Supplemental Fig. S6 – The same analysis as Fig. 5A-B was performed for *POLR2A*. Data from four cell lines (GM12878, HeLa-S3, HepG2 and HUVEC) and three laboratories (HAIB, SYDH, UTA) was analyzed here. By removing GC-content bias, improvements are observed: tighter clustering of samples for the same cell lines.