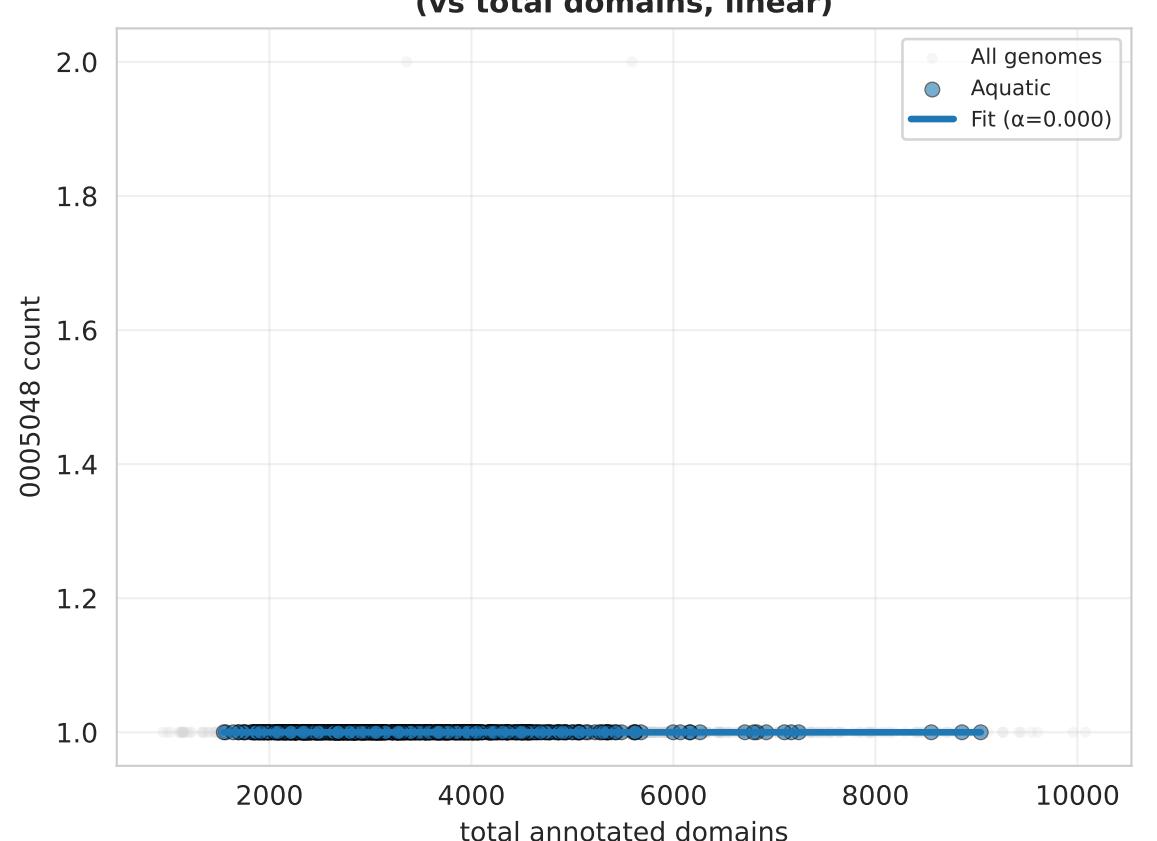
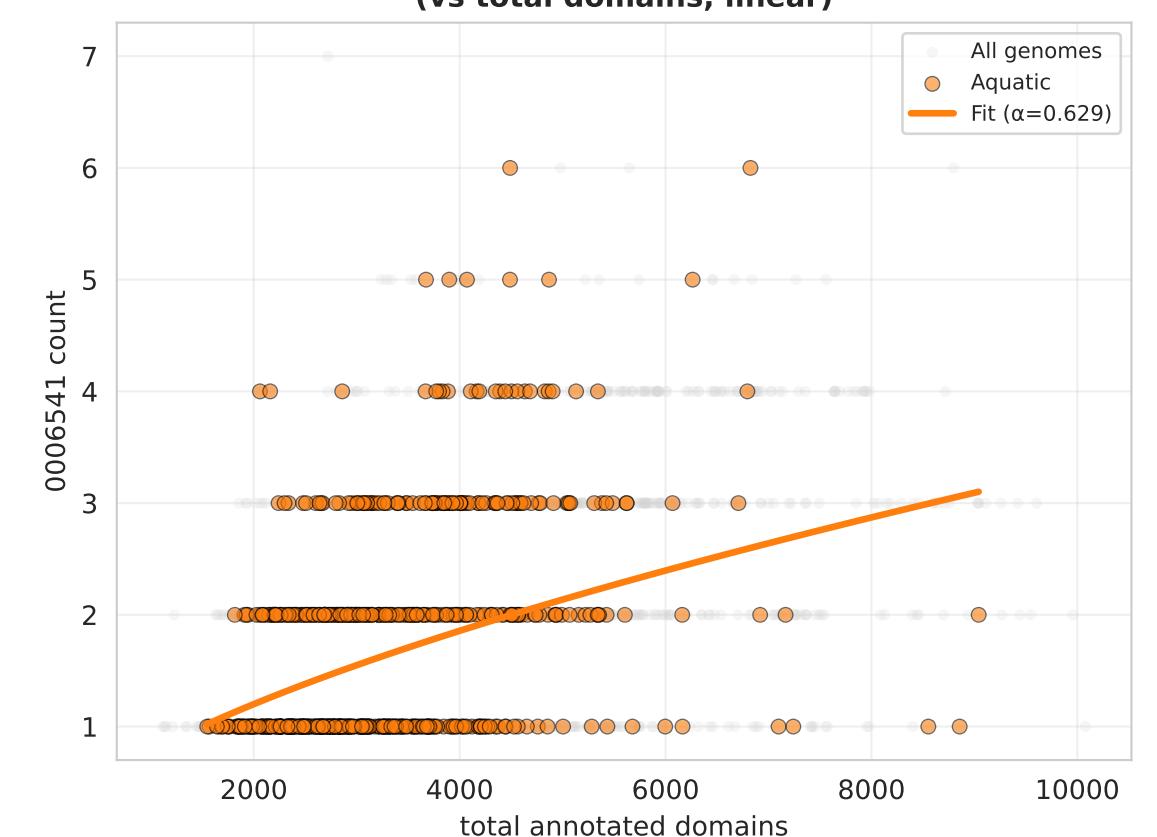


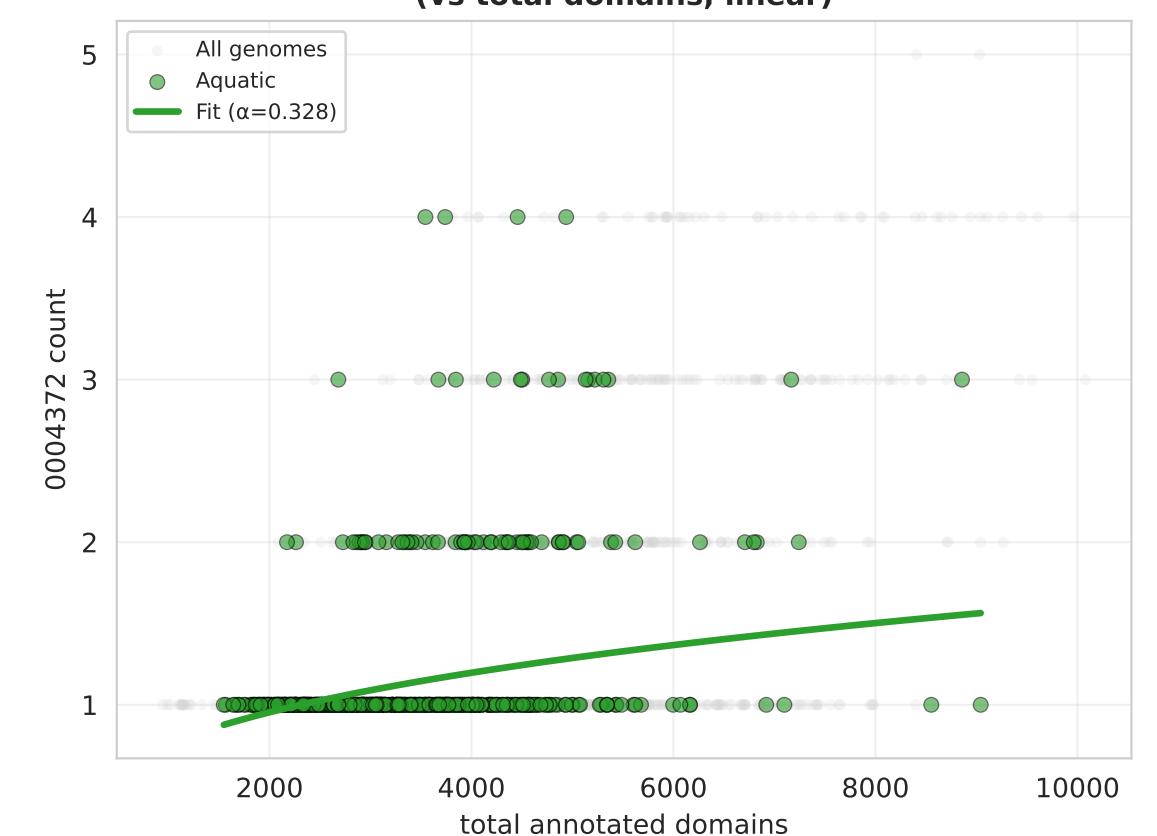
Panel t: Aquatic signal sequence binding (GO:0005048) (vs total domains, linear)



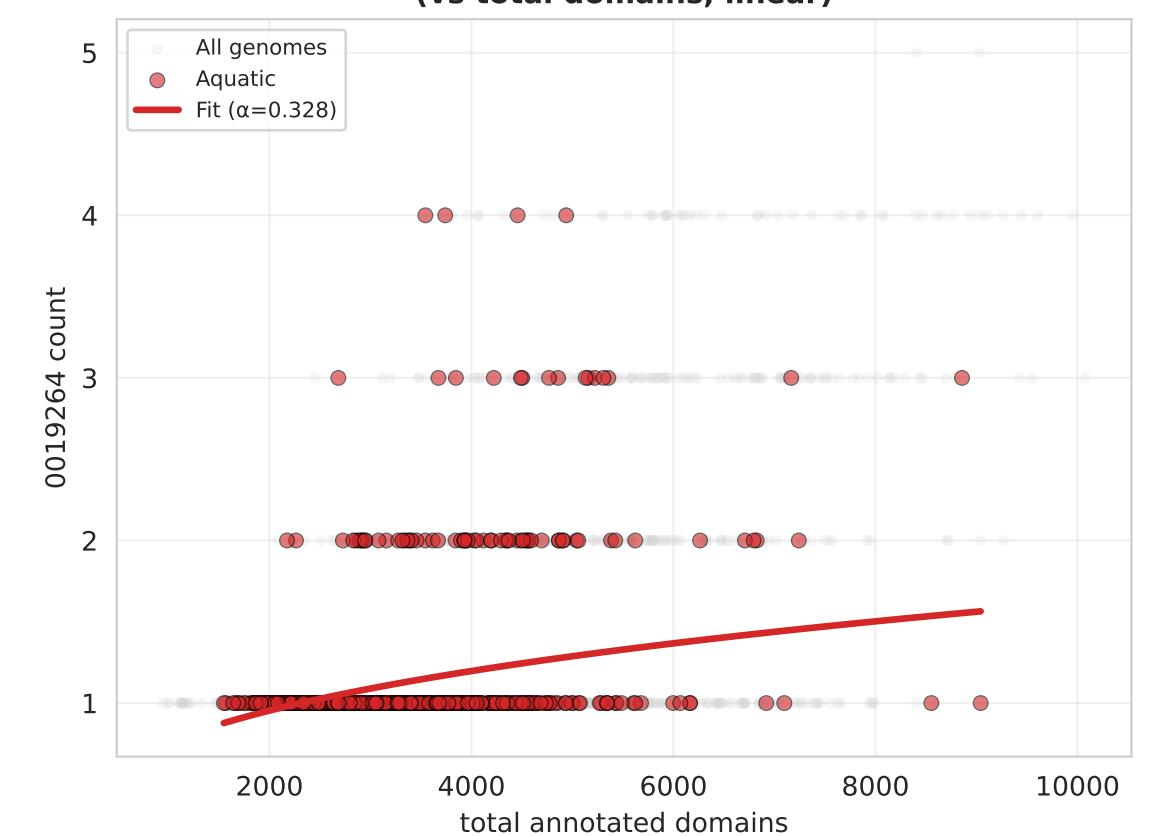
Panel t: Aquatic glutamine metabolic process (GO:0006541) (vs total domains, linear)



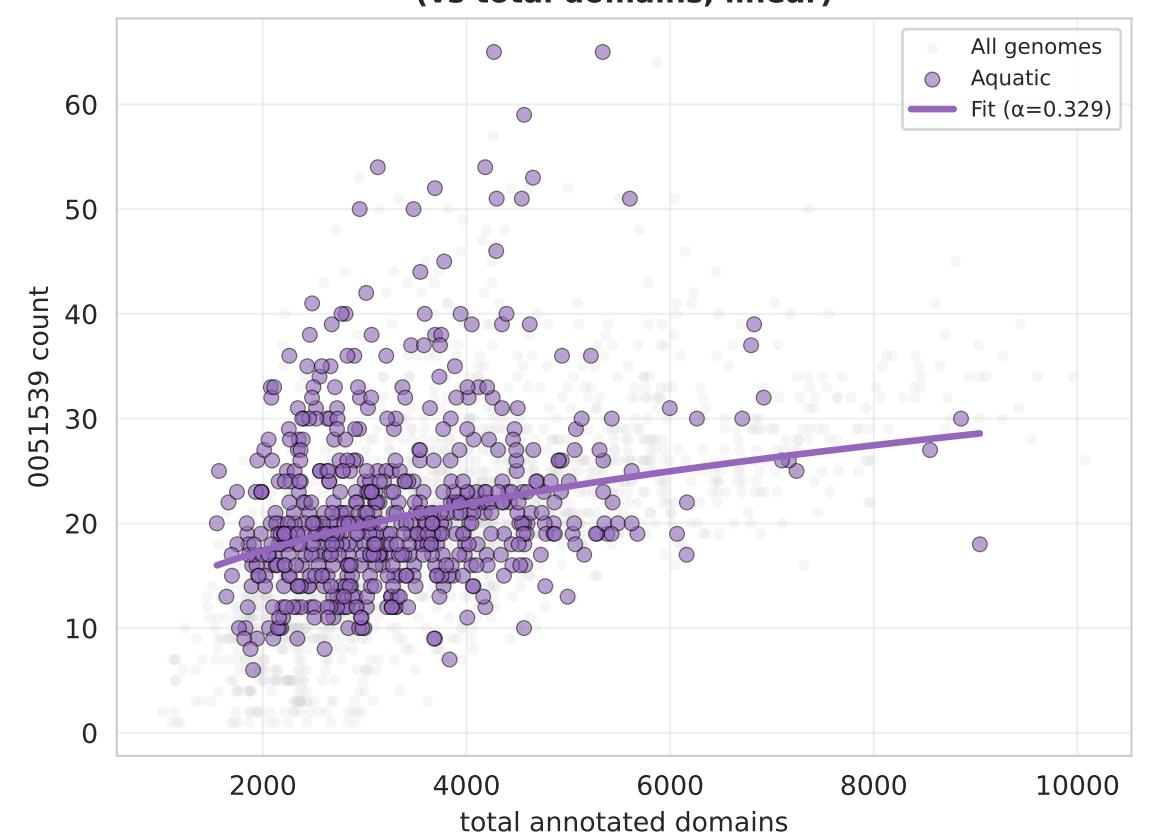
Panel t: Aquatic glycine hydroxymethyltransferase activity (GO:0004372) (vs total domains, linear)



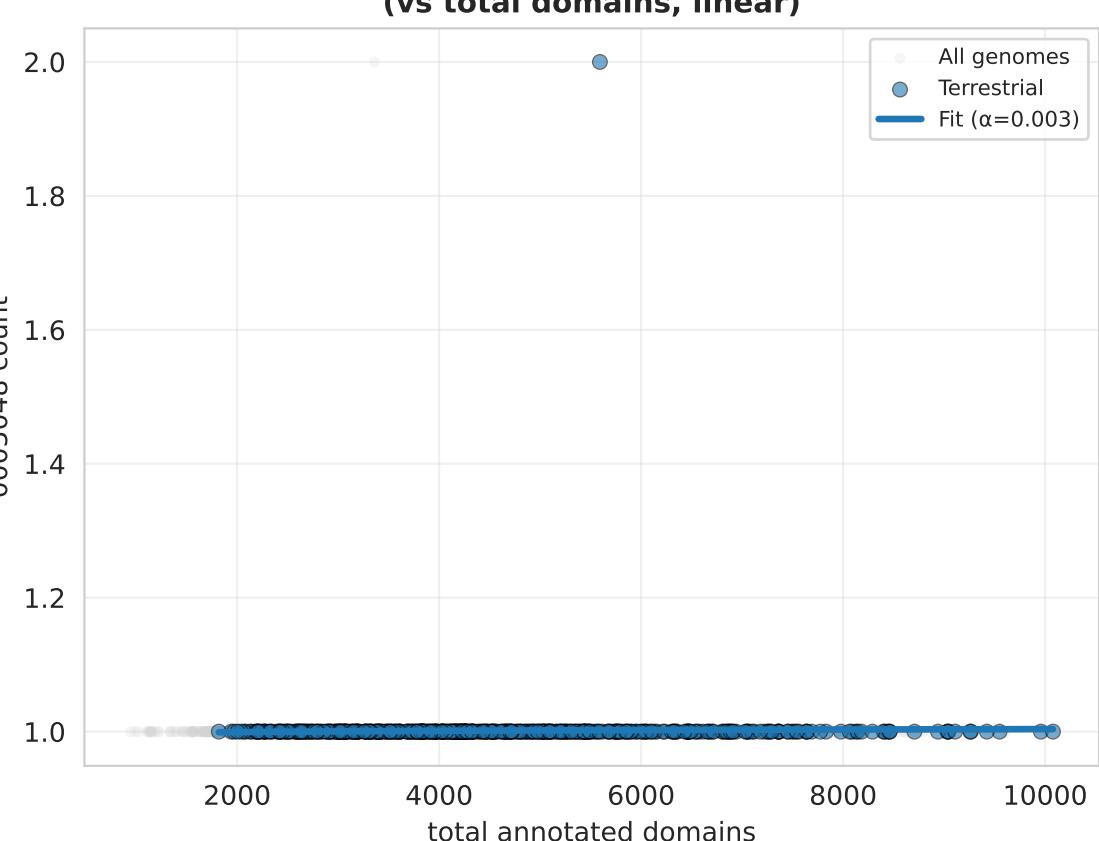
Panel t: Aquatic glycine biosynthetic process from serine (GO:0019264) (vs total domains, linear)



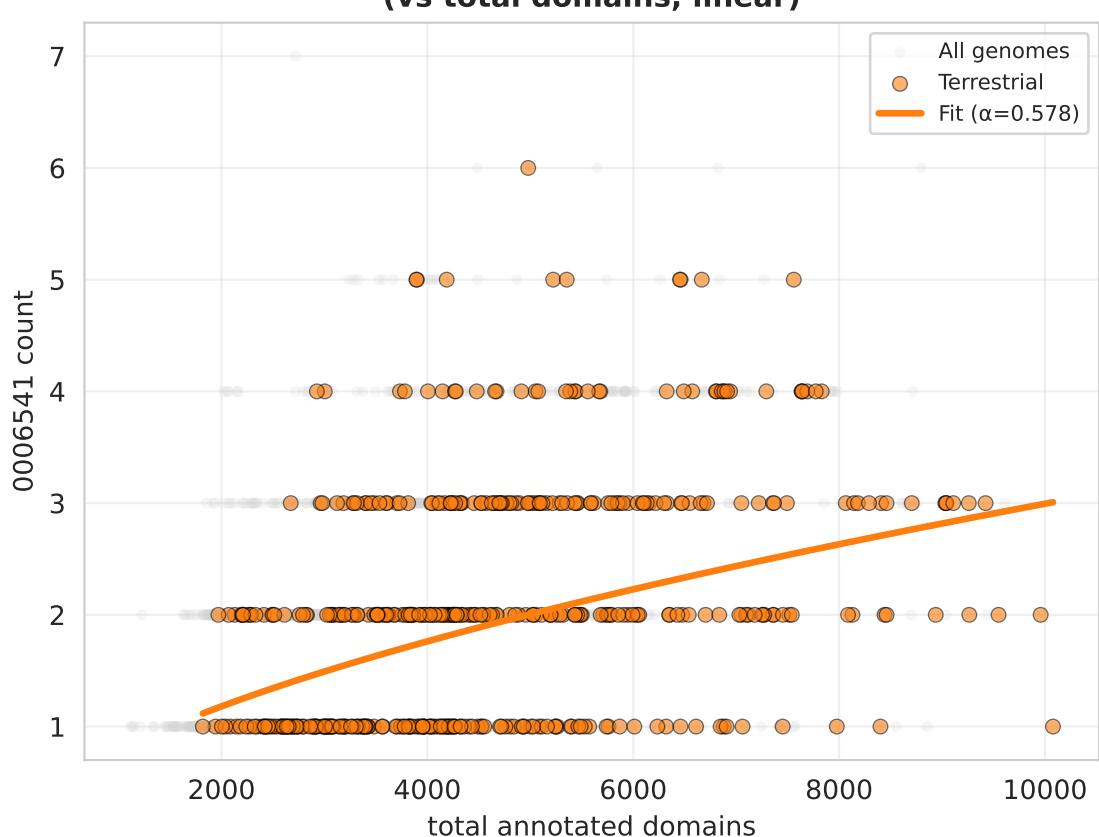
Panel t: Aquatic 4 iron, 4 sulfur cluster binding (GO:0051539) (vs total domains, linear)



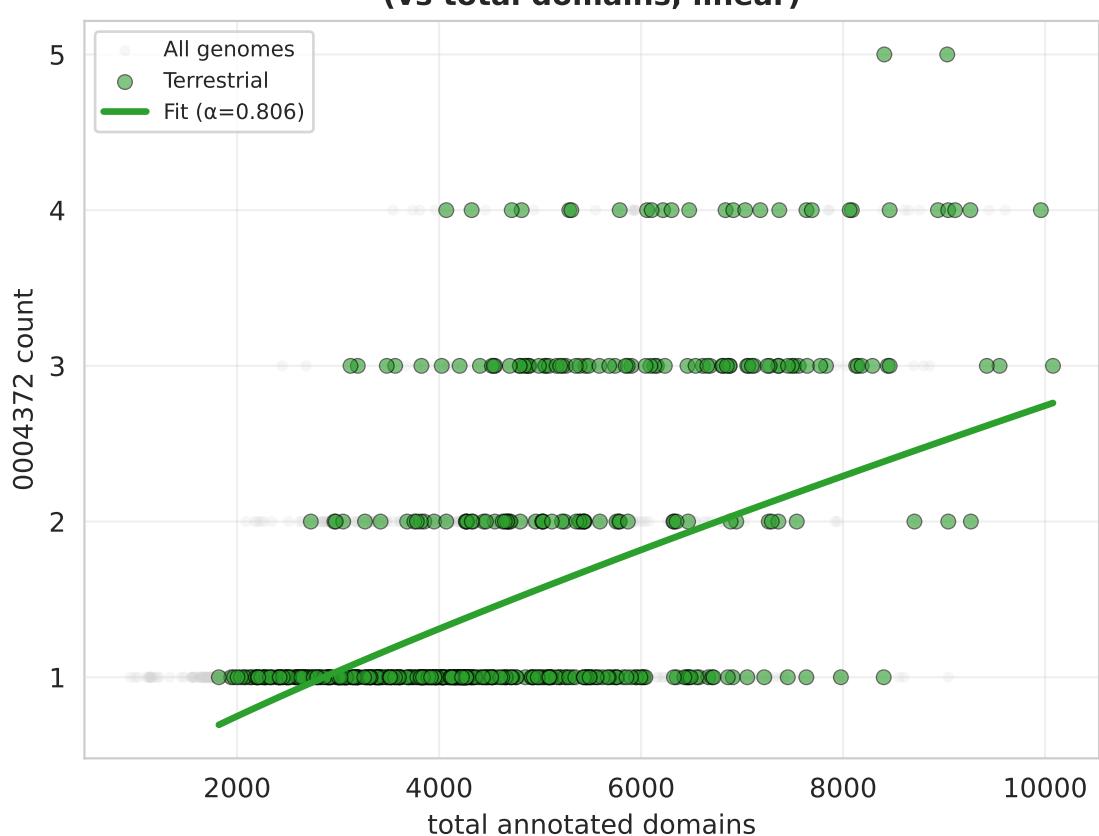
Panel t: Terrestrial signal sequence binding (GO:0005048) (vs total domains, linear)



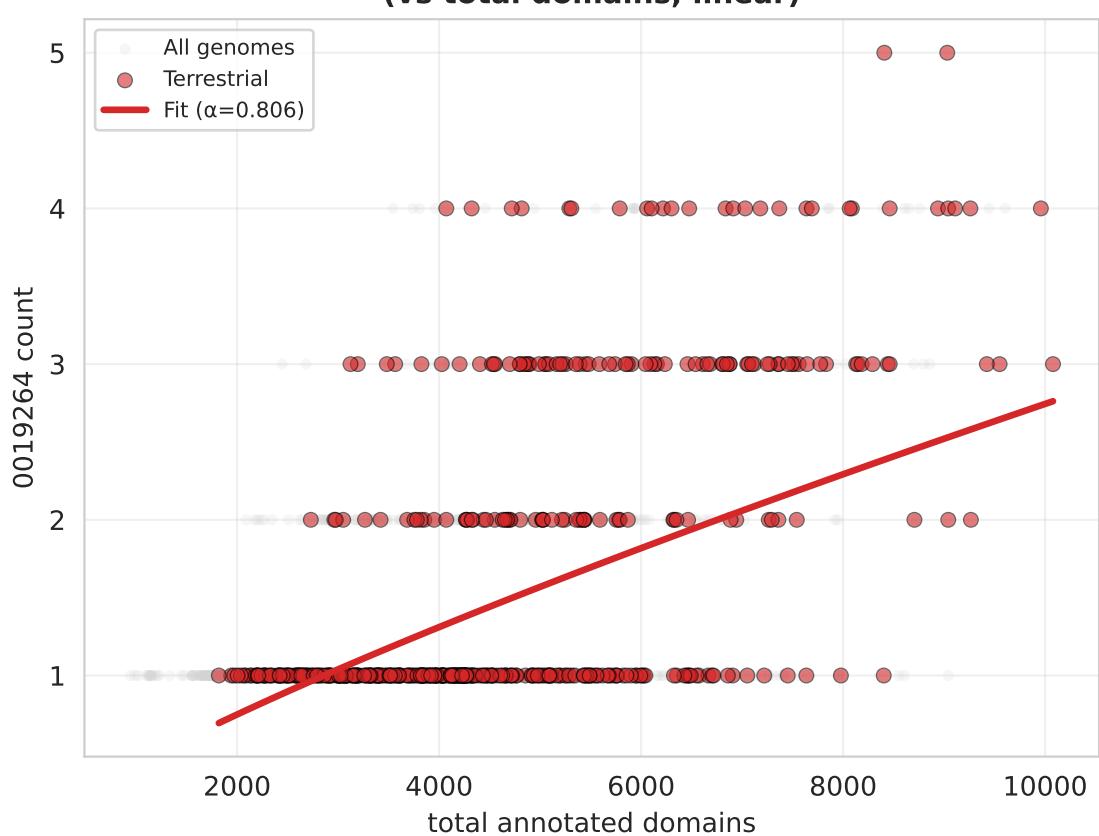
Panel t: Terrestrial glutamine metabolic process (GO:0006541) (vs total domains, linear)



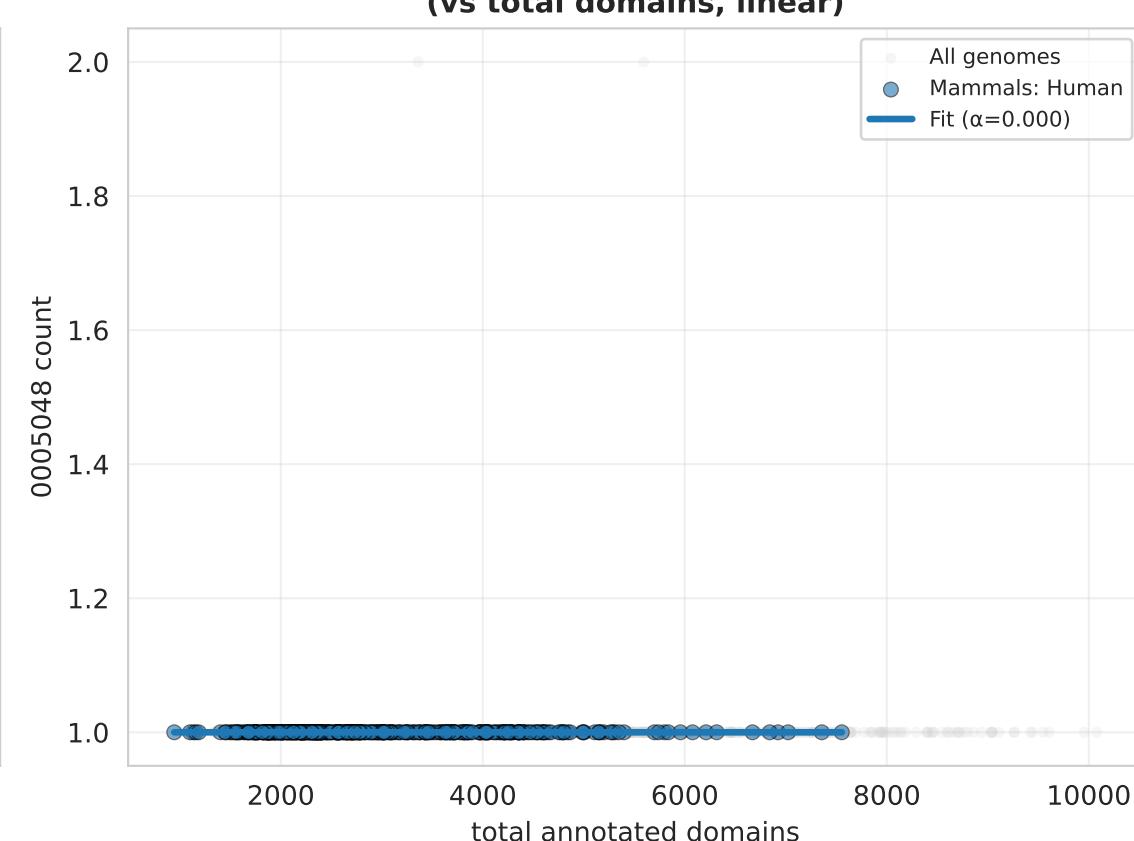
Panel t: Terrestrial glycine hydroxymethyltransferase activity (GO:0004372) (vs total domains, linear)



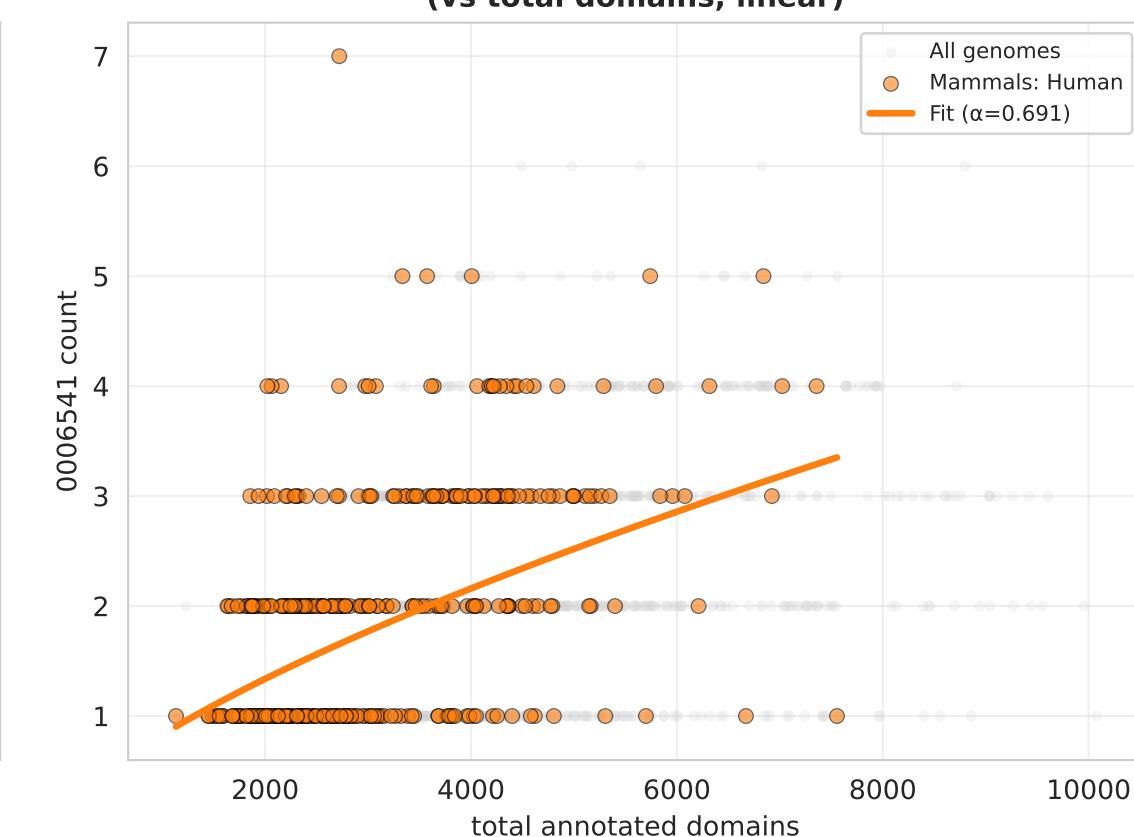
Panel t: Terrestrial glycine biosynthetic process from serine (GO:0019264) (vs total domains, linear)



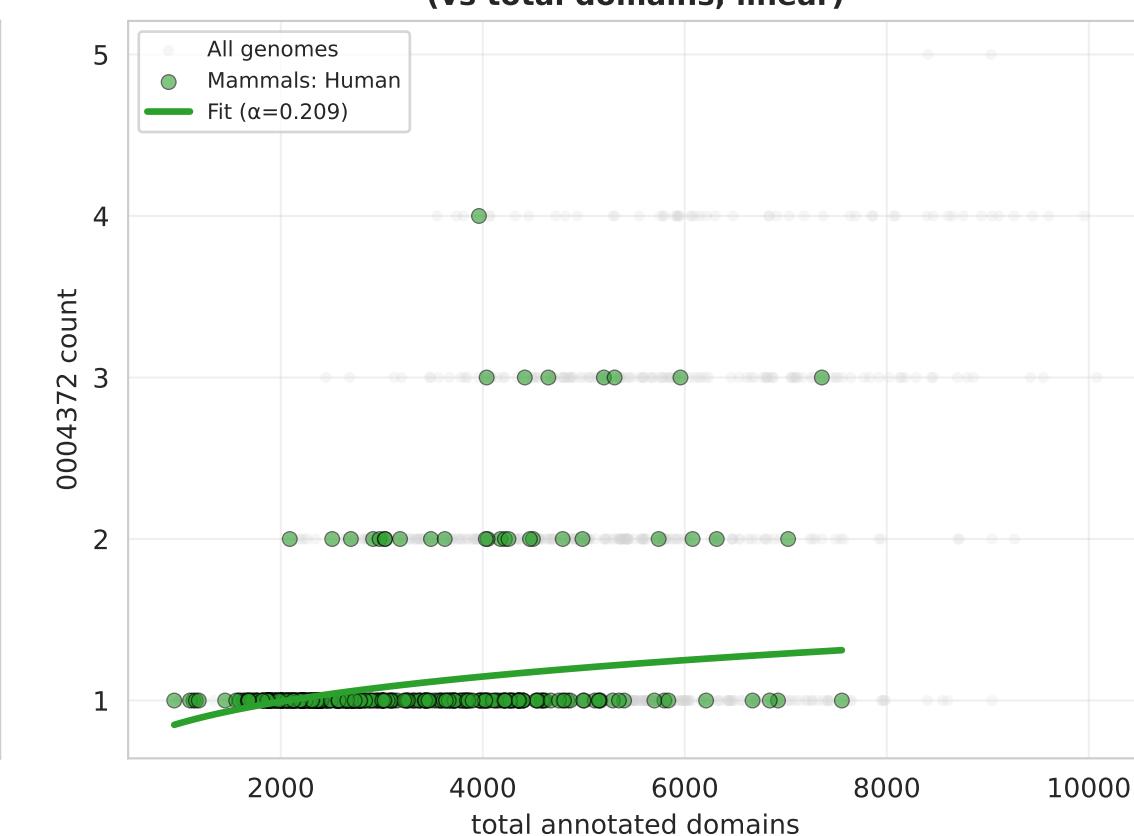
Panel t: Mammals: Human signal sequence binding (GO:0005048) (vs total domains, linear)



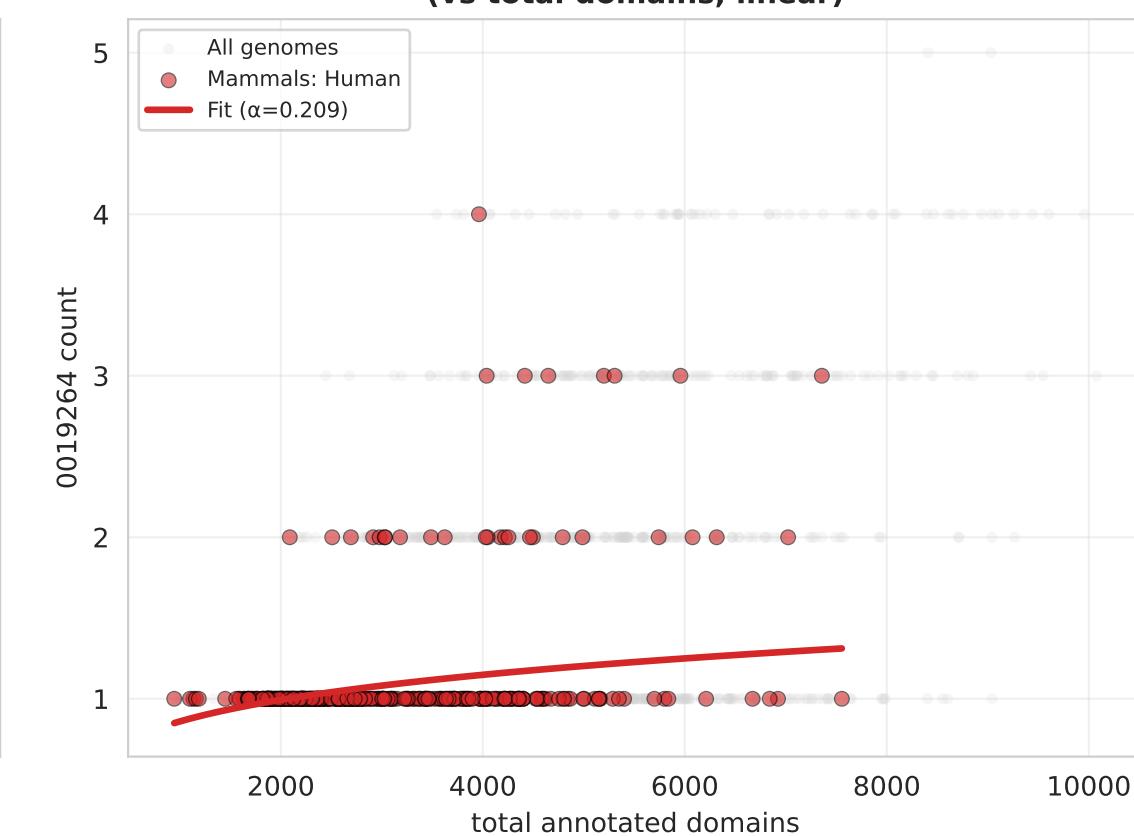
Panel t: Mammals: Human glutamine metabolic process (GO:0006541) (vs total domains, linear)



Panel t: Mammals: Human glycine hydroxymethyltransferase activity (GO:0004372) (vs total domains, linear)



Panel t: Mammals: Human glycine biosynthetic process from serine (GO:0019264) (vs total domains, linear)



Panel t: Mammals: Human 4 iron, 4 sulfur cluster binding (GO:0051539) (vs total domains, linear)

