																					GOT	ERM_	MF_F	AT																						1
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GO:0000150~recombinase activity	GO:0003676~nucleic acid binding	GO:0003682~chromatin binding	GO:0003723~RNA binding	O:000452	GO:0004527~exonuclease activity	GO:0004722~protein serine/threonine phosphatase activity	GO:0004842~ubiquitin-protein transferase activity	GO:0005198~structural molecule activity	GO:0008134~transcription factor binding	GO:0016779~nucleotidyltransferase activity	GO:0019787∼ubiquitin−like protein transferase activity	GO:0019899~enzyme binding	:001990	GO:001990z~priospriatase binding GO:0019901~protein kinase binding	:0031490~chromatin DNA bindi	GO:0031491~nucleosome binding	GO:0031492~nucleosomal DNA binding	GO:0031625∼ubiquitin protein ligase binding	GO:0035257~nuclear hormone receptor binding	GO:0042802∼identical protein binding		:0044389~ubiquitin−like protein	O:0044822~poly(A) RNA bir	O:0045296~cadherin binding	O:0046966~thyroid hormone rec	839~cell adhesion molec	GO:0051427~hormone receptor binding	GO:0097159~organic cyclic compound binding	GO:0098631~protein binding involved in cell adhesion	GO:0098632~protein binding involved in cell–cell adhesion	098641∼cadherin binding inv	O:1901363~heterocyclic compound binding	GO:0016791~phosphatase activity GO:0016651~oxidoreductase activity, acting on NAD(P)H	:003	GO:0000166~nucleotide binding	GO:0003729~mRNA binding	GO:0003730~mRNA 3'-UTR binding	GO:0016817~hydrolase activity, acting on acid anhydrides	GO:0016818~hydrolase activity, acting on acid anhydrides, in phosphorus–containing anhyd	GO:0017016~Ras GTPase binding	GO:0017111~nucleoside-triphosphatase activity	GO:0017137~Rab GTPase binding	iO:0031267∼small GTPase bindinç	0:0036	O:005	