Mean log2 difference in coef estimate (to other proteins)

-2 0 2

MWU p-val (B/H): <0.01 <0.05 n.s

Column headers (left to right):
H2A.Z-Strongly recruited, H2A.Z-Strongly excluded, meDNA-Strongly recruited, meDNA-Strongly excluded, H3K4me1-Strongly recruited, H3K4me1-Strongly excluded, H3K4me3-Strongly recruited, H3K4me3-Strongly excluded, H3ac-Strongly recruited, H3ac-Strongly excluded, H3K9acK14ac-Strongly recruited, H3K9acK14ac-Strongly excluded, H3K27ac-Strongly recruited, H3K27ac-Strongly excluded, H3K9me2-Strongly recruited, H3K9me2-Strongly excluded, H3K9me3-Strongly recruited, H3K9me3-Strongly excluded, H3K27me2-Strongly recruited, H3K27me2-Strongly excluded, H3K27me3-Strongly recruited, H3K27me3-Strongly excluded, H4ac-Strongly recruited, H4ac-Strongly excluded, H4K16ac-Strongly recruited, H4K16ac-Strongly excluded, H4K20me2-Strongly recruited, H4K20me2-Strongly excluded, H4K20me3-Strongly recruited, H4K20me3-Strongly excluded

Row labels:
H3K4me1vsControl, H3K4me3vsControl, H3K4me3vsH3K4me1