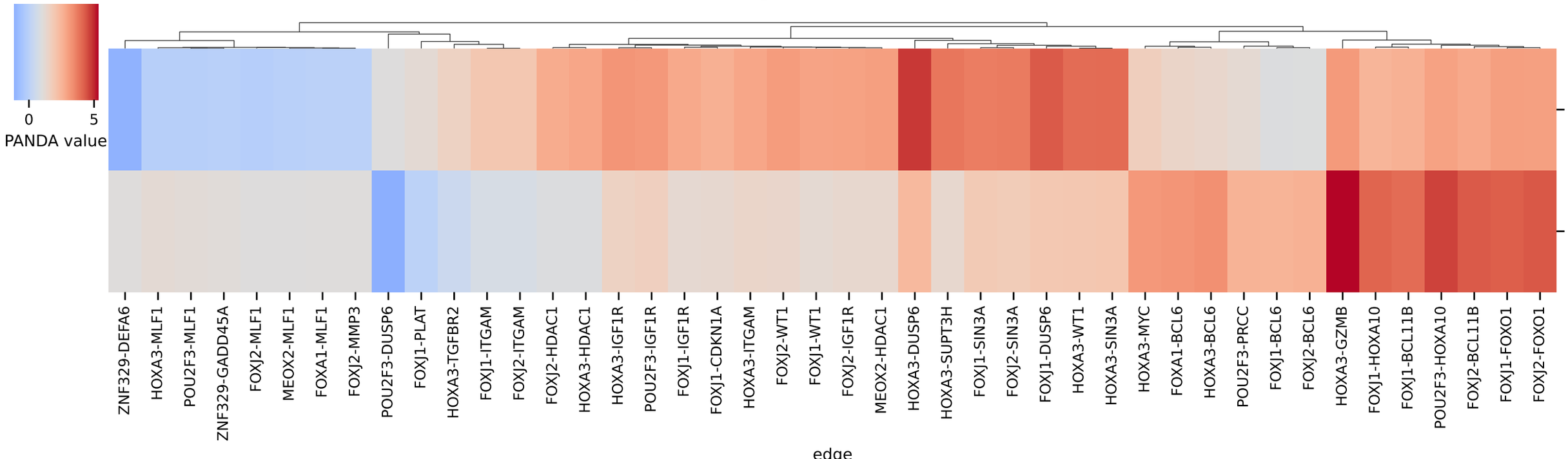
cluster_A: Transcriptional misregulation in cancer (FDR = 0.142)
0
5
PANDA value
cms2
cms4
ZNF329-DEFA6
HOXA3-MLF1
POU2F3-MLF1
ZNF329-GADD45A
FOXJ2-MLF1
MEOX2-MLF1
FOXA1-MLF1
FOXJ2-MMP3
POU2F3-DUSP6
FOXJ1-PLAT
HOXA3-TGFBR2
FOXJ1-ITGAM
FOXJ2-ITGAM
FOXJ2-HDAC1
HOXA3-HDAC1
HOXA3-IGF1R
POU2F3-IGF1R
FOXJ1-IGF1R
FOXJ1-CDKN1A
HOXA3-ITGAM
FOXJ2-WT1
FOXJ1-WT1
FOXJ2-IGF1R
MEOX2-HDAC1
HOXA3-DUSP6
HOXA3-SUPT3H
FOXJ1-SIN3A
FOXJ2-SIN3A
FOXJ1-DUSP6
HOXA3-WT1
HOXA3-SIN3A
HOXA3-MYC
FOXA1-BCL6
HOXA3-BCL6
POU2F3-PRCC
FOXJ1-BCL6
FOXJ2-BCL6
HOXA3-GZMB
FOXJ1-HOXA10
FOXJ1-BCL11B
POU2F3-HOXA10
FOXJ2-BCL11B
FOXJ1-FOXO1
FOXJ2-FOXO1
edge