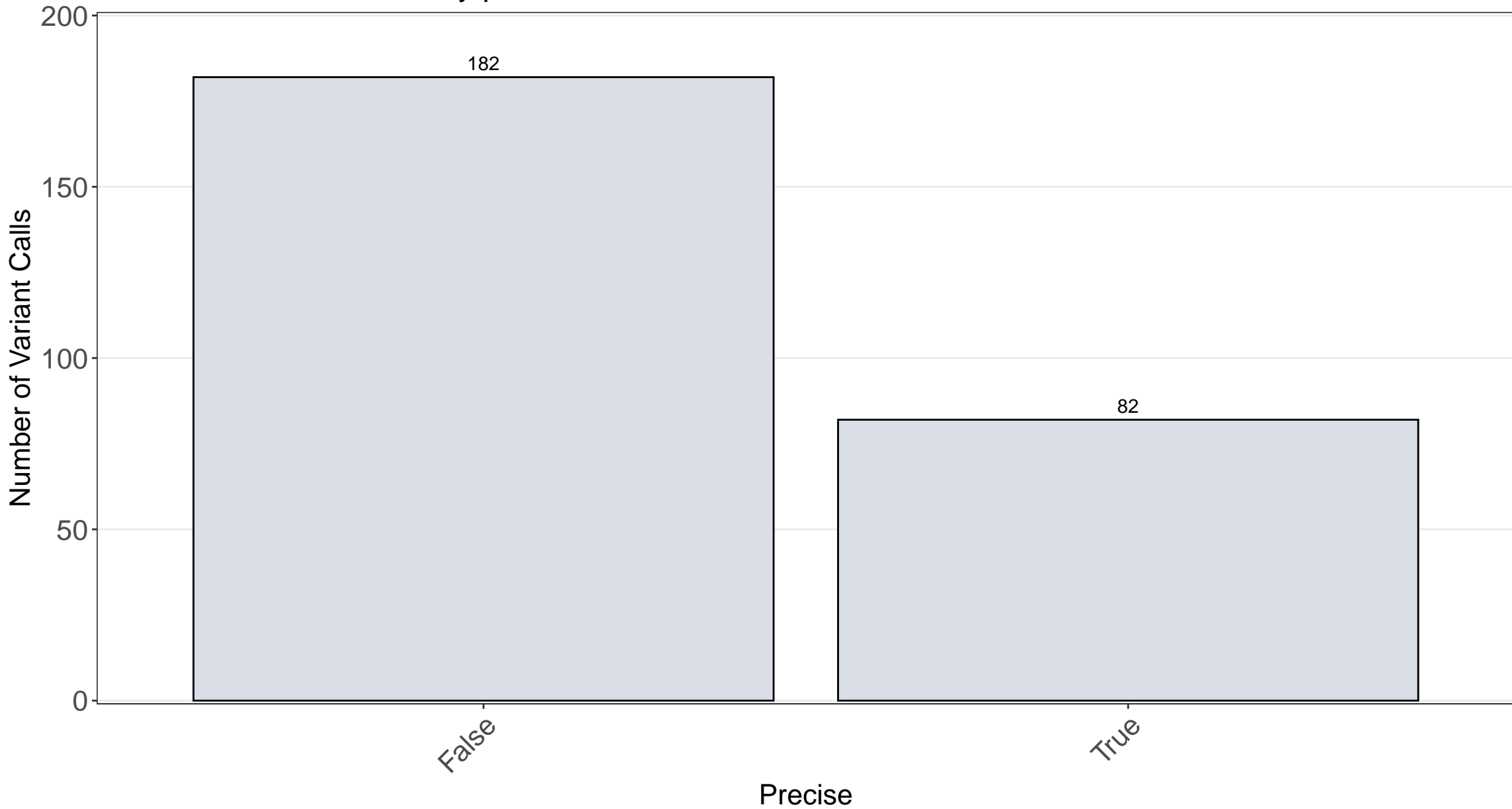
Number of Variant Calls by variant_calling_method

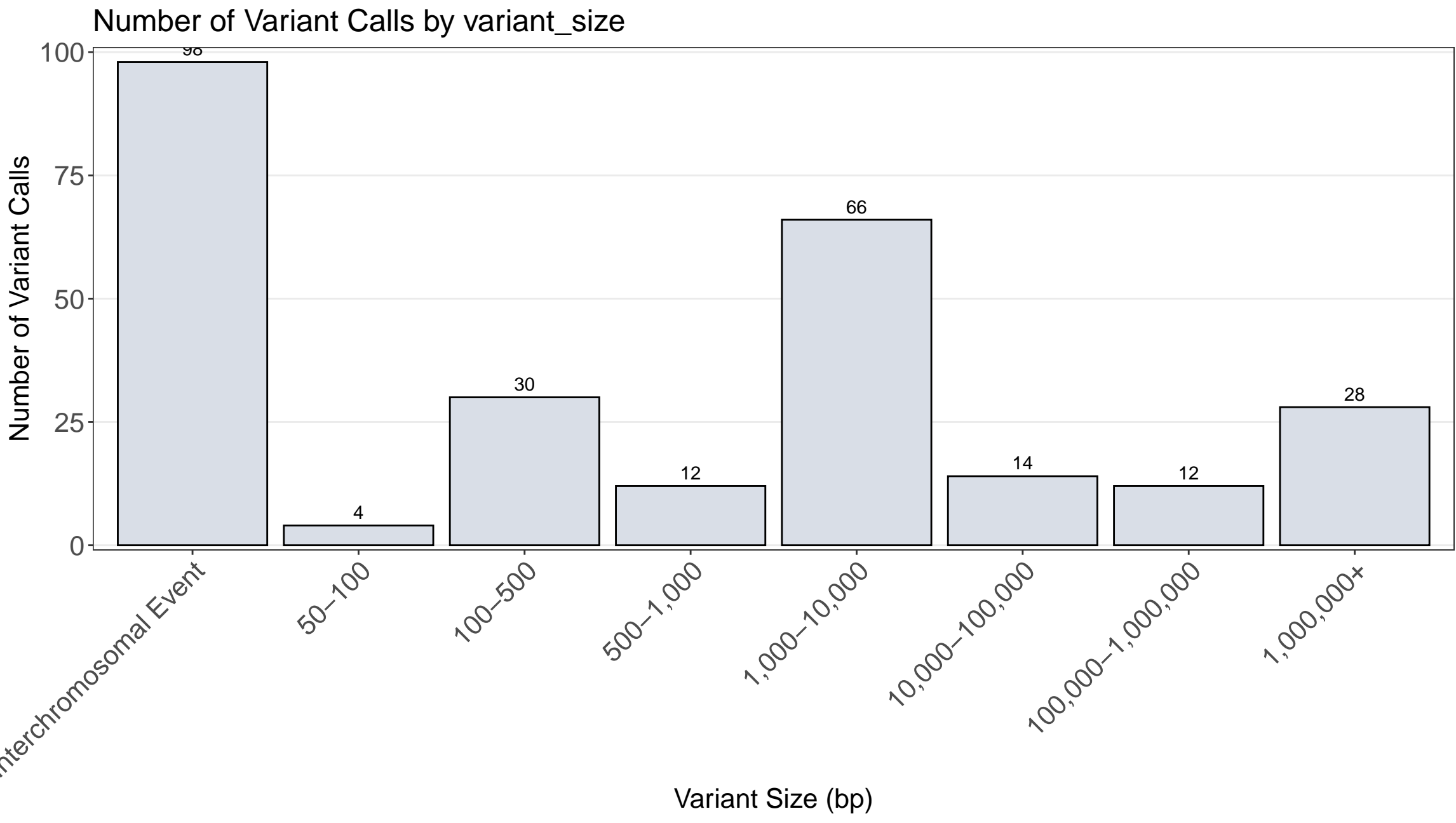


Number of Variant Calls by filter



Number of Variant Calls by precise





Histogram of alternate_allele_read_count

