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6/12 proton-transporting two-sector ATPase complex, proton-transporting domain
11/29 proton-transporting two-sector ATPase complex
4/12 proton-transporting two-sector ATPase complex, catalytic domain
125/606 mitochondrial part
4/11 mitochondrial respiratory chain complex III
40/145 mitochondrial membrane part
16/55 respiratory chain
63/304 organelle inner membrane
6/28 mitochondrial respiratory chain complex I
19/72 oxidoreductase complex
8/23 cytochrome complex
4/11 respiratory chain complex IV
4/11 respiratory chain complex IV
96/425 mitochondrial envelope
4/6 mitochondrial proton-transporting ATP synthase complex, catalytic core F
9/13 mitochondrial proton-transporting ATP synthase complex
 180/685 organelle envelope
 42/289 mitochondrial matrix
 4/12 aminoacyl-tRNA synthetase multienzyme complex 4/6 WASH complex
 65/213 early endosome
 6/11 pseudopodium
 5/12 guanyl-nucleotide exchange factor complex
8/8 endocytic vesicle lumen
 2/5 outer acrosomal membrane
6/12 endoplasmic reticulum chaperone complex
16/27 smooth endoplasmic reticulum
 4/6 alveolar lamellar body 5/9 lamellar body
 10/33 azurophil granule membrane
 125/460 vacuole
0/6 vacuolar proton–transporting V–type ATPase, V1 domain 37/167 vacuolar membrane 83/296 vacuolar part 64/172 extracellular matrix
 43/114 extracellular matrix part
167/662 endoplasmic reticulum part
7/7 collagen trimer
6/9 oligosaccharyltransferase complex
 37/202 intrinsic component of organelle membrane
21/92 intrinsic component of endoplasmic reticulum membrane
 4/6 integral component of lumenal side of endoplasmic reticulum membrane
 3/10 endoplasmic reticulum quality control compartment
2/11 COPI–coated vesicle membrane
73/299 Golgi membrane
9/49 Golgi–associated vesicle membrane
3/18 COPI–coated vesicle
 27/119 Golgi–associated vesicle 9/32 vesicle coat
 16/53 membrane coat
 16/62 endoplasmic reticulum—Golgi intermediate compartment 54/141 basolateral plasma membrane 19/45 basal plasma membrane 212/583 plasma membrane region
 36/77 basal part of cell
7/8 myelin sheath abaxonal region
22/40 lateral plasma membrane
10/17 muscle tendon junction
54/105 cell-substrate junction
8/15 hemidesmosome
29/47 cell-cell adherens junction
101/207 cell-cell junction
6/8 zonula adherens
92/167 anchoring junction
4/5 uropod
 4/5 uropod
163/371 cell junction
7/8 cell trailing edge
20/40 cell-cell contact zone
14/28 intercalated disc
 5/5 desmosome
122/231 cell leading edge
50/86 ruffle
 8/14 invadopodium
 4/5 SCAR complex
67/114 lamellipodium
36/65 filopodium
8/21 axolemma

180/449 axon

121/269 axon part

17/44 main axon

6/12 node of Ranvier
 72/145 site of polarized growth
 16/40 neuron projection membrane
73/202 cell projection membrane
42/90 leading edge membrane
191/513 synapse
 21/61 presynaptic membrane
47/127 synaptic membrane
 30/79 postsynaptic membrane
 50/102 neuron spine
  176/481 cell body
 10/20 costamere
4/11 dystrophin–associated glycoprotein complex
                                  plasma membrane
 23/56 apical cortex
 102/318 apical part of cell
    /10 cytoplasmic side of apical plasma membrane
2/80 apical junction complex
5/27 septate junction
 22/54 occluding junction
 131/508 intrinsic component of plasma membrane 4/15 potassium channel complex
 111/215 actin cytoskeleton 21/31 actin filament bundle
 24/38 actomyosin
 11/20 podosome
7/9 Arp2/3 protein complex
24/43 actin filament
40/60 cortical cytoskeleton
11/14 actomyosin contractile ring
15/24 contractile ring
 7/10 actin cortical patch
37/73 cell division site
131/277 cell cortex
 6/8 exocyst
77/141 cell cortex part
6/7 actin cap
81/152 contractile fiber
8/13 myofilament
43/76 I band
 37/59 A band
26/44 M band
13/28 striated muscle dense body
16/29 myosin complex
5/13 myosin Il complex
 11/17 unconventional myosin complex
  12/22 rhabdomere
 46/102 extrinsic component of plasma membrane
 4/5 heterotrimeric G-protein complex 71/182 extrinsic component of membrane 29/53 extrinsic component of cytoplasmic side of plasma membrane
   15/36 stereocilium
 8/12 microtubule plus-end
 51/125 microtubule

20/35 kinesin complex

6/11 mitotic spindle midzone
 19/40 spindle midzone
 72/215 spindle 35/92 spindle pole
 145/451 microtubule organizing center 6/14 pericentriolar material
 6/14 pericentriolar material
34/108 microtubule organizing center part
24/76 centriole
 7/7 ciliary transition fiber
13/56 nonmotile primary cilium
 136/378 cilium
 15/51 ciliary transition zone 34/104 ciliary basal body
 9/15 axonemal dynein complex
51/110 microtubule associated complex
6/9 inner dynein arm
             intermediate filament cytoskeleton
 11/23 DNA packaging complex
 4/8 condensin complex
7/15 nucleosome
114/315 chromatin
        nuclear nucleosome
 8/17 centromeric heterochromatin
 8/24 nuclear heterochromatin
66/196 nuclear chromatin
3/8 nuclear centromeric heterochromatin
3/8 nuclear centromeric neterochromatin
12/38 condensed chromosome, centromeric region
46/106 chromosome, centromeric region
63/170 protein–DNA complex
3/11 condensed nuclear chromosome kinetochore
100/326 nuclear chromosome
6/17 condensed nuclear chromosome, centromeric region
23/70 condensed nuclear chromosome
203/617 chromosome
 203/617 chromosome
11/23 synaptonemal complex
43/120 condensed chromosome
8/12 lateral element
 4/5 cohesin complex 6/14 female germ cell nucleus
 14/37 polytene chromosome puff
 3/5 Rb-E2F complex
 1/9 methylosome
4/20 U12-type sp
 4/20 U12-type spliceosomal complex
4/24 spliceosomal tri-snRNP complex
 17/67 small nuclear ribonucleoprotein complex
17/67 Small nuclear Houndeleoprotein.
4/17 U5 snRNP
8/38 precatalytic spliceosome
43/135 spliceosomal complex
4/8 commitment complex
21/59 U2-type spliceosomal complex
208/600 ribonucleoprotein complex
10/21 prespliceosome
7/17 U1 snRNP
26/75 catalytic step 2 spliceosome
5/22 U2 snRNP
14/34 cytoplasmic stress granule
4/32 90S preribosome
10/74 preribosome
5/33 small-subunit processome
 2/6 box C/D snoRNP complex
164/611 nucleolus
 3/19 small nucleolar ribonucleoprotein complex
2/24 preribosome, large subunit precursor
29/38 cytosolic small ribosomal subunit
31/62 small ribosomal subunit
99/202 cytosolic part
 0/8 chaperonin-containing T-complex
50/85 large ribosomal subunit
87/165 ribosome
48/54 cytosolic large ribosomal subunit
77/96 cytosolic ribosome
20/22 polysomal ribosome
39/62 polysome
19/23 rough endoplasmic reticulum membrane
26/66 rough endoplasmic reticulum
15/15 cytoplasmic side of rough endoplasmic reticulum membrane
16/23 cytoplasmic side of endoplasmic reticulum membrane
5/9 eukarvotic translation initiation factor 3 complex, eIF3m
 5/9 eukaryotic translation initiation factor 3 complex, eIF3m
5/15 eukaryotic translation initiation factor 3 complex
 2/5 multi-elF complex
7/20 proteasome regulatory particle
15/48 proteasome complex
3/9 proteasome regulatory particle, base subcomplex
2/7 nuclear proteasome complex
3/11 cytosolic proteasome complex
3/9 proteasome regulatory particle, lid subcomplex
3/7 proteasome core complex, alpha–subunit complex
5/16 proteasome core complex
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2/13 cytoplasmic ubiquitin ligase complex

2/5 prefoldin complex

p < 0.01 p < 0.05 p < 0.1