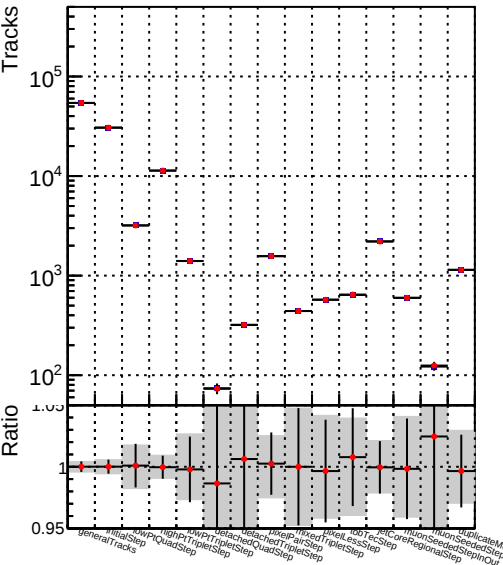
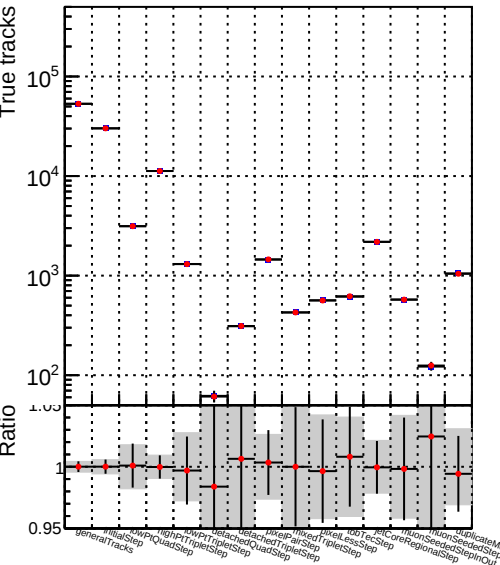


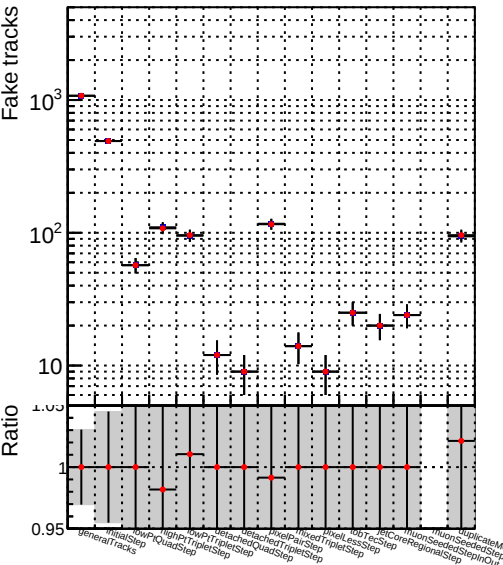
Number of tracks vs collection



Number of true tracks vs collection



Number	Label
1	DQM_V0001_R000000001_Global_mkFitbase2_RECO



Genomic Feature	Ratio (approx.)
GeneralTracks	1.005
MissedStep	1.002
HighPrTripletStep	1.003
DetachedTripletStep	1.004
MissedPairStep	1.006
MissedTripletStep	1.007
MissedCellStep	1.008
MissedRecStep	1.009
MissedCoreStep	1.010
MissedSeedStep	1.011
MissedRegionalStep	1.012
DuplicateTracks	1.025

Number of pileup tracks vs collection

