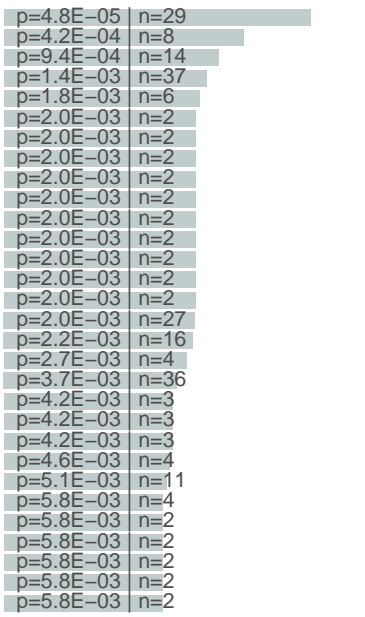


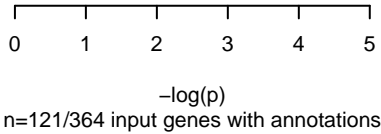
GO:BP  
Elav\_Nvec\_vc1.1\_XM\_032366225.2

fraction genes in fg and expected value

neuron projection development	GO:0031175
regulation of Ras protein signal transdu...	GO:0046578
protein complex oligomerization	GO:0051259
cell surface receptor signaling pathway	GO:0007166
positive regulation of cell–cell adhesio...	GO:0022409
positive regulation of myotube different...	GO:0010831
optic vesicle morphogenesis	GO:0003404
guanosine–containing compound biosynthes...	GO:1901070
plasma membrane raft assembly	GO:0044854
GTP biosynthetic process	GO:0006183
mesectoderm development	GO:0048383
cellular response to exogenous dsRNA	GO:0071360
lens induction in camera–type eye	GO:0060235
positive regulation of myoblast fusion	GO:1901741
establishment or maintenance of cytoskel...	GO:0003380
growth	GO:0040007
chemical synaptic transmission	GO:0007268
adenylate cyclase–activating G protein–c...	GO:0007189
response to external stimulus	GO:0009605
regulation of triglyceride biosynthetic ...	GO:0010866
regulation of intracellular estrogen rec...	GO:0033146
activin receptor signaling pathway	GO:0032924
negative regulation of protein depolymer...	GO:1901880
nematode larval development	GO:0002119
negative regulation of mitochondrion org...	GO:0010823
positive regulation of interferon–gamma ...	GO:0032729
lumen formation, open tracheal system	GO:0035149
post–embryonic body morphogenesis	GO:0040032
microvillus assembly	GO:0030033
negative regulation of triglyceride meta...	GO:0090209



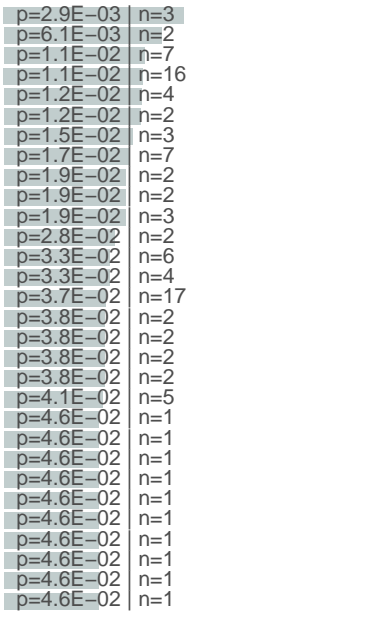
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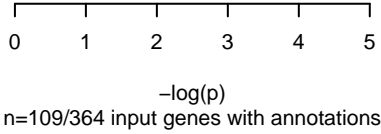
GO:MF  
Elav\_Nvec\_vc1.1\_XM\_032366225.2

fraction genes in fg and expected value

phosphatidylcholine binding	GO:0031210
ligase activity, forming phosphoric este...	GO:0016886
enzyme inhibitor activity	GO:0004857
cytoskeletal protein binding	GO:0008092
phosphoric diester hydrolase activity	GO:0008081
phosphatidylethanolamine binding	GO:0008429
disordered domain specific binding	GO:0097718
microtubule binding	GO:0008017
glucosyltransferase activity	GO:0046527
AP–2 adaptor complex binding	GO:0035612
ammonium ion binding	GO:0070405
BH domain binding	GO:0051400
actin binding	GO:0003779
UDP–glucosyltransferase activity	GO:0008194
signaling receptor binding	GO:0005102
UDP–galactose:beta–N–acetylglucosamine b...	GO:0008499
UDP–galactosyltransferase activity	GO:0035250
phosphatase inhibitor activity	GO:0019212
spectrin binding	GO:0030507
hexosyltransferase activity	GO:0016758
neuregulin binding	GO:0038132
phosphatidylinositol phosphate kinase ac...	GO:0016307
1–phosphatidylinositol–4–phosphate 5–kin...	GO:0016308
1–phosphatidylinositol–5–phosphate 4–kin...	GO:0016309
oxidative DNA demethylase activity	GO:0035516
sodium:dicarboxylate symporter activity	GO:0017153
MHC protein complex binding	GO:0023023
MHC class Ib protein complex binding	GO:0023025
MHC class Ib protein binding	GO:0023029
glutaminase activity	GO:0004359



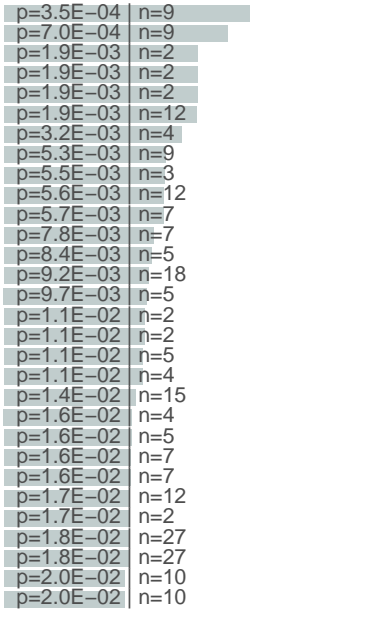
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GO:CC  
Elav\_Nvec\_vc1.1\_XM\_032366225.2

fraction genes in fg and expected value

lamellipodium	GO:0030027
synaptic membrane	GO:0097060
photoreceptor outer segment membrane	GO:0042622
uropod	GO:0001931
collagen and cuticulin–based cuticle ext...	GO:0060102
cell projection membrane	GO:0031253
transcription elongation factor complex	GO:0008023
plasma membrane protein complex	GO:0098797
clathrin–coated pit	GO:0005905
presynapse	GO:0098793
growth cone	GO:0030426
cytoplasmic side of plasma membrane	GO:0009898
plasma membrane raft	GO:0044853
somatodendritic compartment	GO:0036477
extrinsic component of plasma membrane	GO:0019897
filamentous actin	GO:0031941
growth cone lamellipodium	GO:1990761
vacuolar lumen	GO:0005775
extrinsic component of cytoplasmic side ...	GO:0031234
secretory vesicle	GO:0099503
caveola	GO:0005901
brush border	GO:0005903
basolateral plasma membrane	GO:0016323
basal plasma membrane	GO:0009925
side of membrane	GO:0098552
sensory dendrite	GO:0071683
cytoplasmic vesicle	GO:0031410
intracellular vesicle	GO:0097708
anchoring junction	GO:0070161
cell–cell junction	GO:0005911



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