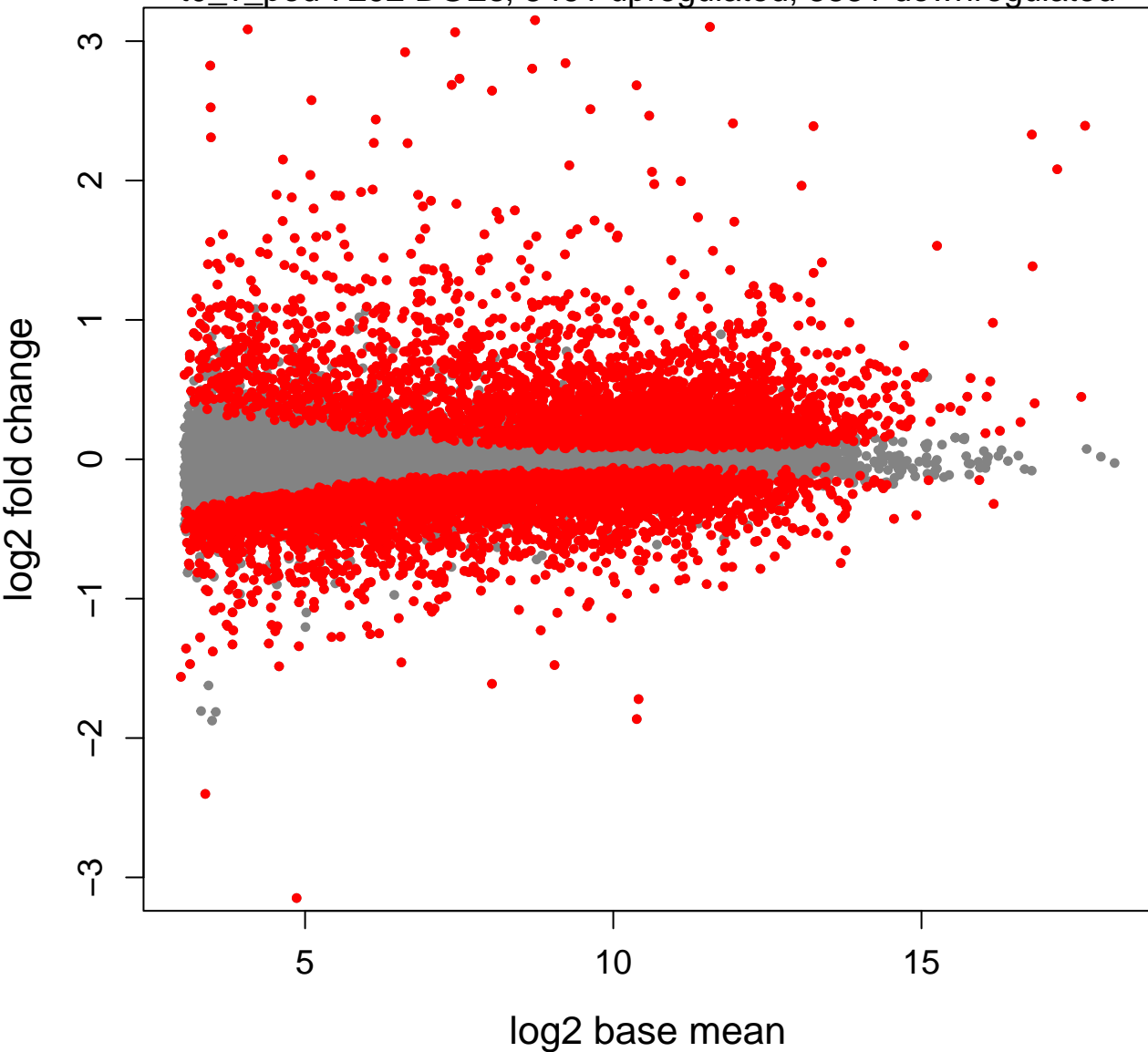
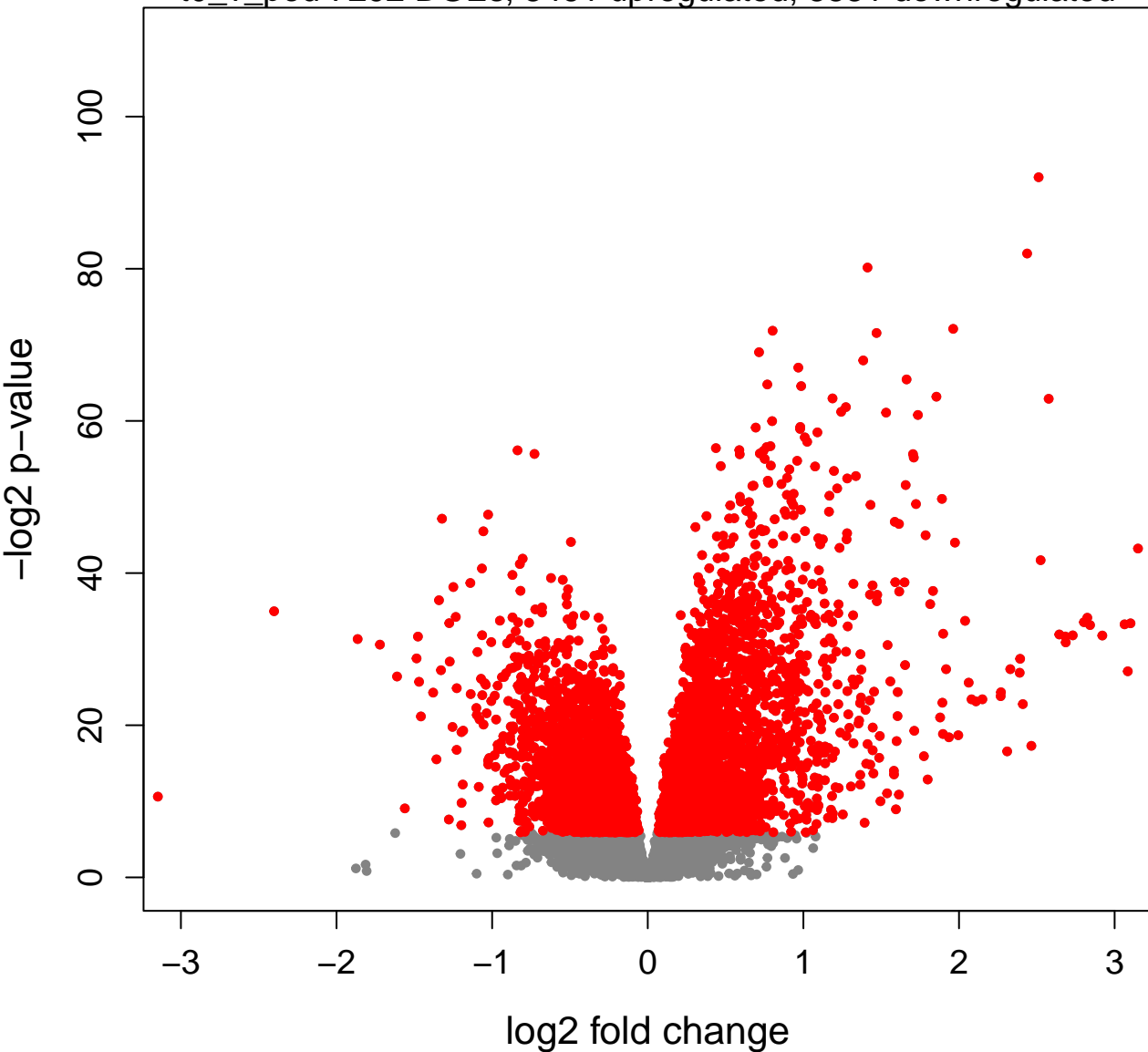


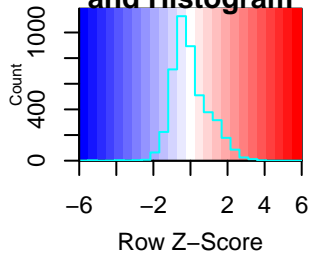
t0_v_pod 7292 DGEs, 3461 upregulated, 3831 downregulated



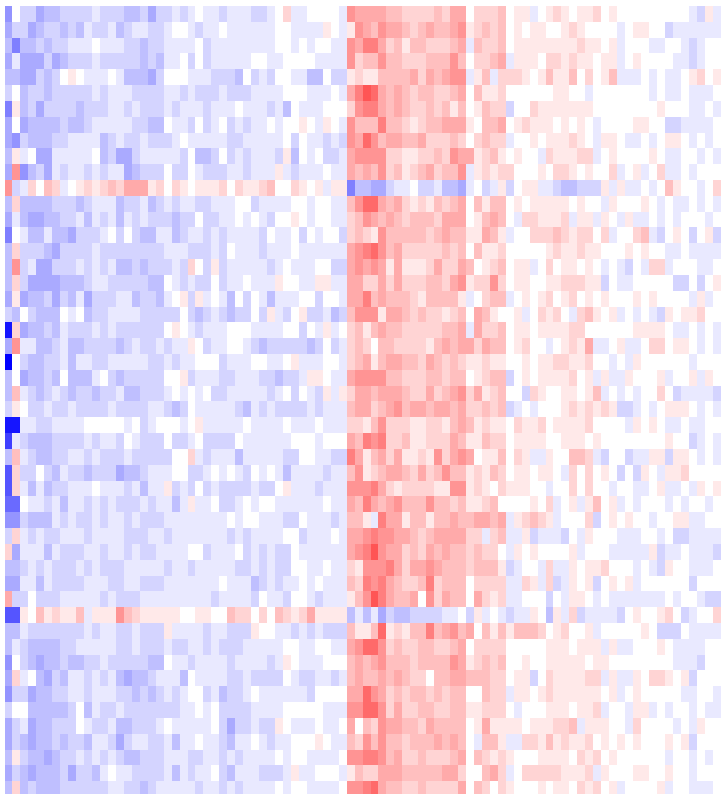
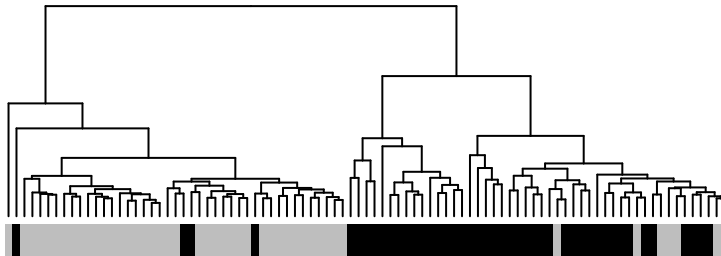
t0_v_pod 7292 DGEs, 3461 upregulated, 3831 downregulated



Color Key and Histogram



Top 50 genes



ENSG00000107745.19 MICU1
 ENSG00000138413.13 IDH1
 ENSG00000141401.12 IMPA2
 ENSG00000135838.13 NPL
 ENSG00000161944.16 ASGR2
 ENSG00000183019.7 MCMP1
 ENSG00000168615.12 ADAM9
 ENSG00000186205.13 MARC1
 ENSG00000163803.13 PLB1
 ENSG00000159348.13 CYB5R1
 ENSG00000140848.17 CPNE2
 ENSG00000148019.13 CEP78
 ENSG00000197006.14 METTL9
 ENSG00000138061.12 CYP1B1
 ENSG00000164125.15 GASK1B
 ENSG00000163221.9 S100A12
 ENSG00000130830.15 MPP1
 ENSG00000143162.9 CREG1
 ENSG00000146592.17 CREB5
 ENSG00000184557.4 SOCS3
 ENSG00000008394.13 MGST1
 ENSG00000110013.13 SIAE
 ENSG0000014257.16 ACP
 ENSG00000156414.19 TDRD9
 ENSG00000169385.3 RNASE2
 ENSG00000108950.12 FAM20A
 ENSG00000103876.12 FAH
 ENSG00000256751.6 PLBD1-AS1
 ENSG00000187837.3 H1-2
 ENSG00000112290.13 WASF1
 ENSG00000131187.9 F12
 ENSG00000163251.4 FZD5
 ENSG00000166033.13 HTRA1
 ENSG00000170439.7 METTL7B
 ENSG00000132170.21 PPARG
 ENSG00000136160.17 EDNRB
 ENSG00000157064.11 NMNAT2
 ENSG00000137869.15 CYP19A1
 ENSG00000136052.9 SLC41A2
 ENSG000000087116.16 ADAMTS2
 ENSG00000142657.21 PGD
 ENSG00000121316.11 PLBD1
 ENSG000000064601.18 CTSA
 ENSG00000198053.11 SIRPA
 ENSG00000135636.14 DYSF
 ENSG0000010327.10 STAB1
 ENSG00000125810.10 CD93
 ENSG00000163220.11 S100A9
 ENSG000000038427.16 VCAN
 ENSG00000143546.10 S100A8

RG046-T0
 RG866-T0
 RG071-T0
 RG002-T0
 RG818-T0
 RG057-T0
 RG019-T0
 RG049-T0
 RG086-T0
 RG076-T0
 RG031-T0
 RG034-POD
 RG820-T0
 RG050-T0
 RG054-T0
 RG048-T0
 RG801-T0
 RG949-T0
 RG940-T0
 RG408-T0
 RG867-T0
 RG827-T0
 RG048-POD
 RG846-POD
 RG073-POD
 RG052-POD
 RG860-POD
 RG012-POD
 RG013-POD
 RG866-POD
 RG066-POD
 RG822-POD
 RG057-POD
 RG831-POD
 RG048-POD
 RG003-POD
 RG834-POD
 RG827-POD
 RG818-POD
 RG847-POD
 RG862-POD
 RG822-T0
 RG087-T0
 RG023-POD
 RG019-POD
 RG813-POD