

ACETYLTRANSFERASE\_COMPLEX, GOCC\_PROTEIN\_ACETYLTRANSFERASE\_COMPLEX

- GOCC\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_REGULATOR\_COMPLEX, GOCC\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_REGULATOR\_COMPLEX
- HP\_HIGHLY\_ARCHED\_EYEBROW, HP\_HIGHLY\_ARCHED\_EYEBROW
- GOMF\_NUCLEAR\_RECEPTOR\_COACTIVATOR\_ACTIVITY, GOMF\_NUCLEAR\_RECEPTOR\_COACTIVATOR\_ACTIVITY
- GOCC\_HISTONE\_METHYLTRANSFERASE\_COMPLEX, GOCC\_HISTONE\_METHYLTRANSFERASE\_COMPLEX
- GOMF\_TRANSCRIPTION\_COREPRESSOR\_ACTIVITY, GOMF\_TRANSCRIPTION\_COREPRESSOR\_ACTIVITY
- HP\_DELAYED\_ABILITY\_TO\_WALK, HP\_DELAYED\_ABILITY\_TO\_WALK
- GOBP\_NEGATIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC, GOBP\_NEGATIVE\_REGULATION\_OF\_GENE\_EXPRESSION\_EPIGENETIC
- GOCC\_TRANSCRIPTION\_FACTOR\_TFIID\_COMPLEX, GOCC\_TRANSCRIPTION\_FACTOR\_TFIID\_COMPLEX
- HP\_FULL\_CHEEKS, HP\_FULL\_CHEEKS
- HP\_DEVIATION\_OF\_TOES, HP\_DEVIATION\_OF\_TOES
- GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX, GOCC\_SWI\_SNF\_SUPERFAMILY\_TYPE\_COMPLEX
- HP\_DEVIATION\_OF\_THE\_HALLUX, HP\_DEVIATION\_OF\_THE\_HALLUX
- GOBP\_TELOMERASE\_RNA\_LOCALIZATION, GOBP\_TELOMERASE\_RNA\_LOCALIZATION
- HP\_ABNORMALITY\_OF\_THE\_HALLUX, HP\_ABNORMALITY\_OF\_THE\_HALLUX
- GOCC\_HISTONE\_DEACETYLASE\_COMPLEX, GOCC\_HISTONE\_DEACETYLASE\_COMPLEX
- GOCC\_INO80\_TYPE\_COMPLEX, GOCC\_INO80\_TYPE\_COMPLEX
- HP\_PROMINENT\_FOREHEAD, HP\_PROMINENT\_FOREHEAD
- HP\_LOW\_HANGING\_COLUMELLA, HP\_LOW\_HANGING\_COLUMELLA
- HP\_ABNORMAL\_DISTAL\_PHALANX\_MORPHOLOGY\_OF\_FINGER, HP\_ABNORMAL\_DISTAL\_PHALANX\_MORPHOLOGY\_OF\_FINGER
- HP\_TELECANTHUS, HP\_TELECANTHUS
- GOCC\_MLL1\_2\_COMPLEX, GOCC\_MLL1\_2\_COMPLEX
- GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERASE\_RNA\_LOCALIZATION\_TO\_CAJAL\_BODY, GOBP\_POSITIVE\_REGULATION\_OF\_TELOMERASE\_RNA\_LOCALIZATION\_TO\_CAJAL\_BODY
- HP\_PATENT\_FORAMEN\_OVALE, HP\_PATENT\_FORAMEN\_OVALE
- GOBP\_HISTONE\_DEUBIQUITINATION, GOBP\_HISTONE\_DEUBIQUITINATION
- HP\_ABNORMALITY\_OF\_INCISOR\_MORPHOLOGY, HP\_ABNORMALITY\_OF\_INCISOR\_MORPHOLOGY
- GOBP\_CHROMATIN\_SILENCING, GOBP\_CHROMATIN\_SILENCING
- HP\_ABNORMAL\_PROXIMAL\_PHALANX\_MORPHOLOGY\_OF\_THE\_HAND, HP\_ABNORMAL\_PROXIMAL\_PHALANX\_MORPHOLOGY\_OF\_THE\_HAND
- GOBP\_HISTONE\_H2A\_ACETYLATION, GOBP\_HISTONE\_H2A\_ACETYLATION
- GOCC\_H4\_H2A\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX, GOCC\_H4\_H2A\_HISTONE\_ACETYLTRANSFERASE\_COMPLEX
- GOBP\_POSITIVE\_REGULATION\_OF\_HISTONE\_METHYLATION, GOBP\_POSITIVE\_REGULATION\_OF\_HISTONE\_METHYLATION
- HP\_EARLY\_ONSET\_OF\_SEXUAL\_MATURATION, HP\_EARLY\_ONSET\_OF\_SEXUAL\_MATURATION
- GOCC\_SWR1\_COMPLEX, GOCC\_SWR1\_COMPLEX
- HP\_ABNORMALLY\_FOLDED\_HELIX, HP\_ABNORMALLY\_FOLDED\_HELIX
- GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_RESPONSE\_TO\_TRANSFORMING\_GROWTH\_FACTOR\_BETA\_STIMULUS, GOBP\_POSITIVE\_REGULATION\_OF\_CELLULAR\_RESPONSE\_TO\_TRANSFORMING\_GROWTH\_FACTOR\_BETA\_STIMULUS
- HP\_DOWN\_SLOPING\_SHOULDERS, HP\_DOWN\_SLOPING\_SHOULDERS
- GOCC\_DNA\_HELICASE\_COMPLEX, GOCC\_DNA\_HELICASE\_COMPLEX
- HP\_ABSENT\_TOE, HP\_ABSENT\_TOE
- GOBP\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE\_BY\_P53\_CLASS\_MEDIATOR, GOBP\_REGULATION\_OF\_INTRINSIC\_APOPTOTIC\_SIGNALING\_PATHWAY\_IN\_RESPONSE\_TO\_DNA\_DAMAGE\_BY\_P53\_CLASS\_MEDIATOR
- GOCC\_SAGA\_TYPE\_COMPLEX, GOCC\_SAGA\_TYPE\_COMPLEX