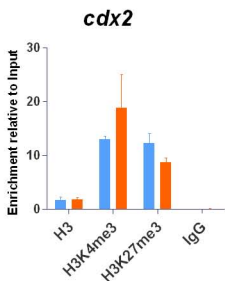
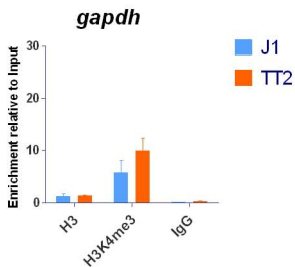


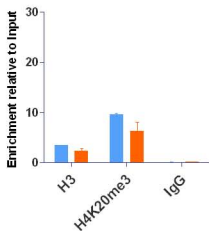
H3K4me3 and H3K27me3 positive control



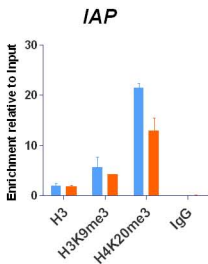
H3K4me3 positive control



H4K20me3 positive control



H3K9me3 and H4K20me3 positive control



Negative region for all histone modifications analyzed

