Clarify of the promestile helpfrior development is the promestile helpfrior development in the promestile helpfrior developmen		cytoskeletal DNA protein binding binding	-myosin light chain binding -sequence–specific single stranded DNA binding -sequence–specific DNA binding -RNA polymerase II cis–regulatory region sequence–specific DNA binding -RNA polymerase II transcription regulatory region sequence–specific DNA binding
mal ubiquitin-dependent protein catabolic process death		DNA-binding transcription factor ac	
nuscle myoblast proliferation  I in micturition mission rgic n aling		enzyme binding	-DNA topoisomerase binding
n aling seudopodium assembly body assembly ille cilium assembly	itium assembly assembly  assembly  assembly  assembly  assembly  assembly  assembly  assembly  over_represented_pvalue  out  out  out  out  out  out  out  o	enzyme regulator activity	- guanylate cyclase activator activity - adenylate cyclase activator activity - acetyltransferase activator activity
cess olyadenylation strand break repair via nonhomologous end joining DNA fragmentation		hydrolase activity, acting on carbo	- N–formylglutamate deformylase activity
olyubiquitination		hydrolase activity, acting on glyco	hydrolase activity, hydrolyzing N–glycosyl compounds
		ion binding	- FAD binding - phosphatidylinositol phosphate binding - spermidine binding
oH reduction ocess ling n remodeling		kinase activity	- glutamate 5-kinase activity - insulin-activated receptor activity - mannokinase activity - hexokinase activity - glucokinase activity - fructokinase activity
chemotaxis to cAMP tion on		ligase activity	-RNA-3'-phosphate cyclase activity -methionine-tRNA ligase activity
one-mediated protein folding		lyase activity	- carbonate dehydratase activity - glutathione specific gamma–glutamylcyclotransferase activity - tryptophanase activity - gamma–glutamylcyclotransferase activity
		oxidoreductase activity	sulfide: quinone oxidoreductase activity Deloxidase activity Service of the control of the contr
way e-activating G protein-coupled receptor signaling pathway		transcription factor binding	- Tat protein binding - cAMP response element binding protein binding
g pathway etic process eess cess cutamate		transferase activity, transferring 	- 1-acylglycerol-3-phosphate O-acyltransferase activity - alpha-1,6-mannosylglycoprotein 4-beta-N-acetylglucosaminyltransferase activity - mycocerosate synthase activity - diamine N-acetyltransferase activity - UDP-alpha-D-glucose:glucosyl-glycogenin alpha-D-glucosyltransferase activity - glycogenin glucosyltransferase activity - methionine adenosyltransferase activity
cess :quinone oxidoreductase :geretion :sertion into mitochondrial outer membrane :geretion :sertion into mitochondrial outer membrane :certion into mitoc		transmembrane transporter activity	fructose transmembrane transporter activity carbohydrate:proton symporter activity dehydroascorbic acid transmembrane transporter activity aromatic amino acid transmembrane transporter activity glucose transmembrane transporter activity thyroid hormone transmembrane transporter activity acetylcholine—gated cation—selective channel activity potassium ion antiporter activity monocarboxylic acid transmembrane transporter activity amino acid transmembrane transporter activity amino acid transmembrane transporter activity
transmembrane transport nembrane t ng growth factor beta2 production test to the second transport test	<u></u>		Tamino acid transmembrane transporter activity  activity control of the control o