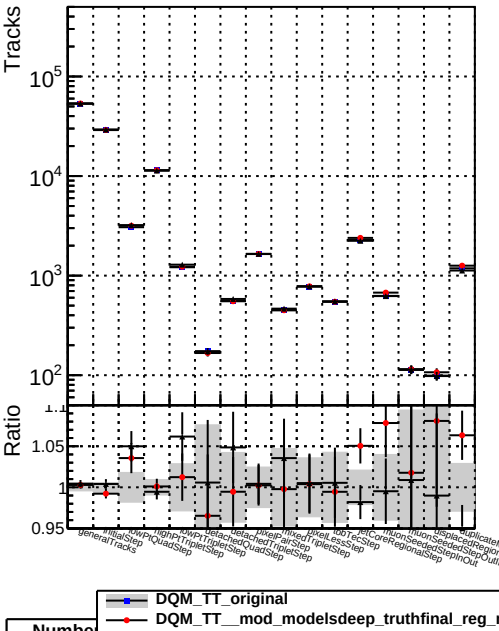
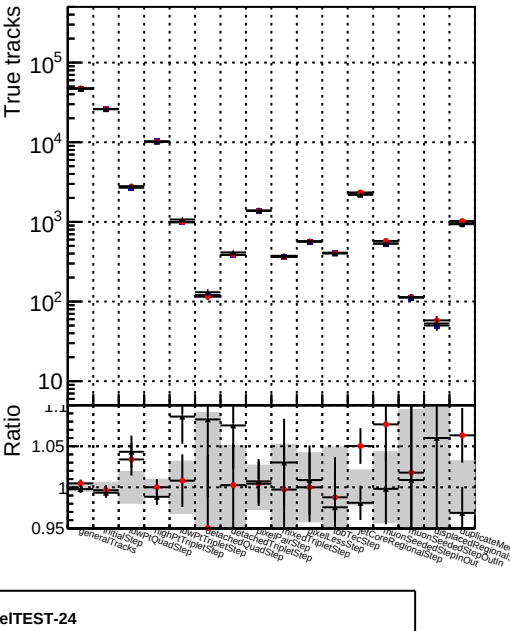


Number of tracks vs collection



Number of true tracks vs collection



Number

- DQM_TT_original
- DQM_TT_mod_modelsdeep_truthfinal_reg_modelTEST-24
- DQM_TT_mod_modelsdeep_something3

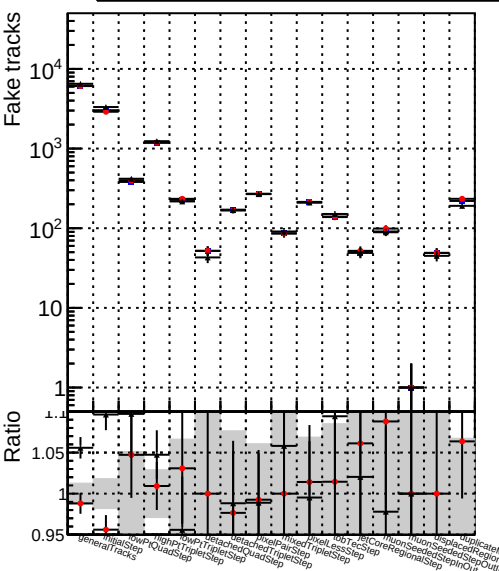


Figure 1: A dot plot showing the ratio of duplicate tracks to total tracks for various genomic features. The y-axis is 'Ratio' on a log scale from 0.8 to 10². The x-axis lists genomic features. A horizontal line is at Ratio = 1.0. A grey shaded region is between Ratio = 0.8 and 1.0. A red dot is at Ratio = 1.0 for 'General tracks'. A black dot is at Ratio = 1.0 for 'Duplicate tracks'. A black dot is at Ratio = 1.0 for 'Duplicate tracks'.

Number of pileup tracks vs collection

