

UNKNOWN, TMTGCGGANR_UNKNOWN

- TOX4_TARGET_GENES, TOX4_TARGET_GENES
- GSE21546_ELK1_KO_VS_SAP1A_KO_AND_ELK1_KO_ANTI_CD3_STIM_DP_THYMOCYTES_UP, GSE21546_ELK1_KO_VS_SAP1A_KO_AND_ELK1_KO_ANTI_CD3_STIM_DP_THYMOCYTES_UP
- HP_WIDE_MOUTH, HP_WIDE_MOUTH
- HP_ABNORMAL_NIPPLE_MORPHOLOGY, HP_ABNORMAL_NIPPLE_MORPHOLOGY
- HP_SHORT_LONG_BONE, HP_SHORT_LONG_BONE
- TAANNYSGCG_UNKNOWN, TAANNYSGCG_UNKNOWN
- GOMF_PROTEIN_PHOSPHATASE_BINDING, GOMF_PROTEIN_PHOSPHATASE_BINDING
- PARK_HSC_AND_MULTIPOTENT_PROGENITORS, PARK_HSC_AND_MULTIPOTENT_PROGENITORS
- HP_SMALL_NAIL, HP_SMALL_NAIL
- HP_ABNORMALITY_OF_SALIVATION, HP_ABNORMALITY_OF_SALIVATION
- HP_ABNORMALITY_OF_THE_PULMONARY_VASCULATURE, HP_ABNORMALITY_OF_THE_PULMONARY_VASCULATURE
- HP_WIDE_INTERMAMILLARY_DISTANCE, HP_WIDE_INTERMAMILLARY_DISTANCE
- GOCC_MOTILE_CILIUM, GOCC_MOTILE_CILIUM
- GACAGGG_MIR339, GACAGGG_MIR339
- HP_BULBAR_SIGNS, HP_BULBAR_SIGNS
- HP_HYPOPLASTIC_FINGERNAIL, HP_HYPOPLASTIC_FINGERNAIL
- HP_FACIAL_HEMANGIOMA, HP_FACIAL_HEMANGIOMA
- HP_INCREASED_NUMBER_OF_TEETH, HP_INCREASED_NUMBER_OF_TEETH
- HP_INCREASED_URINARY_NON_PROTEINOGENIC_AMINO_ACID_LEVEL, HP_INCREASED_URINARY_NON_PROTEINOGENIC_AMINO_ACID_LEVEL
- GOBP_POSITIVE_REGULATION_OF_PROTEIN_POLYUBIQUITINATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_POLYUBIQUITINATION
- GOBP_ACTIVATION_OF_NF_KAPPAB_INDUCING_KINASE_ACTIVITY, GOBP_ACTIVATION_OF_NF_KAPPAB_INDUCING_KINASE_ACTIVITY
- HP_ABNORMALITY_OF_THE_PULMONARY_VEINS, HP_ABNORMALITY_OF_THE_PULMONARY_VEINS