Genes gained in Anomura MF TreeMap

protein binding protein binding		binding		cellular matrix tein binding	protein disulfide isomerase activity		L–lysine transmembrane transporter activity		Y–form DNA binding		
		intermediate filament binding	polyubiquitin modification–dependent protein binding				structural constituent of nuclear pore ctural constituent of n		DNA binding		acetyltransferase activity
							proton–transporting ATP synthase activity,		rRNA binding		DNA-binding transcription repressor activity, RNA
		ubiquitin–like protein binding	p	rotein kinase binding	RNA polymerase II		rotational mechanism				polymerase II–specific RNA polymerase II
		ubiquitin binding			transcription regulatory region sequence–specific DNA binding		proton channel activity		DNA-binding transcription repressor activity		cis–regulatory region sequence–specific DNA binding
WW domain binding	protein homodimerization activity		kinase binding	melanin–concentrating hormone receptor binding	protein phosphatase	nhosnh	atase	prote serine/thre		protein	Innacanataca
		HECT domain binding	type 1 melanin–concentrating hormone receptor binding		regulator i	proteil regulator	n phosp activity	phatase regulator a inhibitor activity		tivity activator activity	