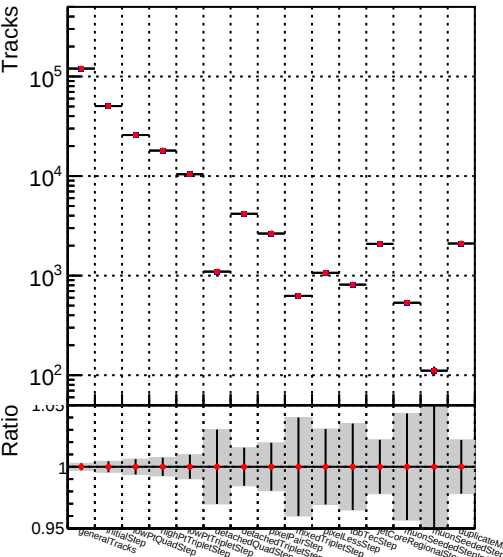
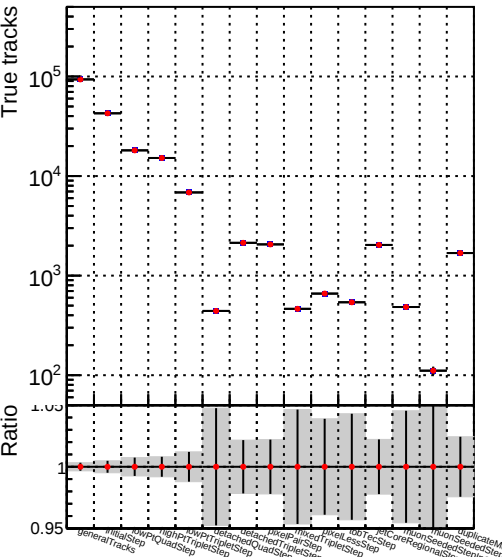


Number of tracks vs collection



Number of true tracks vs collection



Number	—●—	DQM_V0001_R0000000001_Global_mkFiti1000mod_RECO
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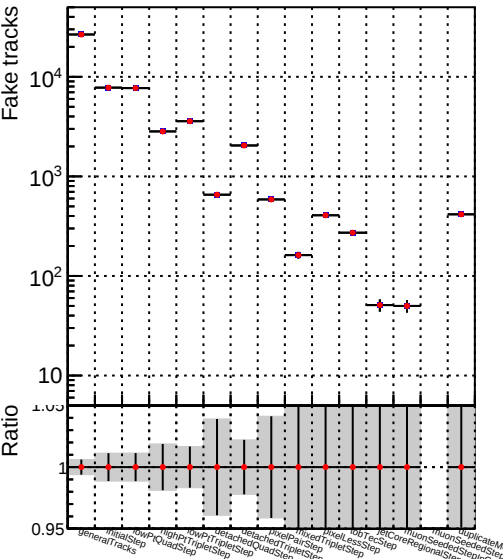


Figure 1 is a plot showing the ratio of duplicate tracks to total tracks for various genomic features. The y-axis is labeled 'Ratio' and ranges from 0.95 to 100 on a logarithmic scale. The x-axis lists genomic features: general tracks, chr1, chr2, chr3, chr4, chr5, chr6, chr7, chr8, chr9, chr10, chr11, chr12, chr13, chr14, chr15, chr16, chr17, chr18, chr19, chr20, chr21, chr22, chr23, chr24, chr25, chr26, chr27, chr28, chr29, chr30, chrX, chrY, chrMT, chrUn, chrUn1, chrUn2, chrUn3, chrUn4, chrUn5, chrUn6, chrUn7, chrUn8, chrUn9, chrUn10, chrUn11, chrUn12, chrUn13, chrUn14, chrUn15, chrUn16, chrUn17, chrUn18, chrUn19, chrUn20, chrUn21, chrUn22, chrUn23, chrUn24, chrUn25, chrUn26, chrUn27, chrUn28, chrUn29, chrUn30, chrUn31, chrUn32, chrUn33, chrUn34, chrUn35, chrUn36, chrUn37, chrUn38, chrUn39, chrUn40, chrUn41, chrUn42, chrUn43, chrUn44, chrUn45, chrUn46, chrUn47, chrUn48, chrUn49, chrUn50, chrUn51, chrUn52, chrUn53, chrUn54, chrUn55, chrUn56, chrUn57, chrUn58, chrUn59, chrUn60, chrUn61, chrUn62, chrUn63, chrUn64, chrUn65, chrUn66, chrUn67, chrUn68, chrUn69, chrUn70, chrUn71, chrUn72, chrUn73, chrUn74, chrUn75, chrUn76, chrUn77, chrUn78, chrUn79, chrUn80, chrUn81, chrUn82, chrUn83, chrUn84, chrUn85, chrUn86, chrUn87, chrUn88, chrUn89, chrUn90, chrUn91, chrUn92, chrUn93, chrUn94, chrUn95, chrUn96, chrUn97, chrUn98, chrUn99, chrUn100. A horizontal line at Ratio = 1 indicates the expected ratio. Features with ratios significantly above 1 are highlighted in grey.

Number of pileup tracks vs collection

