

Human expressed differentially from the non-human primates: p-value < 0.05.				
PANTHER				
Other nucleoside, nucleotide and nucleic acid metabolism	0.04854	1	1	
Oxidative phosphorylation	0.04854	1	1	
RNA catabolism	0.04854	1	1	
GO				
protein transport	0.03063	2	6	
RNA interference, production of siRNA	0.05	1	1	
RNA interference, targeting of mRNA for destruction	0.05	1	1	
branching morphogenesis of a tube	0.05	1	1	
electron transport chain	0.05	1	1	
embryonic hindlimb morphogenesis	0.05	1	1	
exocytosis	0.05	1	1	
gene silencing by RNA	0.05	1	1	
gene silencing by miRNA, production of miRNAs	0.05	1	1	
lung development	0.05	1	1	
mitochondrial electron transport, NADH to ubiquinone	0.05	1	1	
oligopeptide transport	0.05	1	1	
positive regulation of long-term neuronal synaptic plasticity	0.05	1	1	
stem cell maintenance	0.05	1	1	
All differential p-values < 0.05.				
PANTHER				
Hearing	0.007015	2	3	
Protein targeting	0.01429	3	11	
Protein targeting and localization	0.04777	3	17	
Action potential propagation	0.04967	1	1	
Other mRNA transcription	0.04967	1	1	
Phosphate transport	0.04967	1	1	
GO				
regulation of Rab GTPase activity	0.002009	3	6	
ion transport	0.005597	6	35	
mitochondrial electron transport, NADH to ubiquinone	0.01332	2	4	
electron transport chain	0.01391	3	11	
sensory perception of sound	0.01792	3	12	
potassium ion transport	0.0225	3	13	
protein modification process	0.03971	3	16	
translational elongation	0.04658	3	17	
DNA-dependent DNA replication	0.04913	1	1	
L-glutamate transport	0.04913	1	1	
asparaginyl-tRNA aminoacylation	0.04913	1	1	
aspartyl-tRNA aminoacylation	0.04913	1	1	
base-excision repair, gap-filling	0.04913	1	1	
clustering of voltage-gated potassium channels	0.04913	1	1	
cobalamin biosynthetic process	0.04913	1	1	
complement activation, classical pathway	0.04913	1	1	
dTDP biosynthetic process	0.04913	1	1	
dTTP biosynthetic process	0.04913	1	1	
long-term strengthening of neuromuscular junction	0.04913	1	1	
maturity of SSU-rRNA	0.04913	1	1	
metal ion transport	0.04913	1	1	
middle ear morphogenesis	0.04913	1	1	
nucleotide biosynthetic process	0.04913	1	1	
outer ear morphogenesis	0.04913	1	1	
phosphate transport	0.04913	1	1	
positive regulation of phagocytosis	0.04913	1	1	
positive regulation of type Ila hypersensitivity	0.04913	1	1	
protein import into mitochondrial inner membrane	0.04913	1	1	
protein targeting to mitochondrion	0.04913	1	1	
ribosomal small subunit assembly and maintenance	0.04913	1	1	
sequestering of neurotransmitter	0.04913	1	1	
skeletal system morphogenesis	0.04913	1	1	