

# Dynamics of Cullin-RING Ubiquitin Ligase Network Revealed by Proteomics

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Profiling of Protein Interaction Networks of Protein Complexes Using Affinity Purification and Quantitative Mass Spectrometry. <i>Molecular and Cellular Proteomics</i> , 2010, 9, 1650-1665.	2.5	89
2	The cullin protein family. <i>Genome Biology</i> , 2011, 12, 220.	13.9	391
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5	The Molecular Basis of CRL4DDB2/CSA Ubiquitin Ligase Architecture, Targeting, and Activation. <i>Cell</i> , 2011, 147, 1024-1039.	13.5	372
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10	Ubiquitination of E3 ligases: self-regulation of the ubiquitin system via proteolytic and non-proteolytic mechanisms. <i>Cell Death and Differentiation</i> , 2011, 18, 1393-1402.	5.0	239
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15	SCF<sup>FBXO22</sup> Regulates Histone H3 Lysine 9 and 36 Methylation Levels by Targeting Histone Demethylase KDM4A for Ubiquitin-Mediated Proteasomal Degradation. <i>Molecular and Cellular Biology</i> , 2011, 31, 3687-3699.	1.1	81
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18	Simply quantifying ubiquitin complexity. <i>Nature Methods</i> , 2011, 8, 630-631.	9.0	0

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