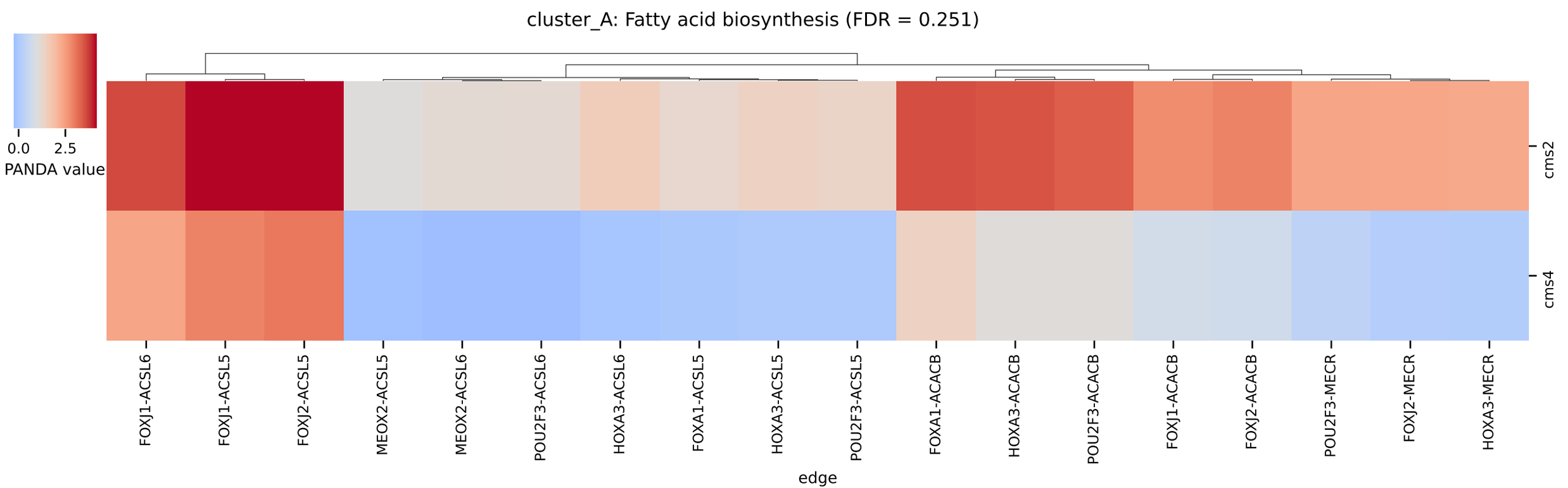
cluster_A: Fatty acid biosynthesis (FDR = 0.251)
0.0
2.5
PANDA value
cms2
cms4
FOXJ1-ACSL6
FOXJ1-ACSL5
FOXJ2-ACSL5
MEOX2-ACSL5
MEOX2-ACSL6
POU2F3-ACSL6
HOXA3-ACSL6
FOXA1-ACSL5
HOXA3-ACSL5
POU2F3-ACSL5
FOXA1-ACACB
HOXA3-ACACB
POU2F3-ACACB
FOXJ1-ACACB
FOXJ2-ACACB
POU2F3-MECR
FOXJ2-MECR
HOXA3-MECR
edge