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UNIQUE hCAD PROTRUSION TERMS	hCAD PROTRUSION IN 2 TERMS	EXCLUSIVE hCAD PROTRUSION TERMS
myelin sheath (GO:0043209)	cytosolic ribosome (GO:0022626)	calcium- and calmodulin-dependent protein kinase complex (GO:0005954)
extracellular exosome (GO:0070062)	myelin sheath (GO:0043209)	RAVE complex (GO:0043291)
extracellular vesicle (GO:1903561)	cytosolic small ribosomal subunit (GO:0022627)	FHF complex (GO:0070695)
extracellular organelle (GO:0043230)	cytosolic part (GO:0044445)	dendritic spine neck (GO:0044326)
ribonucleoprotein complex (GO:1990904)	chaperonin-containing T-complex (GO:0005832)	sodium:potassium-exchanging ATPase complex (GO:0005890)
intracellular ribonucleoprotein complex (GO:0030529)	ribosomal subunit (GO:0044391)	neurofilament (GO:0005883)
cytosolic part (GO:0044445)	chaperone complex (GO:0101031)	cell tip (GO:0051286)
cytosolic ribosome (GO:0022626)	ribosome (GO:0005840)	cation-transporting ATPase complex (GO:0090533)
proteasome complex (GO:0000502)	zona pellucida receptor complex (GO:0002199)	dendritic spine head (GO:0044327)
eukaryotic translation initiation factor 3 complex (GO:0005852)	small ribosomal subunit (GO:0015935)	ATPase dependent transmembrane transport complex (GO:0098533)
endopeptidase complex (GO:1905369)	cytosolic large ribosomal subunit (GO:0022625)	sperm fibrous sheath (GO:0035686)
vesicle (GO:0031982)	extracellular exosome (GO:0070062)	eukaryotic translation initiation factor 3 complex (GO:0005852)
focal adhesion (GO:0005925)	extracellular vesicle (GO:1903561)	dendritic spine (GO:0043197)
extracellular space (GO:0005615)	extracellular organelle (GO:0043230)	neuron spine (GO:0044309)
extracellular region part (GO:0044421)	focal adhesion (GO:0005925)	sperm principal piece (GO:0097228)
cell-substrate adherens junction (GO:0005924)	cell-substrate adherens junction (GO:0005924)	mitotic spindle pole (GO:0097431)
macromolecular complex (GO:0032991)	cell-substrate junction (GO:0030055)	postsynapse (GO:0098794)
cell-substrate junction (GO:0030055)	adherens junction (GO:0005912)	growth cone (GO:0030426)
proteasome accessory complex (GO:0022624)	anchoring junction (GO:0070161)	site of polarized growth (GO:0030427)
proteasome regulatory particle (GO:0005838)	VCP-NSFL1C complex (GO:1990730)	synapse part (GO:0044456)
proteasome core complex (GO:0005839)	mitotic spindle microtubule (GO:1990498)	ciliary part (GO:0044441)
nuclear proteasome complex (GO:0031595)	extracellular matrix (GO:0031012)	neuronal cell body (GO:0043025)
cytosol (GO:0005829)	ribonucleoprotein complex (GO:1990904)	postsynaptic membrane (GO:0045211)
extracellular region (GO:0005576)	intracellular ribonucleoprotein complex (GO:0030529)	dendrite (GO:0030425)
zona pellucida receptor complex (GO:0002199)	large ribosomal subunit (GO:0015934)	dendritic tree (GO:0097447)
proteasome regulatory particle, base subcomplex (GO:0008540)	extracellular region part (GO:0044421)	cell projection part (GO:0044463)
adherens junction (GO:0005912)	extracellular space (GO:0005615)	plasma membrane bounded cell projection part (GO:0120038)
peptidase complex (GO:1905368)	eukaryotic translation elongation factor 1 complex (GO:0005853)	somatodendritic compartment (GO:0036477)
chaperonin-containing T-complex (GO:0005832)	Myb complex (GO:0031523)	synapse (GO:0045202)
ribosome (GO:0005840)	vesicle (GO:0031982)	cell body (GO:0044297)
anchoring junction (GO:0070161)	extracellular region (GO:0005576)	cilium (GO:0005929)
intracellular organelle part (GO:0044446)	cytosol (GO:0005829)	neuron projection (GO:0043005)
chaperone complex (GO:0101031)	proteasome core complex, alpha-subunit complex (GO:0019773)	plasma membrane bounded cell projection (GO:0120025)
cytoplasm (GO:0005737)	type III intermediate filament (GO:0045098)	neuron part (GO:0097458)
organelle part (GO:0044422)	macromolecular complex (GO:0032991)	integral component of plasma membrane (GO:0005887)
cytoplasmic part (GO:0044444)	endoplasmic reticulum chaperone complex (GO:0034663)	intrinsic component of plasma membrane (GO:0031226)
protein complex (GO:0043234)	polysome (GO:0005844)	cell projection (GO:0042995)
cytosolic large ribosomal subunit (GO:0022625)	cytoplasmic part (GO:0044444)	plasma membrane (GO:0005886)
ribosomal subunit (GO:0044391)	cell junction (GO:0030054)	
proteasome core complex, alpha-subunit complex (GO:0019773)	intracellular non-membrane-bounded organelle (GO:0043232)	
eukaryotic translation initiation factor 3 complex, eIF3m (GO:0071541)	non-membrane-bounded organelle (GO:0043228)	
membrane-bounded organelle (GO:0043227)	smooth endoplasmic reticulum (GO:0005790)	
intracellular part (GO:0044424)	glycogen granule (GO:0042587)	
cytosolic proteasome complex (GO:0031597)	eukaryotic translation initiation factor 3 complex (GO:0005852)	
intracellular (GO:0005622)	tubulin complex (GO:0045298)	
organelle (GO:0043226)	mitochondrial proton-transporting ATP synthase complex, catalytic core F(1) (GO:0000275)	
intracellular organelle (GO:0043229)	proton-transporting ATP synthase complex, catalytic core F(1) (GO:0045261)	
prefoldin complex (GO:0016272)	cytoplasm (GO:0005737)	
eukaryotic translation elongation factor 1 complex (GO:0005853)	proton-transporting two-sector ATPase complex, catalytic domain (GO:0033178)	
nucleocytoplasmic transport complex (GO:0031074)	protein complex (GO:0043234)	

Figure S5