

ES = 0.543 NES = 1.74 Nom. p-val = 0.00375 FWER = 0.362 FDR = 0.217

Heatmap visualization showing the expression of 100 genes across two clusters: **clus2** and **NON.clus2**. The y-axis lists the genes, and the x-axis represents the two clusters. The color scale ranges from blue (low expression) to red (high expression).

Genes listed on the y-axis (from top to bottom):

- Class
- NC21
- FCGR2B
- FCGR2D
- FCGR2A
- FCGR2B
- ICK
- FCGR3A
- AV2
- MARCKS
- PRKCA
- PLA2G3A
- PRKCD
- RN1
- ARF1
- PIK3R5
- MAP3K1
- VASP
- LYN
- ARPC2
- ASAP
- BYK
- ARPC4
- GSN
- WAS
- ARF1
- LMN1
- PTPRC
- ARPC2
- ARF6
- CNK
- AMPH
- PRKCB
- CEL1
- DOCK2
- PIK3CD
- ACT1
- MAP3K
- MAP3K
- DOCK2
- SPR1
- RAC2
- VAV1
- PIK3IC
- ACT1
- MAPK1
- RAF1
- WASP2
- DNM2
- PRKCA
- ARPC2L
- QSOX
- INPP5D
- PIK3G
- WASL
- SPR2
- PLC2
- PIK3A
- LMN2
- DNM1
- MYO1B
- MYO1B
- RPRKCC
- PIK3B
- PIK3CA
- PIK3B
- AKT2
- PIK3T
- SAV3
- ASAP3
- PIK3R1
- CLT
- LOL
- PRKCB
- WASP
- PRAP2
- PRAP2A
- PLC1
- SON
- PIK3R1
- PRAP2
- PRAP2B
- MARCKSL1
- PLA2G3A
- DNM3
- DNM3
- WASP1