Genes lost in ECRY2 MF TreeMap

cell-cell adhesion mediator activity		myosin binding		vinculin binding				cyclic–GMP–AMP synthase activity		DNA-directed DNA polymerase activity		G protein-coupled receptor activity	
dystroglycan binding	transformi growth fac beta bindi	factor Interleukin-		nitric–oxide 1 synthase binding	RNA-directed DNA polycyclin-dependent protein kir			nase activity brane receptor protein tyrosine kinase	serine/three	otein nine/tyrosine activity	cyclin–dependent protein kinase activity	G protein–coupled peptide receptor activity	
lamin binding	protein kin	ase	death recept	transforming				activity MAP kinase kinase kinase kinase activity	serin kina e calmodu	-depender protein e/threoning se activity lin-depend	myosin light chain kinase	interleukin–1 receptor activity signaling rec	cytokine receptor activity eptor activity
PDZ domain binding	cadherir binding				RNA polymerase II cis–regulatory region sequence–specific DNA	bubble DNA binding	RNA polyme transcript regulatory r sequence-s	ion DNA egion	I DNA binding		cyclic_di_GMP		olfactory receptor activity
integrin binding	thyroid hormone receptor binding	e receptor r binding		CARD domain binding	binding	DN	DNA bind		ıg	zinc ion binding	neuropeptide binding	. ocopie.	death receptor activity
kinase binding	protein kina C binding		owth factor activity	phosphatidylinositol 3-kinase binding scaffold protein binding	metal ion binding	peptidoglycan binding	heparin bir	nding cobalamin			inding	activity tumor necrosis factor–activated receptor activity	neuropeptide Y receptor activity