97 Coka_g15545. 100 Esil_CBN745 Gonavo_ Sappar_KD0 Pinf_PITG_ Alblai_CC.	Phatr3_EG02211.p1 SET.HG4.0:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA t1 SET.HG4.0:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA t7 SET.HG4.0:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA t7 SET.HG4.0:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA t7 SET.HG4.0:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA t8 t8 SET.HG4.0:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA t8 t8 t8 t8 t8 t8 t8 t
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78 64 59 89 89 89 89 89 89 89 89 89 89 89 89 89	-Rall_EPZ35236 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA
59 96 90 100	**Symmic_Smic39131 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA *Vitbra_CEL99874.1 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA *Manpla_Gene.26883 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA *Phatri_Phatr3_J44204.p1 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA *Tpse_EED86239 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA *Nk52_evm1s193 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA *Cowc_KJE88959 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA *Morvir_scaffold269.g2.t1 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA
98 100 100 100 97 Potfil ETO3	 Morvir_scaffold322.g20.t1 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA Pinf_PITG_00530T0 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA Mcir_OAD06300 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA Crei_PNW73786 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA Bralan_BL09079_cuf1 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA Nvec_NVE22957 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA Spis_XP_022805134_1 SET.HG4.1:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA
Retfil_ETO2 99 -Retfil_ETO2 -Mp -Klenit	S918 SET.HG4.2:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA 8807 SET.HG4.2:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA 901 SET.HG4.2:like:ASH1L/KMT2E/NSD1/NSD2/NSD3/SETD2/SETD5:likeclu:5/9/10/11 NA Cvel_10645.t1-p1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA us_XP_003062753.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Otau_XP_022839927.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA _GAQ86441.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA abra_GBG92028 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Marpol_PTQ44251 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA
99	
100	95 Smoe_EFJ08743 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Atri_ERN12742 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Bdis_KQJ95145 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Sbic_EES14647 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Atha_AT4G15180.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Mgut_XP_012852433.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Acoe_PIA58436.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Acoe_PIA58439.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA
99	Aurlim_84477 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Plabra_CEO99192 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA
80 96 100 100 100 100 100 100 100 100 100 10	Retfil_ETO06287 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Retfil_ETO04525 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Clim_evm11s134 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Cfra_CFRG5212T1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Sarc_Sarc4_g23026T SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Aplker_53819 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Schagg_62231 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA
92 99 100	Aurlim_137975 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Aplker_99154 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Coka_g11223.t1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Esil_CBN77807 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Tpse_EED92469 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Phatri_Phatr3_J44935.p1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA Caulen_g245.t1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA
99	-Crei_PNW71011 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Vcar_XP_002951499.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Otau_XP_003080081.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Mpus_XP_003061887.1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Caulen_g5387.t1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Caulen_g5387.t1 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Ptet_CAK79050 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA -Ptet_CAK92634 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA
31 98 100 56 56	Ptet_CAK61316 SET.HG4.3:like:KMT2E/SETD5:likeclu:5 NA a_CEO95538 SET.HG4.4:like:KMT2E/SETD5:likeclu:5 NA —Ncra_EAA30745 SET.HG4.4:like:KMT2E/SETD5:likeclu:5 NA —Gthe_EKX50551 SET.HG4.4:like:KMT2E/SETD5:likeclu:5 NA —Aurlim_63251 SET.HG4.4:like:KMT2E/SETD5:likeclu:5 NA —Honfer_GBG34253 SET.HG4.4:like:KMT2E/SETD5:likeclu:5 NA —Retfil_ETO08498 SET.HG4.4:like:KMT2E/SETD5:likeclu:5 NA —Spimul_27337_TR13611_c0_g1 SET.HG4.4:like:KMT2E/SETD5:likeclu:5 NA
99 99 99 99 99 99 99 99 99 99 99 99 99	-Nutlon_comp34182_c0_seq1.p3 SET.HG4.4:like:KMT2E/SETD5:likeclu:5 NA -Plamic_ANCSI_03482.mRNA.1 SET.HG4.4:like:KMT2E/SETD5:likeclu:5 NA -Symmic_Smic31890 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Ppat_Pp3c10_12470V3.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Sphfal_Sphfalx18G021700.1.p SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Pinf_PITG_01162T0 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Pinf_PITG_03984T0 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Spimul_11975_TR6327_c0_g1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5
56	 Pavlsp_850672 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Chrtob_KOO25319.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Ehux_EOD38084 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Ehux_EOD22319 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Ehux_EOD20363 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Acas_ELR17287 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Plabra_CEO94779 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Hetalb_XP_020438443.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5
93 63 65 72 99 100 1	Mbre_EDQ89227 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Sman_Smp_161010.1_pep SET.HG4.5:KMT2E/SETD5 islike:KMT2E/SETD5 Helrob_HelroP70550 SET.HG4.5:KMT2E/SETD5 islike:KMT2E/SETD5 Helrob_HelroP161339 SET.HG4.5:KMT2E/SETD5 islike:KMT2E/SETD5 Clim_evm9s183 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Dpul_EFX64531 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Dmel_FBpp0075510 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 Tcas_XP_015838651.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5
82 89 89 89 89 89 89 89 89 89 89 89 89 89	**Clyhem_TCONS_00005305-protein SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Clyhem_TCONS_00005305-protein SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Tadh_P61865 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Mlei_ML0113_g2_t1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Sman_Smp_210660.1_pep SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5
93	-Cowc_KJE91036 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Nk52_evm1s136 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Spom_SPAC22E12.11c.1_pep SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Rall_EPZ35345 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Maljak_MALJA_17586.mRNA.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Maljak_MALJA_06000.mRNA.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Coerev_PIA13995.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Scer_YKR029C SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5
77	**Scer_YJL105W SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Rirr_GBC20234 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Rhidel_EIE79055 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Mcir_OAD09197 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Mcir_OAD03829 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Rhidel_EIE92519 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Pbla_XP_018291544.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 **Plebac_Pbla_mRNA_8400 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5
70	-Lcom_lcpid11936 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Scil_scpid9219 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Scil_scpid12689 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Aque_Aqu2.1.42711_001 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Tetwil_Twilhelma_g3394.t1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Emue_0001g505a SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Emue_0001g538a SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 -Auraur_scaffold20.g20.t2 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5
96	*Morvir_scaffold174.g1.t1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 *Nvec_NVE13168 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 *Exapal_XP_028516061.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 *Adig_XP_015765686.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 *Spis_XP_022797365_1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 *Adig_XP_015780794.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 *Adig_XP_015771949.1 SET.HG4.5:KMT2E/SETD5 KMT2E/SETD5 *Ptyfla_pfl_40v0_9_20150316_1g3212.t1 SET.HG4.5:KMT2E/SETD5 SETD5 *Ptyfla_pfl_40v0_9_20150316_1g3212.t1 SET.HG4.5:KMT2E/SETD5 SETD5
9	Lpol_XP_013791181.2 SET.HG4.5:KMT2E/SETD5 KMT2E Lpol_XP_022234954.1 SET.HG4.5:KMT2E/SETD5 KMT2E Lpol_XP_022234955.1 SET.HG4.5:KMT2E/SETD5 KMT2E
83	Apla_XP_022108494.1 SET.HG4.5:KMT2E/SETD5 KMT2E Spur_XP_011676547.2 SET.HG4.5:KMT2E/SETD5 KMT2E Spur_XP_011676552.2 SET.HG4.5:KMT2E/SETD5 KMT2E Spur_XP_011676552.2 SET.HG4.5:KMT2E/SETD5 KMT2E Hsap_ENSP00000257745.4 SET.HG4.5:KMT2E/SETD5 KMT2E Drer_ENSDARP00000112275.2 SET.HG4.5:KMT2E/SETD5 KMT2E Phoaus_g17028.t1 SET.HG4.5:KMT2E/SETD5 KMT2E Linana_g2085.t1 SET.HG4.5:KMT2E/SETD5 KMT2E Notgen_g1146.t1 SET.HG4.5:KMT2E/SETD5 KMT2E
96 100 Ch	**Cgig_EKC30029 SET.HG4.5:KMT2E/SETD5 KMT2E **Cgig_EKC30029 SET.HG4.5:KMT2E/SETD5 KMT2E **Lgig_LotgiP139900 SET.HG4.5:KMT2E/SETD5 KMT2E **Helrob_HelroP151907 SET.HG4.5:KMT2E/SETD5 KMT2E **pur_KAA8500022.1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA **pori_XP_005716732.1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA **racho_PXF48826.1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA **racho_PXF39871.1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA **Aano_EGB11035 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA
100	Coka_g11793.t1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Esil_CBN76570 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA nf_PITG_05564T0 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Pinf_PITG_08614T0 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Pinf_PITG_03320T0 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Pinf_PITG_12923T0 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Pinf_PITG_09695T0 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Pinf_PITG_19210T0 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA
84	Tpse_EED89570 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Pavlsp_571480 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Pavlsp_576813 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Pavlsp_1043453 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Pavlsp_576221 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Tetwil_Twilhelma_g3663.t1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Tetwil_Twilhelma_g24320.t1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Tetwil_Twilhelma_g24879.t1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA
83	Tetwil_Twilhelma_g24879.t1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA -Retfil_ETO06432 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA -Ulvmut_UM019_01651 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA -Chrtob_KOO26547.1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA -Chrtob_KOO26787.1 SET.HG4.6:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA
23	er_28962 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA —Ngad_EWM21040 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Glam_EDO79379 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA —Carmem_5932.p1 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA —Carmem_2347.p1 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA —Carmem_b1468.p1 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA —Tpse_EED89667 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA —Oxytri_EJY66278 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA
100	Tthe_EAR98905 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Ptet_CAK66947 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Ptet_CAK63244 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Ppat_Pp3c24_14110V3.1 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Sphfal_Sphfalx15G103400.1.p SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Sphfal_Sphfalx15G103400.2.p SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Sphfal_Sphfalx15G103400.2.p SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Asacu_v1.1_s0290.g027166 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Azfi_s0030.g024356 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA
	Marpol_PTQ29863 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Sacu_v1.1_s0031.g010677 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Azfi_s0123.g048409 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Acoe_PIA47765.1 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Mgut_XP_012853733.1 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Atha_AT5G09790.2 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Sbic_EES02119 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Bdis_KOK11697 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA
	Bdis_KQJ93013 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Sbic_OQU84251 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Atha_AT5G24330.1 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Mgut_XP_012852024.1 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Acoe_PIA48877.1 SET.HG4.7:like:ASH1L/NSD1/NSD2/NSD3/SETD2:likeclu:9/10/11 NA Whakir_PHAKIR_09359.mRNA.1 SET.HG4.8:like:NSD1/NSD2/NSD3:likeclu:9 NA Whakir_PHAKIR_09359.mRNA.2 SET.HG4.8:like:NSD1/NSD2/NSD3:likeclu:9 NA
73 100 100 73 87	Symmic_Smic32862 SET.HG4.8:like:NSD1/NSD2/NSD3:likeclu:9 NA Pavlsp_595377 SET.HG4.8:like:NSD1/NSD2/NSD3:likeclu:9 NA
60 62 87 Jak	Tmel_CAZ80586 SET.HG4.8:like:NSD1/NSD2/NSD3:likeclu:9 NA Umay_KIS71975 SET.HG4.8:like:NSD1/NSD2/NSD3:likeclu:9 NA
77° 100° 100° 100° 100° 100° 100° 100° 1	■Maljak_MALJA_19997.mRNA.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 ■Maljak_MALJA_04523.mRNA.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 —■Malcal_MALCA_08315.mRNA.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 —Gefoke_MALSP_09708.mRNA.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 —Gefoke_MALSP_09708.mRNA.2 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 —Styinc_Gene47033_m47033 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 —Andgod_ANDGO_07359.mRNA.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 —Jaklib_JAKLI_23089.mRNA.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3
100	akbah_JAKBA_10103.mRNA.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 ecame_RECAM_00230.mRNA.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Secula_SECUL_09573.mRNA.2 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Secula_SECUL_09573.mRNA.3 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 100
78 90 100 100 100 100 100 100 100 100 100	 Honfer_GBG32395 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Aurlim_141495 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Sappar_KDO24656 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Pinf_PITG_05692T0 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Alblai_CCA15310 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Ngad_EWM29679 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Esil_CBJ25830 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Coka_g11301.t1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3
72	Phatri_Phatr3_J6093.p1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Tpse_EED88618 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Cpar_Cpar_evm1s59 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Otau_XP_003082786.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Gracho_PXF45272.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Chocri_XP_005710598.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Ulvmut_UM149_00291 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3
100	-Cvar_XP_005846950.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Cocsub_XP_005651892.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Vcar_XP_002951402.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Crei_PNW80554 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Sappar_KDO32124 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Pinf_PITG_00474T0 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Klenit_GAQ81469.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Chabra_GBG89943 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3
100	**Ppat_Pp3c2_35860V3.10 SE1.HG4.9:NSD1/NSD2/NSD3 Islike:NSD1/NSD2/NSD3 **Sphfal_Sphfalx02G020500.1.p SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 **Atha_AT4G30860.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 **Mgut_XP_012837814.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 **Acoe_PIA46038.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 **Atri_ERN07128 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3
90	-Sbic_KXG30618 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Bdis_KQK00294 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Bdis_KQJ98371 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Sbic_OQU80550 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Chabra_GBG86446 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Smoe_EFJ19814 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Smoe_EFJ04637 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 -Marpol_PTQ48324 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3
	Sphfal_Sphfalx17G078500.1.p SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Ppat_Pp3c4_29810V3.10 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Ppat_Pp3c4_29810V3.10 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Ppat_Pp3c26_12150V3.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Sacu_v1.1_s0122.g021471 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Azfi_s0090.g042624 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Sacu_v1.1_s1053.g027927 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Sacu_v1.1_s0001.g000771 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Atri_ERN04535 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3
	Sbic_OQU89246 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Bdis_PNT64346 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Bdis_KQK15084 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Acoe_PIA52930.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Atha_AT2G44150.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Atha_AT3G59960.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Mgut_XP_012830182.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Mgut_XP_012829063.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3
93	**Plabra_CEO95530 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 SET.HG4.9:NSD1/NSD2/NSD3 SET.HG4.9:NSD1/NSD2/NSD3 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 SET.HG4.9:NSD1/NSD2/NSD3 SET.HG4.9:NSD1/NSD2/NSD3 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 NK52_evm33s24 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Rigram_Gene.26561 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Sarc_Sarc4_g31219T SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3
74 70 70 100 100 100 100 100 100 100 100 1	Cfra_CFRG0207T1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Nutlon_comp17141_c0_seq1.p1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Nutlon_comp17141_c1_seq1.p1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Sros_EGD78296 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Mbre_EDQ87845 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Spimul_18950_TR10098_c0_g1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Aque_Aqu2.1.38449_001 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Tetwil_Twilhelma_g23087.t1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3
	 Emue_0013g1106a SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Lcom_lcpid32057 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Scil_scpid16325 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Ocar_g3171.t1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Mvib_comp10766_c1_seq1_fr4 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Emul_CDS41338.1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Sman_Smp_137060.1_pep SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Mlei ML0336 g25 t1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3
91 96 21 93	■Tadh_P31338 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 ■Tcas_XP_008198269.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 ■Dpul_EFX71479 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 ■Dmel_FBpp0084636 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 ■Hduj_OQV21199.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 ■Ramvar_GAV05823.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 ■Cpar_Cpar_evm1s5587 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3
17	Sman_Smp_160700.1_pep SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 =Emul_CDS39953.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 =Hydvul_Sc4wPfr_138.g13930.t1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 =Morvir_scaffold171.g5.t1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 =Auraur_scaffold1618.g1.t1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 =Spis_XP_022779411_1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 =Adig_XP_015754820.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3
93	Adig_XP_015754821.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Nvec_NVE19383 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Exapal_XP_020897074.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Exapal_XP_020897075.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Bralan_BL17376_evm0 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Cscu_XP_023226639.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Lpol_XP_013784275.2 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Lpol_XP_022242981.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3
	Apla_XP_022091753.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Spur_XP_030841455.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Spur_XP_030841823.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Ptyfla_pfl_40v0_9_20150316_1g14997.t1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Skow_XP_06812994.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Skow_XP_06812995.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1/NSD2/NSD3 Ctel_CapteP93578 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3 Notgen_g2935.t1 SET.HG4.9:NSD1/NSD2/NSD3 islike:NSD1/NSD2/NSD3
	100
	#Helrob_HelroP110454 SET.HG4.9:NSD1/NSD2/NSD3 NSD3 #Hsap_ENSP00000313983.7 SET.HG4.9:NSD1/NSD2/NSD3 NSD3 #Drer_ENSDARP00000151686.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD3 #Hsap_ENSP00000395929.2 SET.HG4.9:NSD1/NSD2/NSD3 NSD1 #Drer_ENSDARP00000156812.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1 #Drer_ENSDARP00000106822.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD1 #Hsap_ENSP00000372347.5 SET.HG4.9:NSD1/NSD2/NSD3 NSD2 #Drer_ENSDARP00000002944.8 SET.HG4.9:NSD1/NSD2/NSD3 NSD2
100	Too Drer_ENSDARP00000150128.1 SET.HG4.9:NSD1/NSD2/NSD3 NSD2
97 94 95 95	Bden_EGF77213 SET.HG4.10:ASH1L ASH1L Spun_KND00271 SET.HG4.10:ASH1L ASH1L
73 90 100 100	Pbla_XP_018286368.1 SET.HG4.10:ASH1L ASH1L 100 Plebac_Pbla_mRNA_13201 SET.HG4.10:ASH1L ASH1L EKX31239 SET.HG4.10:ASH1L ASH1L ———————————————————————————————————
42	-Esil_CBN74814 SET.HG4.10:ASH1L ASH1L -Coka_g9352.t1 SET.HG4.10:ASH1L ASH1L -Coka_g9352.t2 SET.HG4.10:ASH1L ASH1L -Coka_g9352.t2 SET.HG4.10:ASH1L ASH1L -Coka_g9352.t2 SET.HG4.10:ASH1L ASH1L Mbre_EDQ84292 SET.HG4.10:ASH1L ASH1L Mbre_EDQ84292 SET.HG4.10:ASH1L ASH1L Ngru_EFC42671 SET.HG4.10:ASH1L ASH1L Ramvar_GAV08813.1 SET.HG4.10:ASH1L ASH1L
71	
60 60 57 66 57 66 60 66 60 66 60 66 60 66 60 66 60 66 60 66 60 60	Ptyfla_pfl_40v0_9_20150316_1g5965.t1 SET.HG4.10:ASH1L ASH1L -Apla_XP_022105810.1 SET.HG4.10:ASH1L ASH1L -Spur_XP_030832467.1 SET.HG4.10:ASH1L ASH1L -Spur_XP_030832471.1 SET.HG4.10:ASH1L ASH1L -Bralan_BL25105_evm0 SET.HG4.10:ASH1L ASH1L -Hsap_ENSP00000357330.3 SET.HG4.10:ASH1L ASH1L -Drer_ENSDARP00000052914.6 SET.HG4.10:ASH1L ASH1L -Cscu_XP_023217116.1 SET.HG4.10:ASH1L ASH1L
	Lpol_XP_013779563.1 SET.HG4.10:ASH1L ASH1L Linana_g1428.t1 SET.HG4.10:ASH1L ASH1L Cgig_EKC31816 SET.HG4.10:ASH1L ASH1L Lgig_LotgiP143999 SET.HG4.10:ASH1L ASH1L Phoaus_g4680.t1 SET.HG4.10:ASH1L ASH1L Ctel_CapteP176480 SET.HG4.10:ASH1L ASH1L Helrob_Helrob70629 SET.HG4.10:ASH1L ASH1L Notgen_g13215.t1 SET.HG4.10:ASH1L ASH1L
95	## Notgen_g14626.t1 SET.HG4.10:ASH1L ASH1L ## gram_Gene.38110 SET.HG4.11:SETD2 SETD2 ## Bnat_JGI_V11_85862 SET.HG4.11:SETD2 SETD2 ## Bnat_JGI_V11_72270 SET.HG4.11:SETD2 SETD2 ## Plabra_CEP03120 SET.HG4.11:SETD2 SETD2 ## Acas_ELR22513 SET.HG4.11:SETD2 islike:SETD2 ## Carmem_b10521.p2 SET.HG4.11:SETD2 islike:SETD2 ## Spimul_22204_TR10936_c0_g2 SET.HG4.11:SETD2 islike:SETD2
77° 22° 96° 70° 100° 100° 100° 100° 100° 100° 100°	■Plamic_ANCSI_10688.mRNA.1 SET.HG4.11:SETD2 islike:SETD2 ■Meteor_CRO19MET_NODE_5809.p1 SET.HG4.11:SETD2 islike:SETD2 ■Meteor_CRO19MET_NODE_8184.p1 SET.HG4.11:SETD2 islike:SETD2 ■Ngru_EFC46377 SET.HG4.11:SETD2 islike:SETD2 ■Amobb2_AMBB2_12527.mRNA.1 SET.HG4.11:SETD2 islike:SETD2 ■Andgod_ANDGO_02265.mRNA.1 SET.HG4.11:SETD2 islike:SETD2 ■Styinc_Gene46551_m46551 SET.HG4.11:SETD2 islike:SETD2 ■Phakir_PHAKIR_05159.mRNA.1 SET.HG4.11:SETD2 islike:SETD2
74 100 99	Recame_RECAM_16147.mRNA.1 SET.HG4.11:SETD2 islike:SETD2
97 100 100 71 99 97	■Tielac_KYQ92214 SET.HG4.11:SETD2 islike:SETD2 ■Hetalb_XP_020433149.1 SET.HG4.11:SETD2 islike:SETD2 ■Cavfas_EGG13662 SET.HG4.11:SETD2 islike:SETD2 ■Diprot_Gene.29586 SET.HG4.11:SETD2 islike:SETD2 ■Meteor_CRO19MET_NODE_594.p1 SET.HG4.11:SETD2 islike:SETD2 —Gonavo_g682.t1 SET.HG4.11:SETD2 islike:SETD2 —Chabra_GBG73012 SET.HG4.11:SETD2 islike:SETD2 —Klenit_GAQ81302.1 SET.HG4.11:SETD2 islike:SETD2
100	Sphfal_Sphfalx10G032300.1.p SET.HG4.11:SETD2 islike:SETD2 Sphfal_Sphfalx17G060000.1.p SET.HG4.11:SETD2 islike:SETD2 Marpol_PTQ41050 SET.HG4.11:SETD2 islike:SETD2 Sacu_v1.1_s0001.g000376 SET.HG4.11:SETD2 islike:SETD2 Azfi_s0345.g065775 SET.HG4.11:SETD2 islike:SETD2 Mgut_XP_012839082.1 SET.HG4.11:SETD2 islike:SETD2 Mgut_XP_012856728.1 SET.HG4.11:SETD2 islike:SETD2 Set.HG4.11:SETD2 islike:SETD2 Set.HG4.11:SETD2 islike:SETD2 Set.HG4.11:SETD2 islike:SETD2 Set.HG4.11:SETD2 islike:SETD2
39	### Acoe_PIA32317.1 SET.HG4.11:SETD2 islike:SETD2 ###################################
75 100 <u>100</u>	-Ulvmut_UM003_02701 SET.HG4.11:SETD2 SETD2
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66 98 100 66 87 94 1	■Ihof_evm3s600 SET.HG4.11:SETD2 SETD2 ■Sarc_Sarc4_g33662T SET.HG4.11:SETD2 SETD2 ■Cfra_CFRG7962T1 SET.HG4.11:SETD2 SETD2 ■Falb_KCV70881 SET.HG4.11:SETD2 SETD2 ■Acas_ELR15294 SET.HG4.11:SETD2 SETD2 ■Fabtro_Gene.55848 SET.HG4.11:SETD2 SETD2 ■Ancsig_Gene.21664 SET.HG4.11:SETD2 SETD2 ■Nutlon_comp18930_c0_seq1.p1 SET.HG4.11:SETD2 SETD2
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68 [•]	**Spun_KNC98643 SET.HG4.11:SETD2 SETD2 **Bden_EGF78935 SET.HG4.11:SETD2 SETD2 **Gpro_KXS11000 SET.HG4.11:SETD2 SETD2 **Cang_ORZ31119 SET.HG4.11:SETD2 SETD2 **Amac_KNE70080 SET.HG4.11:SETD2 SETD2 **Morelo_OAQ28409 SET.HG4.11:SETD2 SETD2 **Morelo_OAQ32941 SET.HG4.11:SETD2 SETD2 **Morelo_OAQ32941 SET.HG4.11:SETD2 SETD2 **Morelo_OAQ32941 SET.HG4.11:SETD2 SETD2
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78 93 95	■Plamic_ANCSI_03774.mRNA.1 SET.HG4.11:SETD2 SETD2 ■Nk52_evm50s230 SET.HG4.11:SETD2 islike:SETD2 ■Ramvar_GAV09639.1 SET.HG4.11:SETD2 islike:SETD2 ■Sman_Smp_210650.1_pep SET.HG4.11:SETD2 islike:SETD2 ■Emul_CUT99633.1 SET.HG4.11:SETD2 islike:SETD2 ■Sros_EGD77782 SET.HG4.11:SETD2 SETD2 ■Mbre_EDQ87823 SET.HG4.11:SETD2 SETD2 ■Mlei_ML0966_g3_t1 SET.HG4.11:SETD2 SETD2
91	#Aque_Aqu2.1.27840_001 SET.HG4.11:SETD2 SETD2 #Tetwil_Twilhelma_g14211.t1 SET.HG4.11:SETD2 SETD2 #Auraur_scaffold330.g7.t1 SET.HG4.11:SETD2 SETD2 #Morvir_scaffold6.g150.t1 SET.HG4.11:SETD2 SETD2 #Clyhem_TCONS_00016153-protein SET.HG4.11:SETD2 SETD2 #Hydvul_Sc4wPfr_1040.g7654.t1 SET.HG4.11:SETD2 SETD2 #Lcom_lcpid35039 SET.HG4.11:SETD2 SETD2 #Scil_scpid27866 SET.HG4.11:SETD2 SETD2
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