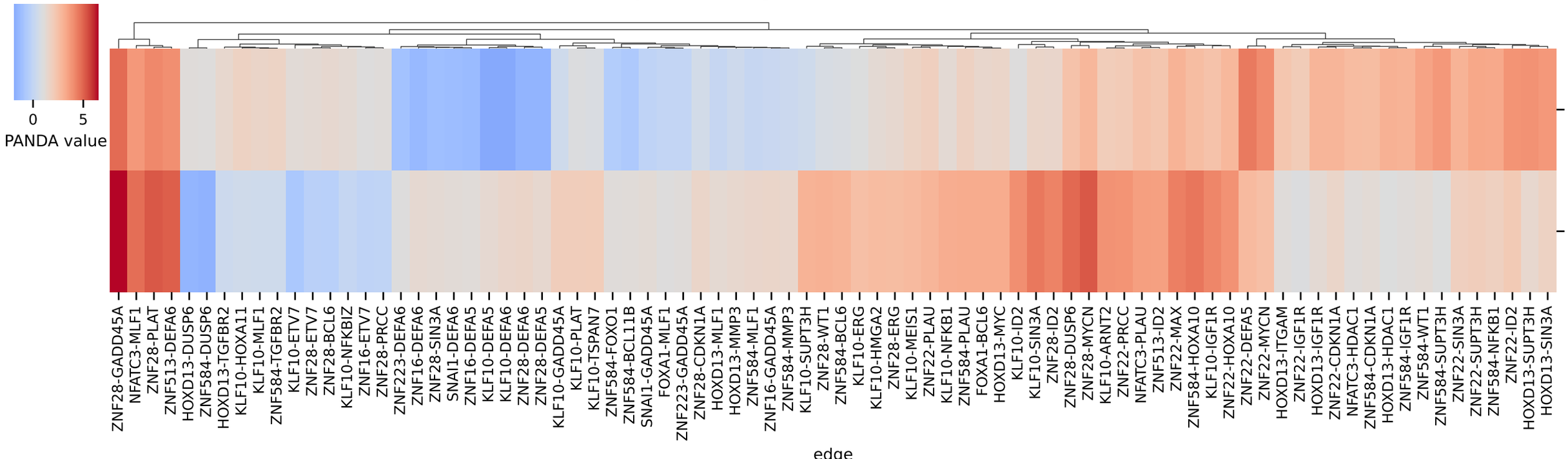
cluster_D: Transcriptional misregulation in cancer (FDR = 0.018)
0
5
PANDA value
cms2
cms4
ZNF28-GADD45A
NFATC3-MLF1
ZNF28-PLAT
ZNF513-DEFA6
HOXD13-DUSP6
ZNF584-DUSP6
HOXD13-TGFBR2
KLF10-HOXA11
KLF10-MLF1
ZNF584-TGFBR2
KLF10-ETV7
ZNF28-ETV7
ZNF28-BCL6
KLF10-NFKBIZ
ZNF16-ETV7
ZNF28-PRCC
ZNF223-DEFA6
ZNF16-DEFA6
ZNF28-SIN3A
SNAI1-DEFA6
ZNF16-DEFA5
KLF10-DEFA5
KLF10-DEFA6
ZNF28-DEFA6
ZNF28-DEFA5
KLF10-GADD45A
KLF10-PLAT
KLF10-TSPAN7
ZNF584-FOXO1
ZNF584-BCL11B
SNAI1-GADD45A
FOXA1-MLF1
ZNF223-GADD45A
ZNF28-CDKN1A
HOXD13-MLF1
HOXD13-MMP3
ZNF584-MLF1
ZNF16-GADD45A
ZNF584-MMP3
KLF10-SUPT3H
ZNF28-WT1
ZNF584-BCL6
KLF10-ERG
KLF10-HMGA2
ZNF28-ERG
KLF10-MEIS1
ZNF22-PLAU
KLF10-NFKB1
ZNF584-PLAU
FOXA1-BCL6
HOXD13-MYC
KLF10-ID2
KLF10-SIN3A
ZNF28-ID2
ZNF28-DUSP6
ZNF28-MYCN
KLF10-ARNT2
ZNF22-PRCC
NFATC3-PLAU
ZNF513-ID2
ZNF22-MAX
ZNF584-HOXA10
KLF10-IGF1R
ZNF22-HOXA10
ZNF22-DEFA5
ZNF22-MYCN
HOXD13-ITGAM
ZNF22-IGF1R
HOXD13-IGF1R
ZNF22-CDKN1A
NFATC3-HDAC1
ZNF584-CDKN1A
HOXD13-HDAC1
ZNF584-IGF1R
ZNF584-WT1
ZNF584-SUPT3H
ZNF22-SIN3A
ZNF22-SUPT3H
ZNF584-NFKB1
ZNF22-ID2
HOXD13-SUPT3H
HOXD13-SIN3A
edge