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Huang, Y. J.; Tejero, R.; Powers, R.; Montelione, G.T. A topology-constrained distance network algorithm for protein structure determination from NOESY data. *PROTEINS: Struct. Funct. Bioinformatics* 15, 587-603 (2006)

Huang, Y. J.; Powers, R. & Montelione, G. T. Protein NMR recall, precision, and F-measure scores (RPF scores): structure quality assessment measures based on information retrieval statistics. *J Am Chem Soc* 127, 1665-74 (2005)

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