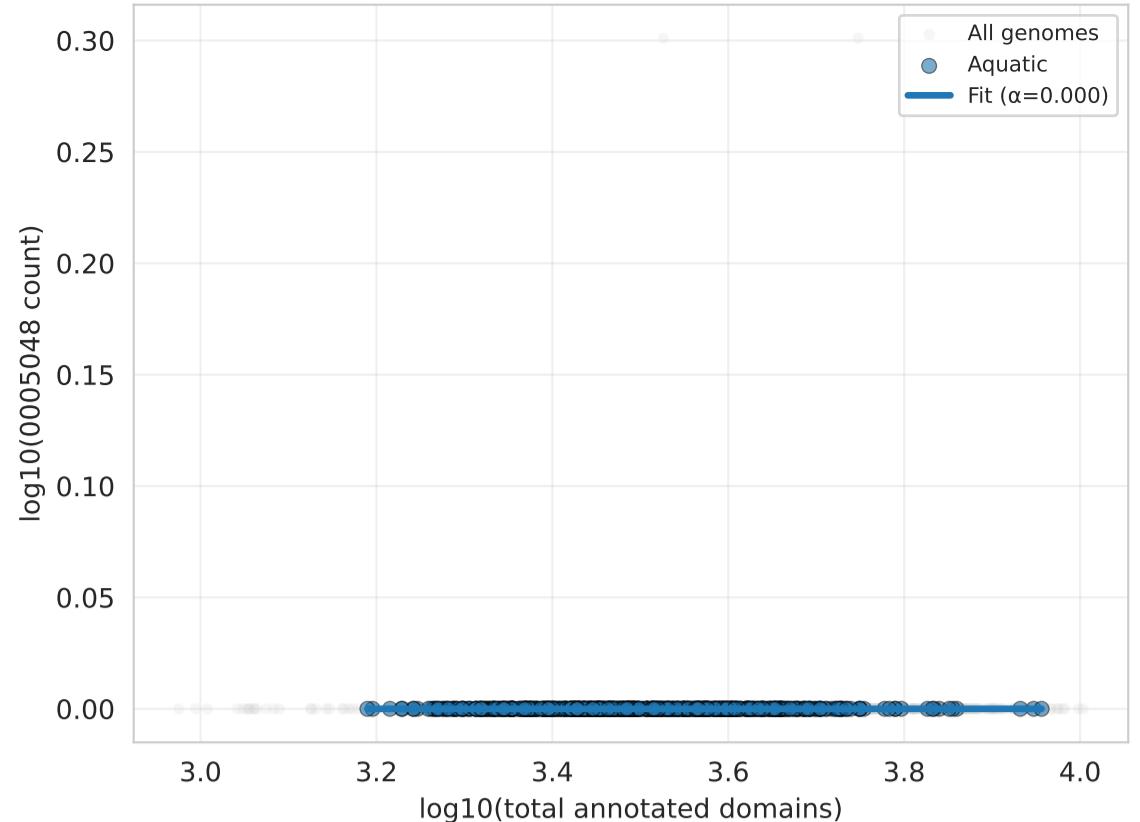
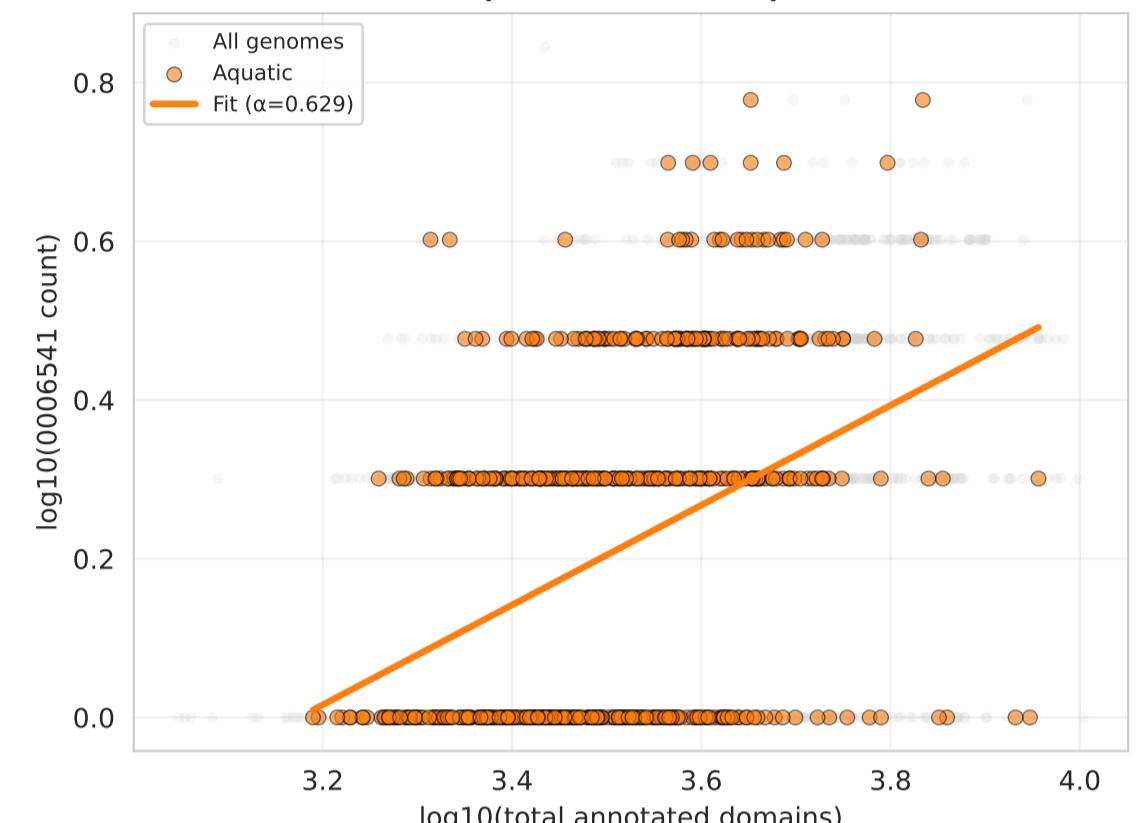


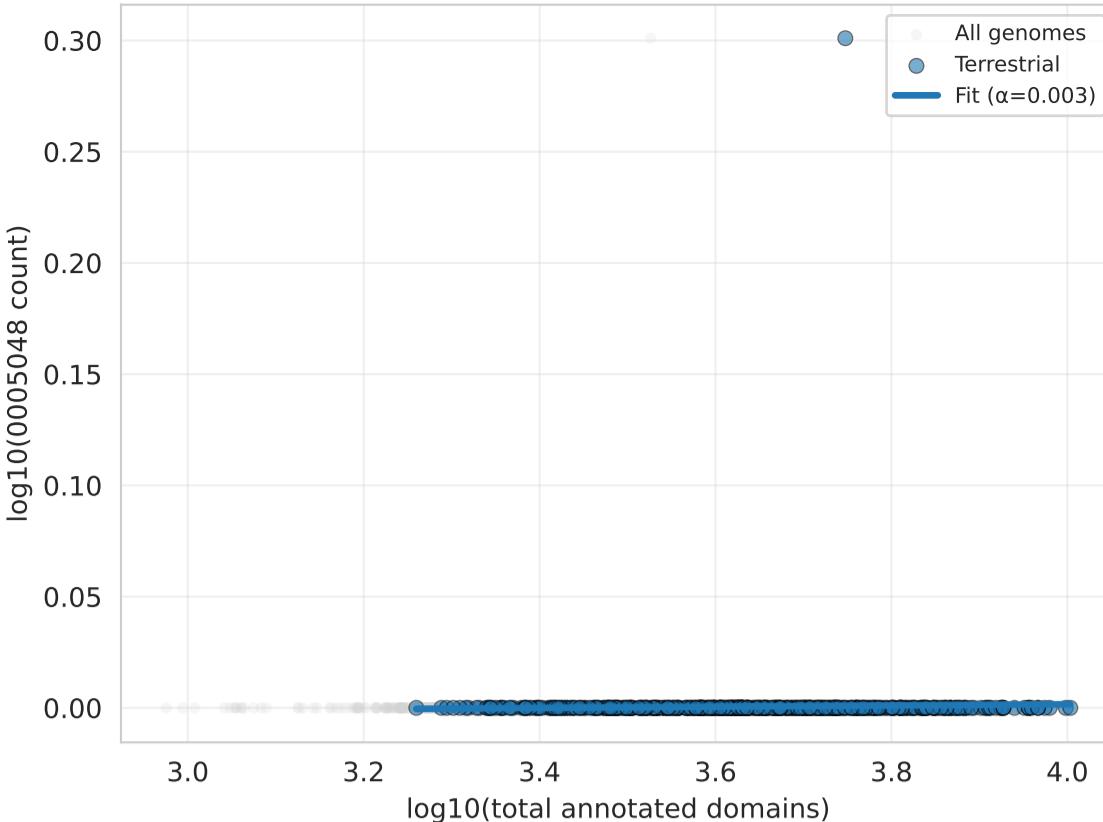
Panel f: Aquatic signal sequence binding (GO:0005048) (vs total domains)



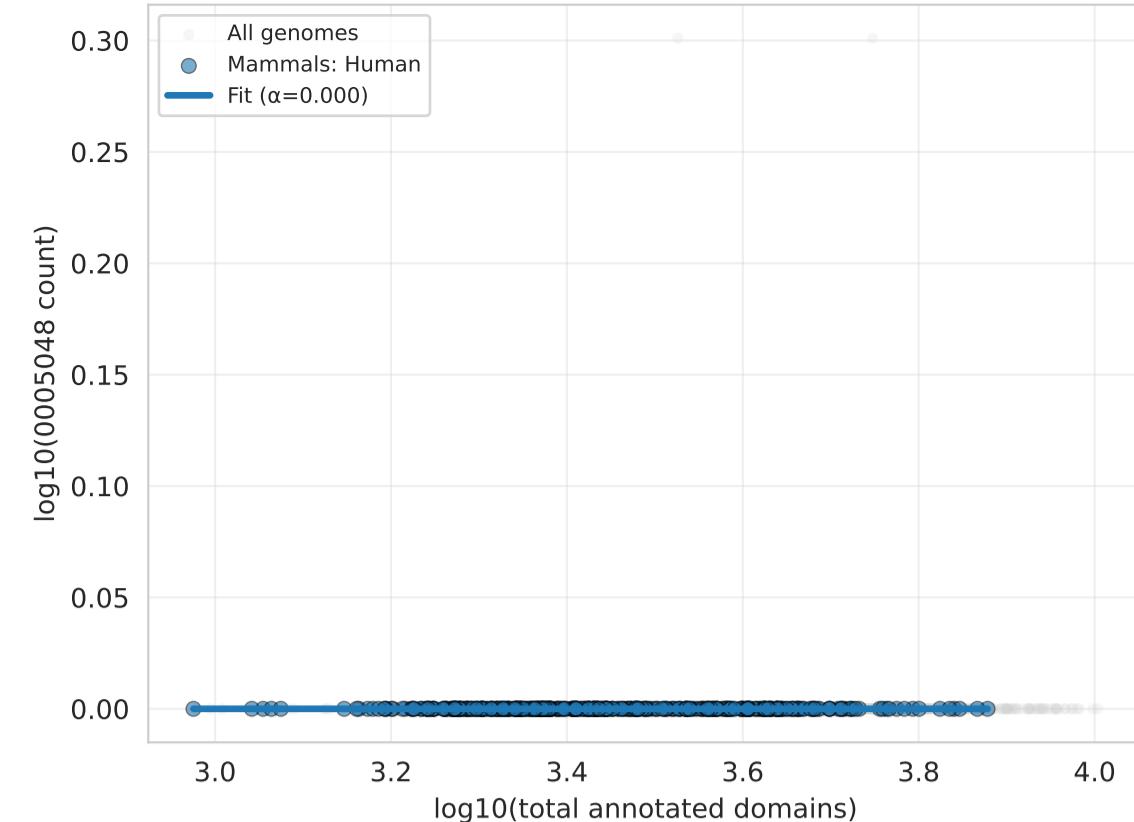
Panel i: Aquatic glutamine metabolic process (GO:0006541) (vs total domains)



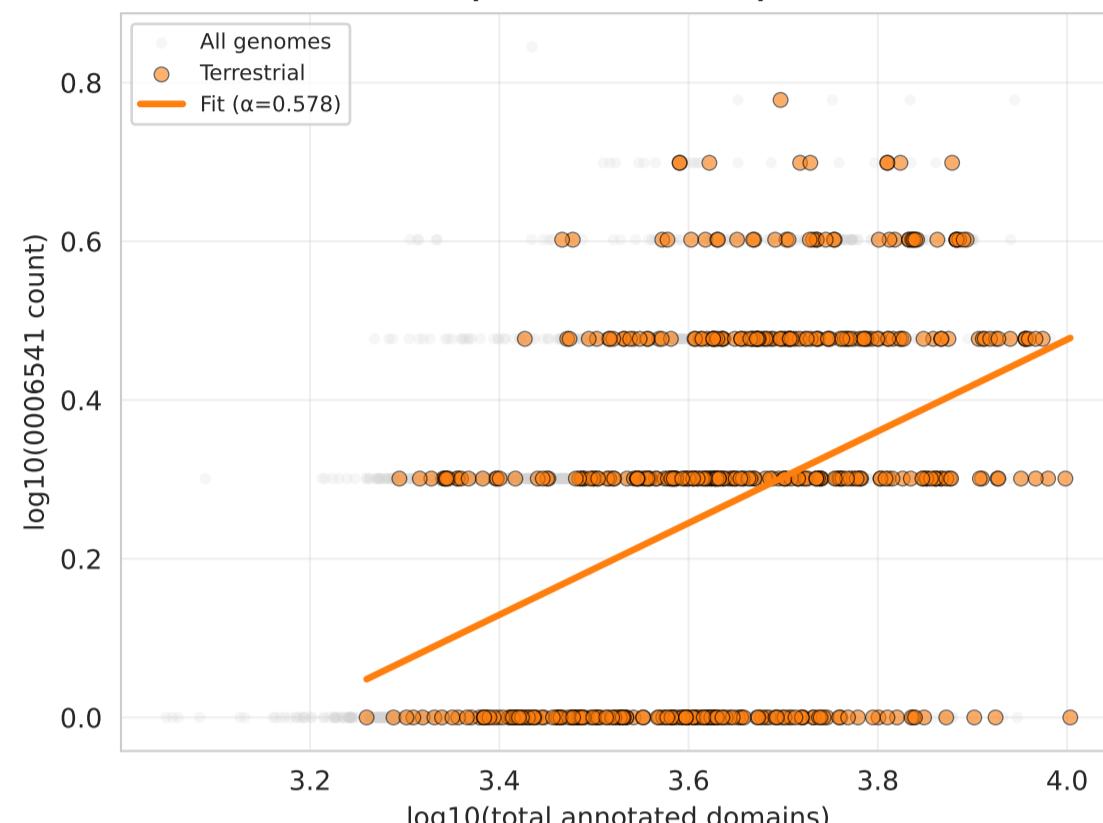
Panel g: Terrestrial signal sequence binding (GO:0005048) (vs total domains)



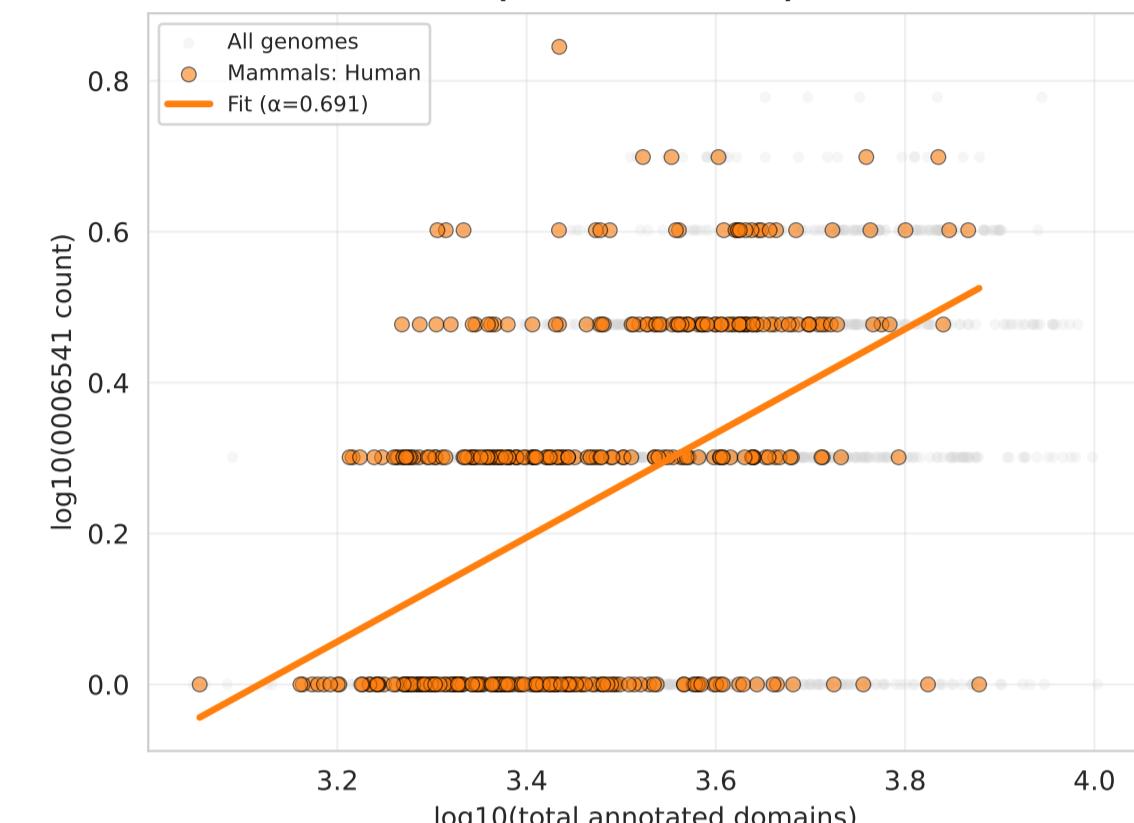
Panel h: Mammals: Human signal sequence binding (GO:0005048) (vs total domains)



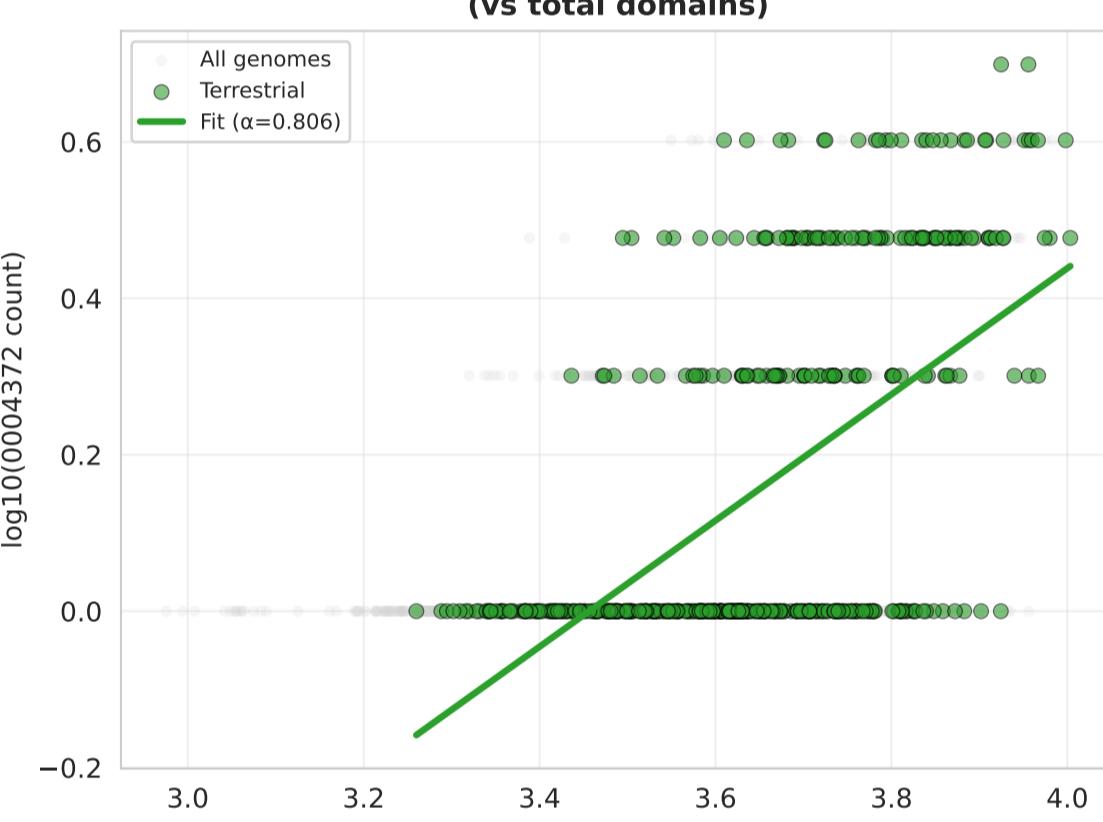
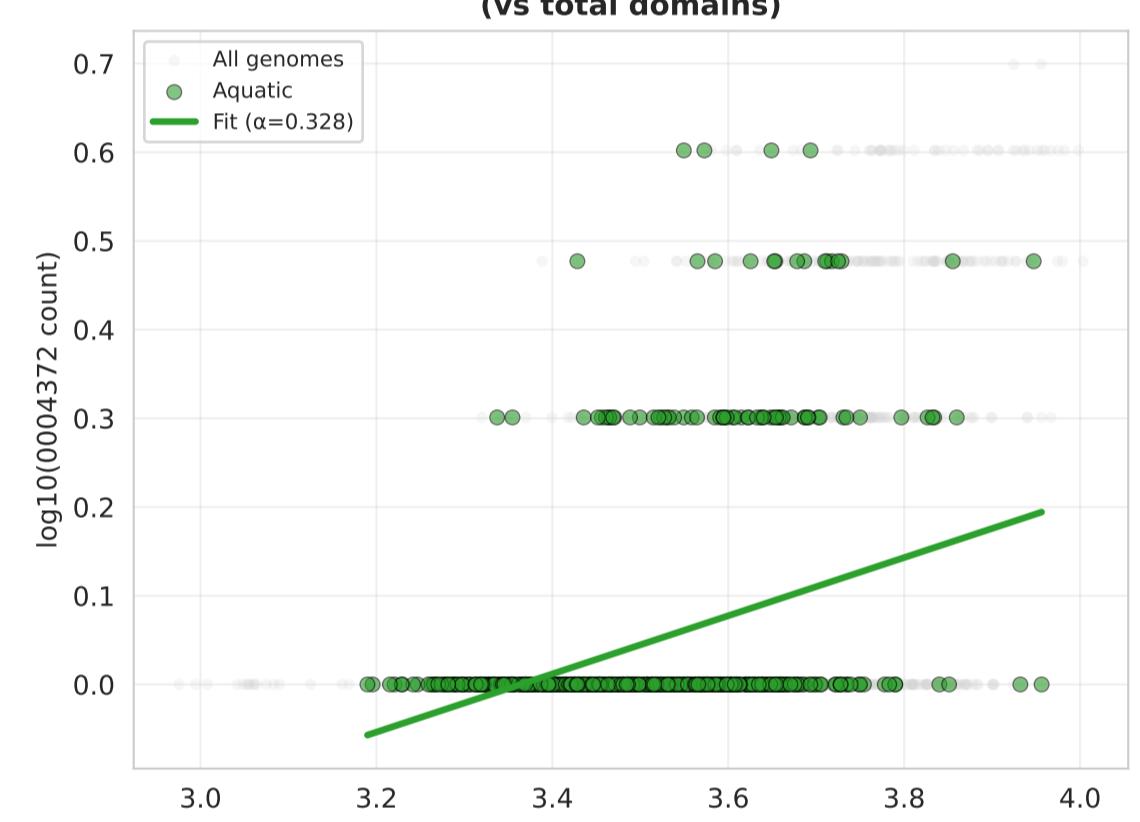
Panel j: Terrestrial glutamine metabolic process (GO:0006541) (vs total domains)



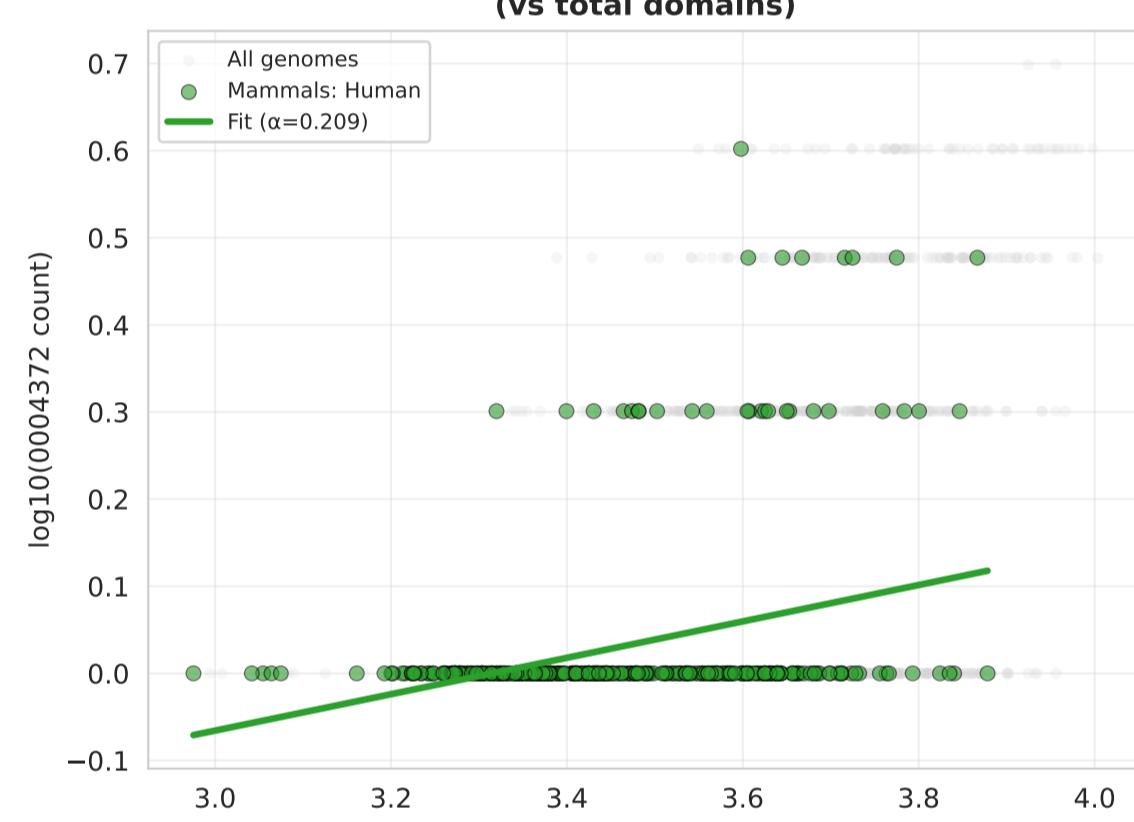
Panel k: Mammals: Human glutamine metabolic process (GO:0006541) (vs total domains)



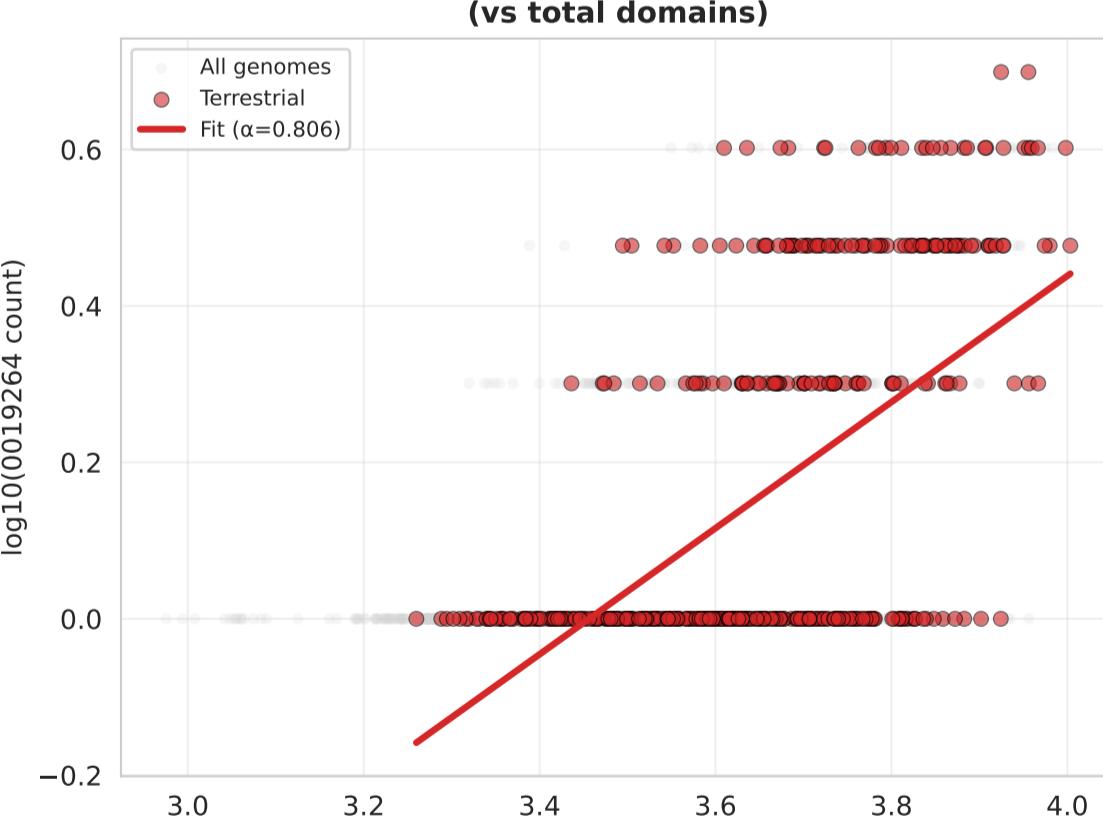
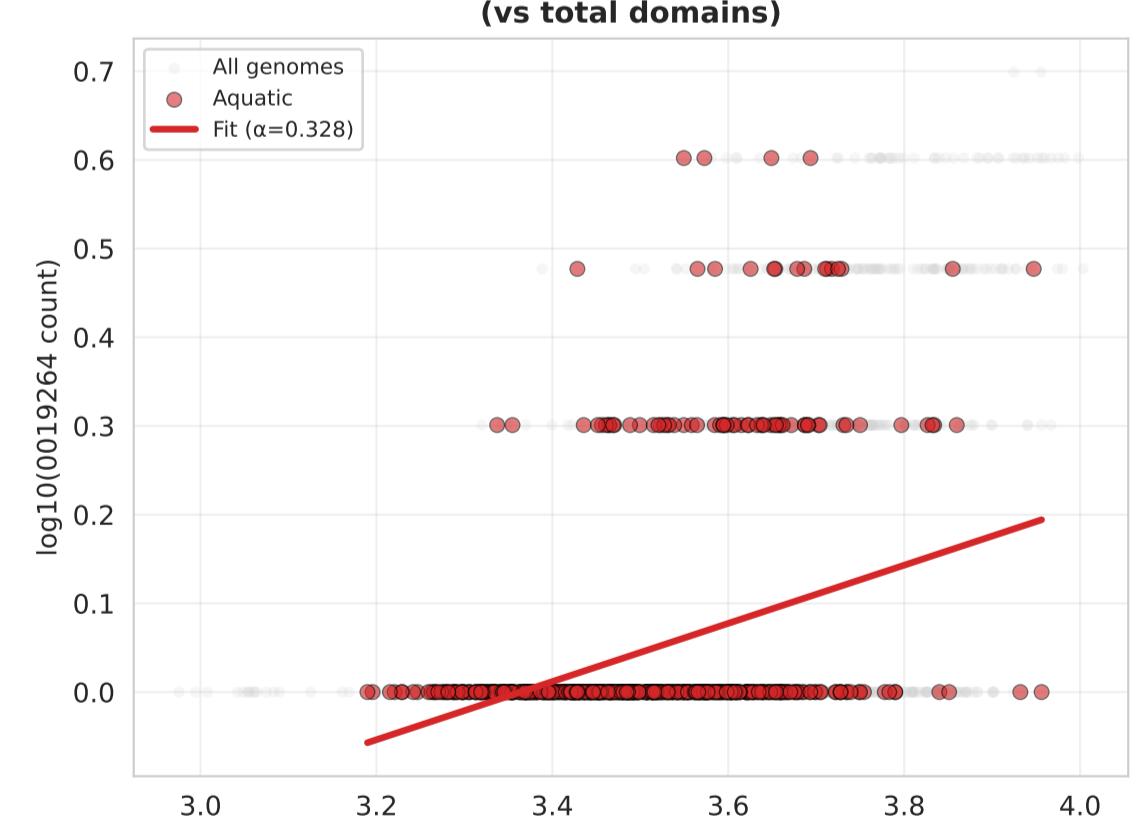
Panel l: Aquatic glycine hydroxymethyltransferase activity (GO:0004372) (vs total domains)



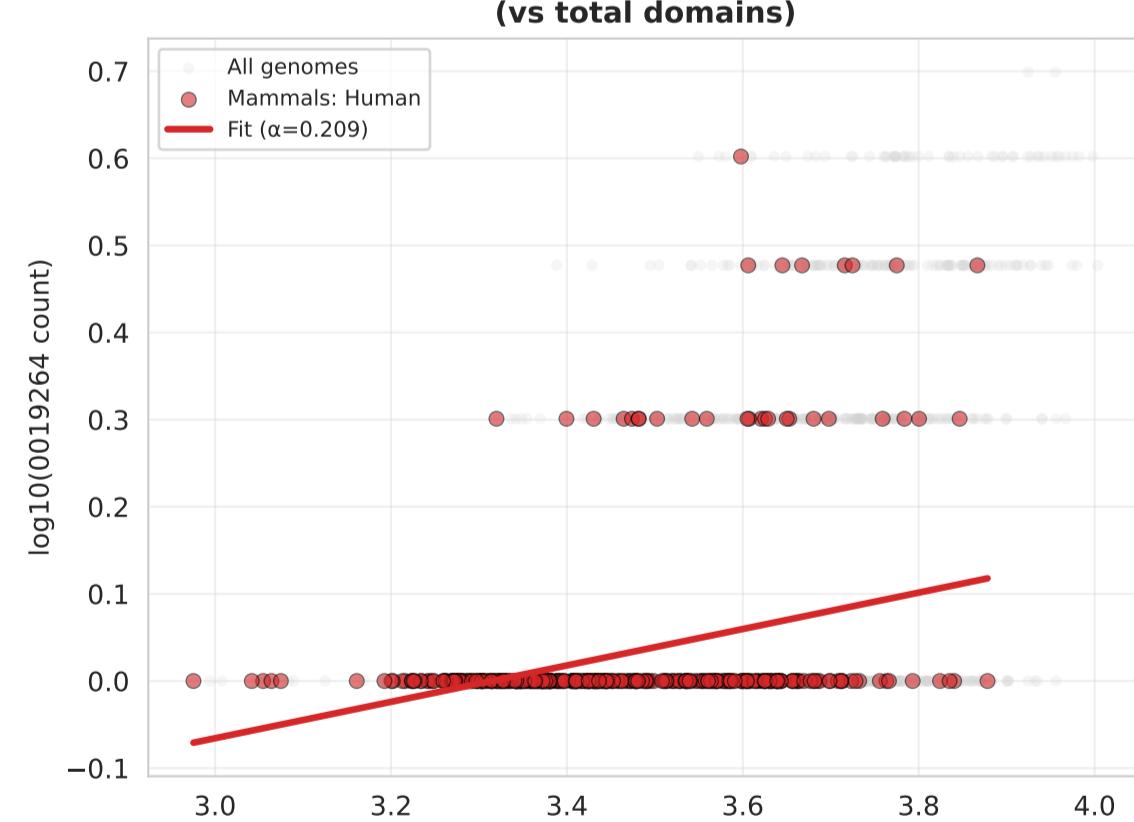
Panel n: Mammals: Human glycine hydroxymethyltransferase activity (GO:0004372) (vs total domains)



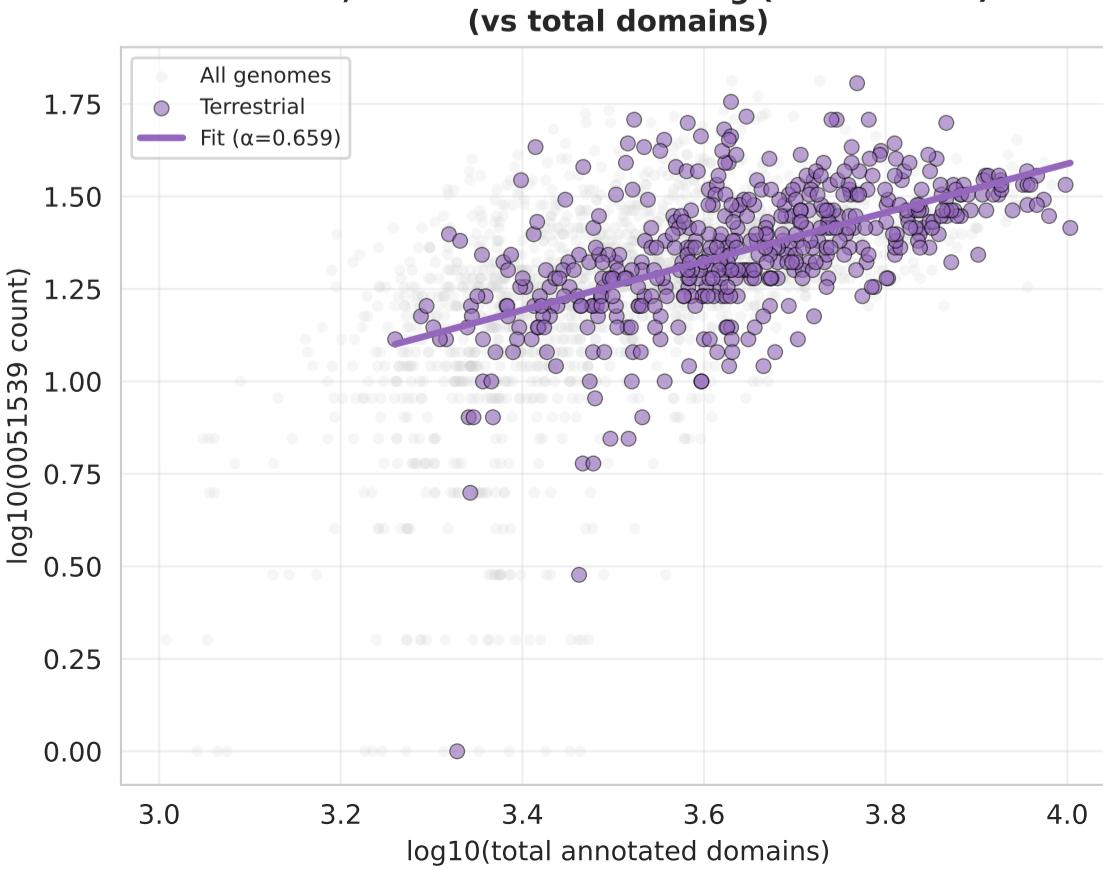
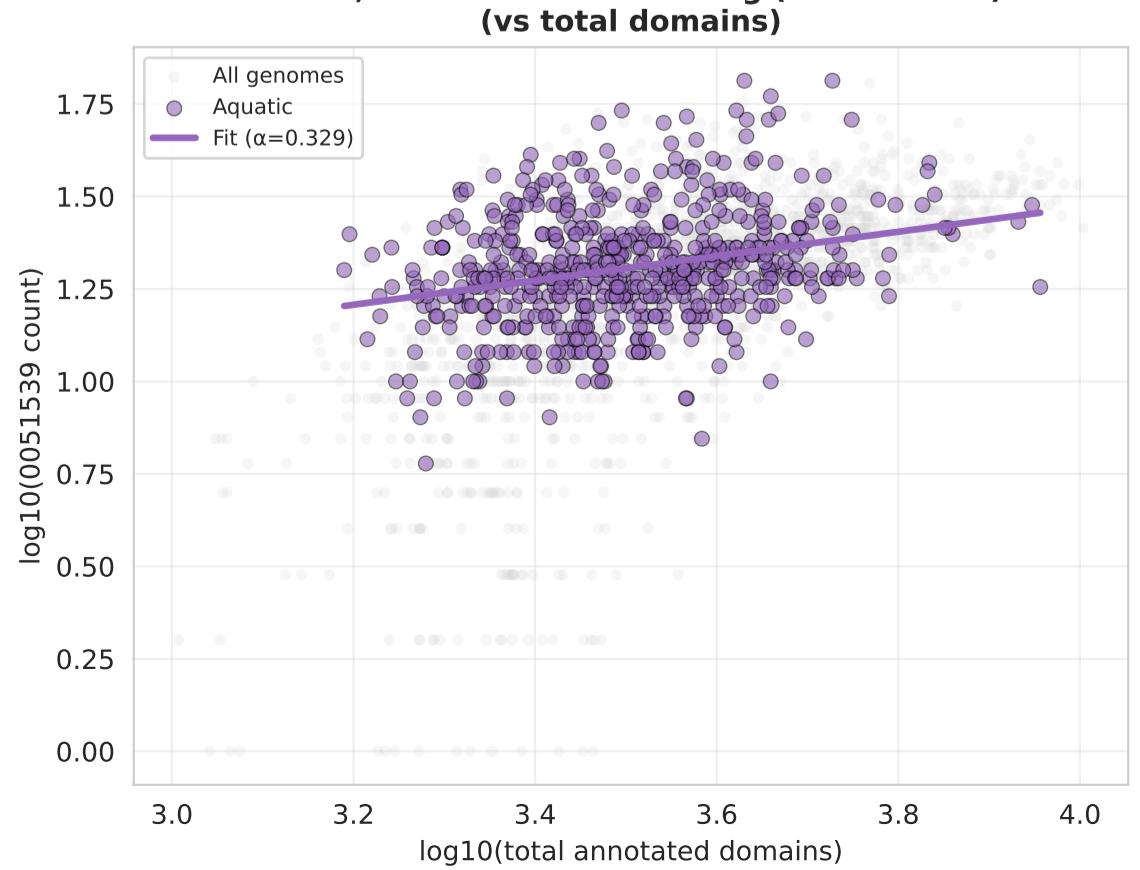
Panel o: Aquatic glycine biosynthetic process from serine (GO:0019264) (vs total domains)



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



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



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

