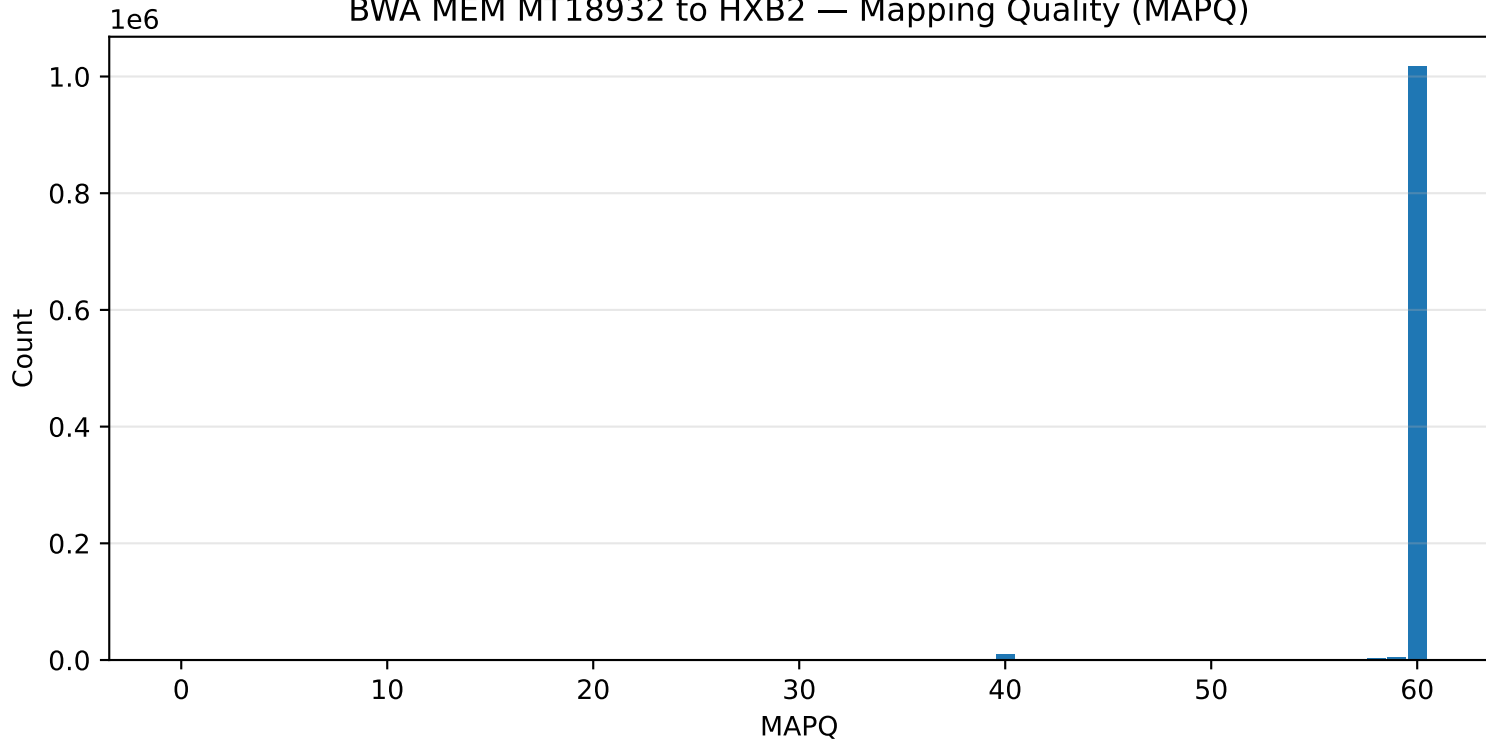
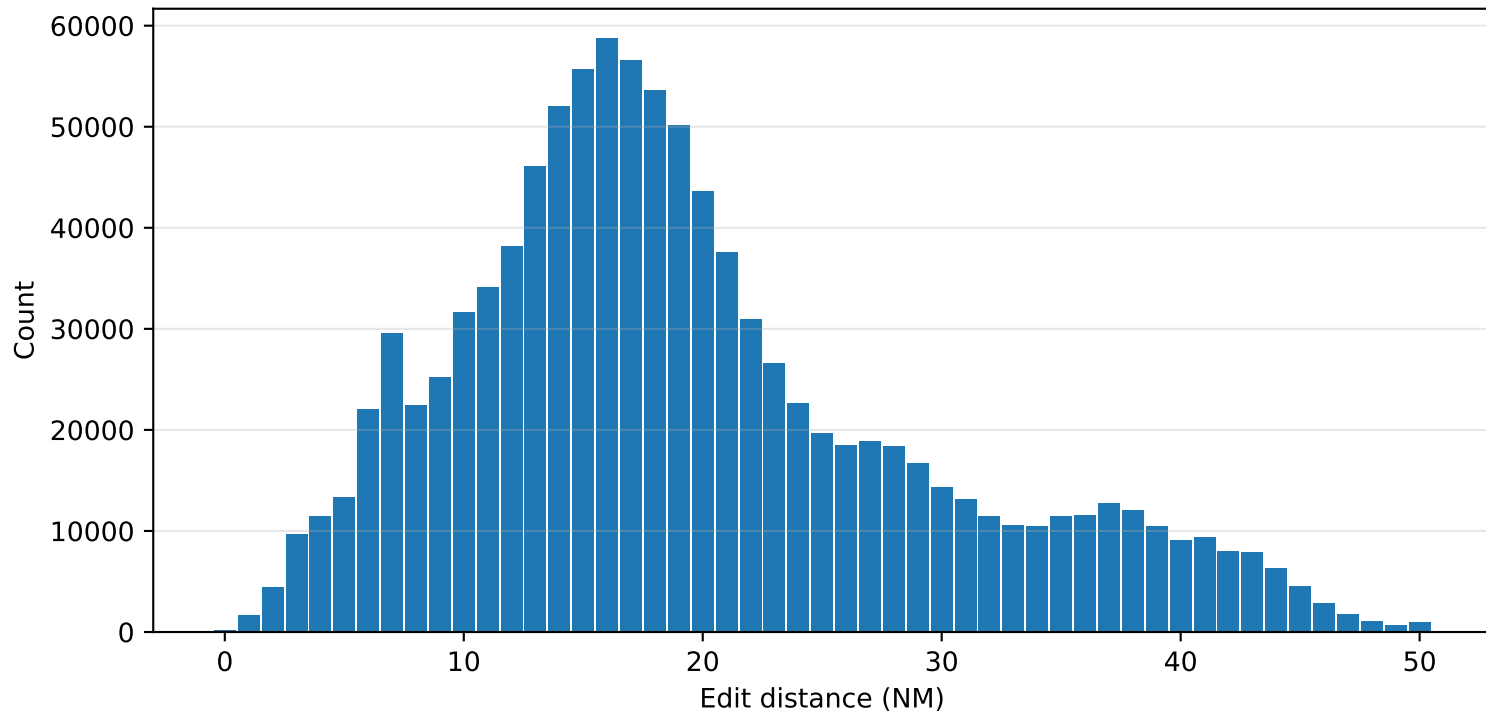


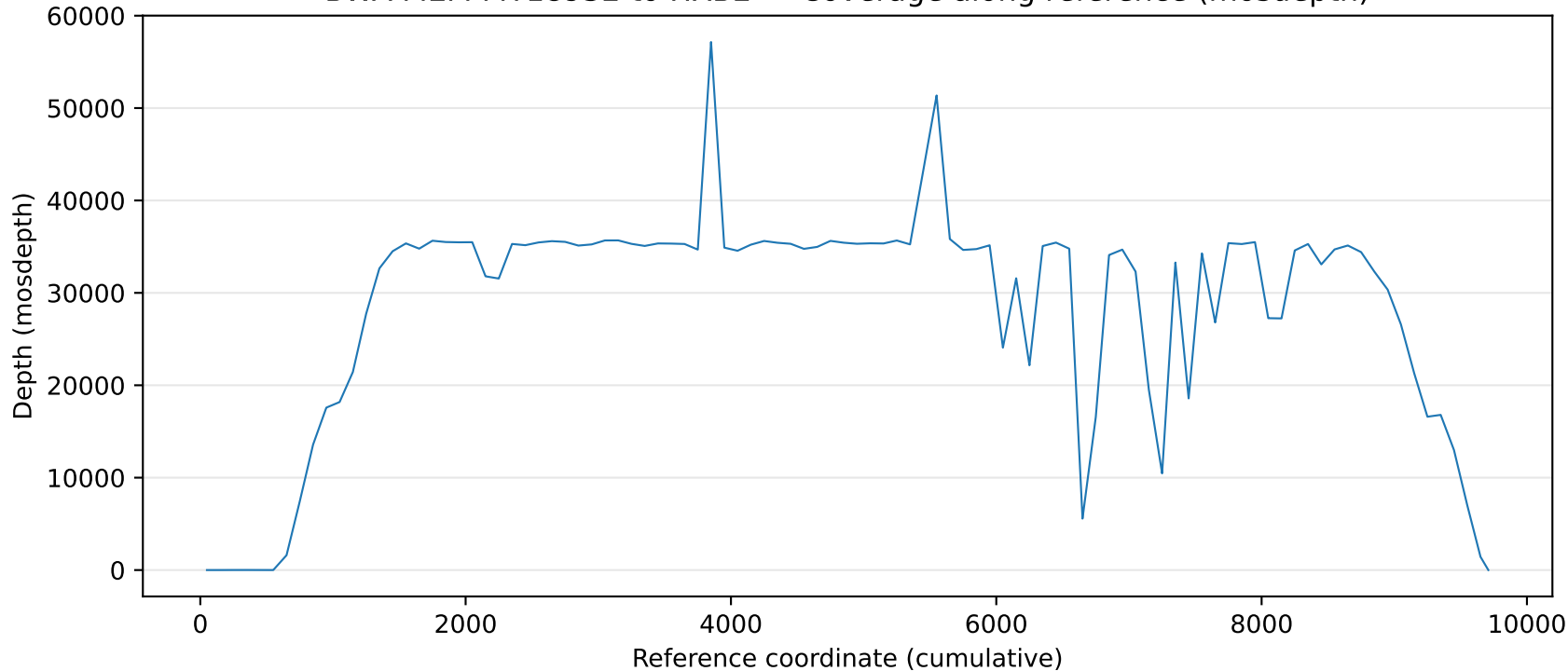
BWA MEM MT18932 to HXB2 — Mapping Quality (MAPQ)



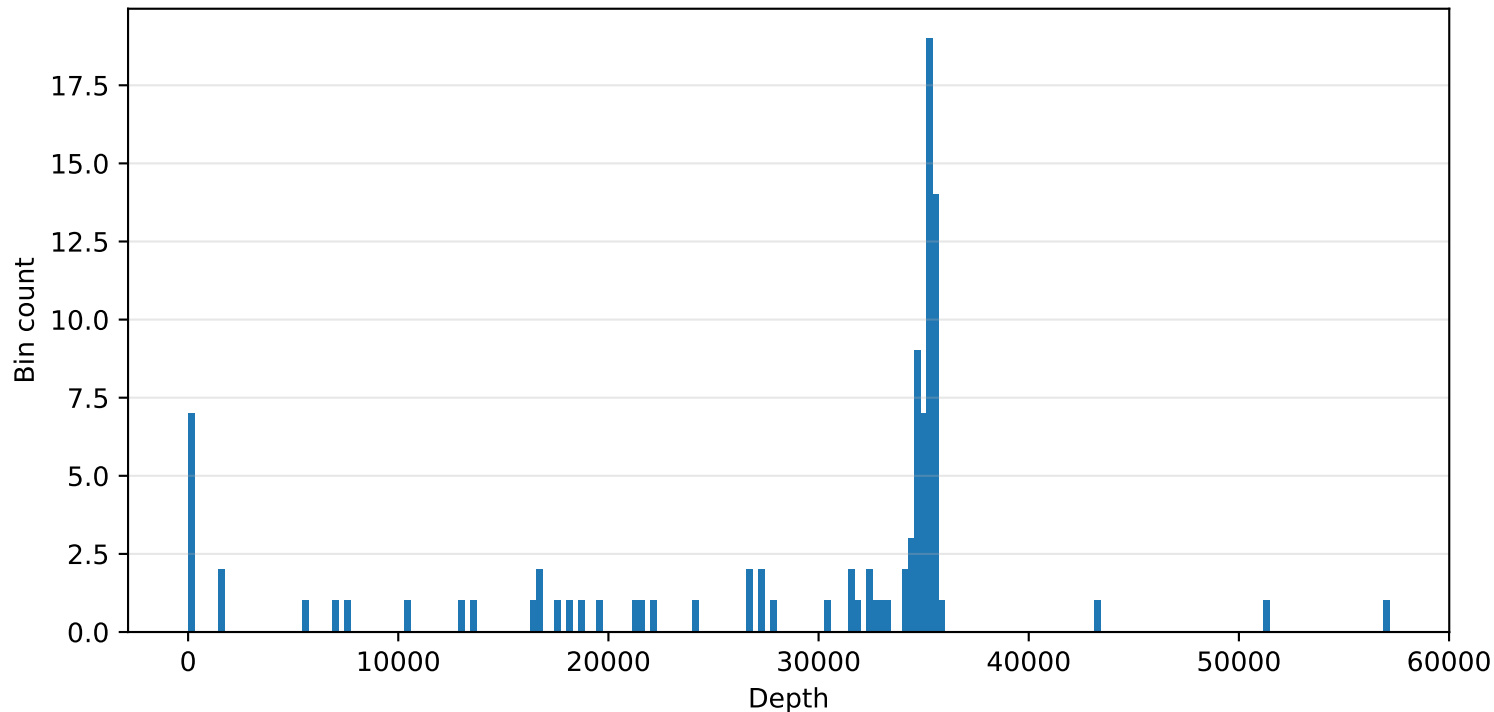
BWA MEM MT18932 to HXB2 — Edit Distance (NM)



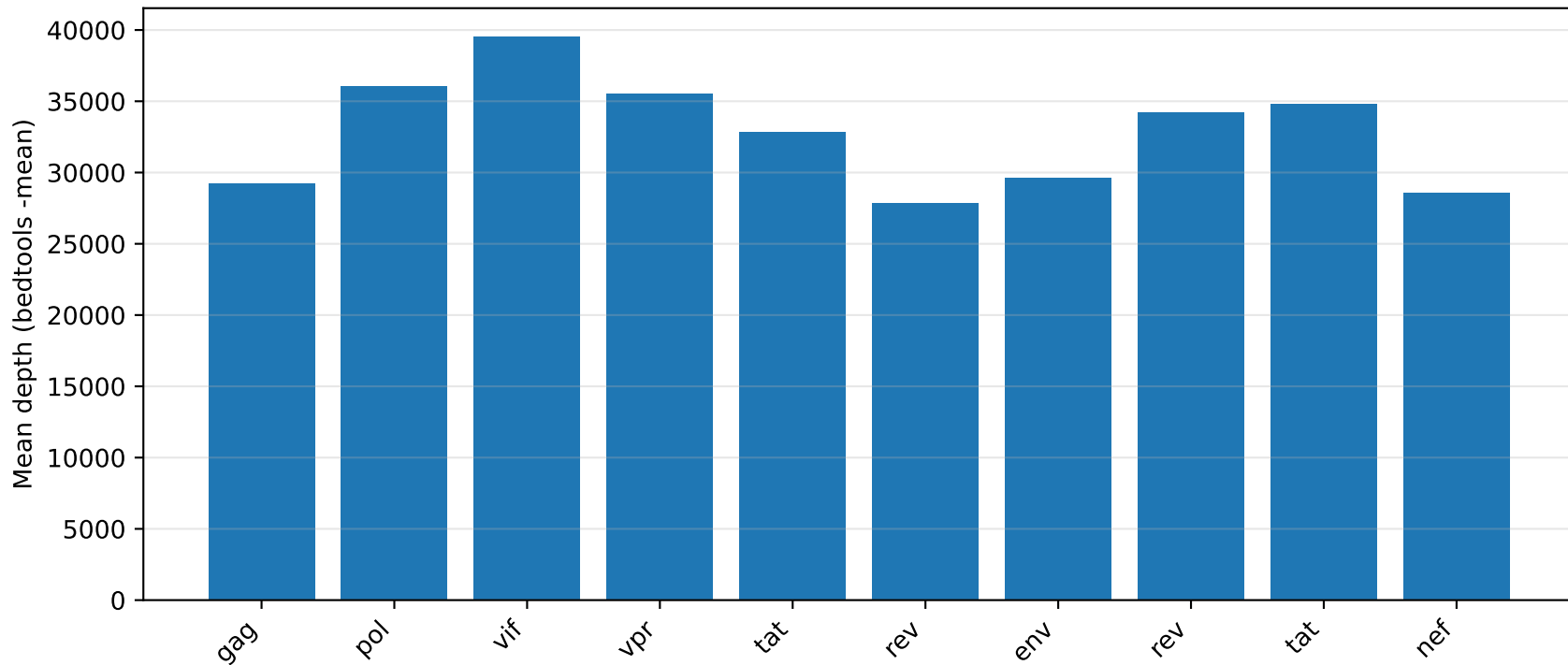
BWA MEM MT18932 to HXB2 — Coverage along reference (mosdepth)



BWA MEM MT18932 to HXB2 — Coverage distribution (mosdepth)



BWA MEM MT18932 to HXB2 — Per-gene mean depth



plot_align_stats.py – summary

Title: BWA MEM MT18932 to HXB2

MAPQ hist: stats/B.US.2016.5104_CMV_NEG_F21.MT18932_sim_miseq_hbx2.sorted.mapq_hist.tsv

NM hist: stats/B.US.2016.5104_CMV_NEG_F21.MT18932_sim_miseq_hbx2.sorted.nm_hist.tsv

mosdepth regions: stats/B.US.2016.5104_CMV_NEG_F21.MT18932_sim_miseq_hbx2.sorted.regions.bed.gz

mosdepth per-base: (none)

gene coverage: stats/B.US.2016.5104_CMV_NEG_F21.MT18932_sim_miseq_hbx2.sorted.gene_coverage.tsv

Max MAPQ: 60

Max NM: 50

Max depth (hist clip): (none)