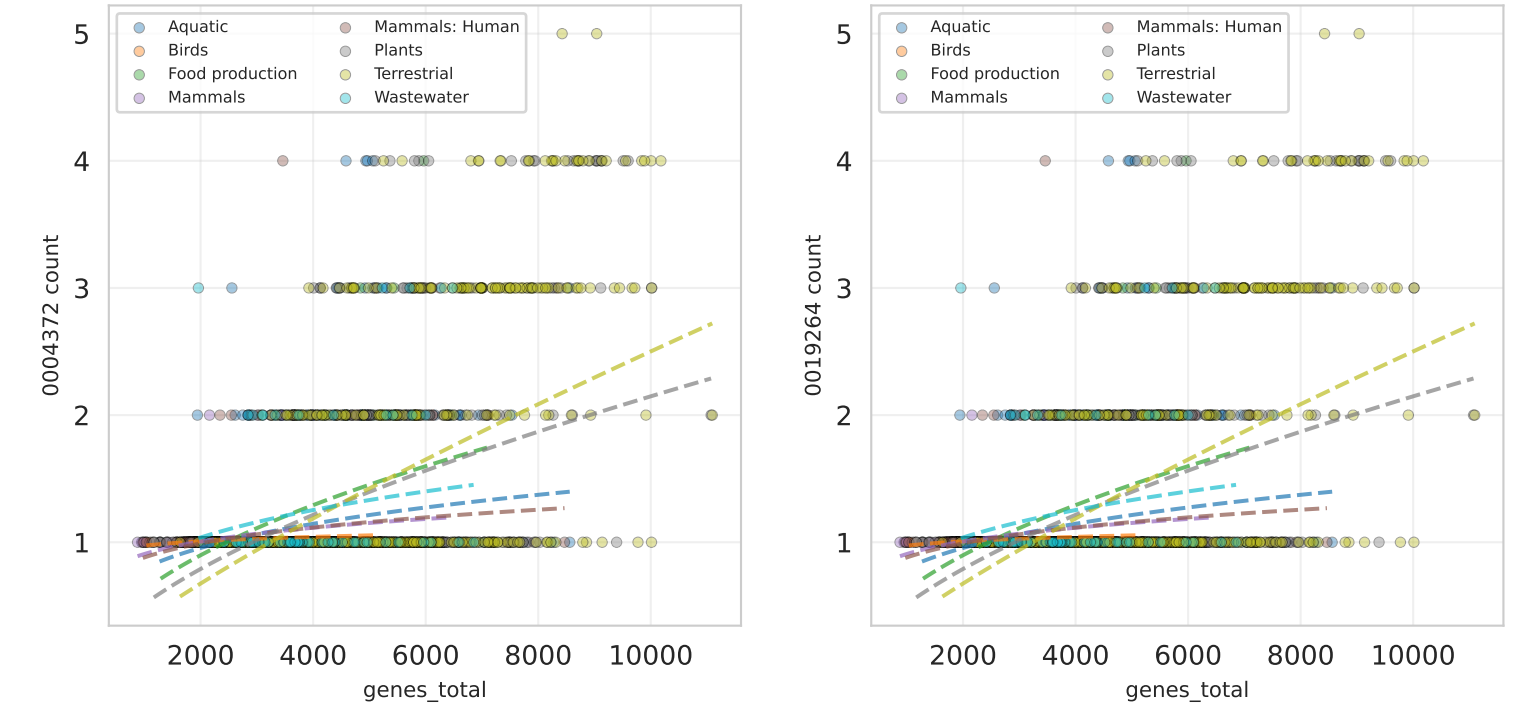
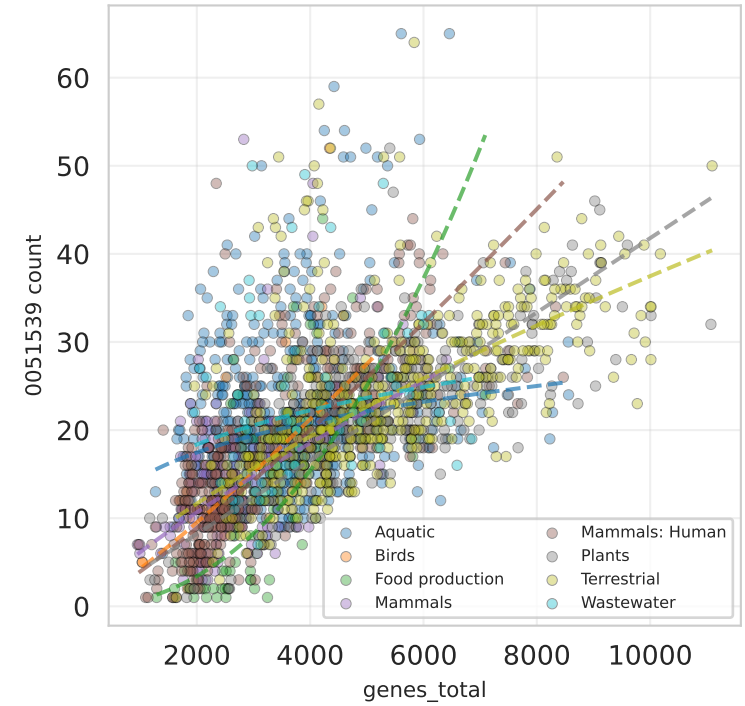


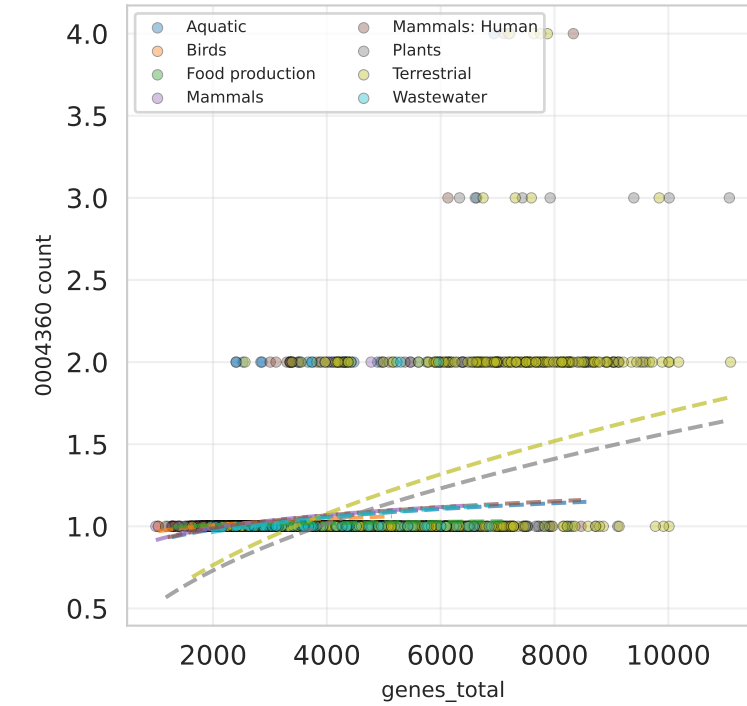
glycine hydroxymethyltransferase activity (GO:0004372) ~~glycine~~ biosynthetic process from serine (GO:0019264) (



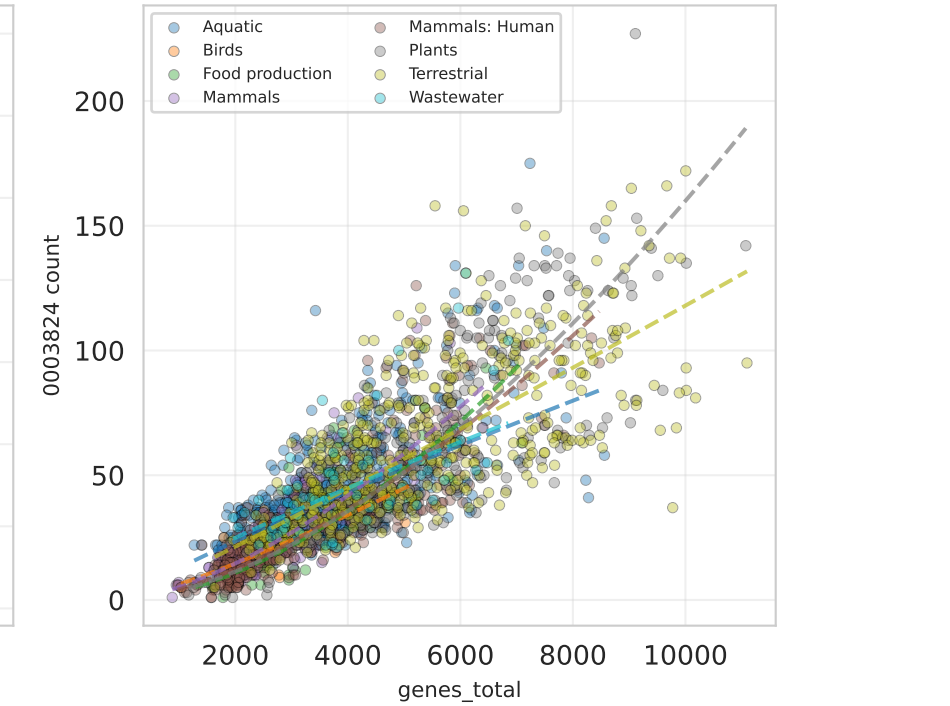
linea4)iron, 4 sulfur cluster binding (GO:0051539) (light blue)



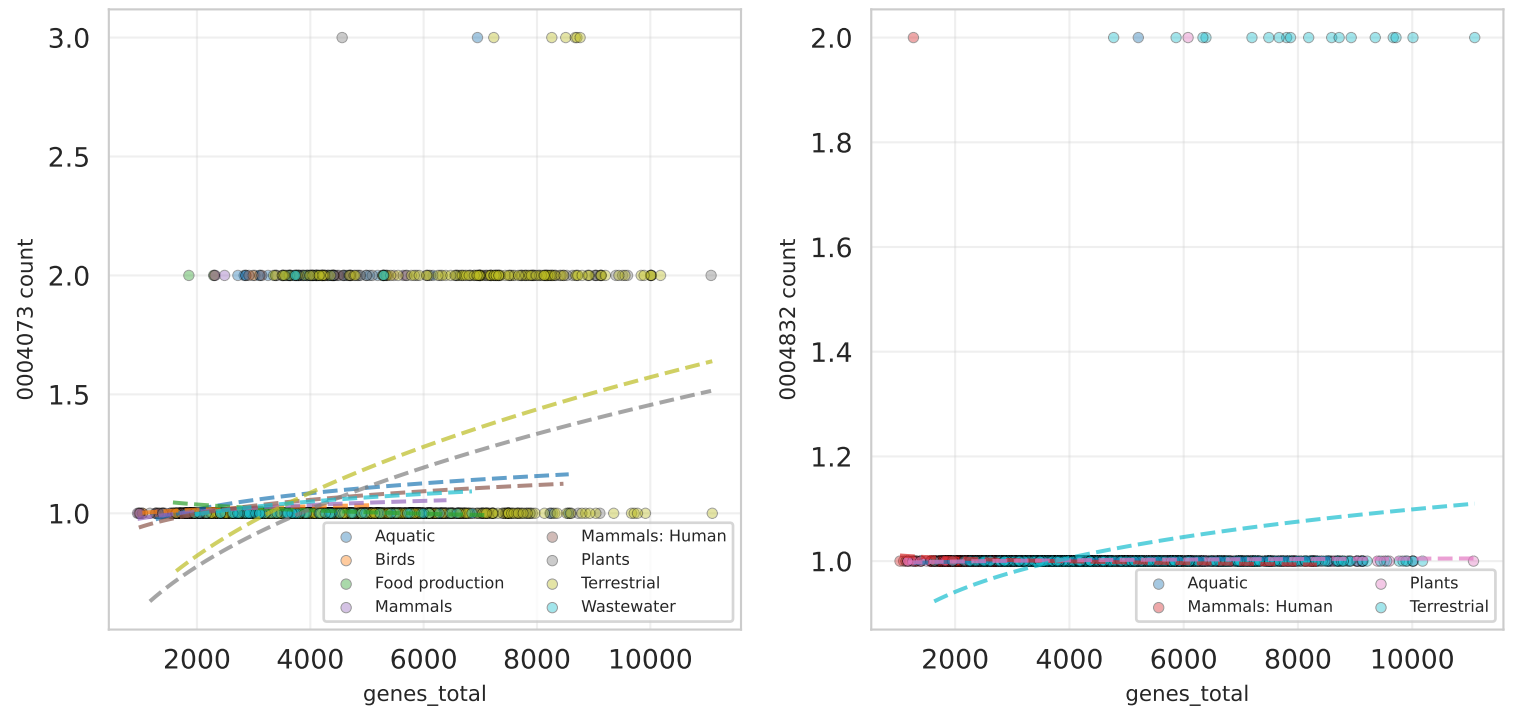
Glucose-6-phosphate transaminase (isomerizing)



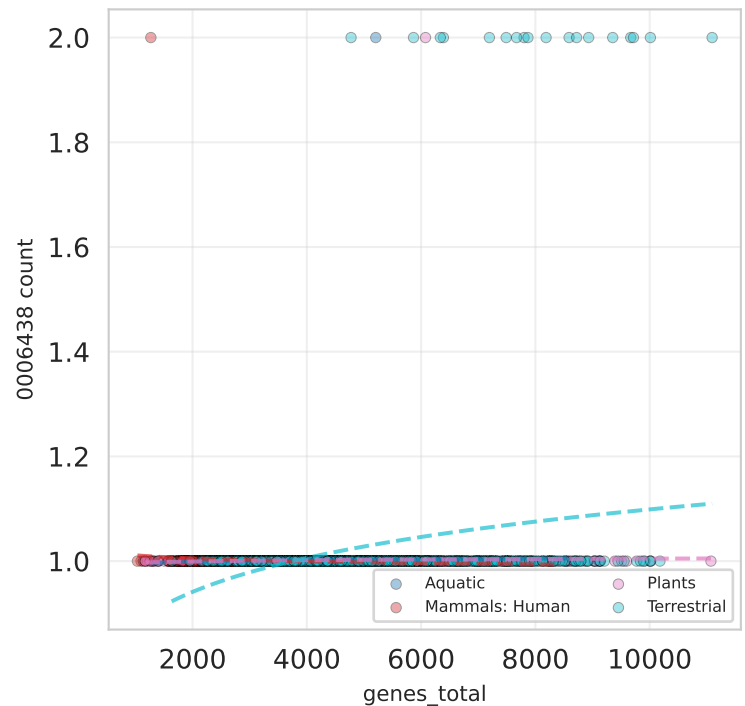
ng)... (linear) catalytic activity (GO:0003824) (linear)



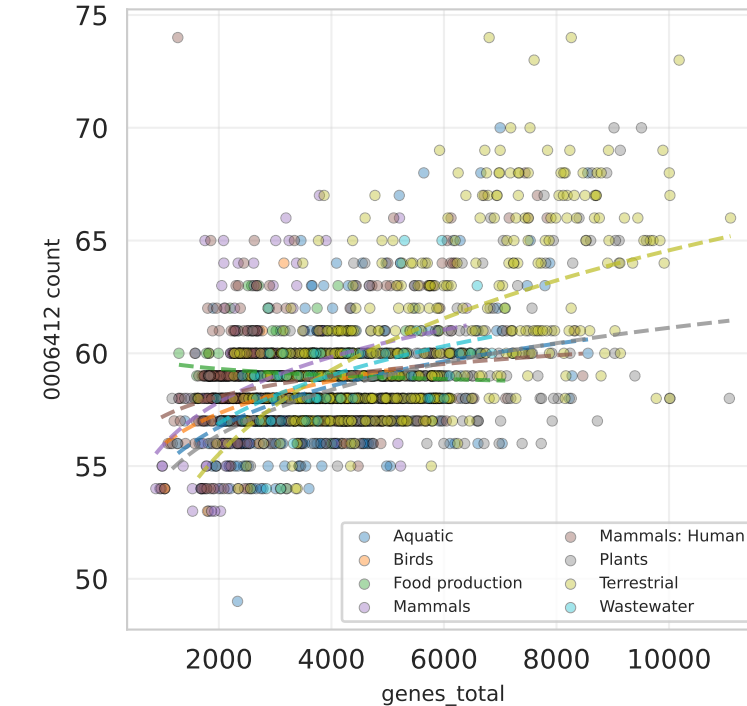
aspartate-semialdehyde dehydrogenase activity (GO:0004073) (linear) **line-tRNA ligase activity (GO:0004832) (linear)**



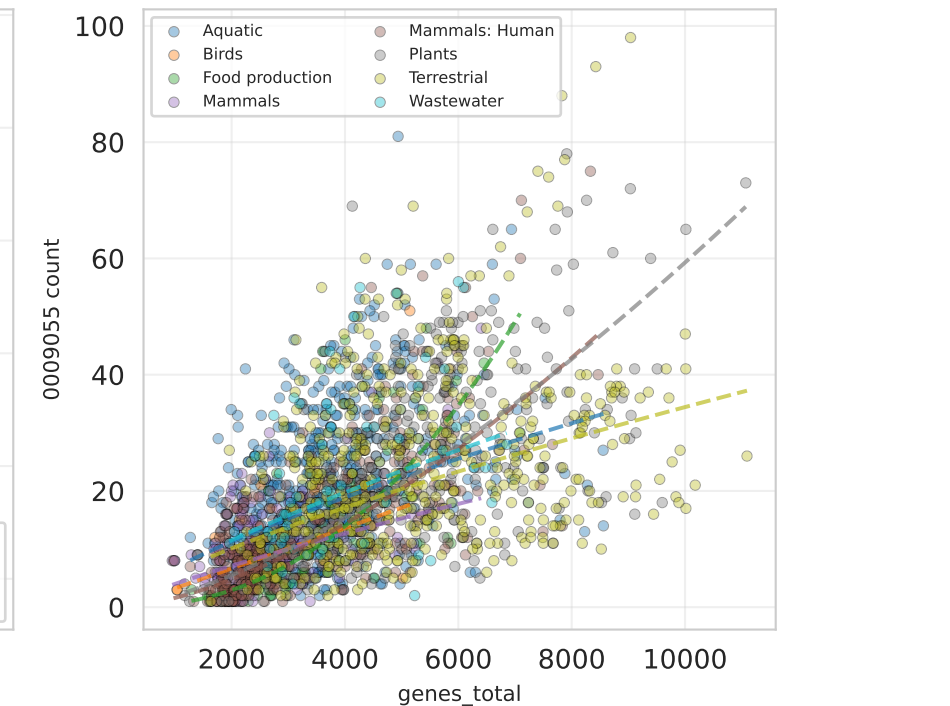
valyl-tRNA aminoacylation (GO:0006438) (linear



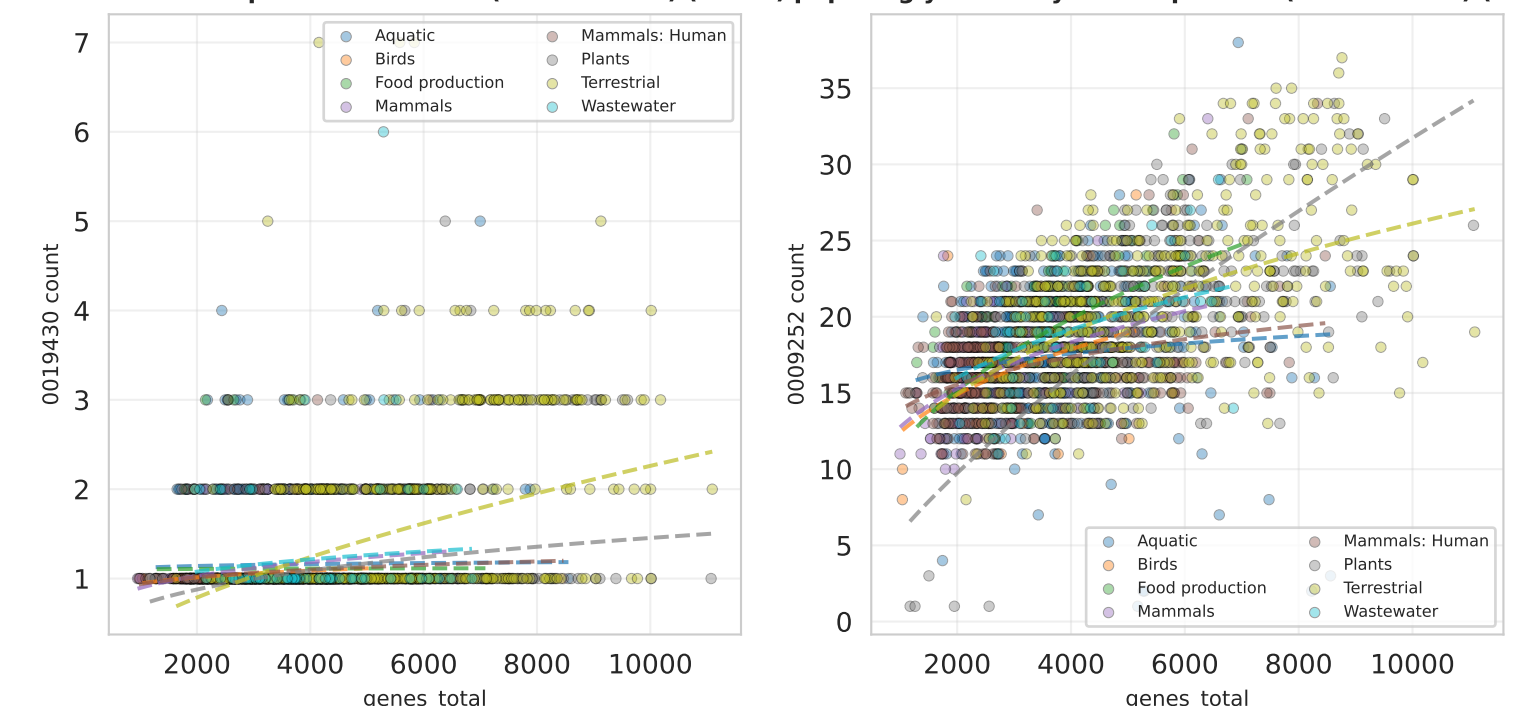
) translation (GO:0006412) (linear)



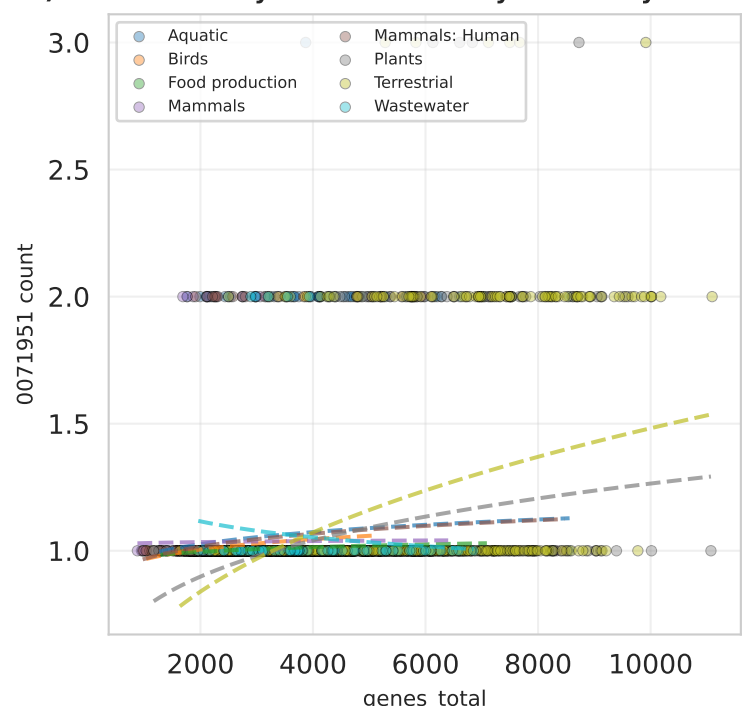
electron transfer activity (GO:0009055) (linear)



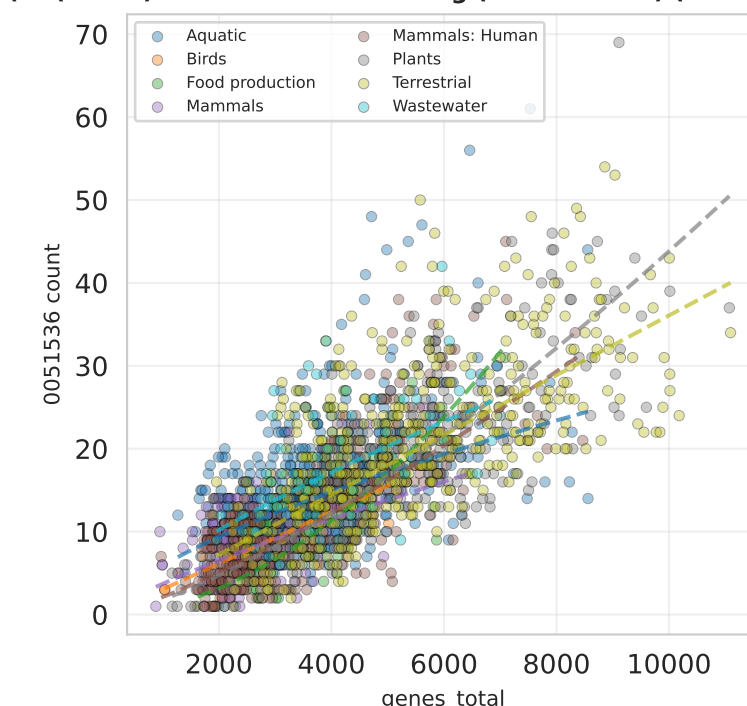
removal of superoxide radicals (GO:0019430) (linear) peptidoglycan biosynthetic process (GO:0009252) (lin



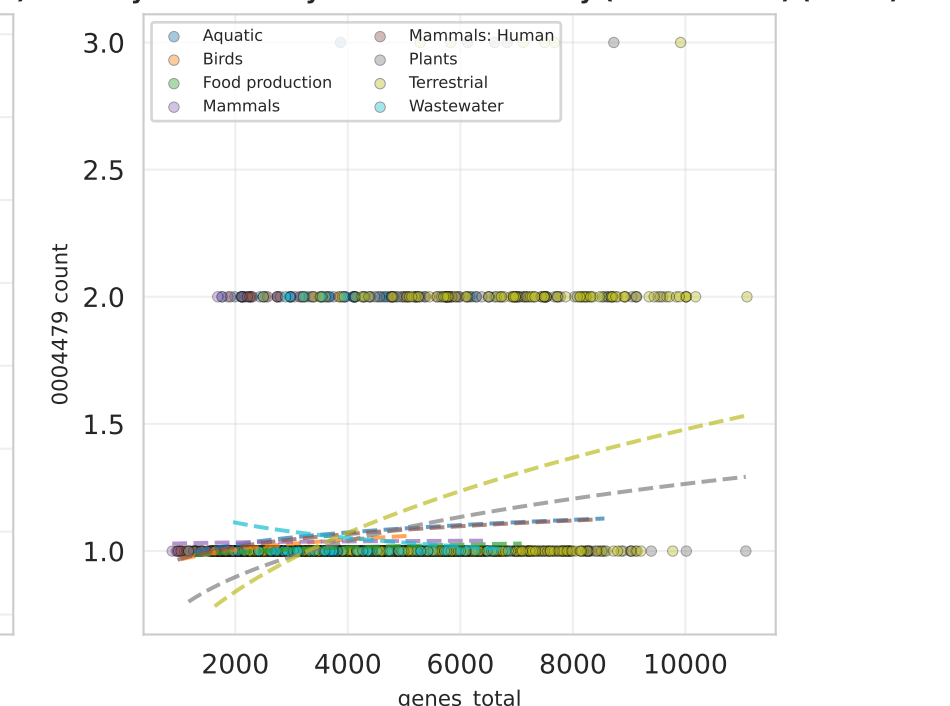
Conversion of methionyl-tRNA to N-formyl-methionyl-tRNA



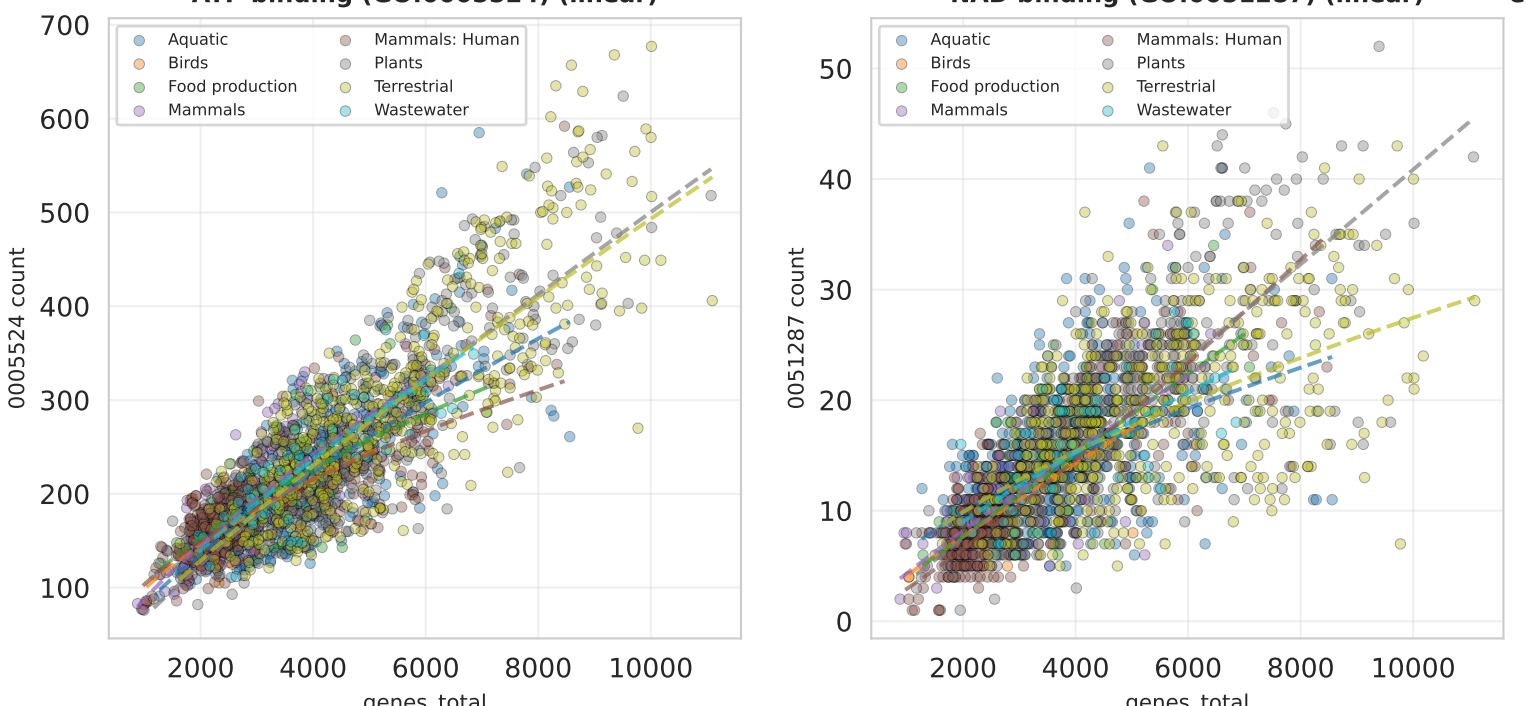
(... (line)iron-sulfur cluster binding (GO:0051536) (line



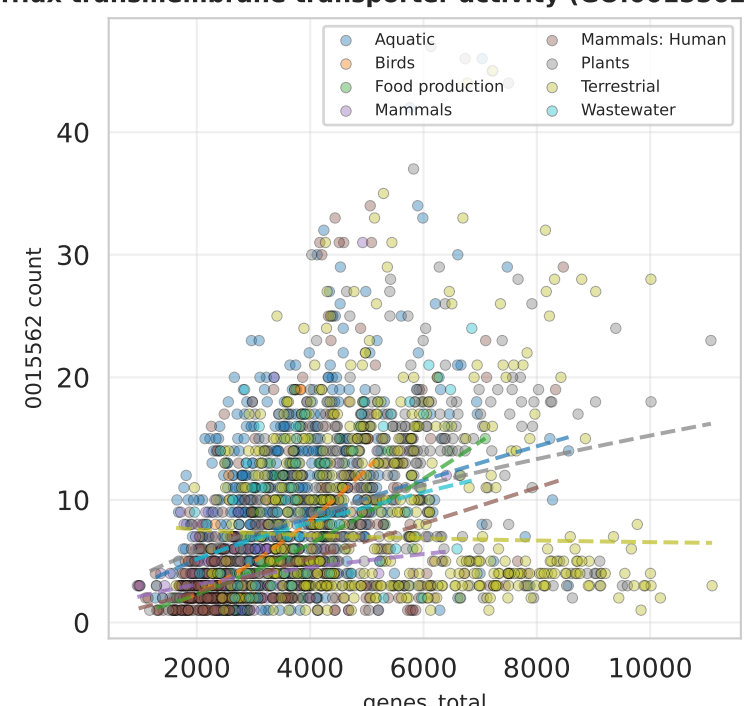
methionyl-tRNA formyltransferase activity (GO:0004479) (linear)



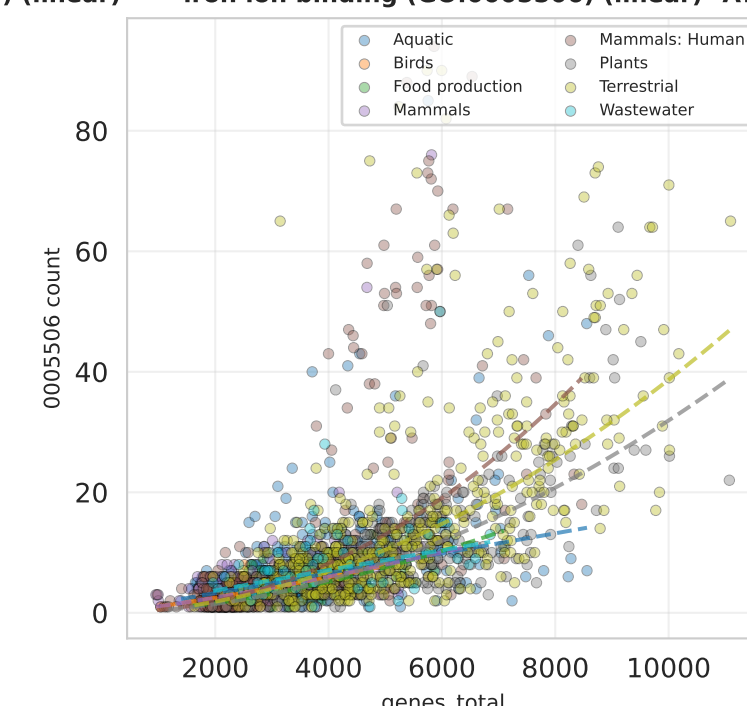
ATP binding (GO:0005524) (linear) NAD binding (GO:0051287) (linear) e



fflux transmembrane transporter activity (GO:0015562)



) (linear) iron ion binding (GO:0005506) (linear) AT



TPase-coupled transmembrane transporter activity (GO:004... (linear)

