Genes lost in Crassiclitellata MF TreeMap

thioes	tumor necrosis factor receptor binding			RNA–directed DNA polymerase activity	ubiquitin protein ligase	ubiquitin protein ligase activity		tric–oxide nase activity	DNA-binding transcription repressor activity, RNA polymerase II-specific transcrip cytoskeleton-nuclea membrane anchor	transcri initiat factor ac tion regula	eral ption ion ctivity utor acti	transcription regulator activity	
					ligase activity pe endopeptidase activity	3–galactosyl–N–ac 4–alpha–L–fucc activ	osyltransferase	ubiquitin–protein transferase activity	activity DNA binding		ntromeri DNA	cis–regulatory region	
interleukin–1 i	dent protein bi	kine activ nding m-deper		endonuclease activity	RNA-directed 5'-3' RNA polymerase activity	RNA-DNA hybrid ribonuclease activity ubiquitin-like		non–membrane spanning protein tyrosine kinase activity	RNA polymeras transcription regulatory regions sequence-spect DNA binding	ONA bindii e II on		sequence-specific DNA binding NA binding	
		ein bindi			dolichyl-diphosphooligosaccharide-protein glycotransferase activity	prot transfe acti	tein erase	ligase activity	rib	nucleoside-diphosphat			
interleukin–1 binding	peptide hormone receptor binding	microtubule plus-end bindii		l adhesion iator activity		copper ion binding		ohatidylinosi oinding	tol galactoside binding	serine/threc <mark>ribonucl</mark> kinase inhi reducta activity cyc	voltage-gat		voltage-gated potassium channel activity
cell-cell adhesion mediator activity	calmodulin binding			ubiquitin protein	zinc ion binding	galactoside binding	₽			nattern recognition			transmembrane
		hormone red binding	•	ligase binding		intraciliary transport particle A binding	phosphatidylinositol–5-phospl binding		nate phosphalidylinositol-3,4-bisphosphali binding	transmembrand receptor activity	signaling activ	signaling receptor active receptor activity	