ULTIPOTENT_VS_PRO_BCELL_DN, GSE15330_LYMPHOID_MULTIPOTENT_VS_PRO_BCELL_DN GOBP ALANINE CATABOLIC PROCESS, GOBP ALANINE CATABOLIC PROCESS

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GSE13738_RESTING_VS_BYSTANDER_ACTIVATED_CD4_TCELL_DN, GSE13738_RESTING_VS_BYSTANDER_ACTIVATED_CD4_TCELL_DN
 GSE2770 TGFB AND IL4 VS IL12 TREATED ACT CD4 TCELL 6H UP, GSE2770 TGFB AND IL4 VS IL12 TREATED ACT CD4 TCELL 6H UP
 GSE21678_WT_VS_FOXO1_FOXO3_KO_TREG_UP, GSE21678_WT_VS_FOXO1_FOXO3_KO_TREG_UP
GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4HIGH_BCELL_DN, GSE46606_DAY1_VS_DAY3_CD40L_IL2_IL5_STIMULATED_IRF4HIGH_BCELL
GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_12H_DN, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_12H_DN
GSE17721_POLYIC_VS_GARDIQUIMOD_2H_BMDC_DN, GSE17721_POLYIC_VS_GARDIQUIMOD_2H_BMDC_DN
GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP, GSE11961_GERMINAL_CENTER_BCELL_DAY7_VS_PLASMA_CELL_DAY7_UP
GSE46606_IRF4HIGH_VS_WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_DN, GSE46606_IRF4HIGH_VS_WT_CD40L_IL2_IL5_DAY1_STIMULATED_BCELL_DN
 GSE34006_WT_VS_A2AR_KO_TREG_UP, GSE34006_WT_VS_A2AR_KO_TREG_UP
GRABARCZYK_BCL11B_TARGETS_UP, GRABARCZYK_BCL11B_TARGETS_UP
GOBP_POSITIVE_REGULATION_OF_AUTOPHAGY, GOBP_POSITIVE_REGULATION_OF_AUTOPHAGY
HP_NEURODEGENERATION, HP_NEURODEGENERATION
 MIR1226_3P, MIR1226_3P
WP_IL6_SIGNALING_PATHWAY, WP_IL6_SIGNALING_PATHWAY
 GSE37605_NOD_VS_C57BL6_IRES_GFP_TREG_UP, GSE37605_NOD_VS_C57BL6_IRES_GFP_TREG_UP
 VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN, VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN
 GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MEDULLA_THYMUS_DN, GSE18281_PERIMEDULLARY_CORTICAL_REGION_VS_WHOLE_MI
MIR4471, MIR4471
BURTON_ADIPOGENESIS_PEAK_AT_0HR, BURTON_ADIPOGENESIS_PEAK_AT_0HR
HP_INCREASED_LACTATE_DEHYDROGENASE_LEVEL, HP_INCREASED_LACTATE_DEHYDROGENASE_LEVEL
GSE13229_IMM_VS_INTMATURE_NKCELL_UP, GSE13229_IMM_VS_INTMATURE_NKCELL_UP
PRC2_EED_UP.V1_UP, PRC2_EED_UP.V1_UP
CTCNANGTGNY_UNKNOWN, CTCNANGTGNY_UNKNOWN
GSE13411_NAIVE_VS_SWITCHED_MEMORY_BCELL_UP, GSE13411_NAIVE_VS_SWITCHED_MEMORY_BCELL_UP
GOBP_CHLORIDE_TRANSPORT, GOBP_CHLORIDE_TRANSPORT
 WP_PHOSPHODIESTERASES_IN_NEURONAL_FUNCTION, WP_PHOSPHODIESTERASES_IN_NEURONAL_FUNCTION
 VALK_AML_WITH_11Q23_REARRANGED, VALK_AML_WITH_11Q23_REARRANGED
 GOBP_RESPIRATORY_GASEOUS_EXCHANGE_BY_RESPIRATORY_SYSTEM, GOBP_RESPIRATORY_GASEOUS_EXCHANGE_BY_RESPIRATORY_SYSTEM
GOBP_ONE_CARBON_COMPOUND_TRANSPORT, GOBP_ONE_CARBON_COMPOUND_TRANSPORT
 CUI_TCF21_TARGETS_UP, CUI_TCF21_TARGETS_UP
GOMF_CHLORIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_CHLORIDE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GOBP_POSITIVE_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_STAT_PROTEIN, GOBP_POSITIVE_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHORYLATION_OF_TYROSINE_PHOSPHO
BIOCARTA_EPONFKB_PATHWAY, BIOCARTA_EPONFKB_PATHWAY
 chr2q34, chr2q34
HP_FOLLICULAR_HYPERPLASIA, HP_FOLLICULAR_HYPERPLASIA
GOBP_RESPIRATORY_SYSTEM_PROCESS, GOBP_RESPIRATORY_SYSTEM_PROCESS
chr13q32, chr13q32
HP_IRIDODONESIS, HP_IRIDODONESIS
GOBP_INTERCELLULAR_TRANSPORT, GOBP_INTERCELLULAR_TRANSPORT
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