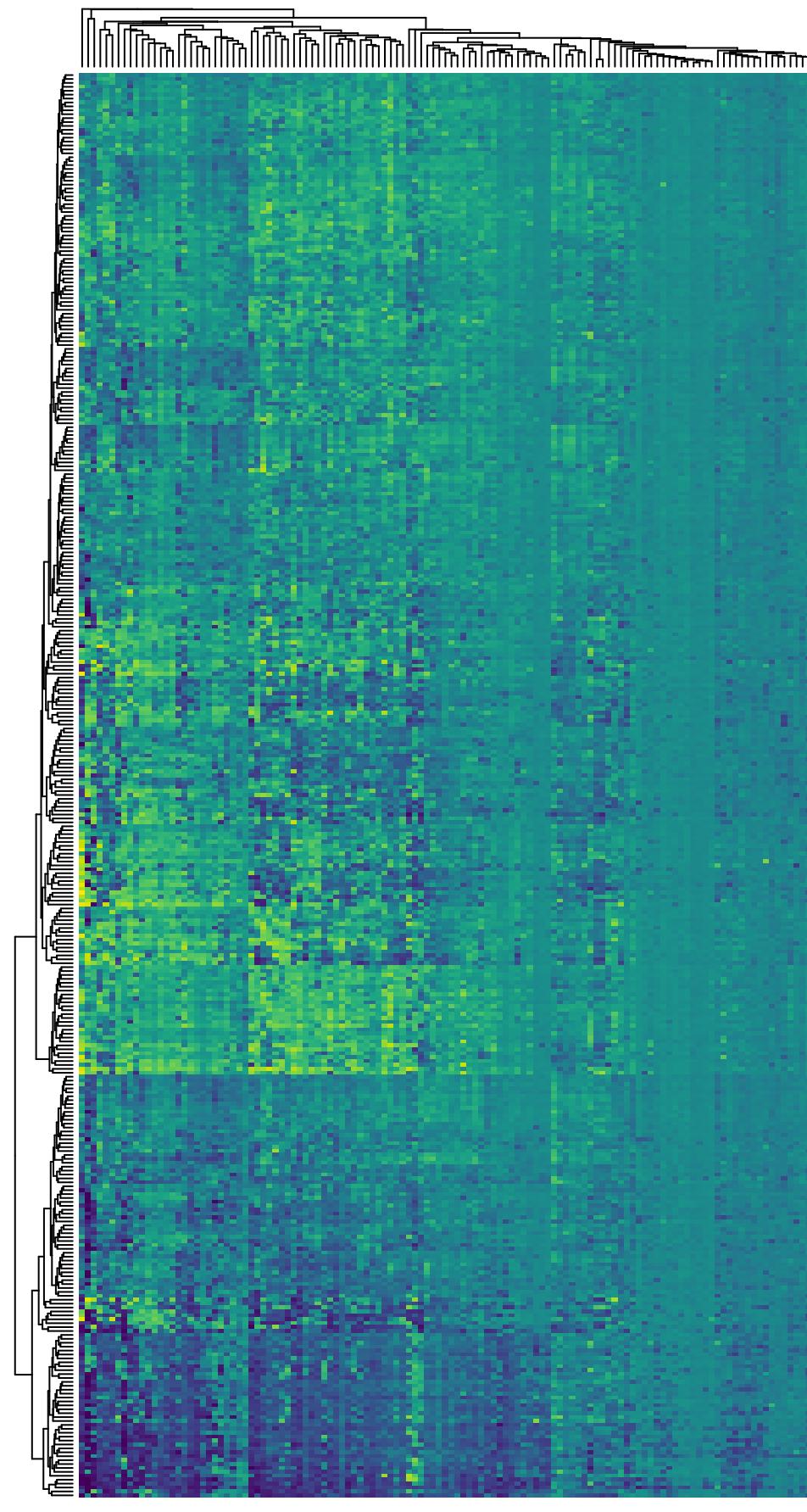
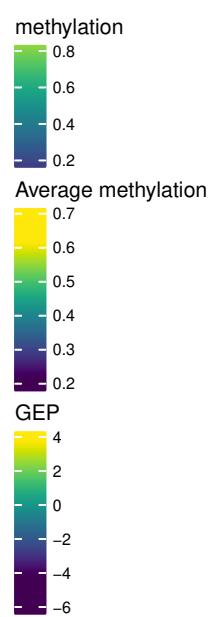
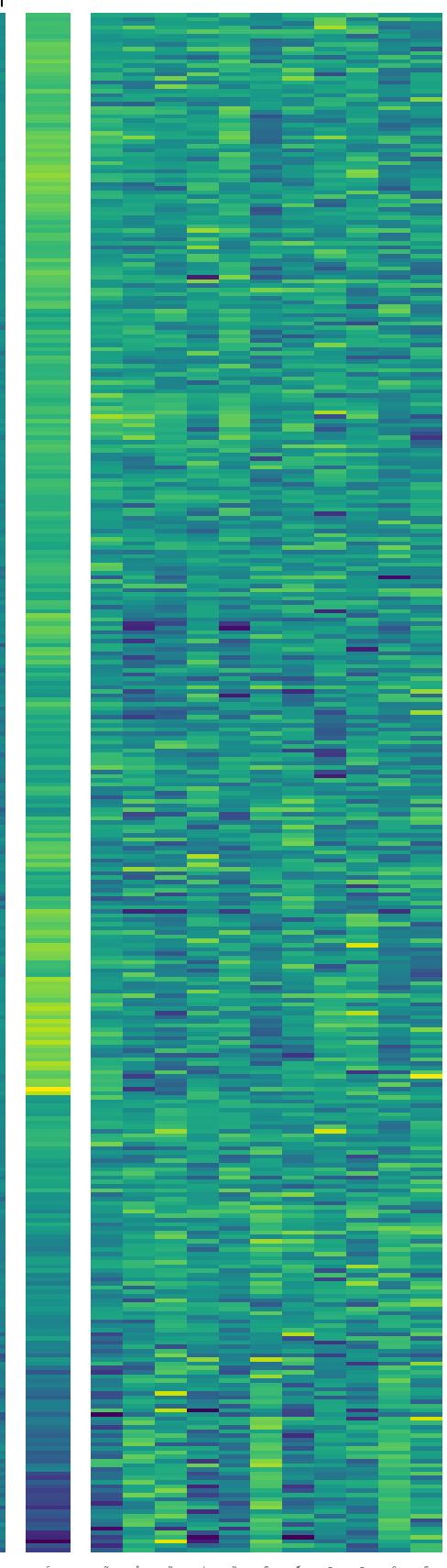


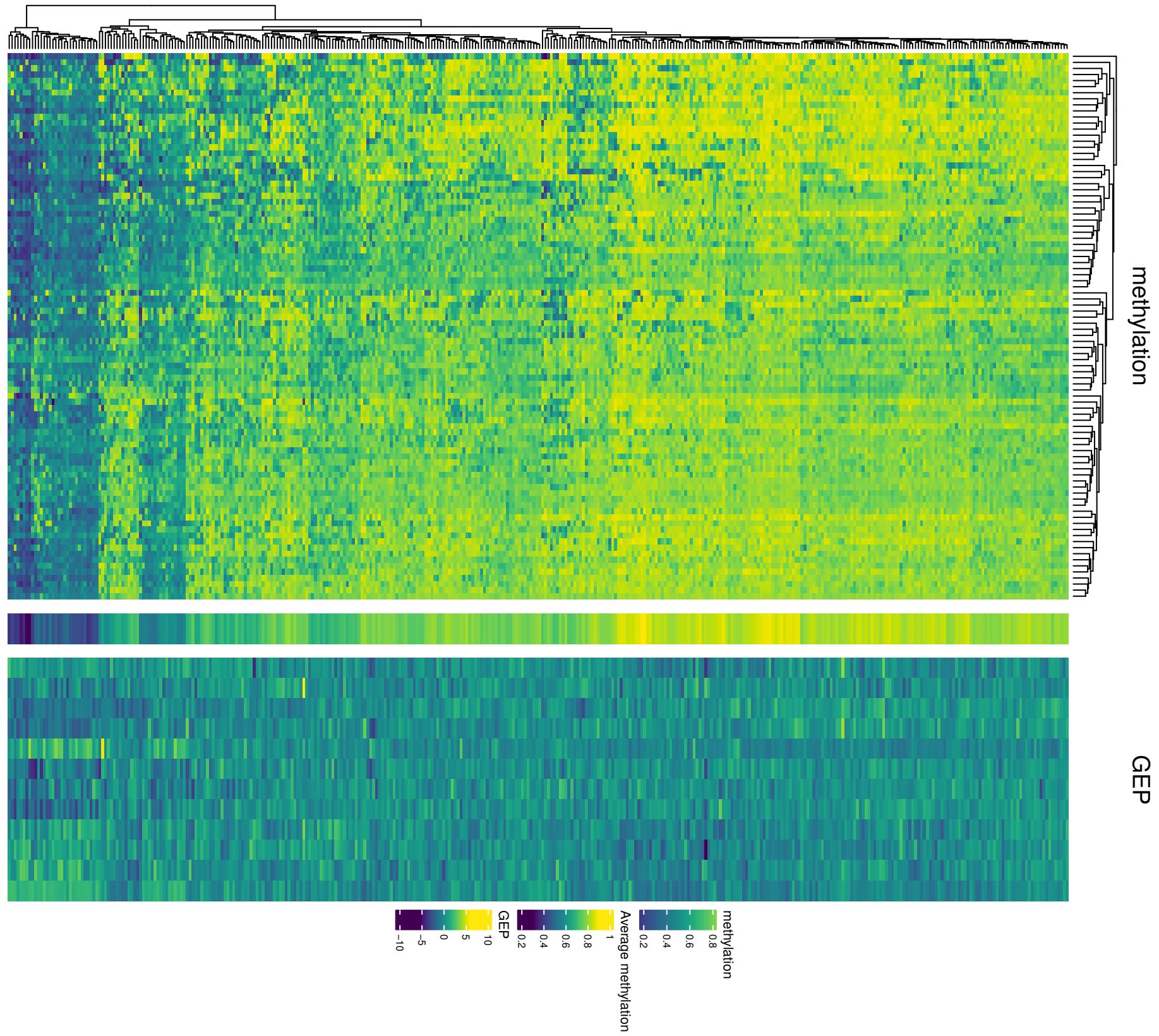
methylation

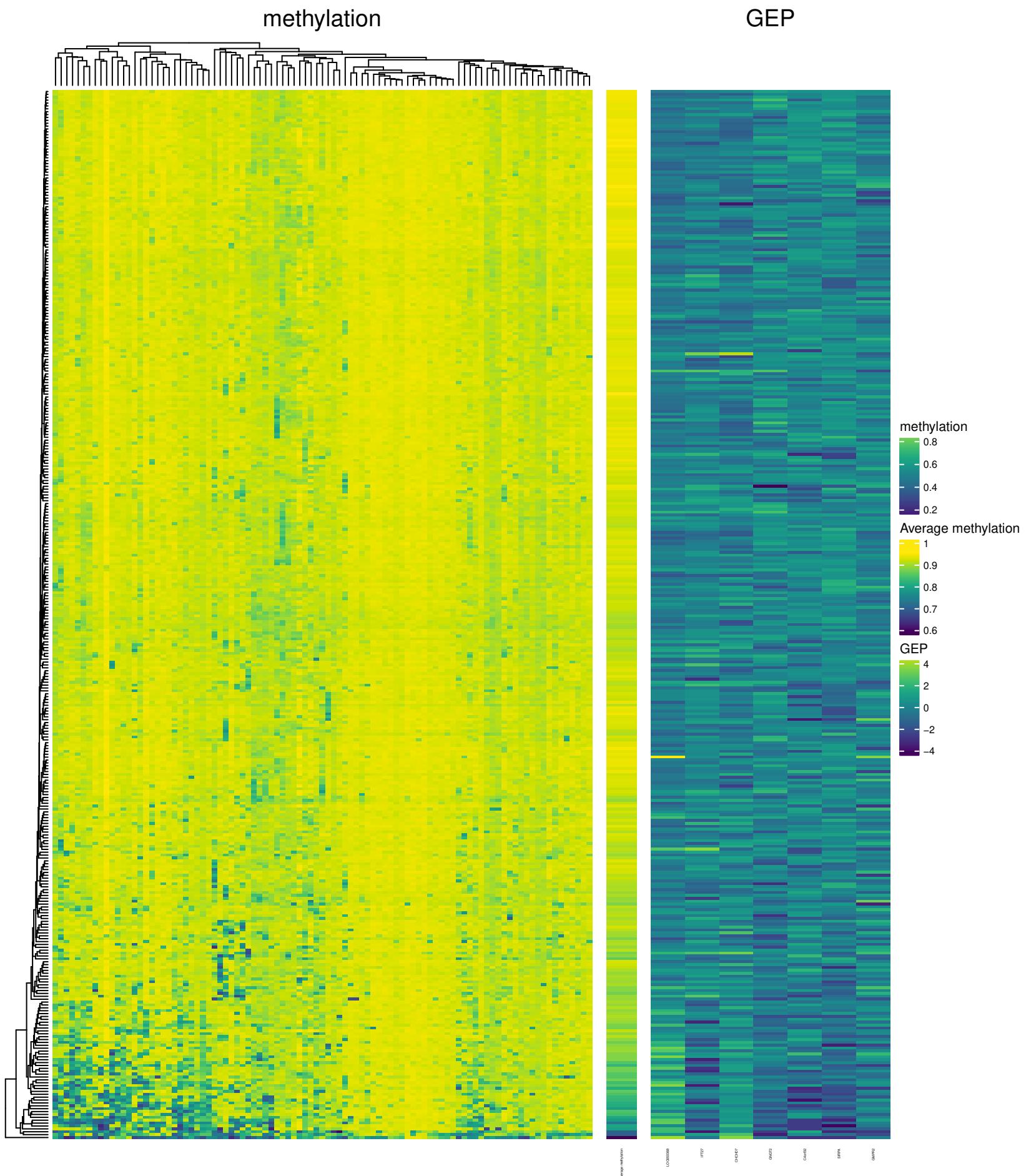


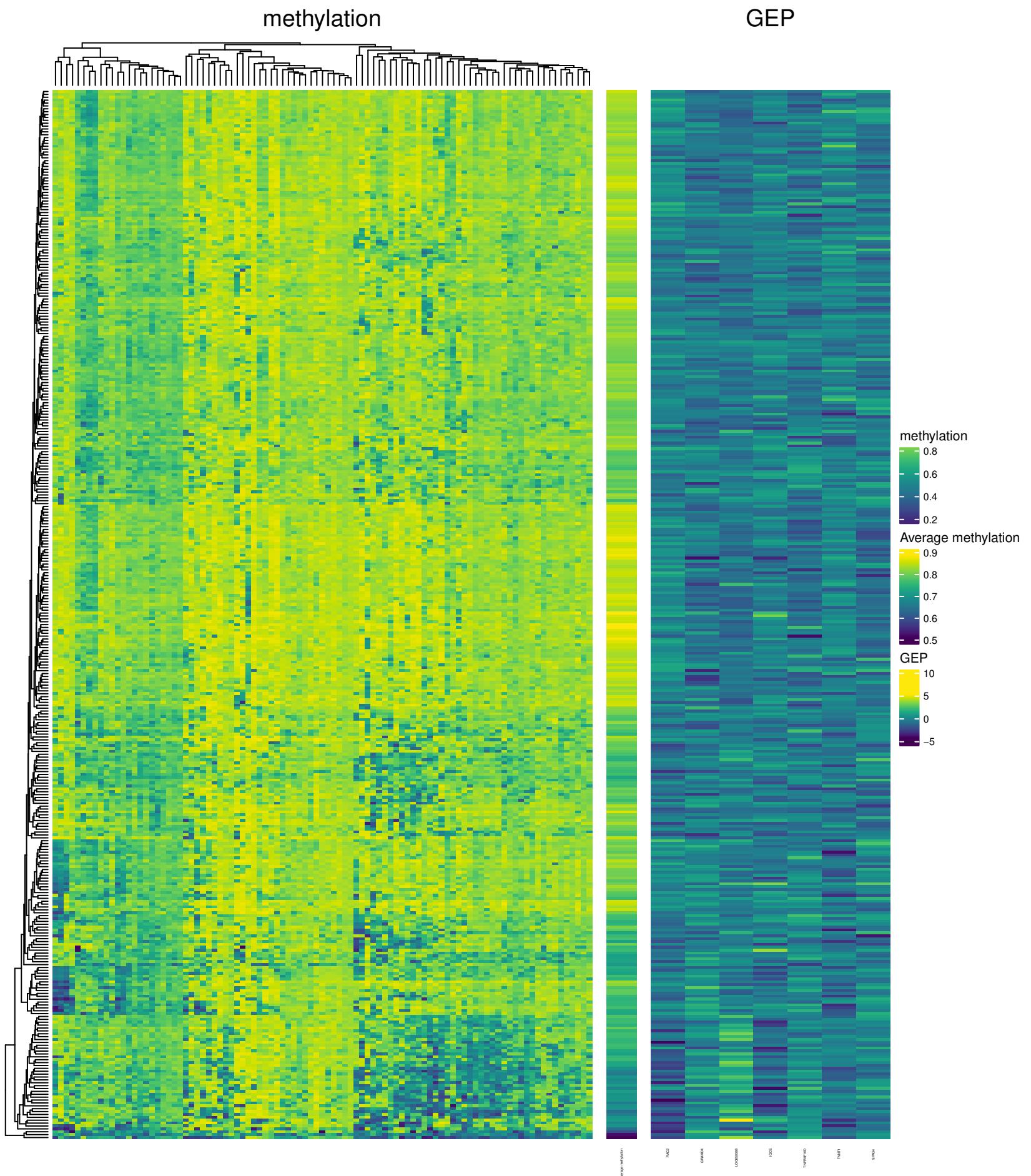
GEP



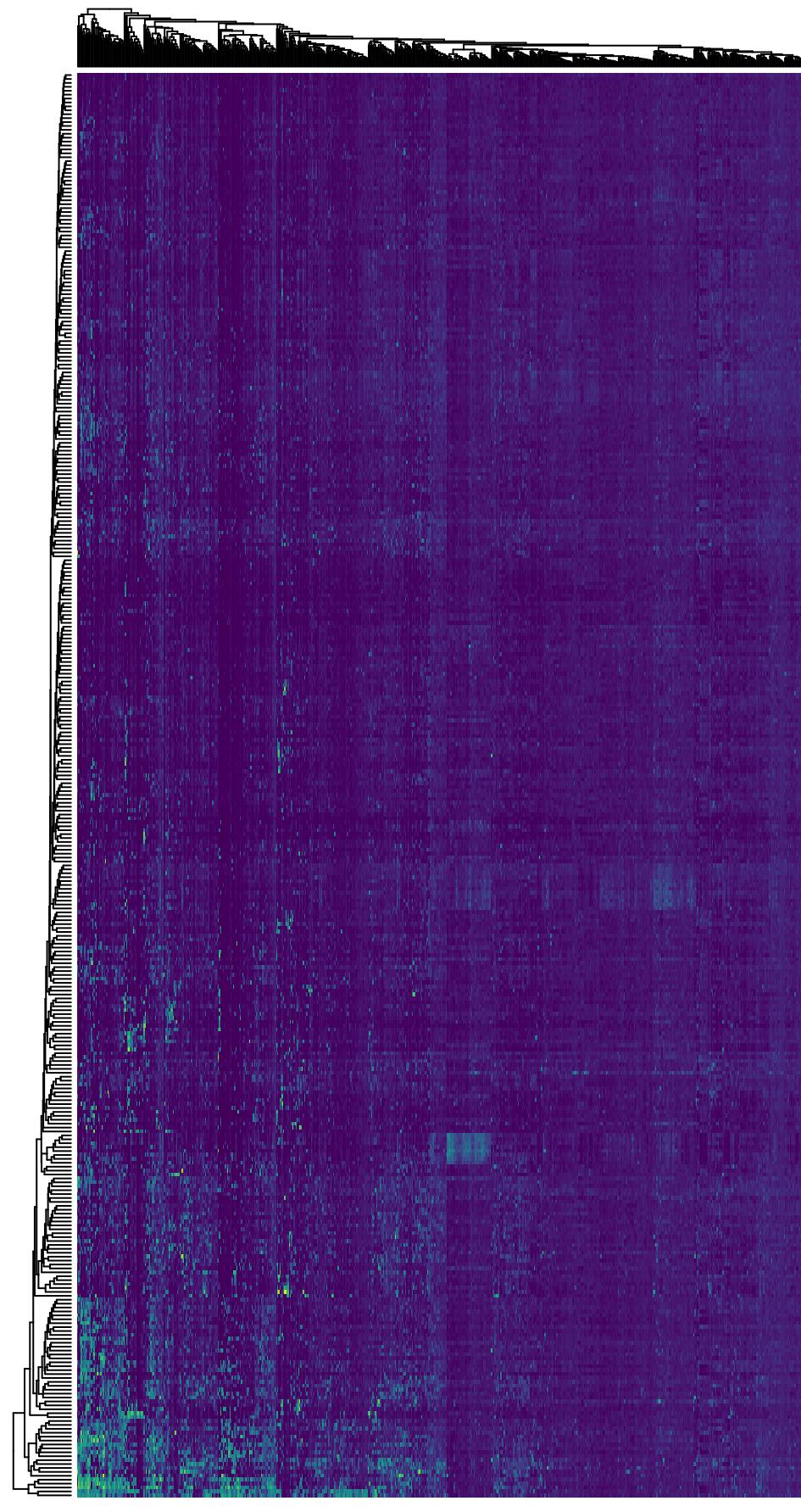
PAC2 F3MAP ACAN D1SL HMG2 SMC3 OLA TSHZ PPT1 GRIN2D IFRD



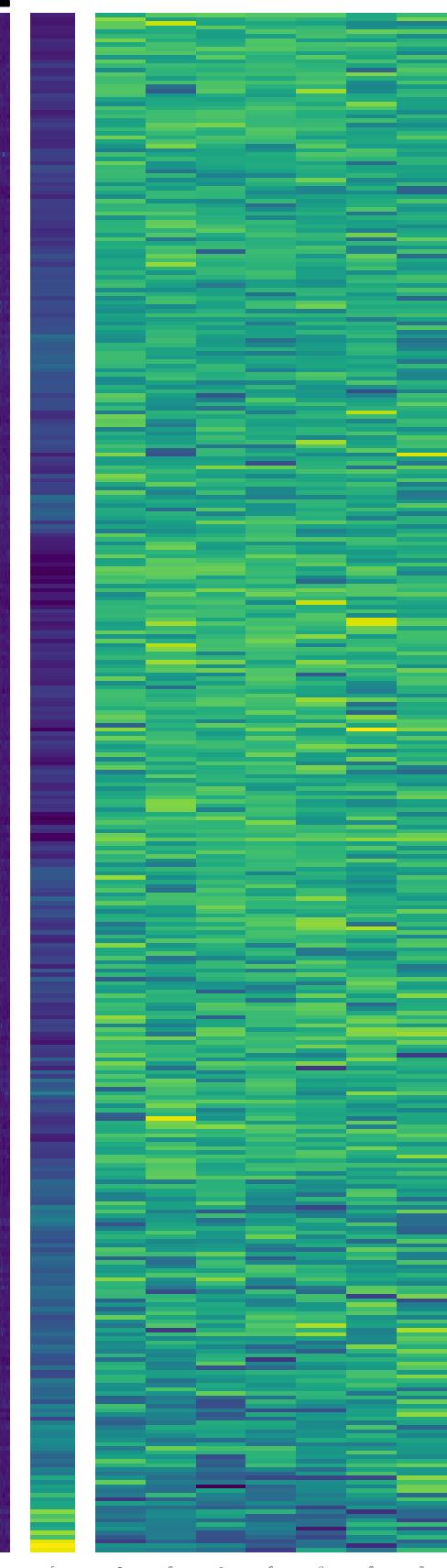




methylation



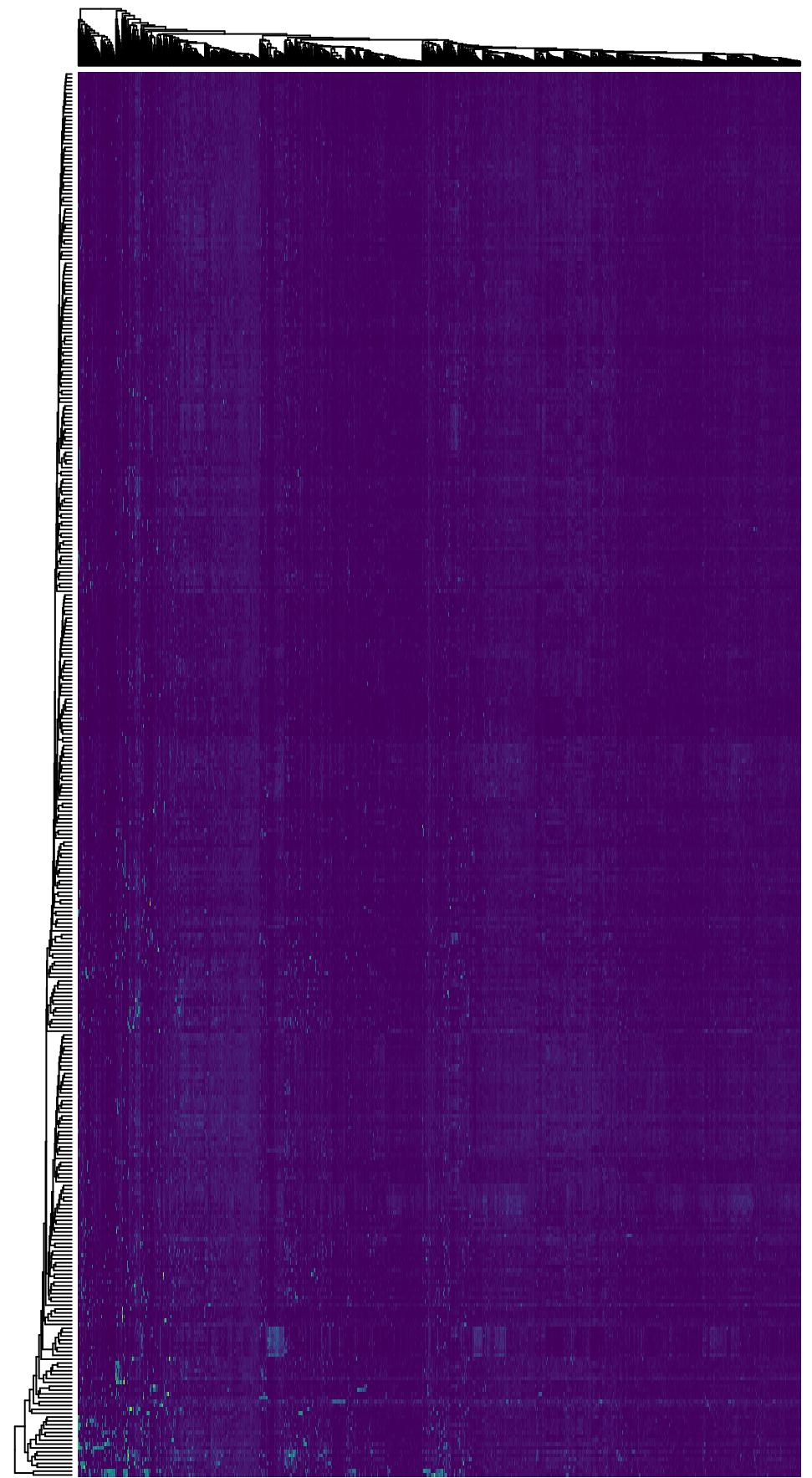
GEP



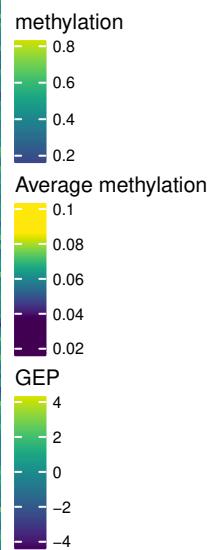
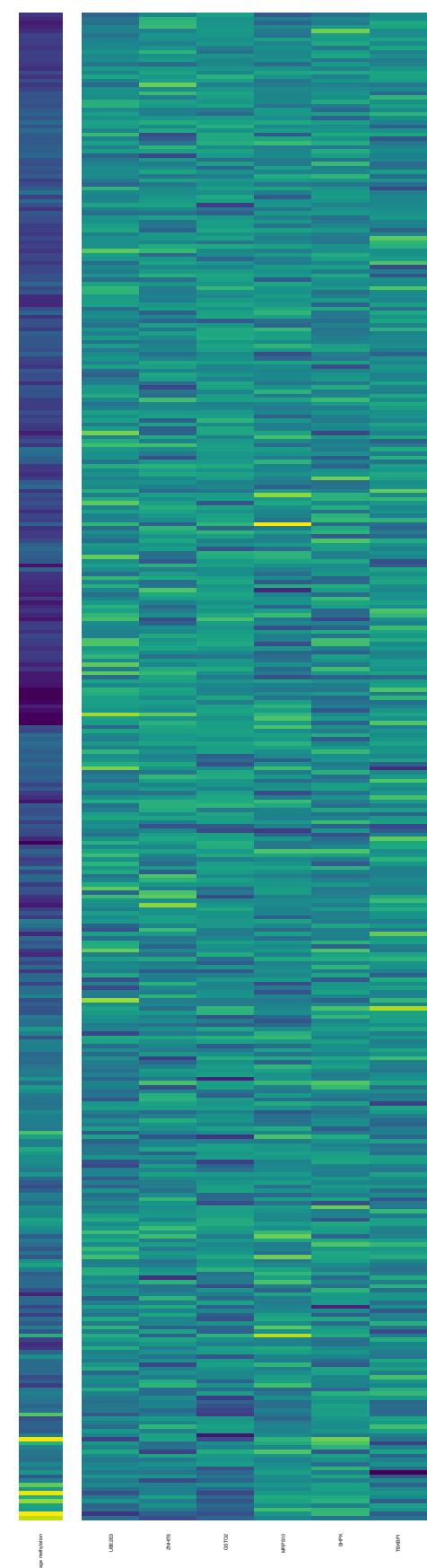
methylation
0.8
0.6
0.4
0.2
0.05
Average methylation
0.2
0.15
0.1
0.05
GEP
4
2
0
-2
-4
-6

F1401 H1405 S1401 P1405 P1407 Z1403 CY1402

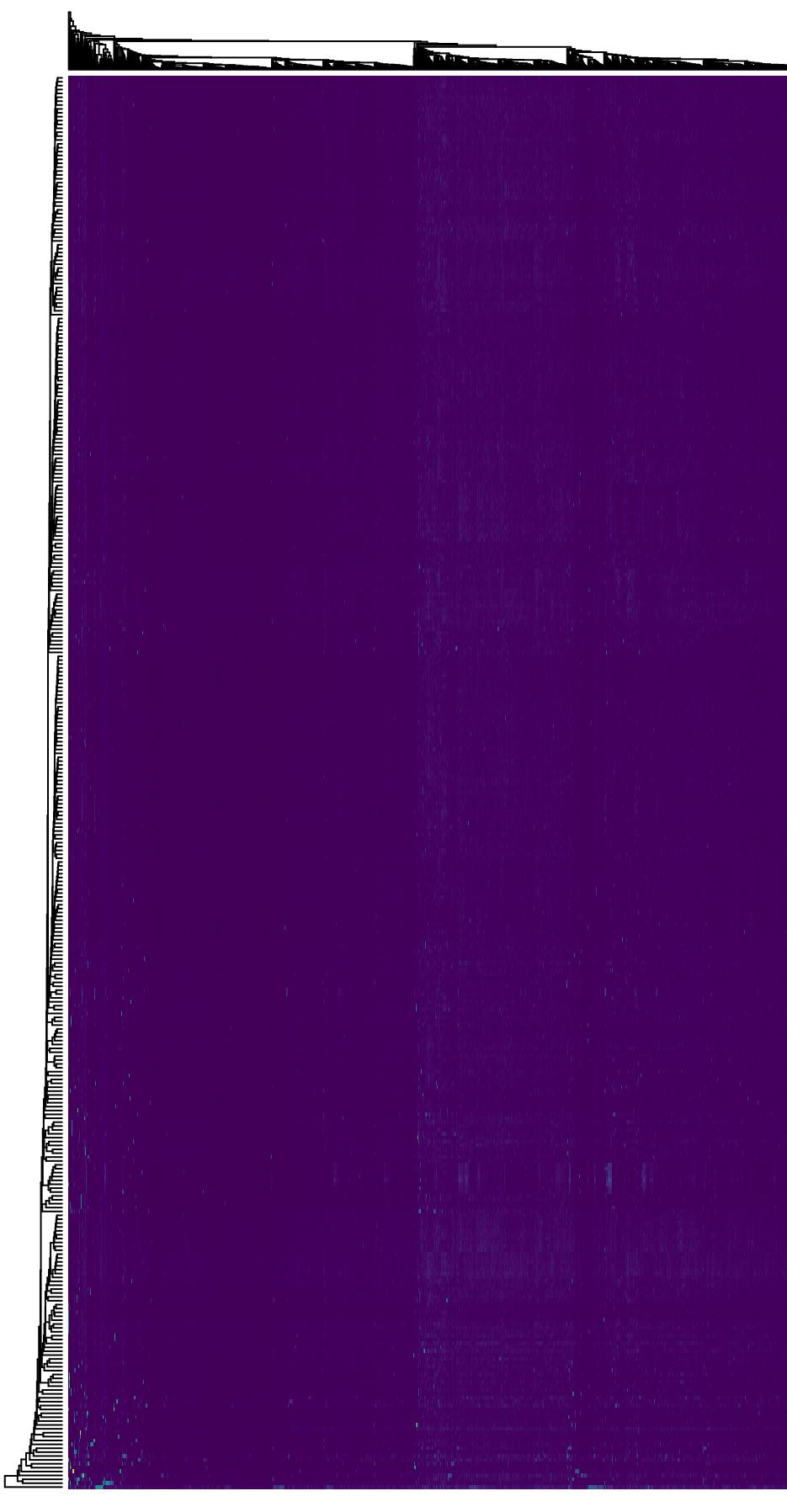
methylation



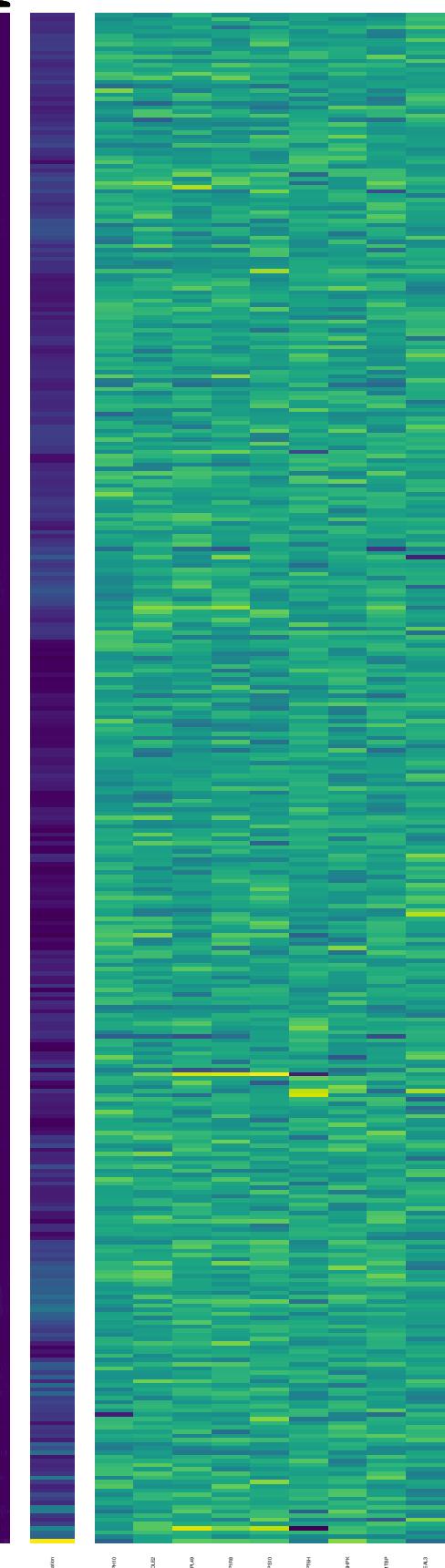
GEP



methylation



GEP



methylation

0.2
0.4
0.6
0.8
1

Average methylation

0.02
0.03
0.04
0.05
0.06
0.07

GEP

-10
-5
0
5

