PR002110 Ankyrin repeat

IPR008949 Isoprenoid syMR939 348 Ellingah N-Remogati (to path on seperal-containing doma	number of	gene
IPR000092 Polyprenyl synthetase IPR020683 Donam of unknown function DUF	\bigcirc 10	
IPR010736 Heterokaryon incompatibility IPR007++1 NACHT nucleoside triphosphatas	() 20	
IPR031352 NACHT-NTPase and P-loop NTPase		
PR041698 Methyltransferase domain 25 IPR013026 None	() 30	
IPR019734 Tetratrice peptide repeat		
IPR035992 Ricin B-like lectins IPR009078 Ferritin-like superfamily		
IPR029063 S-adenosyl-L-methionine-depend IPR001173 Glycosyltransferase 2-like	p.adjust	
IPR016461 O-methyltransferase COMT-type		
IPR036259 MFS transporter superfamily	0.003	ł
IPR001077 Ō-methyltransferase domain	0.000	•
IPR013120 Fatty acyl-coenzyme A redult@93629 NAD(P)-binding domain superfam	0.002	,
IPR042099 ANL, N-terminal domain IPR009081 Phosphopantetheine binding ACP	0.002	•
IPP011008 Dimoric alpha, hota pared	0.001	
iproly853 Glycoside nydrolase superfamil	0.001	
IPR000873 AMP-despite the property of the prop		
IPR006162 Phosphopantetheins attachment		