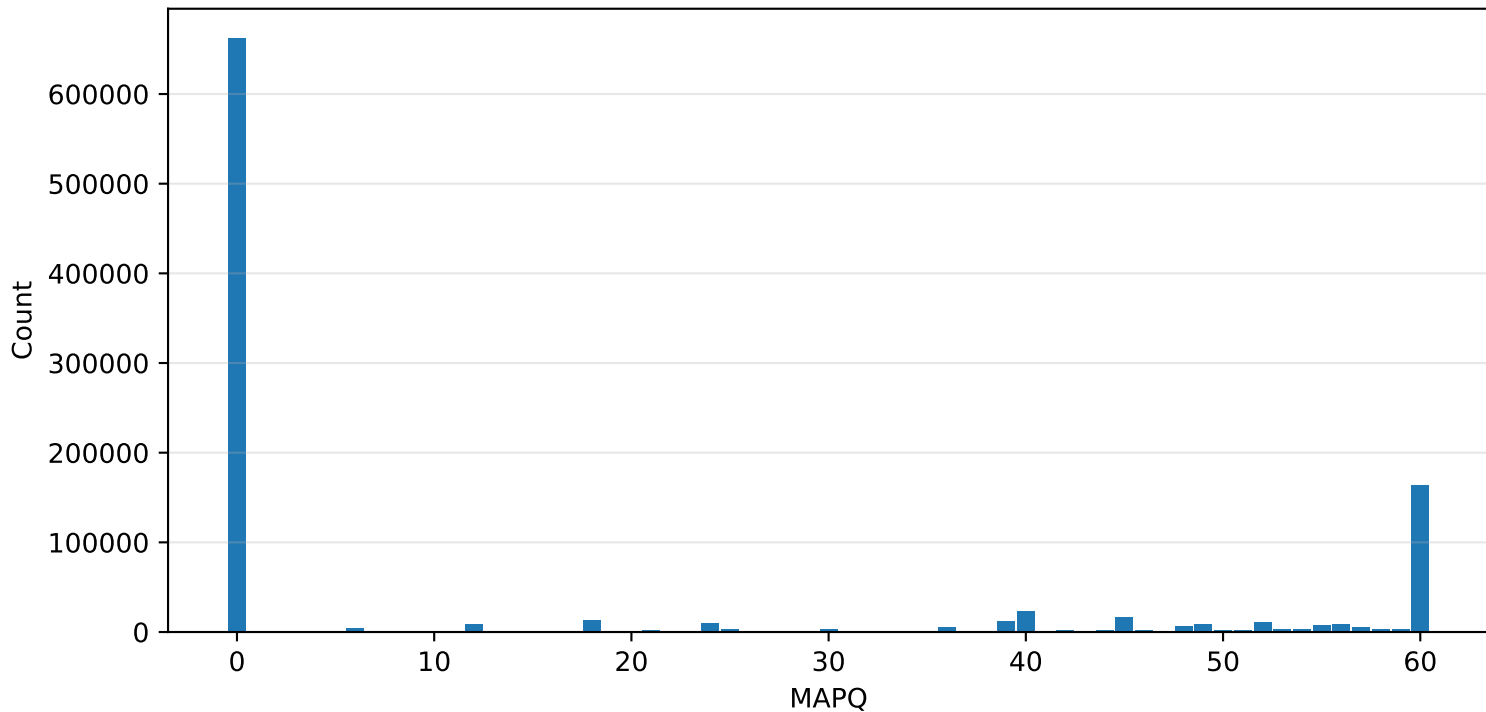
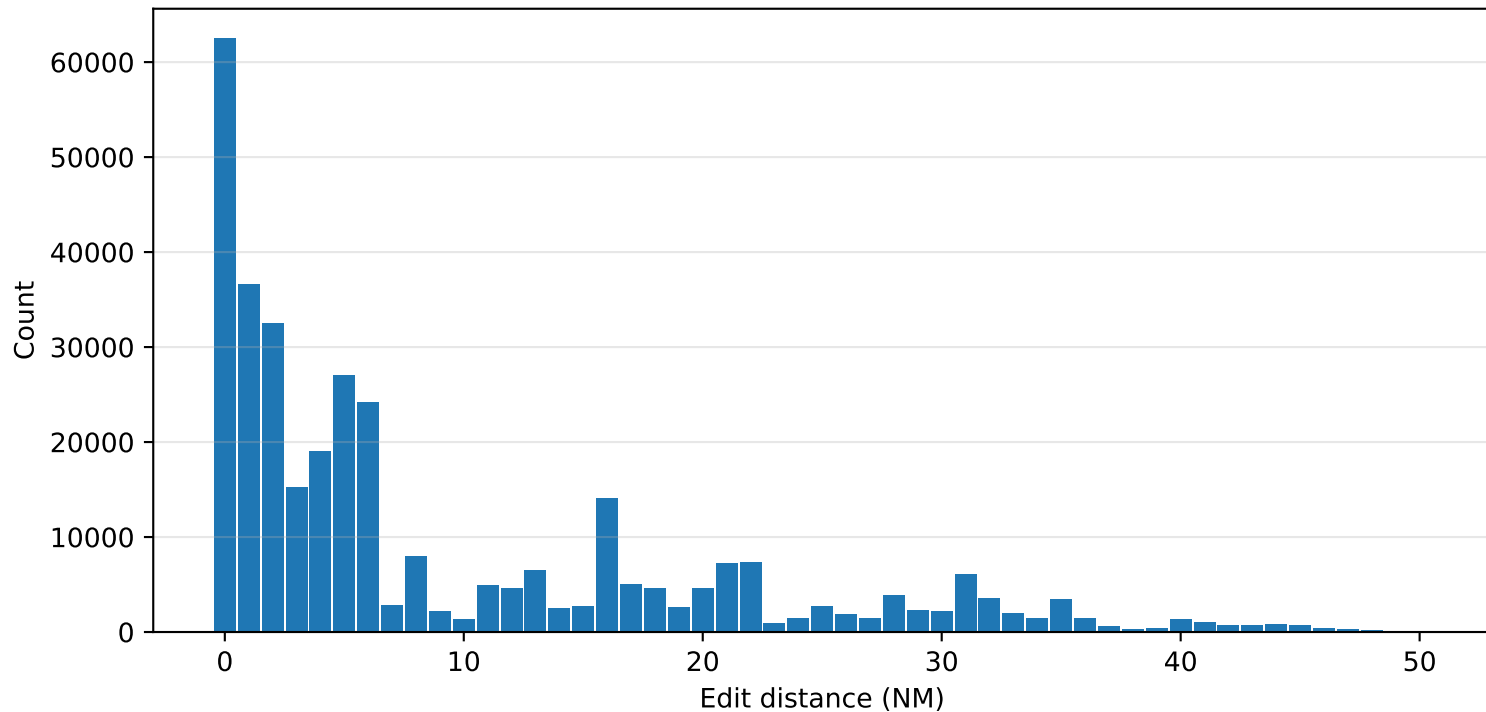


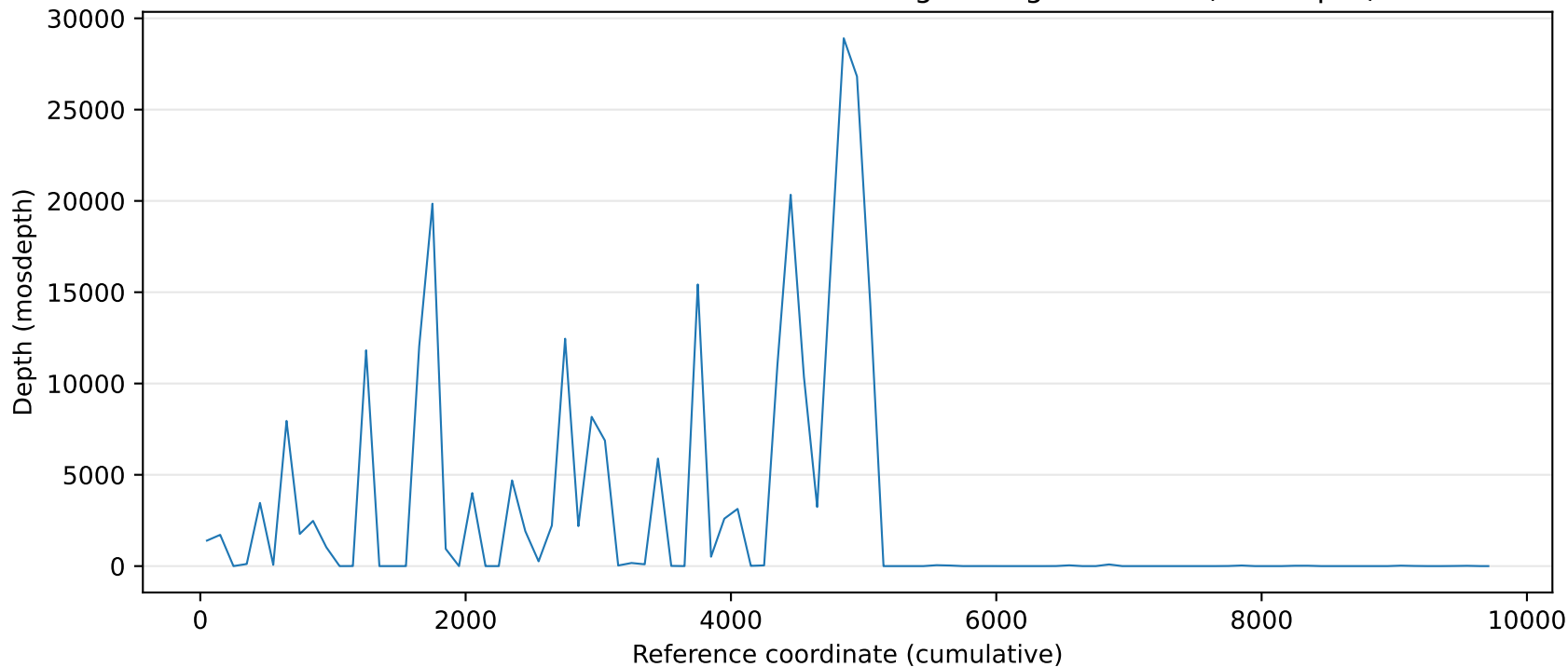
BWA MEM KU168292 to HXB2 — Mapping Quality (MAPQ)



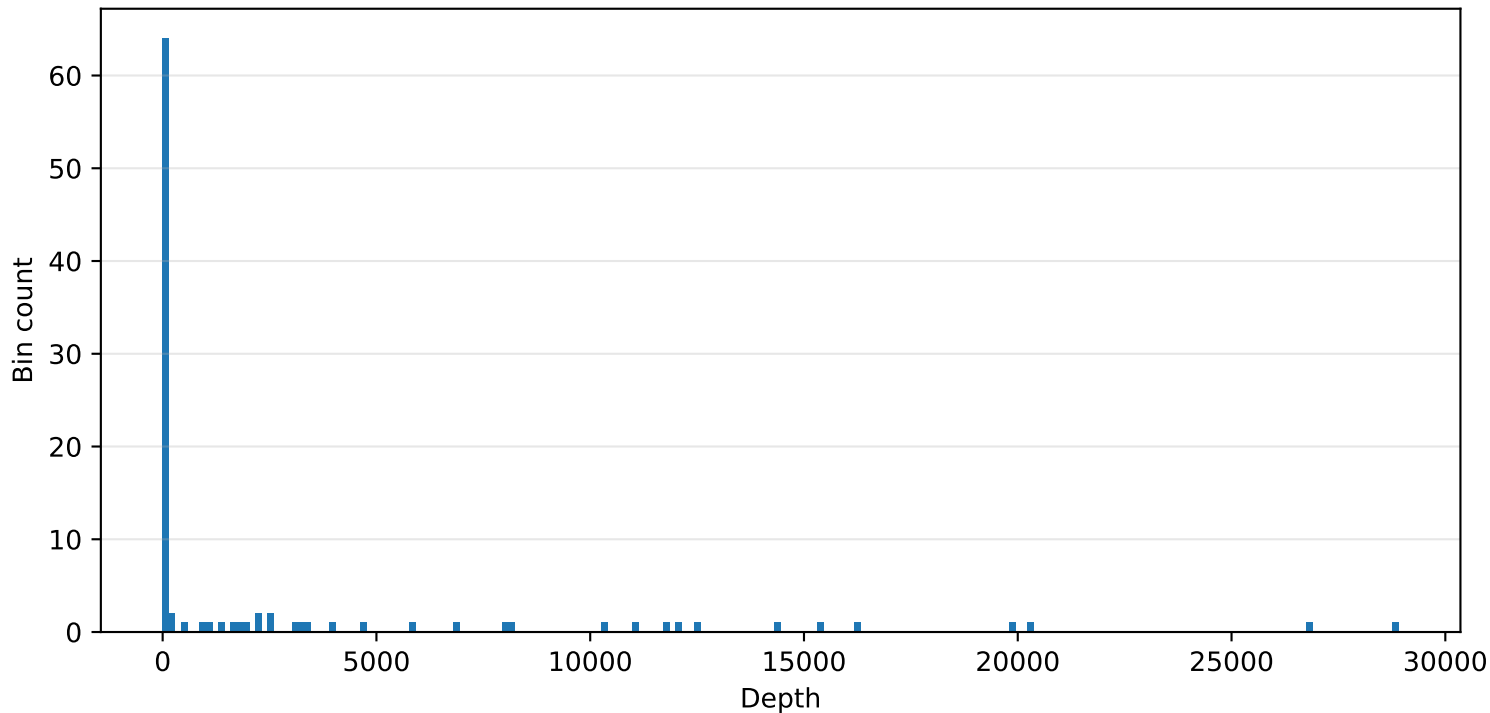
BWA MEM KU168292 to HXB2 — Edit Distance (NM)



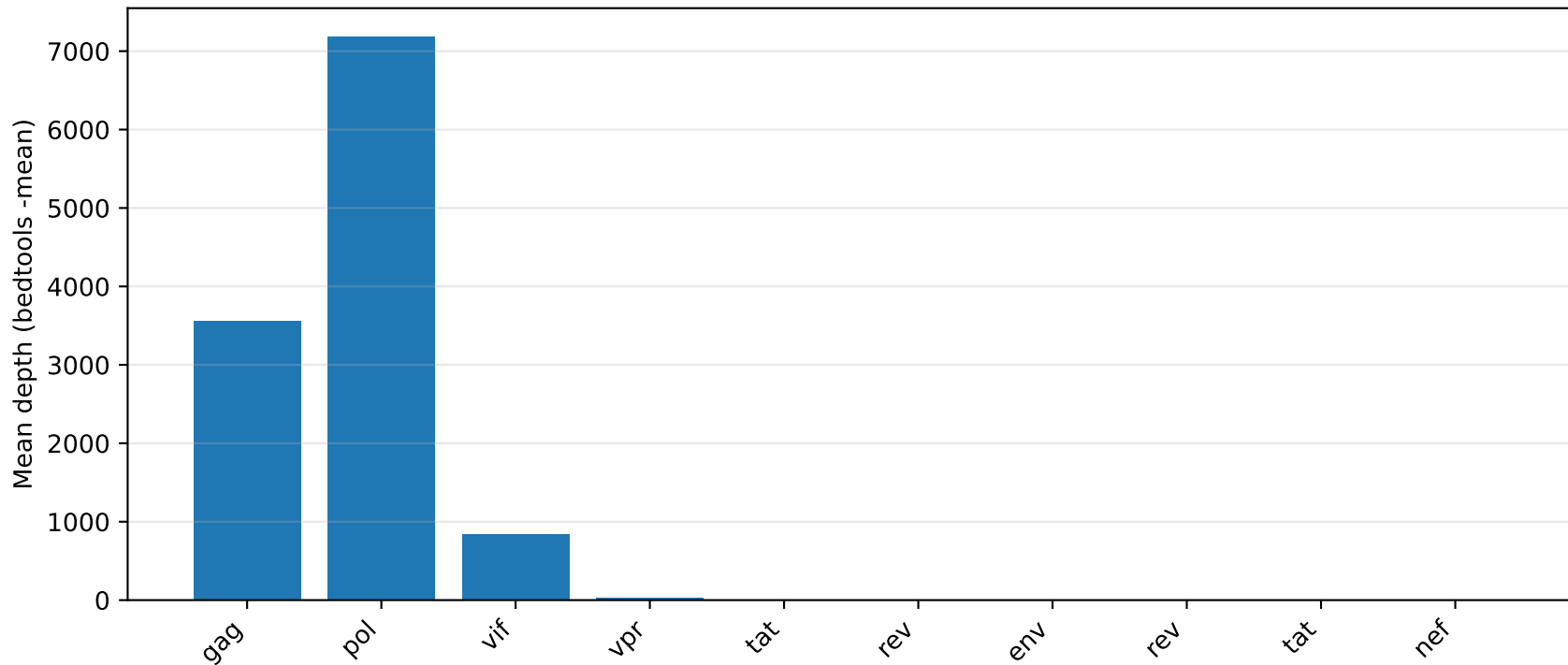
BWA MEM KU168292 to HXB2 — Coverage along reference (mosdepth)



BWA MEM KU168292 to HXB2 — Coverage distribution (mosdepth)



BWA MEM KU168292 to HXB2 — Per-gene mean depth



plot_align_stats.py – summary

Title: BWA MEM KU168292 to HXB2

MAPQ hist: stats/0.FR.2005.LA49RBF189.KU168292_sim_miseq_hxb2.sorted.mapq_hist.tsv

NM hist: stats/0.FR.2005.LA49RBF189.KU168292_sim_miseq_hxb2.sorted.nm_hist.tsv

mosdepth regions: stats/0.FR.2005.LA49RBF189.KU168292_sim_miseq_hxb2.sorted.regions.bed.gz

mosdepth per-base: (none)

gene coverage: stats/0.FR.2005.LA49RBF189.KU168292_sim_miseq_hxb2.sorted.gene_coverage.tsv

Max MAPQ: 60

Max NM: 50

Max depth (hist clip): (none)