carbohydrate derivative metabolic process carbohydrate carbohydrate metabolic process cell cell cytoskeleton detormation motility cell cytoskeleton organization	romithine biosynthetic process methlorine catabolic process crolline catabolic process crolline catabolic process and the proc			DNA binding	sulfide guinone exidoreductase activity Involving a civity. Involving N-glycosyl compounds - High Will Color of the civity of t	
DNA crepair replication DNA transcription	negative regulation of double–strand break repair via nonhomologous end joining DNA double–strand break processing involved in repair via single–strand annealing regulation of DNA endoreduplication positive regulation of transcription of nucleolar large rRNA by RNA polymerase I positive regulation of transcription from RNA polymerase II promoter in response to heat stress complement activation, classical pathway regulation of mast cell activation			lipid binding	-lipoic acid binding	
ular lipid metabolic movement metabolic process movement metabolic process process process process	positive regulation of protein insertion into mitochondrial outer membrane green leaf volatile biosynthetic process positive regulation of rativ acid process positive regulation of colliary planar beating movement pattern positive regulation of mRNA polyadenylation regulation of the force of skeletal muscle contraction process process positive regulation of skeletal muscle fiber contraction process	over_represented_pvalue 0.04 0.03 0.02 0.01		molecular mo function tran regulator activity	-ubiquitin-protein transferase activator activity -guanylate cyclase activator activity -adenylate cyclase activator activity -acetyltransferase activator activity	over_represented_pvalue 0.04 0.03 0.02 0.01
protein catabolic protein protein complex process folding maturation assembly process signaling	negative regulation of hydrogen peroxide—mediated programmed cell death anaphase—promoting complex—dependent catabolic process negative regulation of chaperone—mediated protein folding zymogen activation protein heterooligomerization protein homotrimerization radial spoke assembly protein tetramerization multicellular organismal reproductive process male meiosis I positive regulation of meiotic cell cycle regulation of meiotic nuclear division protein tetramerization of meiotic nuclear division positive regulation of meiotic nuclear division protein heterooligomerization multicellular organismal reproductive process male meiosis I positive regulation of meiotic cell cycle regulation of meiotic nuclear division protein heterooligomerization protein hetero			g transpo	regulatory RNA binding sugar transmembrane transporter activity fructose transmembrane transporter activity gated channel activity dehydroascorbic acid transmembrane transporter activity acetylcholine–gated monoatomic cation–selective channel activity	
telomere transmembrane esicle-med transport	hydrogen sulfide metabolic process sulfide oxidation, using sulfide:quinone oxidoreductase glucosinolate biosynthetic process S-adenosylmethionine biosynthetic process t-circle formation telomeric D-loop disassembly fructose transmembrane transport glucose transmembrane transport carbohydrate transmembrane transport inorganic cation transmembrane transport negative regulation of regulated secretory pathway phagocytosis, engulfment endocytosis vesicle docking vitamin B6 metabolic process		MF	N _A	-D-glucose transmembrane transporter activity -trehalose transmembrane transporter activity -phosphatidic acid transfer activity -extracellular ligand-gated monoatomic ion channel activity -glucose binding -cargo receptor activity -FAD binding -low-density lipoprotein particle binding -amyloid-beta binding -Hsp90 protein binding -D5 dopamine receptor binding -translation elongation factor binding -insulin-like growth factor II binding -insulin-like growth factor I binding -insulin-like growth factor I binding -TORC2 complex binding -insulin-like growth factor receptor binding	