									muscle[M] • vascular endothelial growth factor receptor activity[M]	• sarcomere organization[M][R] • striated muscle cell development[M]
Homo sapiens	TBB5_HUMAN (Uniprot_id_acc)	Q6LC01_HUMAN	**Q6LC01_HUMAN , Q6LC01**	Yes	6:30825252-30830311	203068	Yes	• cytoplasm[M][R][H] • cytoskeleton[H] • microtubule[M][R][H] • protein complex[M][H] • tubulin complex[M]	• GTP binding[M][R][H] • GTPase activity[M][R][H] • MHC class I protein binding[M][R][H] • nucleotide binding[M][R][H] • structural constituent of cytoskeleton[M][R][H] • structural molecule activity[M][H]	• cell motility[H] • microtubule-based movement[M][R][H] • microtubule-based process[M][R] • mitotic spindle assembly[M][R][H] • natural killer cell mediated cytotoxicity[H] • protein polymerization[M][R][H]
Homo sapiens	Q59FC6_HUMAN (Uniprot_id_acc)	Q96GW1_HUMAN	**Q96GW1_HUMAN , Q96GW1**	Yes	12:102848290-102865809	7184	-	• cytosol[H] • endoplasmic reticulum[M][H] • endoplasmic reticulum lumen[H] • endoplasmic reticulum membrane[H] • extracellular space[M] • microsome[H] • perinuclear region of cytoplasm[H]	• ATP binding[M][H] • RNA binding[H] • calcium ion binding[M][H] • low-density lipoprotein receptor binding[M][H] • nucleotide binding[M] • unfolded protein binding[M][H] • virion binding[H]	• anti-apoptosis[H] • protein folding[M][H] • protein transport[H] • response to hypoxia[H] • response to unfolded protein[M] • sequestering of calcium ion[H]
Homo sapiens	UQCR1_HUMAN (Uniprot_id_acc)	UQCR1_HUMAN	**UQCR1_HUMAN , P31930**	Yes	3:48611440-48622474	7384	Yes	• membrane[M][R][H] • mitochondrial electron transport chain[M][R][H] • mitochondrial inner membrane[M][R] • mitochondrion[M][R][H] • ubiquinol-cytochrome-c reductase complex[R]	• metalloendopeptidase activity[M] • oxidoreductase activity[M][R][H] • ubiquinol-cytochrome-c reductase activity[M][R][H]	• aerobic respiration[H] • electron transport[M][R][H] • mitochondrial electron transport, ubiquinol to cytochrome c[R] • oxidative phosphorylation[H] • proteolysis[M] • transport[M][R][H]
Homo sapiens	Q6ZWI2_HUMAN (Uniprot_id_acc)	Q6ZWI2_HUMAN	**Q6ZWI2_HUMAN , Q6ZWI2**	Yes	10:49034607-49054190	143162	-			
Homo sapiens	VIME_HUMAN (Uniprot_id_acc)	Q8N9S6_HUMAN	**Q8N9S6_HUMAN , Q8N9S6**	Yes	10:17311283-17319598	7431	Yes	• cytoplasm[M][R][H] • cytoskeleton[M][R][H] • intermediate filament[M][R][H] • nucleus[R] • type III intermediate filament[M][R]	• protein binding[M][R][H] • structural constituent of cytoskeleton[R][H] • structural molecule activity[M][R][H]	• biological_process[R] • cell motility[R][H] • intermediate filament-based process[M][R]
Homo sapiens	WIF1_HUMAN (Uniprot_id_acc)	WIF1_HUMAN	**WIF1_HUMAN , Q9Y5W5**	Yes	12:63730674-63801383	11197	Yes	• extracellular space[M][R] • integral to membrane[M] • membrane[M]	• Wnt-protein binding[M] • protein-tyrosine kinase activity[M][R][H]	• Wnt receptor signaling pathway[M][R][H] • cell-cell signaling[R][H] • multicellular organismal development[M][R][H] • negative regulation of Wnt receptor signaling pathway[M] • signal transduction[R][H]
Homo sapiens	ZN364_HUMAN (Uniprot_id_acc)	ZN364_HUMAN	**ZN364_HUMAN , Q9Y4L5**	Yes	1:144322393-144402018	27246	-		• metal ion binding[M][H] • protein binding[M][R][H] • zinc ion binding[M][H]	