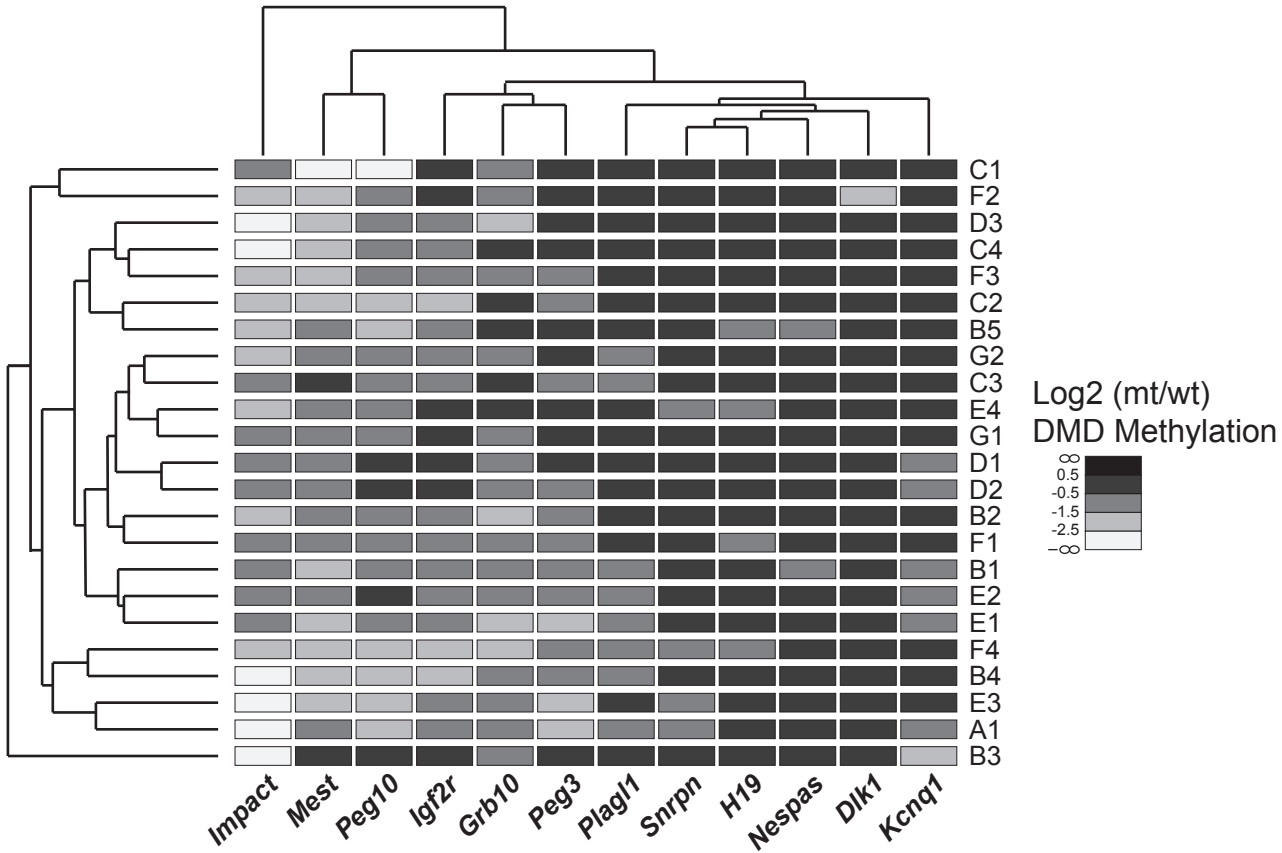
E17.5
C1
F2
D3
C4
F3
C2
B5
G2
C3
E4
G1
D1
D2
B2
F1
B1
E2
E1
F4
B4
E3
A1
B3
Impact
Mest
Peg10
Igf2r
Grb10
Peg3
Plagl1
Snrpn
H19
Nespas
Dlk1
Kcnq1
Log2 (mt/wt)
DMD Methylation
∞
0.5
-0.5
-1.5
-2.5
−∞