Genes gained in Anomura MF TreeMap

protein binding protein binding			protein disulfide isomerase activity		L-lysine transmembrane transporter activity		Y–form DNA binding		WW domain binding		protein homodimerization activity		
			DNA-binding transcription activator activity, RNA polymerase II-specific	thioredoxin–disulfide reductase activity		regulatory region		structural constituent of					
			polymerase II-specific					nuclear pore			omain binding HECT domain		
			acetyltransferase activity	tructural constituent protein serine/threonine phosphatase inhibitor		prote phosph activa	ein atase ator	proton-transporting ATP synthase activity, rotational	intermediate filament binding	te	binding	binding	
				ad	ectivity	activity		mechanism	protois Linear		titin Z domain	melanin-concentrating	
			protein phosphatase regulator activity	proton channel activity		phosphatase activator activity		telomeric DNA binding	protein kinase binding		binding	hormone activity	
binding	extracellular matrix protein binding	ubiquitin–like protein binding										melanin–concentrating hormone receptor	
			phosphatase regulator activity	rRNA bind	oinding polyi	NA-binding transcription † epressor activity, RNA polymerase II-specific		or sequence-specific	modification-depend protein binding		hormo b melanin- hormo	concentrating ne receptor inding -concentrating ne receptor inding	