DNasel count.1M DNasel count.100k *** DNasel count.10k DNasel count.1k CpG_density.1M CpG_density.100k CpG density.10k CpG_counts.10k CpG_counts.1k GC.1M GC.100k *** GC.10k GC.1k *** GC.100 refSeq_counts.1M *** refSeq counts.100k *** refSeq_counts.10k within_refSeq_gene gene.width *** general.width start.dist boundary.dist onco.100k aCFR.1M NFR.1M

