| 2 | FAM117A | 0.0011096316 | SCGB1D2 | 0.022212726 | FAM117A | 0.037322378 |
|---|---------|--------------|---------|-------------|---------|-------------|
| 3 | CSRP1 | 0.0046135985 | FAM117A | 0.026315680 | SCGB1D2 | 0.050938051 |
| 4 | PDPK1 | 0.0064068762 | SYT2 | 0.038265689 | CSRP1 | 0.167752625 |
| 5 | SCGB1D2 | 0.0064658899 | BSDC1 | 0.038674689 | ACHE | 0.167839362 |
| 6 | BSDC1 | 0.0065827091 | NA | 0.038757503 | BSDC1 | 0.179164177 |
| 7 | ARHGEF2 | 0.0066663337 | ACHE | 0.039096858 | ETV5 | 0.183981587 |
| 8 | NA | 0.0069919835 | NA | 0.043563967 | BHLHB9 | 0.188265768 |
| | | | | | | |

mfdr.mfdr

0.045324658

mfdr.cv

0.241818619

feature.cv

0.004235537 ZC2HC1A 0.007887758

0.043636338 ARHGEF2 0.236661507 NA

feature.mfdr

ZC2HC1A

WNK1

NA

feature.uni

ETV5

NA

1 ZC2HC1A

9

10

fdr.uni

0.0007518885

0.0071492096

0.0078690664