Supplementary Data 10. Gene Ontology (GO) enrichment of genes in HOGs lost in Clitellata. Up, treemap visualisation of all terms (CC, Cellular Component; BP, Biological Process; MF, Molecular Function). Down, bar plot of GOs enriched in functions related to cell cycle, chromatin remodelling, DNA repair and genome stability.

TreeMap_Enrichm_all_seqs_in_LOST_HOGs_Select_backg_CC

glycerol-3-phosphate dehydrogenase complex		extracellular region	nuclear RNA-directed RNA polymerase complex	Fanconi SUMO-targeted ubiquitin ligase complex		ATF4-CREB1 transcription factor complex	
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TreeMap_Enrichm_all_seqs_in_LOST_HOGs_Select_backg_MF

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