Homo sapiens	Q6ZNL4_HUMAN (Uniprot_id_acc)	Q6ZNL4_HUMAN	**Q6ZNL4_HUMAN , Q6ZNL4**	Yes	22:35007273-35113958	4627	Yes	cell cortex[M][R][H] • cell-cell adherens junction[M][R][H] • cellular_component[R] • cleavage furrow[M][R][H] • contractile ring[M][R][H] • cortical cytoskeleton[M][R][H] • cytoplasm[M][R][H] • cytosol[M][R][H] • immunological synapse[M][R][H] • integrin complex[M][R][H] • myosin complex[M][R][H] • neuromuscular junction[M][R][H] • nucleus[M][R][H] • plasma membrane[M][R][H] • protein complex[R] • ruffle[M][R][H] • spindle[M][R][H] • stress fiber[M][R][H] • uropod[M][R][H]	• ADP binding[M] • ATP binding[M][R][H] • ATPase activity[M][R][H] • actin binding[M] • actin filament binding[M][R][H] • actin-dependent ATPase activity[M] • calmodulin binding[M][R][H] • hydrolase activity[M] • microfilament motor activity[M][R][H] • molecular_function[R] • motor activity[M][H] • nucleotide binding[M][R][H] • protein anchor[M][R][H] • protein binding[M][R] • protein homodimerization activity[M][R][H] • sodium:potassium-exchanging ATPase activity[R]	• cell motility[M][R][H] • cell-cell adhesion[M][R][H] • cellular morphogenesis during differentiation[M][R][H] • cytokinesis[M][R][H] • establishment of T cell polarity[M][R][H] • establishment of meiotic spindle localization[M][R][H] • in utero embryonic development[M][R][H] • integrin-mediated signaling pathway[H] • leukocyte migration[H] • meiotic metaphase I[M][R][H] • meiotic spindle organization and biogenesis[M][R][H] • membrane protein ectodomain proteolysis[M][R][H] • monocyte differentiation[M][R][H] • myoblast fusion[M][R][H] • platelet formation[M][R][H] • protein transport[M][R][H] • regulation of cell shape[M][R][H] • sensory perception of sound[H]
Homo sapiens	Q9H5Z9_HUMAN (Uniprot_id_acc)	Q9H5Z9_HUMAN	**Q9H5Z9_HUMAN , Q9H5Z9**	Yes	17:26250127-26257379	79736	-	• nucleus[M]	• hydrolase activity[M][R][H] • nuclease activity[M][R][H]	
Homo sapiens	GANAB_HUMAN (Uniprot_id_acc)	GANAB_HUMAN	**GANAB_HUMAN , Q14697-2**	Yes	11:62148882-62170680	23193	Yes	• alpha-glucosidase II complex[M][R] • endoplasmic reticulum[M][R]	• glucosidase activity[M] • hydrolase activity[M] • hydrolase activity, hydrolyzing O-glycosyl compounds[M][R][H] • protein binding[M][R]	• N-glycan processing[M][R] • carbohydrate metabolic process[M][R][H]
Homo sapiens	GRP78_HUMAN (Uniprot_id_acc)	Q2KHP4_HUMAN	**Q2KHP4_HUMAN , Q2KHP4**	Yes	9:127036953-127043430	3309	Yes	• ER-Golgi intermediate compartment[M][R][H] • caspase complex[M][R][H] • cell surface[M][R][H] • endoplasmic reticulum[M][R][H] • endoplasmic reticulum lumen[R][H] • integral to endoplasmic reticulum membrane[M][R][H] • nucleus[M][R][H] • perinuclear region of cytoplasm[M][R][H]	• ATP binding[M][R][H] • calcium ion binding[R][H] • caspase inhibitor activity[M][R][H] • nucleotide binding[M][R][H] • protein binding[M][R] • protein binding, bridging[R][H] • ribosome binding[M][R][H] • unfolded protein binding[R][H]	• ER overload response[M][R][H] • anti-apoptosis[M][R][H] • negative regulation of caspase activity[M][R][H] • protein folding[M] • response to unfolded protein[M]
Homo sapiens	GLU2B_HUMAN (Uniprot_id_acc)	GLU2B_HUMAN	**GLU2B_HUMAN , P14314**	Yes	19:11407269-11422783	5589	-	• alpha-glucosidase II complex[M][R][H] • cellular_component[R] • endoplasmic reticulum[M][R][H] • extracellular space[M][R] • intracellular[M][R][H]	• alpha-glucosidase activity[M][R][H] • calcium ion binding[M][R][H] • kinase activity[M] • molecular_function[R][H] • protein binding[M][R][H]	• N-glycan processing[M][R] • biological_process[R] • protein kinase cascade[M][R][H]
Homo sapiens	GRP75_HUMAN (Uniprot_id_acc)	Q8N1C8_HUMAN	**Q8N1C8_HUMAN , Q8N1C8**	Yes	5:137918924-137939014	3313	-	• cell surface[M][R][H] • cytoplasm[M][R][H] • mitochondrion[M][R][H]	• ATP binding[M][R][H] • nucleotide binding[M][R][H] • protein binding[M][R] • unfolded protein binding[M][R][H]	• anti-apoptosis[R][H] • protein export from nucleus[M][R][H] • protein folding[M][R][H] • response to heat[M] • response to unfolded protein[M][H]
Homo sapiens	Q53GZ6_HUMAN (Uniprot_id_acc)	Q9NZ87_HUMAN	**Q9NZ87_HUMAN , Q9NZ87**	Yes	11:122433411-122438054	3312	Yes	• cell surface[R] • intracellular[R] • nucleus[M][R]	• ATP binding[M][R][H] • ATPase activity, coupled[M][R] • nucleotide binding[M][R][H] • protein binding[M][R] • unfolded protein binding[M][R]	• chaperone cofactor-dependent protein folding[M][R] • protein folding[M][R][H] • regulation of progression through cell cycle[M][R] • response to heat[M] • response to unfolded protein[M][R][H]
Homo sapiens	HSP7C_HUMAN (Uniprot_id_acc)	Q9NZ87_HUMAN	**Q9NZ87_HUMAN , Q9NZ87**	Yes	11:122433411-122438054	3312	Yes	• cell surface[R] • intracellular[R] • nucleus[M][R]	• ATP binding[M][R][H] • ATPase activity, coupled[M][R] • nucleotide binding[M][R][H] • protein binding[M][R] • unfolded protein binding[M][R]	• chaperone cofactor-dependent protein folding[M][R] • protein folding[M][R][H] • regulation of progression through cell cycle[M][R] • response to heat[M] • response to unfolded protein[M][R][H]
									• ATP binding[M][R][H] chaperone binding[R][H]	• cellular protein metabolic process[M][H]