Genes gained in Crassiclitellata MF TreeMap

dynein complex binding dynein complex binding			growth factor activity		kinesin binding			filamin binding	microtubule binding	kinase binding	DNA-binding transcription factor activity
				in protein binding				alpha–tubulin binding protein binding	exogenous protein binding	microtubule plus-end binding	DNA-binding transcription factor activity, RNA polymerase II-specific DNA-binding transcription factor activity
				peptide ne activity	histone deacetylase binding	binding		ubiquitin binding	ubiquitin–like protein binding		
									S	SH2 domain binding	
anaphase-promoting complex binding	IgG binding	inding retinoic acid receptor binding	polysaccharide binding	glycogen binding	p53 binding			eukaryotic initiation factor 4E binding	growth factor receptor binding	fatty acid derivative binding	transcription coregulator activity
								transcription	RNA polymera transcriptio		
ubiquitin conjugating enzyme binding	virus receptor activity	nuclear localizatio sequence binding		transcription corepressor binding	mRNA binding			cis-regulatory region binding mRNA binding	regulatory reg sequence–spe DNA bindin	ion binding cific	DNA-binding transcription activator activity, RNA polymerase II-specific
	cytokine activity	protein phosphatase 1 binding		n-specific se binding				anslation repressor vity, mRNA regulatory element binding	mRNA 3'–U [*] binding	fatty-acyl-CoA binding	DNA-binding transcription activator activity