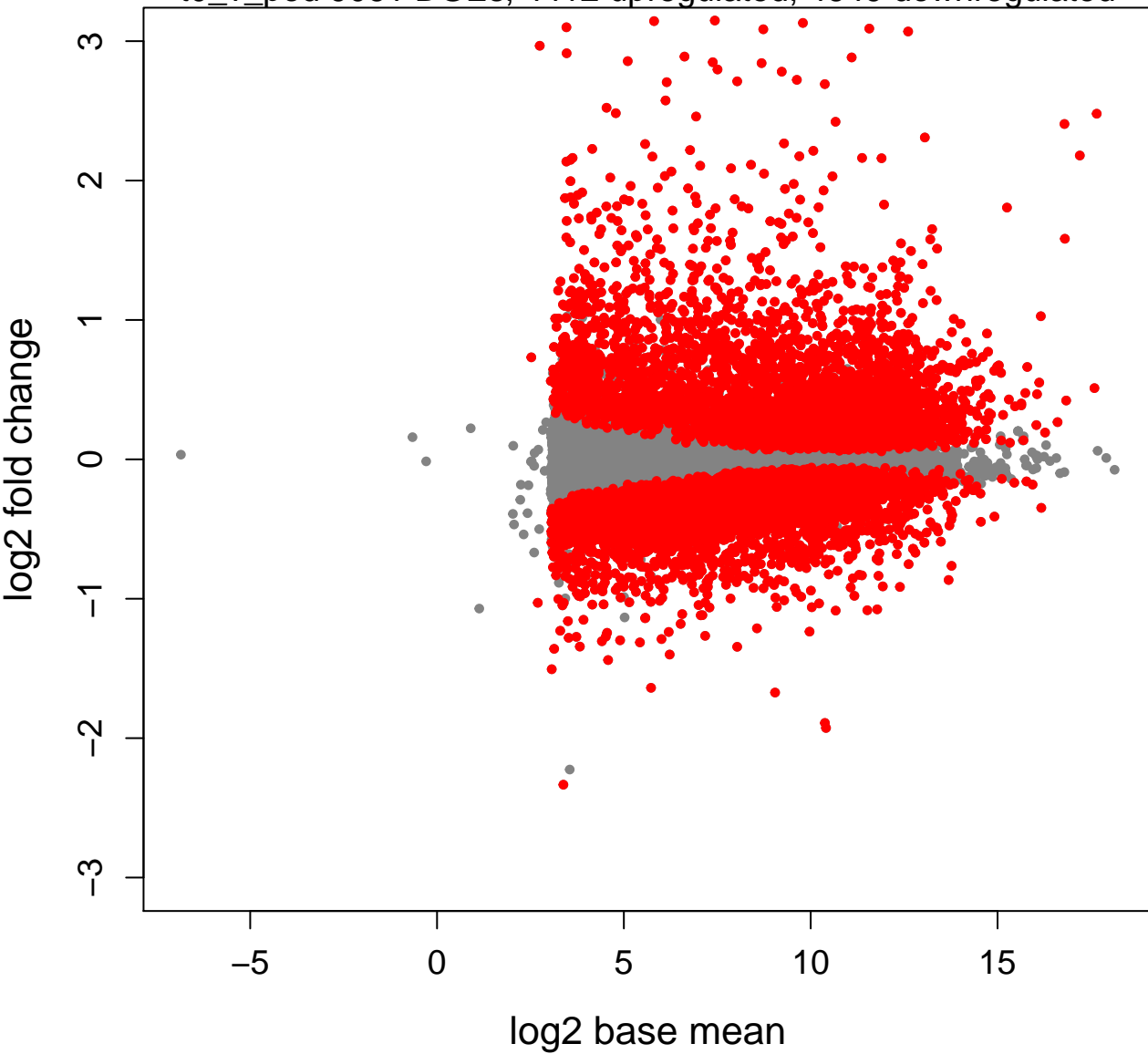
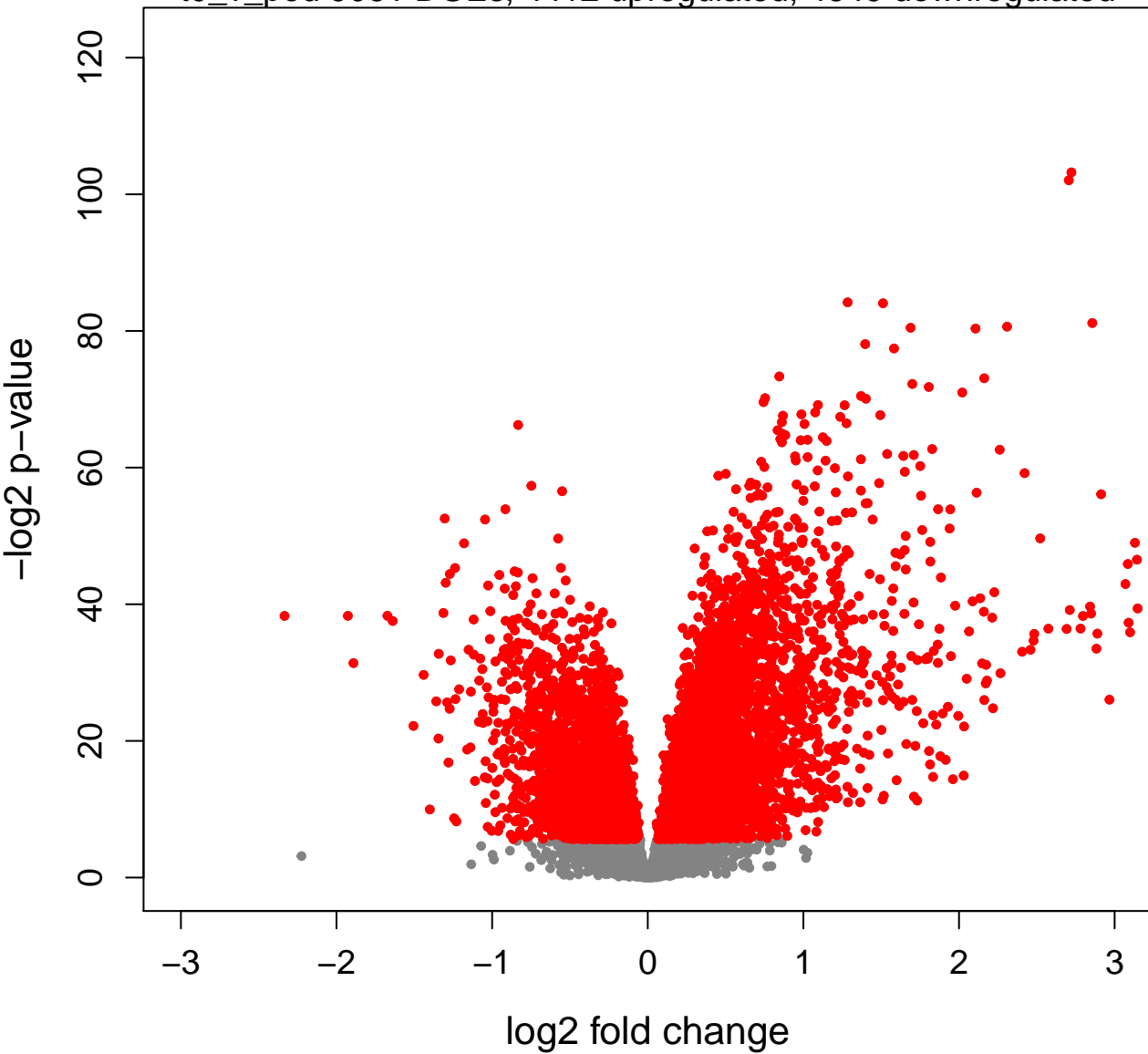


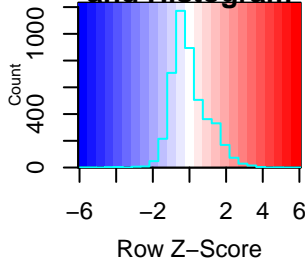
t0_v_pod 9061 DGEs, 4412 upregulated, 4649 downregulated



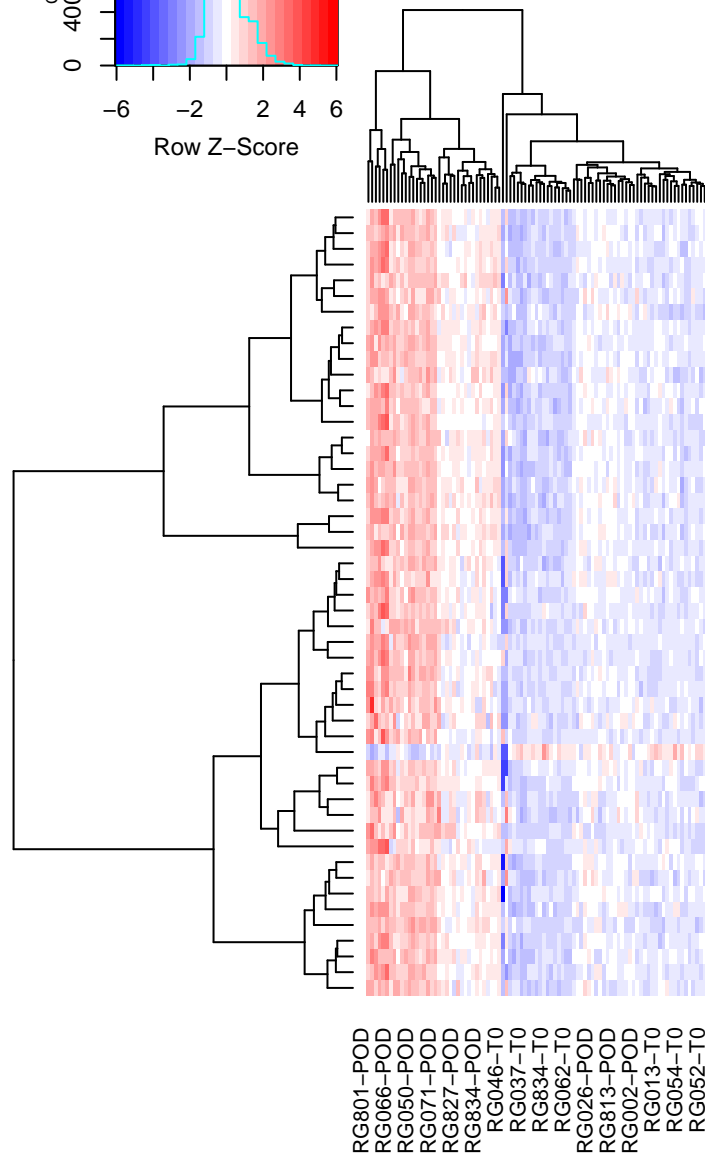
t0_v_pod 9061 DGEs, 4412 upregulated, 4649 downregulated



Color Key and Histogram



Top 50 genes



ENSG00000197006.14 METTL9
 ENSG00000138061.12 CYP1B1
 ENSG00000203710.11 CR1
 ENSG00000163221.9 S100A12
 ENSG00000164125.15 GASK1B
 ENSG00000130830.15 MPP1
 ENSG00000184557.4 SOCS3
 ENSG00000141401.12 IMPA2
 ENSG00000138413.13 IDH1
 ENSG00000135838.13 NPL
 ENSG00000161944.16 ASGR2
 ENSG00000109743.11 BST1
 ENSG00000146592.17 CREB5
 ENSG00000183019.7 MCEMP1
 ENSG00000064601.18 CTSA
 ENSG00000198053.11 SIRPA
 ENSG00000121316.11 PLBD1
 ENSG00000010327.10 STAB1
 ENSG00000125810.10 CD93
 ENSG00000163220.11 S100A9
 ENSG00000038427.16 VCAN
 ENSG00000143546.10 S100A8
 ENSG00000112290.13 WASF1
 ENSG00000131187.9 F12
 ENSG00000163251.4 FZD5
 ENSG00000276644.5 DACH1
 ENSG00000166033.13 HTRA1
 ENSG00000170439.7 METTL7B
 ENSG00000132170.21 PPARG
 ENSG00000136160.17 EDNRB
 ENSG00000157064.11 NMNAT2
 ENSG00000129538.14 RNASE1
 ENSG00000196747.4 H2AC13
 ENSG00000137869.15 CYP19A1
 ENSG00000136052.9 SLC41A2
 ENSG00000232973.13 CYP1B1-AS1
 ENSG00000256751.6 PLBD1-AS1
 ENSG00000187837.3 H1-2
 ENSG00000047597.7 XK
 ENSG00000087116.16 ADAMTS2
 ENSG00000204936.10 CD177
 ENSG00000008394.13 MGST1
 ENSG00000110013.13 SIAE
 ENSG00000014257.16 ACP
 ENSG00000156414.19 TDRD9
 ENSG00000108950.12 FAM20A
 ENSG00000168615.12 ADAM9
 ENSG00000186205.13 MARC1
 ENSG00000163803.13 PLB1
 ENSG00000169385.3 RNASE2