

GO	GO Description	Method
GO:0045095	keratin filament	LMMS, DLMMS, Mean, SME
GO:0005882	intermediate filament	LMMS, DLMMS, Mean, SME
GO:0044822	poly(A) RNA binding	LMMS, DLMMS, Mean, SME
GO:0005634	nucleus	LMMS, DLMMS, Mean, SME
GO:1900740	positive regulation of protein insertion into mitochondrial membrane involved in apoptotic signaling pathway	LMMS, DLMMS, Mean, SME
GO:0005200	structural constituent of cytoskeleton	LMMS, DLMMS, Mean, SME
GO:0005198	structural molecule activity	LMMS, DLMMS, Mean, SME
GO:0005719	nuclear euchromatin	DLMMS, Mean, SME
GO:0005615	extracellular space	DLMMS, Mean, SME
GO:0005515	protein binding	DLMMS, Mean, SME
GO:0005737	cytoplasm	DLMMS, Mean, SME
GO:0005829	cytosol	LMMS, Mean, SME
GO:0031982	vesicle	LMMS, DLMMS, Mean
GO:0030529	ribonucleoprotein complex	LMMS, DLMMS, Mean
GO:0006096	glycolytic process	LMMS, DLMMS, Mean
GO:0031012	extracellular matrix	LMMS, DLMMS, Mean
GO:0004743	pyruvate kinase activity	LMMS, DLMMS, Mean
GO:0023026	MHC class II protein complex binding	LMMS, DLMMS, Mean
GO:0003725	double-stranded RNA binding	LMMS, DLMMS, Mean
GO:0097193	intrinsic apoptotic signaling pathway	Mean, SME
GO:0001895	retina homeostasis	Mean, SME
GO:0031490	chromatin DNA binding	Mean, SME
GO:0021762	substantia nigra development	Mean, SME
GO:0097110	scaffold protein binding	Mean, SME
GO:0005925	focal adhesion	Mean, SME
GO:0070062	extracellular vesicular exosome	Mean, SME
GO:0019904	protein domain specific binding	Mean, SME
GO:0072562	blood microparticle	Mean, SME
GO:0004601	peroxidase activity	Mean, SME
GO:0006605	protein targeting	Mean, SME
GO:0014069	postsynaptic density	Mean, SME
GO:0071901	negative regulation of protein serine/threonine kinase activity	Mean, SME
GO:0042470	melanosome	DLMMS, Mean
GO:0031093	platelet alpha granule lumen	LMMS, Mean
GO:0008285	negative regulation of cell proliferation	LMMS, Mean
GO:0000790	nuclear chromatin	LMMS, Mean
GO:0008544	epidermis development	LMMS, DLMMS
GO:0006006	glucose metabolic process	LMMS, DLMMS
GO:0005929	cilium	LMMS, DLMMS
GO:0030955	potassium ion binding	LMMS, DLMMS
GO:0012501	programmed cell death	LMMS, DLMMS
GO:0060548	negative regulation of cell death	LMMS, DLMMS
GO:0005975	carbohydrate metabolic process	LMMS, DLMMS