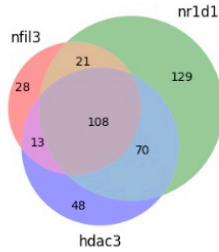


H3K27ac unique genes motif overlap (pval < 0.01)

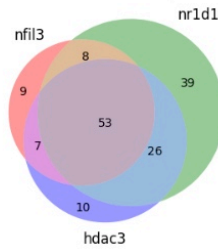
n= hdac3:239 nfil3:170 nr1d1:328 all:847

HOCOMOCO and JASPAR



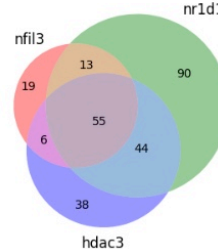
n= hdac3:96 nfil3:77 nr1d1:126 all:269

HOCOMOCO only



n= hdac3:143 nfil3:93 nr1d1:202 all:578

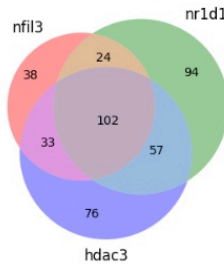
JASPAR only



H3K9ac unique genes motif overlap (pval < 0.01)

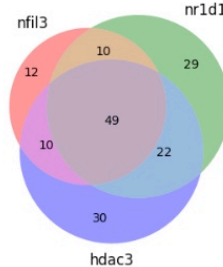
n= hdac3:268 nfil3:197 nr1d1:277 all:870

HOCOMOCO and JASPAR



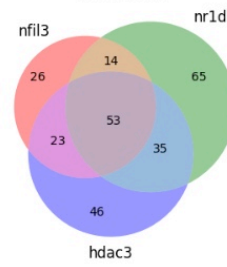
n= hdac3:111 nfil3:81 nr1d1:110 all:277

HOCOMOCO only



n= hdac3:157 nfil3:116 nr1d1:167 all:593

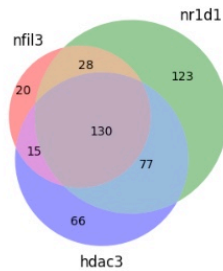
JASPAR only



500bp unique genes motif overlap (pval < 0.01)

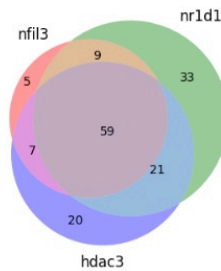
n= hdac3:288 nfil3:193 nr1d1:358 all:874

HOCOMOCO and JASPAR



n= hdac3:107 nfil3:80 nr1d1:122 all:285

HOCOMOCO only



n= hdac3:181 nfil3:113 nr1d1:236 all:589

JASPAR only

