reconstructed (row-normalized) genotype genotype Control cluster ENSMUSG00000107383.1
Atp5a1
Sec61b
ENSMUSG00000114184.1
ENSMUSG00000109610.1
ENSMUSG00000110966.1
ENSMUSG00000110966.1
ENSMUSG000001109762.1
ENSMUSG0000011087593.1
ENSMUSG00000115593.1 Mutant 2 1 SHISTAGE GOOODOO 108772 1
SHISTAGE GOOODOO 115593.1
SHISTAGE GOOODOO 115593.1
SHISTAGE GOOODOO 110469.1
SHISTAGE GOOODOO 110469.1
SHISTAGE GOOODOO 113061.1
SHISTAGE GOOODOO 113061.1
SHISTAGE GOOODOO 113061.1
SHISTAGE GOOODOO 113073.1
SHISTAGE GOOODOO 113073.1
SHISTAGE GOOODOO 113073.1
SHISTAGE GOOODOO 1108073.1
SHISTAGE GOOODOO 1108073.1
SHISTAGE GOOODOO 113070.1
SHISTAGE GOOODOO 110275.1
SHISTAGE GOOODOO 110275.1
SHISTAGE GOOODOO 113075.1
SHISTAGE GOOODOO 110077.1
SHISTAGE GOOODOO 110077.1 0 -1 cluster 2 3 -3 4 5 6 7 ENSMINSG00000116972:1 Gm8730 Gm10076 Wdr89 ENSMUSG00000115194.1 ENSMUSG00000112697.1 ENSMUSG00000112697.1 Msb10 Gm26384 ENSMUSG00000114127.1 ENSMUSG00000114127.1 ENSMUSG00000114127.1 ENSMUSG00000112908.1 Actb Pkm Lrrc58 Oaz1 Atp5g2 Gm10175 Hmgn2 Gm10282 H2afz Ran Nme1 Npm1 Srsf3 Hint1 Hmgb1 Himidal Papinal Funal Gem838398 Gem6743 Sparc Sparc Siln5 Polyrp1 Gatm Pigtrn Tifrc Sicr40a1 Sicr40a1 Beg Fos Eose & Silc646 Silc646 cluster