

deg\_new\_clusters\_6-3

-log<sub>10</sub> adjusted p-value

300

200

100

0

-1.0

-0.5

0.0

0.5

1.0

log<sub>2</sub> FC

NRN1

NDUFA4L2

BNIP3

P4HA1

LDHA

ALDOA

TUBB2A

RGS16

PPP1R15A

SERTAD1

MYLIP

GADD45B

RGS2

DUSP2

MFSD2A

DUSP1

IER5

DNAJB1

JUN

FABP5

CTPS1

CITED4

NR4A1

ARL4D

ARL4A

ID2

BTG2

JUNB

IER2

ARL4A

GADD45G

DNAJB1

JUN

FABP5

CTPS1

ARL4A

GADD45G

IER5

DUSP1

MFSD2A

DUSP2

RGS2

GADD45B

HSPA1A

RASD1

EGR1

CTPS1

ARL4A