calcium ion binding -2.81642969046316e-07 extracellular matrix structural constituent -8.48506754698523e-06 serine-type endopeptidase activity -0.0211994164371091 carbohydrate binding -0.0369089192613595 metalloendopeptidase activity -0.0179004687202011 DNA-binding transcription factor activity, RNA polymerase II-specific -0.000767702340783581 0.0370758975739399 structural constituent of cytoskeleton -0.0395704812691337 scavenger receptor activity -0.0015987360189158 chitin binding -0.00479878127683983 hydrolase activity, hydrolyzing O-glycosyl compounds -0.00589883571925086 cysteine-type endopeptidase activity -0.0479593786362308 copper ion binding -0.0294859587085942 protease binding -0.00665654830351578 carboxylic ester hydrolase activity -0.0295669246247243 SMAD binding -0.0307305142412766 P-type ion transporter activity -0.0054814469052228 oxidoreductase activity, acting on single donors with incorporation of molecular oxygen, incorporation of two atoms of oxygen-1.04700716802029e-05 minor groove of adenine-thymine-rich DNA binding -0.00246663480764292 0.038375912411378 ATPase-coupled inorganic anion transmembrane transporter activity -0.0307305142412766 15-hydroxyprostaglandin dehydrogenase (NAD+) activity -0.00577169523335724 RNA endonuclease activity -0.00103386222541996 myosin heavy chain binding -0.0226474004747081 hydrolase activity, acting on ester bonds -0.0141694859956542 extracellular matrix binding -0.0027559330496216 platelet-derived growth factor binding -0.00170665389625459 nucleosomal DNA binding -0.0159777431406043 lipid transporter activity -0.00234262767002173 dipeptidyl-peptidase activity -0.0349647851066897 core promoter sequence-specific DNA binding -0.00659592144310945 chitinase activity -0.0285473693932665 ribonucleoside binding -0.00703947088111381 phosphopantetheine binding low-density lipoprotein particle binding -0.00834260045599648 ----0.0322409970493545 phosphorus-oxygen lyase activity -0.00374613817294753 nutrient reservoir activity -0.00499253960739584 extracellular matrix structural constituent conferring tensile strength -0.0470024334439132 enzyme regulator activity -0.0206920578934323 betaine-homocysteine S-methyltransferase activity -0.0327290793960894 arachidonate 5-lipoxygenase activity -----0.0488949149103114 acetylcholinesterase activity -0.0482983459918834 wide pore channel activity -0.0410822010058722 rhamnose binding --0.0293059807949241 protein-lysine 6-oxidase activity -----0.0278860475409049 propane-1,3-diamine oxidase activity -----0.0353179845027128 peptidyl-dipeptidase activity -----0.0361270439899008 penicillin amidase activity -----0.0146364182192919 oxidoreductase activity, acting on the CH-NH2 group of donors, oxygen as acceptor -----0.0146362883483593 mycocerosate synthase activity -----0.0478381062166624 modified amino acid binding -0.0278860475409049 methylputrescine oxidase activity -----0.0486023882908976 metallodipeptidase activity -• 0.0410822010058722 melibiose binding -0.00522530288149781 L-amino-acid oxidase activity --0.0278859766536323 kinetochore binding -----0.0278860475409049 histamine oxidase activity -----0.0285027343523419 heme transmembrane transporter activity • 0.033094821687563 GPI anchor binding --0.00512302544682537 glutamine-tRNA ligase activity -0.0056021780606361 ferrochelatase activity -• 0.0400794560822725 dopamine neurotransmitter receptor activity, coupled via Gi/Go -0.0167610410616861 DNA topoisomerase binding -----0.0278860475409049 diamine oxidase activity -----0.039414289693472 7SK snRNA binding

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