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Please use this identifier to cite or link to this item: http://hdl.handle.net/123456789/2968 Title: Numerically optimized band-selective pulses in SOFAST-HMQC experiments for biomolecular Authors: Dogra, S. (/jspui/browse?type=author&value=Dogra%2C+S.) Dorai, K. (/jspui/browse?type=author&value=Dorai%2C+K.) Numerically optimized pulses Keywords: GRAPE algorithm Band-selective excitation SOFAST-HMQC Biomolecular NMR Issue 2014 Date: Publisher: Elsevier Citation: Journal of Molecular Structure, 1063(1), pp.45-50. Abstract: This work demonstrates the efficacy of numerically optimized band-selective pulses in 2D fastpulsing NMR pulse sequences of the SOFAST-HMQC variety. In order to achieve robust band selectivity the amplitude and phase of the shaped RF pulses are modulated according to a numerically optimized function. During the pulse duration, the spin trajectories evolve along complex and often unexpected pathways. The pulses have been designed using the GRAPE algorithm and are experimentally implemented on a model protein ubiquitin (13C, 15N labeled). Signal to noise ratios of peaks have been computed and compared for the different experiments performed using both numerically optimized band-selective pulses and standard pulse shapes. The numerically optimized pulses perform better in terms of signal enhancement and phase, as compared to standard pulse shapes.

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