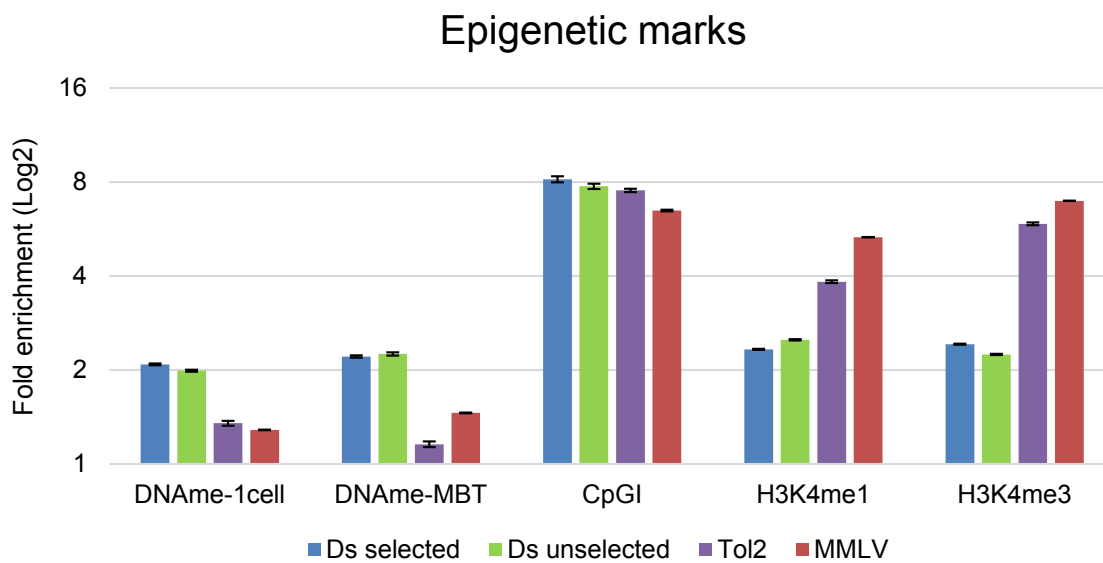


Figure S4



**Figure S4. Chromatin marks are correlated with integrations.** There is a strong preference for CpG islands, as well as CpG DNA methylation at 1-cell and MBT stages. Integration sites overlap with histone methylation marks of enhancer and promoter activity (ADAY *et al.* 2011). Fold enrichment values +/- standard error (n=1000).