Table S4 GO Categories for the genes reactive to Ifi27l2a , Sh3gl3 and Kcmf1

transition metal ion transport 0.001 protein import into nucleus, translocation 0.002 iron ion transport 0.004 protein import into nucleus 0.006 regulation of endocytosis 0.006 positive regulation of endocytosis 0.006 positive regulation of phagocytosis 0.006 nuclear import 0.006 metal ion transport 0.008 protein targeting 0.009 protein localization in nucleus 0.009 negative regulation of multicellular organismal process 0.009 cation transport 0.012 protein localization in organelle 0.015 regulation of vesicle-mediated transport 0.015 B. Sh3gl3 multicellular organismal process 0 mesoderm development 0.003 multicellular organismal development 0.004 cell migration 0.004
protein import into nucleus, translocation iron ion transport protein import into nucleus  regulation of endocytosis positive regulation of endocytosis positive regulation of phagocytosis nuclear import positive regulation of phagocytosis nuclear import protein targeting protein import protein localization in nucleus process cation transport nucleocytoplasmic transport nuclear transport protein localization in organelle protein localization in organelle regulation of vesicle-mediated transport protein localization in organelle regulation of multicellular organismal process  B. Sh3gl3 multicellular organismal process oregulation of multicellular organismal development ocological services ocological service
iron ion transport  protein import into nucleus  regulation of endocytosis  positive regulation of endocytosis  regulation of phagocytosis  positive regulation of phagocytosis  positive regulation of phagocytosis  positive regulation of phagocytosis  nuclear import  nuclear import  protein targeting  protein targeting  protein localization in nucleus  negative regulation of multicellular organismal  process  cation transport  nucleocytoplasmic transport  nuclear transport  protein localization in organelle  protein localization in organelle  regulation of vesicle-mediated transport  B. Sh3gl3  multicellular organismal process  regulation of multicellular organismal process  nucleocytophasmic process  oregulation of multicellular organismal process  output  nuclear transport  output  o
protein import into nucleus  regulation of endocytosis  positive regulation of endocytosis  positive regulation of phagocytosis  positive regulation transport  protein targeting  positive regulation of nucleus  protein import  protein localization in nucleus  process  cation transport  protein localization of multicellular organismal  process  cation transport  protein localization in organelle  protein localization in organelle  protein localization in organelle  protein localization of vesicle-mediated transport  protein localization of multicellular organismal process  pregulation of multicellular organismal development  protein localization in organismal development  protein l
regulation of endocytosis  positive regulation of endocytosis  regulation of phagocytosis  positive regulation of phagocytosis  positive regulation of phagocytosis  nuclear import  nuclear import  protein targeting  protein import  protein localization in nucleus  negative regulation of multicellular organismal  process  cation transport  nucleocytoplasmic transport  nuclear transport  protein localization in organelle  nuclear transport  protein localization in organelle  protein localization in organelle  nuclear transport
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regulation of phagocytosis  positive regulation of phagocytosis  nuclear import  nuclear import  protein targeting  protein import  protein localization in nucleus  negative regulation of multicellular organismal  process  cation transport  nucleocytoplasmic transport  nuclear transport  protein localization in organelle  regulation of vesicle-mediated transport  B. Sh3gl3  multicellular organismal process  mesoderm development  nuclear import  nuclear development
positive regulation of phagocytosis  nuclear import  0.006  metal ion transport  0.008  protein targeting  0.009  protein import  0.009  protein localization in nucleus  negative regulation of multicellular organismal  process  0.009  cation transport  0.012  nucleocytoplasmic transport  0.012  protein localization in organelle  regulation of vesicle-mediated transport  0.015  B. Sh3gl3  multicellular organismal process  0  mesoderm development  0.003  multicellular organismal development  0.004  cell migration  0.004
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protein localization in nucleus  negative regulation of multicellular organismal process  cation transport  nucleocytoplasmic transport  nuclear transport  protein localization in organelle  regulation of vesicle-mediated transport  B. Sh3gl3  multicellular organismal process  regulation of multicellular organismal process  mesoderm development  nuclear transport  0.012  0.015  0.015  0.015  0.005  0.006  0.003  0.004
negative regulation of multicellular organismal process 0.009 cation transport 0.009 nucleocytoplasmic transport 0.012 nuclear transport 0.012 protein localization in organelle 0.015 regulation of vesicle-mediated transport 0.015  B. Sh3gl3 multicellular organismal process 0 regulation of multicellular organismal process 0 mesoderm development 0.003 multicellular organismal development 0.004 cell migration 0.004
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cation transport 0.009 nucleocytoplasmic transport 0.012 nuclear transport 0.012 protein localization in organelle 0.015 regulation of vesicle-mediated transport 0.015  B. Sh3gl3 multicellular organismal process 0 regulation of multicellular organismal process 0 mesoderm development 0.003 multicellular organismal development 0.004 cell migration 0.004
nucleocytoplasmic transport 0.012 nuclear transport 0.012 protein localization in organelle 0.015 regulation of vesicle-mediated transport 0.015  B. Sh3gl3 multicellular organismal process 0 regulation of multicellular organismal process 0 mesoderm development 0.003 multicellular organismal development 0.004 cell migration 0.004
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regulation of vesicle-mediated transport 0.015  B. Sh3gl3  multicellular organismal process 0  regulation of multicellular organismal process 0  mesoderm development 0.003  multicellular organismal development 0.004  cell migration 0.004
B. Sh3gl3  multicellular organismal process 0  regulation of multicellular organismal process 0  mesoderm development 0.003  multicellular organismal development 0.004  cell migration 0.004
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regulation of multicellular organismal process 0 mesoderm development 0.003 multicellular organismal development 0.004 cell migration 0.004
mesoderm development 0.003 multicellular organismal development 0.004 cell migration 0.004
multicellular organismal development 0.004 cell migration 0.004
cell migration 0.004
regulation of developmental process 0.004
negative regulation of cell growth 0.004
negative regulation of cell size 0.004
cellular component movement 0.005
patterning of blood vessels 0.006
olfactory bulb development 0.006
olfactory lobe development 0.006
retinal ganglion cell axon guidance 0.006
positive regulation of protein binding 0.006

regulation of protein binding	0.006
locomotion	0.007
anatomical structure formation involved in	
morphogenesis	0.007
establishment of protein localization	0.007
formation of primary germ layer	0.008
mesoderm formation	0.008
C. Kcmf1	
cholesterol metabolic process	0.007
sterol metabolic process	0.007
steroid metabolic process	0.010
response to radiation	0.020
arginine metabolic process	0.023
xenobiotic metabolic process	0.023
membrane budding	0.023
vesicle coating	0.023
striated muscle contraction	0.023
response to xenobiotic stimulus	0.023
coenzyme A metabolic process	0.023
dendrite development	0.023
cytidine to uridine editing	0.023
mRNA modification	0.023
protein sumoylation	0.023
regulation of protein sumoylation	0.023
positive regulation of protein sumoylation	0.023
nucleoside bisphosphate metabolic process	0.023
lipoprotein transport	0.023

The genes reactive to either *Ifi27l2a* (A) or *Sh3gl3* (B) or *Kcmf1* (C) were entered into a GO analysis focusing on overrepresentation of biological process terms. Shown are the top 20 GO terms and their P-values for all tables.