



- 3/21 proteasome core complex
- 15/108 peptidase complex
- 0/13 proteasome storage granule
- 17/90 protein acetyltransferase complex
- 9/34 endoplasmic reticulum-Golgi intermediate compartment membrane
- 19/79 endoplasmic reticulum-Golgi intermediate compartment**
- 14/95 endoplasmic reticulum protein-containing complex**
- 4/25 inner dynein arm**
- 13/92 dynein complex**
- 24/173 plasma membrane bounded cell projection cytoplasm**
- 21/155 microtubule associated complex**
- 6/41 kinesin complex**
- 131/766 microtubule cytoskeleton**
- 69/451 microtubule organizing center**
- 16/105 ciliary basal body**
- 7/57 ciliary transition zone**
- 4/49 ciliary tip**
- 81/428 cilium**
- 1/8 mitotic spindle pole body
- 7/13 Lewy body
- 35/205 motile cilium**
- 8/12 tubulin complex
- 7/28 chaperone complex**
- 1/8 zona pellucida receptor complex
- 4/21 apoplast**
- 9/44 plasmodesma**
- 5/31 plastid envelope**
- 16/85 plastid**
- 9/52 obsolete chloroplast part**
- 7/22 thylakoid**
- 5/21 fungal-type cell wall**
- 11/53 cell wall**
- 53/222 external encapsulating structure
- 2/8 phragmoplast
- 10/64 cytoplasmic stress granule
- 5/7 amino acid transport complex
- 53/235 plasma membrane protein complex
- 10/33 tertiary granule membrane
- 25/83 leading edge membrane
- 49/213 cell leading edge
- 21/106 apical junction complex
- 26/123 cell-substrate junction
- 3/16 MCM complex
- 2/13 replication fork protection complex**
- 25/114 oxidoreductase complex**
- 4/12 succinate dehydrogenase complex
- 5/19 ribonucleoside-diphosphate reductase complex**
- 8/36 organellar small ribosomal subunit
- 14/83 organellar ribosome**
- 59/160 small ribosomal subunit
- 85/239 large ribosomal subunit**

$p < 0.001$
 $p < 0.005$
 $p < 0.01$