

id	var name	att name	group name	P-val raw	sig bonf	sig fdr	mean class=0	mean class=1	method
1	chromMod_CTCF_overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	8.960	4.282	wilcox
2	chromMod_CTCF_overlapTotalLength	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	106.1	59.93	wilcox
3	chromMod_H2A_Z_overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	47.85	21.63	wilcox
4	chromMod_H2A_Z_overlapTotalLength	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	376.6	236.3	wilcox
5	chromMod_H2BK5me1_overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	7.819	4.772	wilcox
6	chromMod_H2BK5me1_overlapTotalLength	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	132.6	87.38	wilcox
7	chromMod_H3K27me1_overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	6.461	3.774	wilcox
8	chromMod_H3K27me1_overlapTotalLength	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	128.7	79.57	wilcox
9	chromMod_H3K4me2_overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	18.60	11.34	wilcox
10	chromMod_H3K4me2_overlapTotalLength	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	258.5	177.5	wilcox
11	chromMod_H3K4me3_overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	228.4	100.7	wilcox
12	chromMod_H3K9me1_overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	19.46	13.86	wilcox
13	chromMod_PolII_overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	27.24	8.970	wilcox
14	chromMod_PolII_overlapTotalLength	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	294.5	124.2	wilcox
15	overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	0	Yes	Yes	431.3	232.4	wilcox
16	thymus	GNF_Atlas_2	Transcriptome	0	Yes	Yes	0.176	0.010	wilcox
17	overlapRegionsCount	Human_ESTs	Transcriptome	0	Yes	Yes	101.3	37.77	wilcox
18	overlapRegionsCount	Spliced_ESTs	Transcriptome	0	Yes	Yes	93.35	32.82	wilcox
19	chromMod_H3K27me3_overlapRegionsCount	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	4.91e-41	Yes	Yes	2.241	5.902	wilcox
20	chromMod_H3K27me3_overlapTotalLength	NIH_Chromatin_Blood	Epigenome_and_Chromatin_Structure	1.98e-40	Yes	Yes	46.95	113.1	wilcox