Genes lost in RVAR1 MF TreeMap

metalloendopeptidase activity		transmembrane receptor protein tyrosine kinase activity			DEAD/H–box RNA helicase binding	protease binding		misfolded protein binding virion binding		carbohydrate derivative binding epidermal growth factor receptor		lipopolysaccharide binding		zinc ion binding	
		peroxidase activity			mitogen–activated protein kinase kinase kinase binding	cytokine transforming	ig or			bino drate		purine ribonucle binding	mRN	NA nu	purine binding binding purine ucleotide
RNA-directed DNA cal	farnesol con-nitrogen lyaso activity		glutathione activity:idase activity	NAD(P)+ nucleosidase activity	cell-cell adhesion mediator activity	beta bindii TIR doma binding		3MP binding •	unfo pro bino acetylo rece bino	tein ding choline eptor	G-protein beta/gamma-subunit complex binding	phosphatidylcholine binding purine ribonucleoside triphosphate binding	3'-U' bindi ribonucle bindi	ng AT eotide ng pep	binding ATP binding peptidoglycan binding
protein serine/threonine/tyrosine kinase activity	NA nucleo cyc ADP-i gener	otidase, clic ribose	endonuclease activity non-membrane	glutathione transferase activity	urotensin II receptor activity	peptide		in–coupled e receptor ctivity		G protein–coupled receptor activity pattern recognition		ribosomal–protein–al N–acetyltransfera activity	se	e N-acetyltransferase activity	
cholinesterase activity	protein fo chaper serine- endopep activi	n folding perone e-type eptidase	spanning protein tyrosine kinase activity transmembrane receptor protein	carbon–nitrogen lyase activity fibroblast growth factor–activated receptor activity	G prote	im-coupled rec				receptor activity		peptide-serine-N-acet peptide-serine-N-acet		tultraneforaco	
				metalloexopeptidase activity		activ		pheromone receptor activity		neuropeptide Y receptor activity		activity			O-acetyltransferase activity