

Complete linkage, euclidean
cluster distance

21
18
15
12
9
6
3
0

MTOR (H6)

TGFBFR1 (M4)

RIPK4 (G17)

PIPK3RB (K8)

Scrambled (J2)

Scrambled (H2)

Scrambled (G23)

Scrambled (L1)

