Genes lost in ETES1 MF TreeMap

RNA-directed DNA polymerase activity	aspartic-type endopeptidase activity			DNA-directed DNA polymerase activity		iron ion binding	metal ion l	oinding h	nistamine binding	
			Pι			nitric oxide binding	odorant binding odorant bind	binding		
	metalloendopeptidase cargo receptor activity	RNA polymerase II cis—regulatory region sequence—specific DNA binding oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxygen, reduced flavin or flavoprotein as one donor,				zinc ion binding	co-receptor binding	homodimerizat activity		
			acetylcholi cation-s channel	selective	monooxygenase activity		heme binding	adhesion	glycolipid DEAD/H-box RNA helicase binding binding	
						alpha–tubulin binding	receptor antagonist activity	Wnt–protein binding		
endonuclease activity	RNA–DNA hybrid ribonuclease activity		extracellular ligand-gated ion channel activity	oxidoreductase acting acting on paired don with incorporation reduction of molecu oxygen	or sulfotransferase activity	C protoin coupled	acetylchol receptor ac		peptide	
			3',5'-cyclic-AMP phosphodiesterase activity	scavenger receptor activity protein tyrosine kinase activity	G protein–coupled receptor activity	G gnaling receptor	G protein-coup	receptor		
	serine-type endopeptidase activity		DNA-binding transcription activator	fatty acid synthase activity	spanning protein		receptor activity	signalin	g G protein-coupled	
		serine-type endopeptidase inhibitor activity	Activity DNA-binding transcription activator activity, RNA polymerase II-specific	protein-membrane adaptor activity	regulatory region sequence–specific DNA binding	transmembrane signa receptor activity	ling neurotransmitte receptor activit	activity	activity neuropeptide	