

NUCLEOCYTOPLASMIC\_TRANSPORT, GO\_POSITIVE\_REGULATION\_OF\_NUCLEOCYTOPLASMIC\_TRANSPORT

GO\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_FACTOR\_COMPLEX, GO\_RNA\_POLYMERASE\_II\_TRANSCRIPTION\_FACTOR\_COMPLEX  
GO\_INTERLEUKIN\_12\_PRODUCTION, GO\_INTERLEUKIN\_12\_PRODUCTION  
GO\_REGULATION\_OF\_DNA\_DAMAGE\_RESPONSE\_SIGNAL\_TRANSDUCTION\_BY\_P53\_CLASS\_MEDIATOR, GO\_REGULATION\_OF\_DNA\_DAMAGE\_RESPONSE\_SIGNAL\_TRANSD  
GO\_MODIFICATION\_BY\_SYMBIONT\_OF\_HOST\_MORPHOLOGY\_OR\_PHYSIOLOGY, GO\_MODIFICATION\_BY\_SYMBIONT\_OF\_HOST\_MORPHOLOGY\_OR\_PHYSIOLOGY  
GO\_CELL\_DIFFERENTIATION\_INVOLVED\_IN\_KIDNEY\_DEVELOPMENT, GO\_CELL\_DIFFERENTIATION\_INVOLVED\_IN\_KIDNEY\_DEVELOPMENT  
GO\_MULTI\_ORGANISM\_CELLULAR\_PROCESS, GO\_MULTI\_ORGANISM\_CELLULAR\_PROCESS  
GO\_SALIVARY\_GLAND\_DEVELOPMENT, GO\_SALIVARY\_GLAND\_DEVELOPMENT  
GO\_EXOCRINE\_SYSTEM\_DEVELOPMENT, GO\_EXOCRINE\_SYSTEM\_DEVELOPMENT  
GO\_MODULATION\_BY\_VIRUS\_OF\_HOST\_MORPHOLOGY\_OR\_PHYSIOLOGY, GO\_MODULATION\_BY\_VIRUS\_OF\_HOST\_MORPHOLOGY\_OR\_PHYSIOLOGY  
GO\_NUCLEAR\_INNER\_MEMBRANE, GO\_NUCLEAR\_INNER\_MEMBRANE  
GO\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY, GO\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY  
GO\_FIBROBLAST\_APOPTOTIC\_PROCESS, GO\_FIBROBLAST\_APOPTOTIC\_PROCESS  
GO\_POSITIVE\_REGULATION\_OF\_INTERFERON\_BETA\_PRODUCTION, GO\_POSITIVE\_REGULATION\_OF\_INTERFERON\_BETA\_PRODUCTION  
GO\_ASTROCYTE\_DIFFERENTIATION, GO\_ASTROCYTE\_DIFFERENTIATION  
GO\_ACTIN\_FILAMENT\_BUNDLE\_ORGANIZATION, GO\_ACTIN\_FILAMENT\_BUNDLE\_ORGANIZATION  
GO\_POSITIVE\_REGULATION\_OF\_SUPRAMOLECULAR\_FIBER\_ORGANIZATION, GO\_POSITIVE\_REGULATION\_OF\_SUPRAMOLECULAR\_FIBER\_ORGANIZATION  
GO\_CELL\_FATE\_SPECIFICATION, GO\_CELL\_FATE\_SPECIFICATION  
GO\_DORSAL\_VENTRAL\_NEURAL\_TUBE\_PATTERNING, GO\_DORSAL\_VENTRAL\_NEURAL\_TUBE\_PATTERNING  
GO\_REGULATION\_OF\_TELOMERASE\_ACTIVITY, GO\_REGULATION\_OF\_TELOMERASE\_ACTIVITY  
GO\_POSITIVE\_REGULATION\_OF\_PHOSPHOLIPID\_TRANSPORT, GO\_POSITIVE\_REGULATION\_OF\_PHOSPHOLIPID\_TRANSPORT  
GO\_NEGATIVE\_REGULATION\_OF\_CYSSTEINE\_TYPE\_ENDOPEPTIDASE\_ACTIVITY, GO\_NEGATIVE\_REGULATION\_OF\_CYSSTEINE\_TYPE\_ENDOPEPTIDASE\_ACTIVITY  
GO\_RESPONSE\_TO\_STEROL, GO\_RESPONSE\_TO\_STEROL  
GO\_MAP\_KINASE\_KINASE\_ACTIVITY, GO\_MAP\_KINASE\_KINASE\_ACTIVITY  
GO\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS, GO\_REGULATION\_OF\_PROTEASOMAL\_UBIQUITIN\_DEPENDENT\_PROTEIN\_CATABOLIC\_PROCESS  
GO\_REGULATION\_OF\_KIDNEY\_DEVELOPMENT, GO\_REGULATION\_OF\_KIDNEY\_DEVELOPMENT  
GO\_POSITIVE\_REGULATION\_OF\_PROTEIN\_EXPORT\_FROM\_NUCLEUS, GO\_POSITIVE\_REGULATION\_OF\_PROTEIN\_EXPORT\_FROM\_NUCLEUS