

# Tunable alignment of macromolecules by filamentous p interactions

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Citation Report

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2	Biomolecular NMR: recent advances in liquids, solids and screening. <i>Current Opinion in Chemical Biology</i> , 1999, 3, 530-536.	2.8	23
3	NMR structure and mutagenesis of the inhibitor-of-apoptosis protein XIAP. <i>Nature</i> , 1999, 401, 818-822.	13.7	332
4	Induced alignment and measurement of dipolar couplings of an SH2 domain through direct binding with filamentous phage. <i>Journal of Biomolecular NMR</i> , 1999, 14, 175-179.	1.6	18
5	Title is missing!. <i>Journal of Biomolecular NMR</i> , 1999, 14, 333-343.	1.6	122
6	Recognition of protein folds via dipolar couplings. <i>Journal of Biomolecular NMR</i> , 1999, 14, 223-230.	1.6	83
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9	Establishing a degree of order: obtaining high-resolution NMR structures from molecular alignment. <i>Structure</i> , 1999, 7, R205-R211.	1.6	53
10	Order Matrix Analysis of Residual Dipolar Couplings Using Singular Value Decomposition. <i>Journal of Magnetic Resonance</i> , 1999, 138, 334-342.	1.2	544
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15	Solution structure of human GAIP ( $\hat{G}$ interacting protein): a regulator of G protein signaling. <i>Journal of Molecular Biology</i> , 1999, 291, 927-939.	2.0	65
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17	Residual dipolar coupling derived orientational constraints on ligand geometry in a 53 kDa protein-ligand complex. <i>Journal of Molecular Biology</i> , 1999, 293, 107-115.	2.0	89
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