Randomized accession names
TE deletions
TE insertions
Rsq = −0.0206 , p = 0.988
Rsq = 0.0133 , p = 0.195
Rsq = −0.00961 , p = 0.588
Rsq = −0.0126 , p = 0.678
C-DMRs
CG-DMRs
Rare TE variants (count)
DMR DNA methylation level rank (low-high)