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41 40 39	Membrane insertion topology of the central apolipoprotein A-I region. Fluorescence studies using single tryptophan mutants. <i>Biochemistry</i> , 2011 , 50, 466-79 Structure-function relationship in an archaebacterial methionine sulphoxide reductase B. <i>Molecular Microbiology</i> , 2011 , 79, 342-58 Hydrogen exchange during cell-free incorporation of deuterated amino acids and an approach to its inhibition. <i>Journal of Biomolecular NMR</i> , 2011 , 51, 467-76 Zinc(II) complexes of ubiquitin: speciation, affinity and binding features. <i>Chemistry - A European</i>	3.2 4.1 3	9 5 24
41 40 39 38	Membrane insertion topology of the central apolipoprotein A-I region. Fluorescence studies using single tryptophan mutants. <i>Biochemistry</i> , 2011 , 50, 466-79 Structure-function relationship in an archaebacterial methionine sulphoxide reductase B. <i>Molecular Microbiology</i> , 2011 , 79, 342-58 Hydrogen exchange during cell-free incorporation of deuterated amino acids and an approach to its inhibition. <i>Journal of Biomolecular NMR</i> , 2011 , 51, 467-76 Zinc(II) complexes of ubiquitin: speciation, affinity and binding features. <i>Chemistry - A European Journal</i> , 2011 , 17, 11596-603 Interpreting protein chemical shift data. <i>Progress in Nuclear Magnetic Resonance Spectroscopy</i> , 2011	3.2 4.1 3 4.8	9 5 24 29
41 40 39 38 37	Membrane insertion topology of the central apolipoprotein A-I region. Fluorescence studies using single tryptophan mutants. <i>Biochemistry</i> , 2011 , 50, 466-79 Structure-function relationship in an archaebacterial methionine sulphoxide reductase B. <i>Molecular Microbiology</i> , 2011 , 79, 342-58 Hydrogen exchange during cell-free incorporation of deuterated amino acids and an approach to its inhibition. <i>Journal of Biomolecular NMR</i> , 2011 , 51, 467-76 Zinc(II) complexes of ubiquitin: speciation, affinity and binding features. <i>Chemistry - A European Journal</i> , 2011 , 17, 11596-603 Interpreting protein chemical shift data. <i>Progress in Nuclear Magnetic Resonance Spectroscopy</i> , 2011 , 58, 62-87 Structural and functional comparison of the RING domains of two p53 E3 ligases, Mdm2 and Pirh2.	3.2 4.1 3 4.8	9 5 24 29 171

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