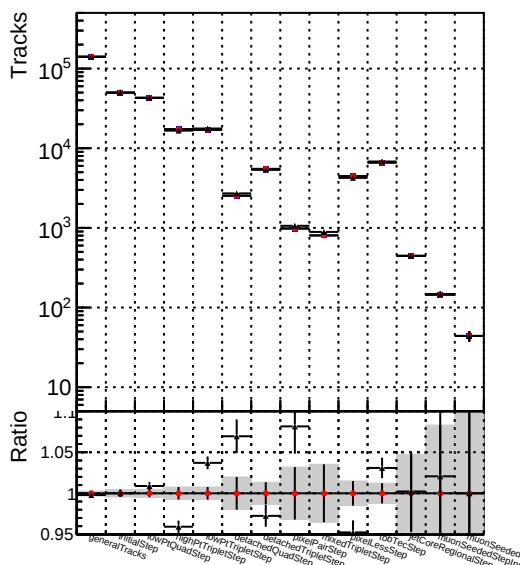
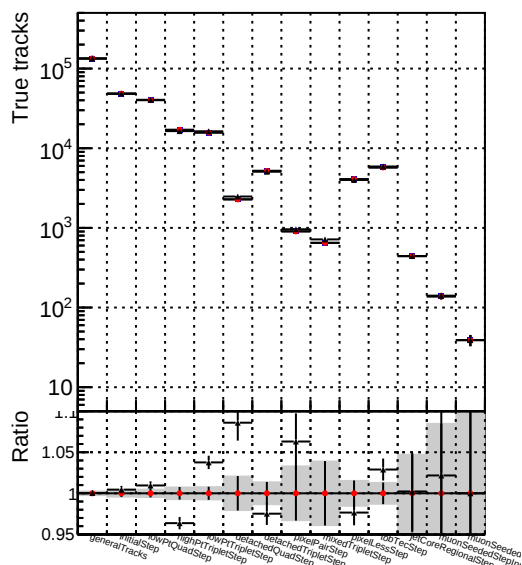


Number of tracks vs collection



Number of true tracks vs collection



Number	Legend
	Blue line with square: ../CMSSW_12_4_0_pre3/src/DQM_V0001_R0000000001_Global_mkFit_batchCAND-adjusted
	Red line with circle: ../CMSSW_12_4_0_pre3/src/DQM_V0001_R0000000001_Global_mkFit_CAND
	Black line with triangle: DQM_V0001_R0000000001_Global_mkFit_CAND-AFTERCONFLICT2

Number

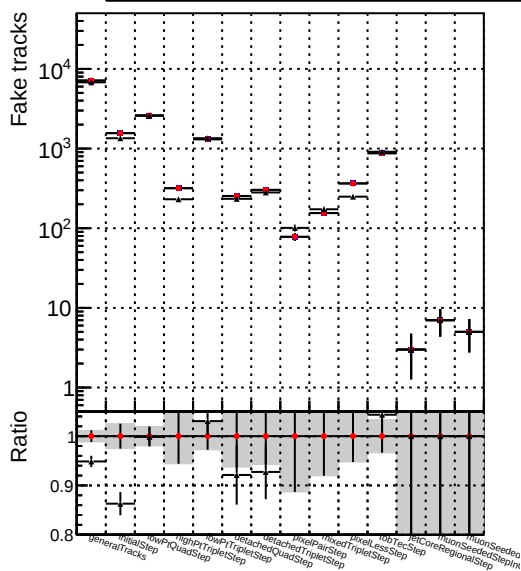


Figure 1 is a log-log plot showing the ratio of duplicate tracks to total tracks for various genomic features. The y-axis is labeled 'Duplicate tracks' and ranges from 10^1 to 10^4 . The x-axis is labeled 'Ratio' and ranges from 10^{-1} to 10^1 . The plot shows that the ratio of duplicate tracks to total tracks is generally high for most features, with some features showing a significant decrease in the ratio as the number of tracks increases.

Number of pileup tracks vs collection

