

InterProScan: protein domains identifier

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Citation Report

#	ARTICLE	IF	CITATIONS
1	InterPro. , 2005, , .		9
2	Genome-wide review of transcriptional complexity in mouse protein kinases and phosphatases. <i>Genome Biology</i> , 2006, 7, R5.	9.6	48
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18	The effects of multiple features of alternatively spliced exons on the K A /K S ratio test. <i>BMC Bioinformatics</i> , 2006, 7, 259.	2.6	8

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1533	Genome Sequences of vB_RleM_RL38JI and vB_RleM_RL2RES, Two Virulent <i>Rhizobium leguminosarum</i> Transducing Phages. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.6	1
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1535	GrAPFI: predicting enzymatic function of proteins from domain similarity graphs. <i>BMC Bioinformatics</i> , 2020, 21, 168.	2.6	8
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1653	Genome-Wide Patterns of Bracovirus Chromosomal Integration into Multiple Host Tissues during Parasitism. <i>Journal of Virology</i> , 2021, 95, e0068421.	3.4	6
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1678	Genome-wide identification studies â€˜ A primer to explore new genes in plant species. <i>Plant Biology</i> , 2022, 24, 9-22.	3.8	5
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1940	The draft chromosome-level genome assembly of tetraploid ground cherry (<i>Prunus fruticosa</i> Pall.) from long reads. Genomics, 2021, 113, 4173-4183.	2.9	14
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1955	Identification of Phosphorus Stress Related Proteins in the Seedlings of Dongxiang Wild Rice (<i>Oryza</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	2.4	7
1956	Genomic insight into the scale specialization of the biological control agent <i>Novius pumilus</i> (Weise,) Tj ETQq1 1 0.784314 rgBT /Overbo	2.8	6

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1961	Genome-Wide Analysis Indicates a Complete Prostaglandin Pathway from Synthesis to Inactivation in Pacific White Shrimp, <i>Litopenaeus vannamei</i> . <i>International Journal of Molecular Sciences</i> , 2022, 23, 1654.	4.1	0
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1986	Selection drives convergent gene expression changes during transitions to co-sexuality in haploid sexual systems. <i>Nature Ecology and Evolution</i> , 2022, 6, 579-589.	7.8	12
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1992	NBS-LRR-WRKY genes and protease inhibitors (PIs) seem essential for cowpea resistance to root-knot nematode. <i>Journal of Proteomics</i> , 2022, 261, 104575.	2.4	4
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1996	Genomic characterization of ZIP genes in pigeonpea (CcZIP) and their expression analysis among the genotypes with contrasting host response to pod borer. <i>Physiology and Molecular Biology of Plants</i> , 2021, 27, 2787-2804.	3.1	0
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2231	Comparative bioinformatics analysis and abiotic stress responses of expansin proteins in Cucurbitaceae members: watermelon and melon. <i>Protoplasma</i> , 2023, 260, 509-527.	2.1	7
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