

Electric Supplementary Information for
**Doubly spin-labeled nanodiscs to improve structural determination of
membrane proteins by ESR**

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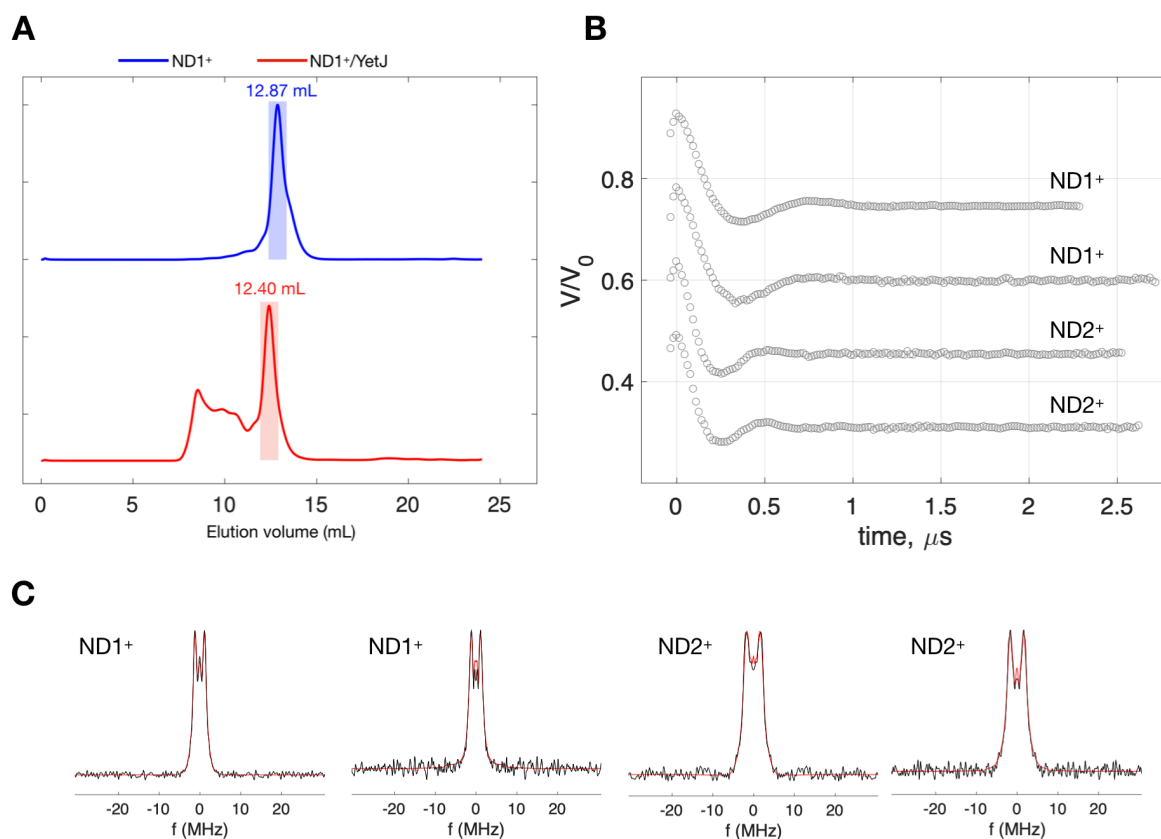


Figure S1. Sample preparation and reproducibility. (A) Representative purification results for empty ND1⁺ and ND⁺/YetJ samples by size exclusion chromatography (SEC-FPLC). The homogeneity, as highlighted by shaded regions, is better for ND⁺/YetJ than ND1⁺ owing to the passage through nickel column after nanodisc formation. The shaded areas also indicate the fractions collected for DEER measurements. (B) Representative background-corrected DEER traces of the samples ND1⁺ and ND2⁺ at pH 7. It shows that these ND⁺ samples can be reliably reproduced with the protocol reported in this study such that the experimental DEER time-domain traces are reproducible in replicate experiments. Same average distances are obtained by TIKR for the replicate measurements of ND1⁺ or ND2⁺. Pake doublet plots of these replicate measurements are shown in (C), confirming that the same dipolar frequency in the Pake doublets for ND1⁺ (or ND2⁺) is obtained in the replicate measurements. The Pake doublets were generated using DeerAnalysis.

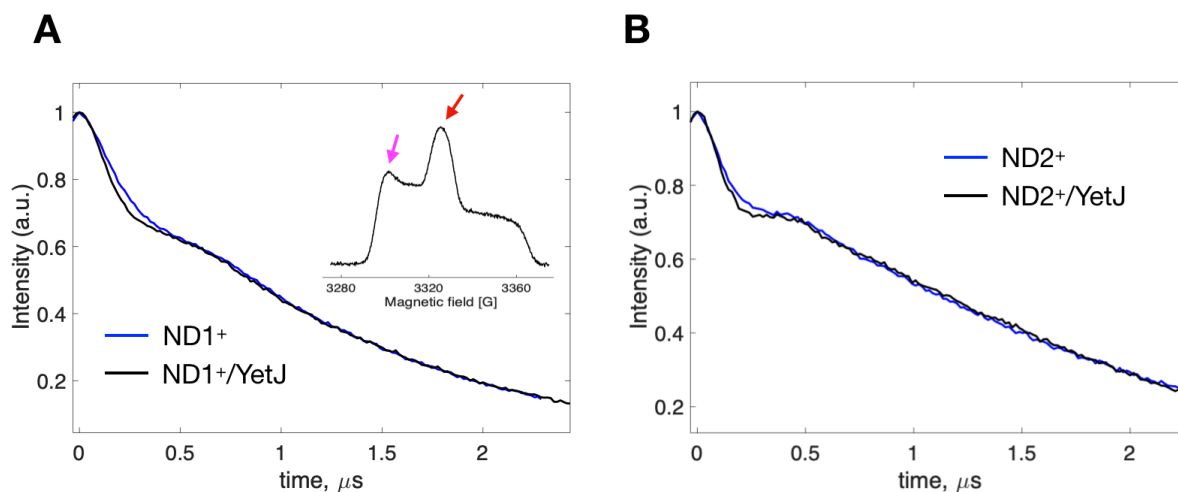


Figure S2. Sensitivity of spin-labeled ND to the change in ND geometry. (A) A comparison of the DEER experimental data of ND1⁺ with/without the incorporated YetJ. For the sake of a clear comparison, the raw data are plotted with the modulation-depth scaling using the DeerAnalysis program. The subtle differences in the DEER traces between the ND1⁺ and the ND1⁺/YetJ are clearly revealed, demonstrating the high sensitivity of ND1⁺ to a change in the ND geometry. Specifically, the time-domain signal for ND1⁺/YetJ decays distinctly faster than that for ND1⁺, supporting a shorter interspin distance in the former. Inset: Hahn echo-detected field swept spectrum of ND1⁺ and the positions of the pump and observe pulses (red and magenta arrows, respectively). The comparison of the DEER experimental data of ND2⁺ with/without the incorporated YetJ is provided in (B), which leads to the same conclusion about the high sensitivity of doubly spin-labeled ND.

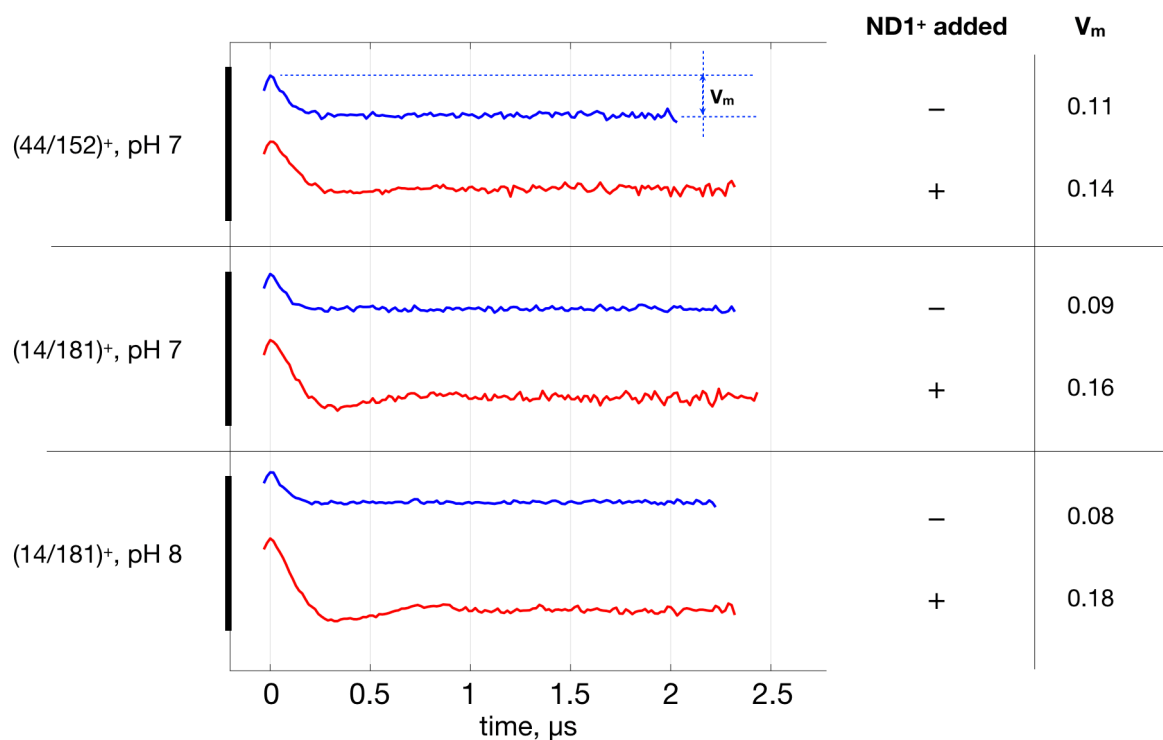


Figure S3. DEER time-domain traces after the removal of background signals. Data are displayed without normalization to the same magnitude so as to emphasize that when ND1⁺ are present in the YetJ⁺ (e.g., (44/152)⁺ or (14/181)⁺) solutions, the overall dipolar modulation depth (V_m , as illustrated) is more clearly revealed in the background-corrected DEER traces. All samples shown have the same spin concentration, as described in Experimental. The presence of ND⁺ samples, whose spin-labeling locations are more solvent-exposed, is useful to enhance the overall modulation depth of DEER signals and thus effectively improve the SNR of DEER data. Note that as the SNR of DEER data is defined by a ratio of the peak intensity to the root-mean-square of noise, it is directly proportional to the peak intensity (i.e., V_m) of the background-corrected data when all of the experimental data have a similar noise level. As shown, we observed an improvement of SNR in the range of 27% and 125%. With the SNR improvement, it directly leads to an enhancement in the distance resolution obtainable by TIKR, hence a distinct reduction in the distance distribution width as compared to the pure YetJ⁺ results (Figs. 3 and 4).