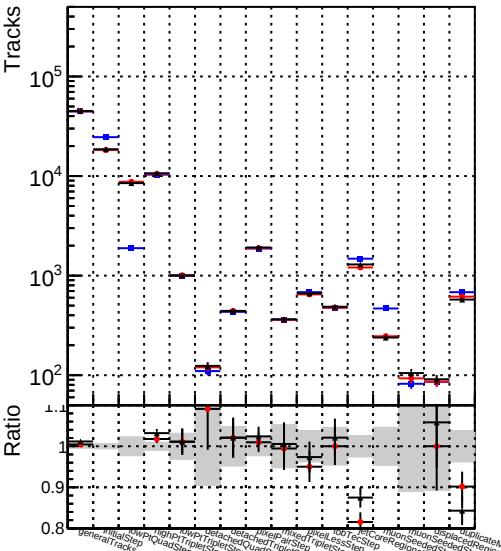
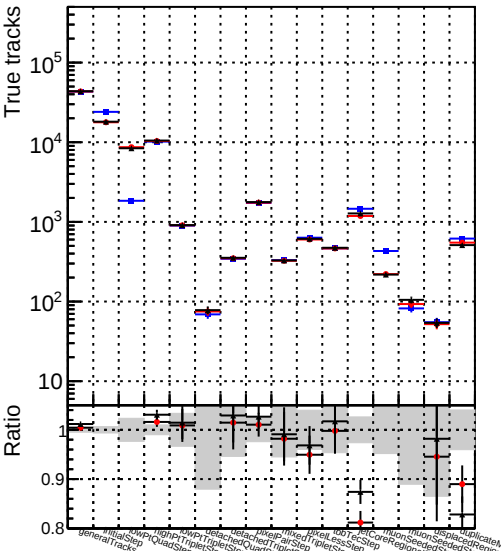


Number of tracks vs collection



Number of true tracks vs collection



 DQM_pixelCPETemplate_step2_ttbar-phase1-newgeom
 DQM_pixelCPETemplateCutChiSquareNew7_step2_ttbar-phase1-newgeom

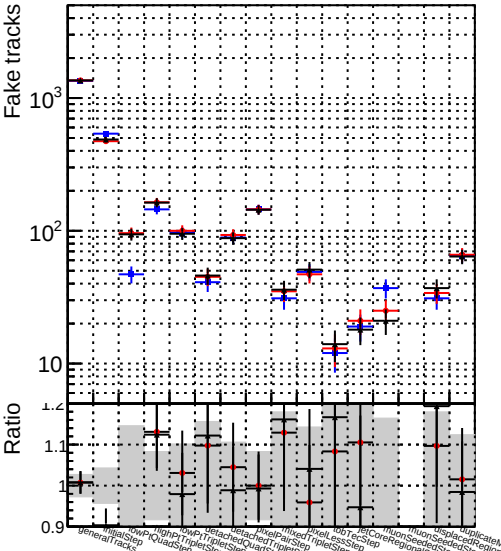


Figure 1 is a plot showing the ratio of duplicate tracks to the total number of tracks for various methods. The y-axis is labeled 'Ratio' and ranges from 0.8 to 10^2 on a logarithmic scale. The x-axis lists 20 methods: General track, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, Haplotype, and Haplotype. A horizontal line at Ratio=1 indicates the expected ratio. Most methods are clustered around 1.0, with some outliers like 'General track' and 'Haplotype' showing higher ratios.

Number of pileup tracks vs collection

