

| MCODE 1mRNA splicing -Major Pathway-12.4MCODE 1mRNA splicing-12.2MCODE 2Regulation of cellular amide metabolic process-6.7Regulation of cellular amide metabolic process-6.7Regulation of translation-6.6Vesicle budding from membrane-6.6Activation of gene expression by SREBF (SREBP)-10.9Regulation of cholesterol biosynthesis by SREBP (SREBP)-10.3MCODE 3Carbohydrate metabolic process-8.6MCODE 4Carbohydrate metabolic process-6.2MCODE 5Mitochondrial protein import-8.8MCODE 6Diabetic cardiomyopathy-8.8MCODE 7Diabetic cardiomyopathy-8.8MCODE 8DNA damage recognition in GG-NER-6.9MCODE 6Formation of incision complex in GG-NER-6.9MCODE 7Negative regulation of intrinsic apoptotic signaling-6.1MCODE 8Cell cycle-6.9MCODE 9Loss of NIp from mitotic centrosomes-6.4MCODE 9Intra-Golgi raffic-13.1MCODE 9Intra-Golgi ind retrograde Golgi-to-ER traffic-13.1MCODE 9Intra-Golgi vesicle mediated transport-10.7MCODE 9Intra-Golgi vesicle mediated transport-10.7MCODE 10PID ERA genomic pathway-7.0 | MCODE subcluster | Downregulated GO Pathways | Log ₁₀ P |
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