



*3/21 proteasome core complex*  
 15/108 peptidase complex  
 0/13 proteasome storage granule  
*17/90 acetyltransferase complex*  
*9/34 endoplasmic reticulum-Golgi intermediate compartment membrane*  
**19/79 endoplasmic reticulum-Golgi intermediate compartment**  
**14/95 endoplasmic reticulum protein-containing complex**  
**69/451 microtubule organizing center**  
 39/329 centrosome  
**131/766 microtubule cytoskeleton**  
**16/105 ciliary basal body**  
**2/32 intraciliary transport particle**  
**81/428 cilium**  
**4/47 ciliary tip**  
**7/57 ciliary transition zone**  
 6/35 photoreceptor connecting cilium  
*12/59 photoreceptor cell cilium*  
 1/8 mitotic spindle pole body  
**8/66 obsolete axoneme part**  
**24/173 plasma membrane bounded cell projection cytoplasm**  
**21/155 microtubule associated complex**  
**5/41 axonemal dynein complex**  
**10/67 dynein complex**  
**4/25 inner dynein arm**  
*7/12 Lewy body*  
**24/154 motile cilium**  
 2/8 phragmoplast  
**6/41 kinesin complex**  
*8/12 tubulin complex*  
**4/21 apoplast**  
**9/44 plasmodesma**  
**5/31 plastid envelope**  
**16/85 plastid**  
**9/52 obsolete chloroplast part**  
**6/36 plastid stroma**  
**7/22 thylakoid**  
**5/21 fungal-type cell wall**  
**11/53 cell wall**  
 53/222 external encapsulating structure  
 3/13 hyphal cell wall  
 10/64 cytoplasmic stress granule  
**4/20 chaperonin-containing T-complex**  
*1/8 zona pellucida receptor complex*  
**7/28 chaperone complex**  
 26/123 cell-substrate junction  
 38/205 adherens junction  
 25/83 leading edge membrane  
 49/213 cell leading edge  
 21/106 apical junction complex  
 10/51 cation channel complex  
 10/33 tertiary granule membrane  
 8/36 organellar small ribosomal subunit  
**14/83 organellar ribosome**  
**145/426 ribosome**  
*59/160 small ribosomal subunit*  
**85/239 large ribosomal subunit**  
 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**  
*5/7 amino acid transport complex*

**p < 0.001**  
 p < 0.005  
 p < 0.01