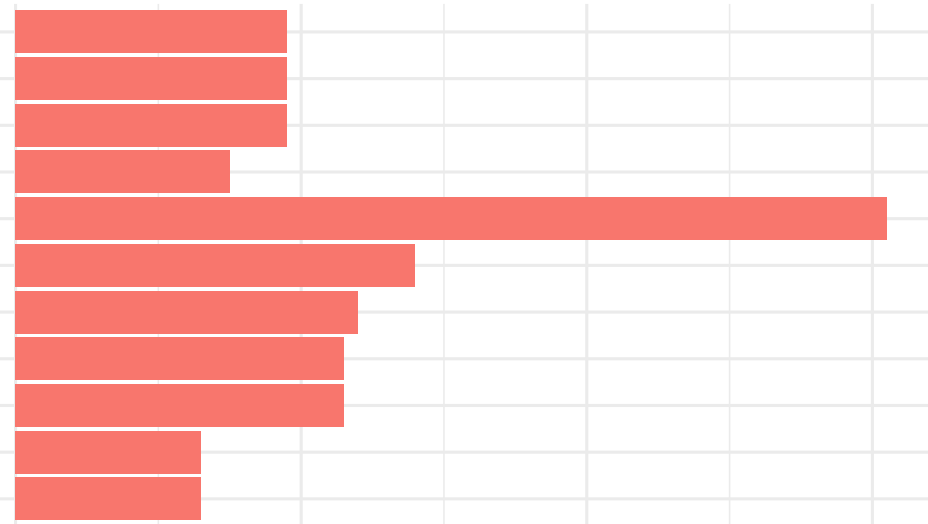


# GO Enrichment by Ontology (BP, MF, CC): AAB\_vs\_DGB

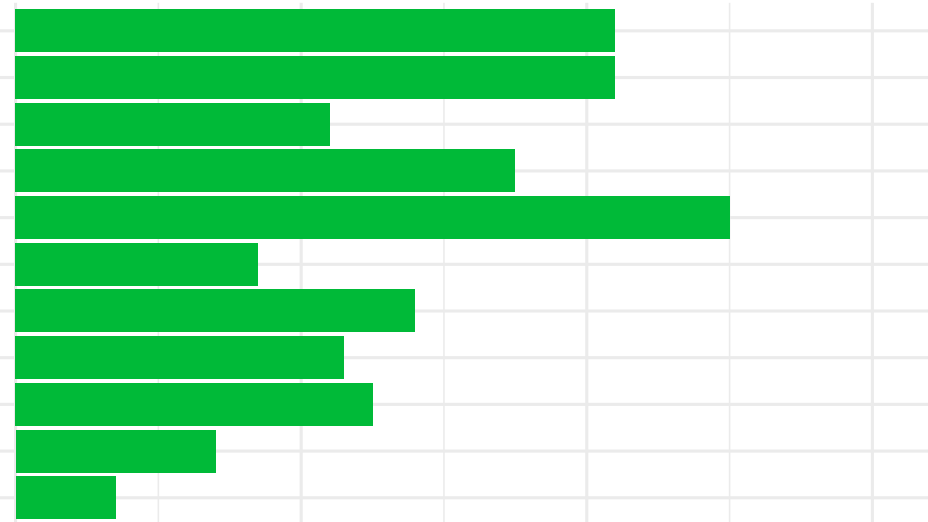
## BP

extracellular matrix organization  
external encapsulating structure organization  
extracellular structure organization  
amino acid biosynthetic process  
cell development  
axon development  
alpha-amino acid metabolic process  
organic acid biosynthetic process  
carboxylic acid biosynthetic process  
L-amino acid biosynthetic process  
proteinogenic amino acid biosynthetic process



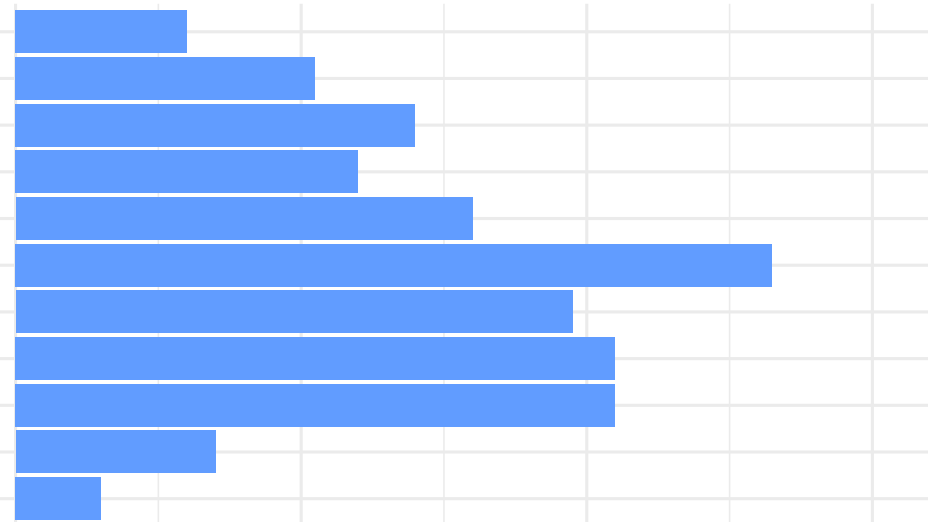
## CC

extracellular matrix  
external encapsulating structure  
collagen-containing extracellular matrix  
cell surface  
neuron projection  
postsynaptic membrane  
receptor complex  
external side of plasma membrane  
postsynapse  
plasma membrane signaling receptor complex  
postsynaptic density membrane



## MF

extracellular matrix structural constituent  
metalloendopeptidase activity  
metallopeptidase activity  
cell adhesion molecule binding  
gated channel activity  
peptidase activity  
endopeptidase activity  
channel activity  
passive transmembrane transporter activity  
amino acid transmembrane transporter activity  
Hsp70 protein binding



## Ontology



Gene Count