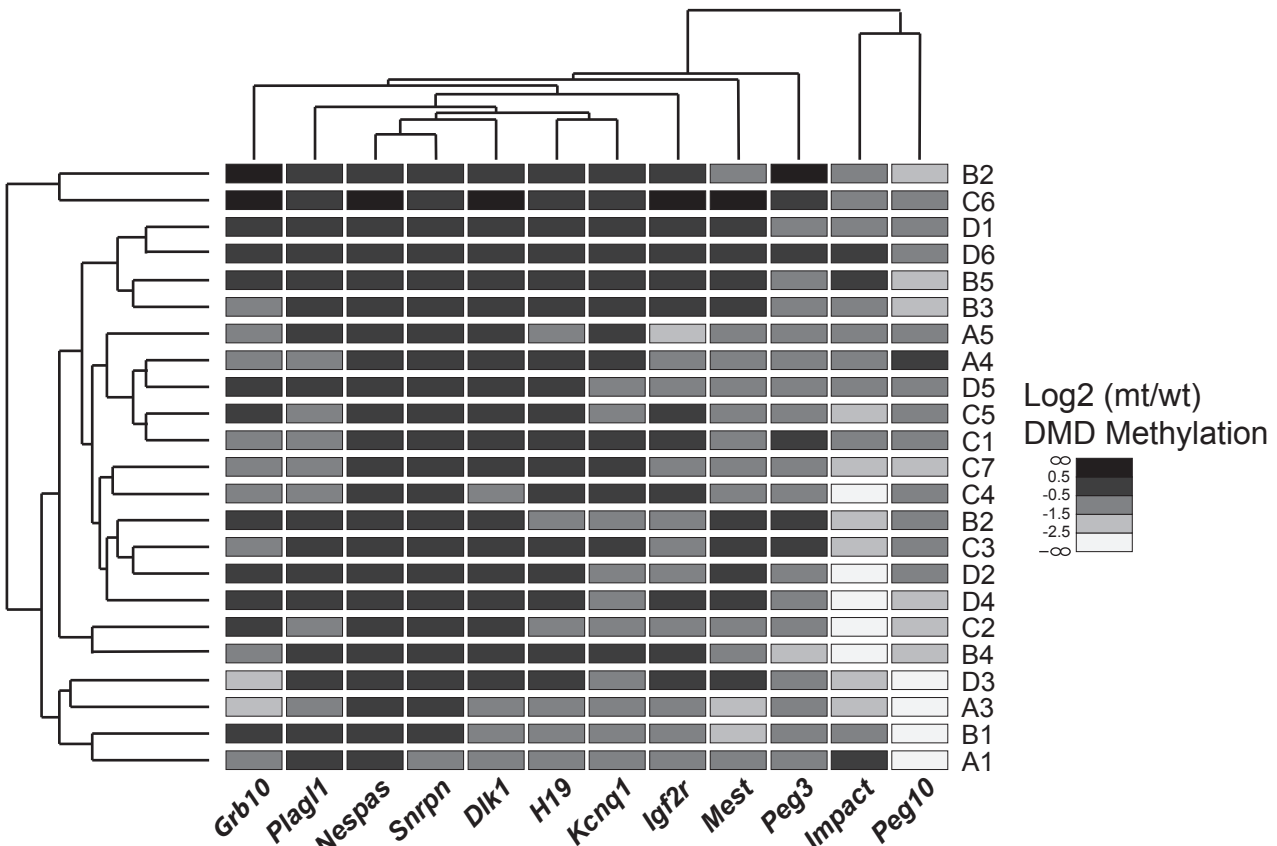
E15.5
B2
C6
D1
D6
B5
B3
A5
A4
D5
C5
C1
C7
C4
B2
C3
D2
D4
C2
B4
D3
A3
B1
A1
Grb10
Plagl1
Nespas
Snrpn
Dlk1
H19
Kcnq1
Igf2r
Mest
Peg3
Impact
Peg10
Log2 (mt/wt)
DMD Methylation
∞
0.5
-0.5
-1.5
-2.5
-∞