

## **SUPPLEMENTARY MATERIAL**

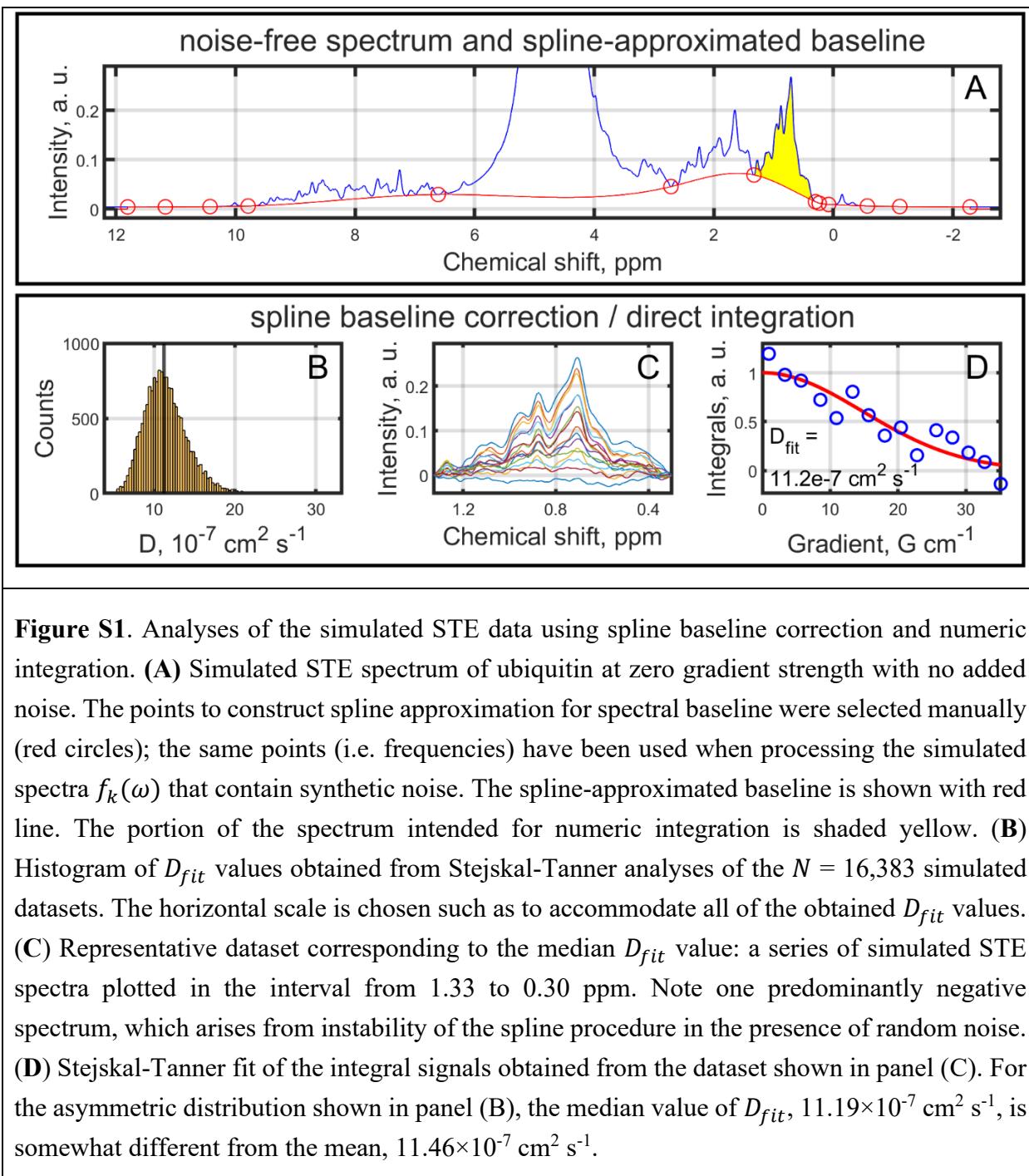
### **Web server DDfit: a new scheme to process PFG NMR diffusion data with improved precision**

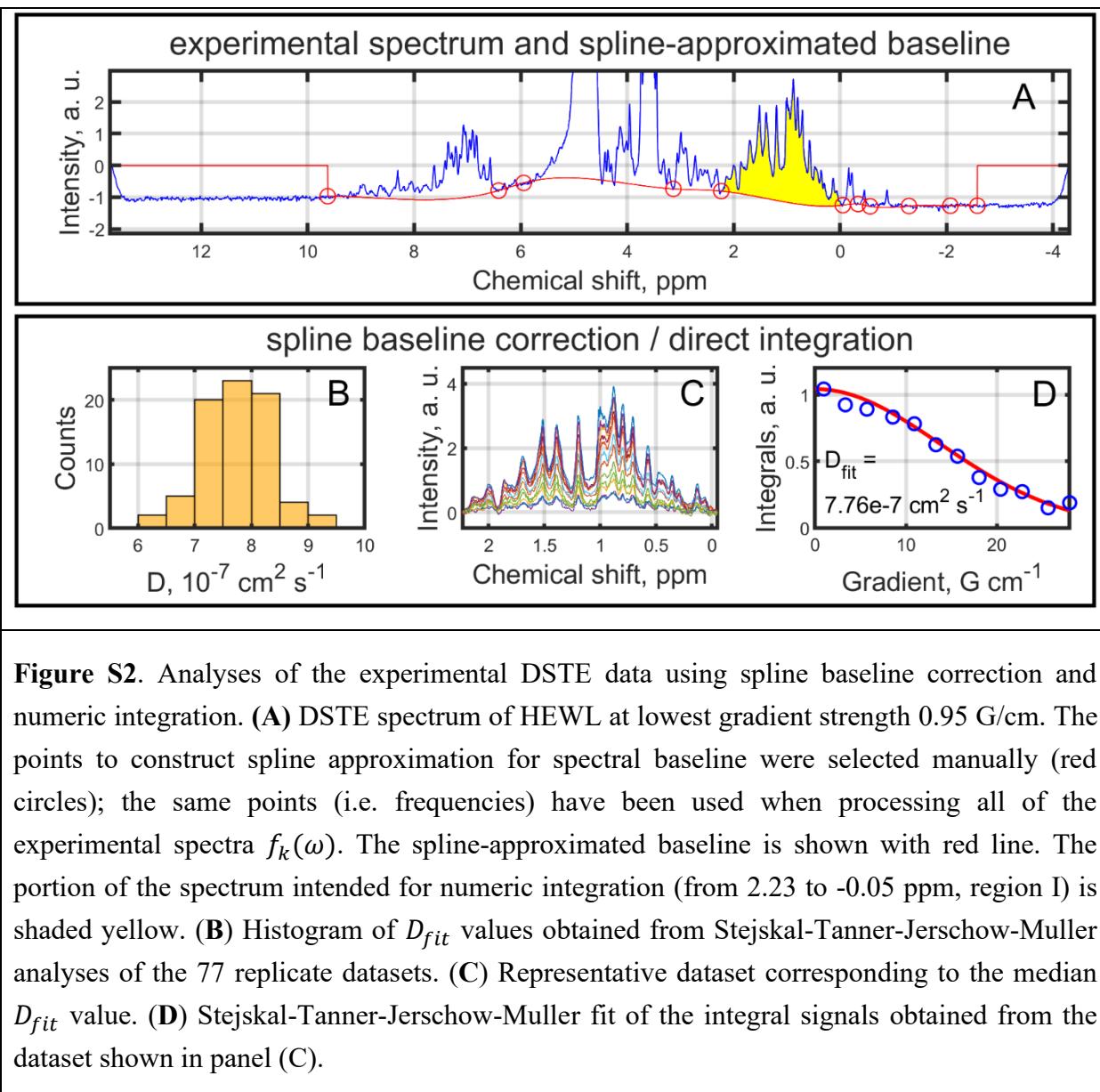
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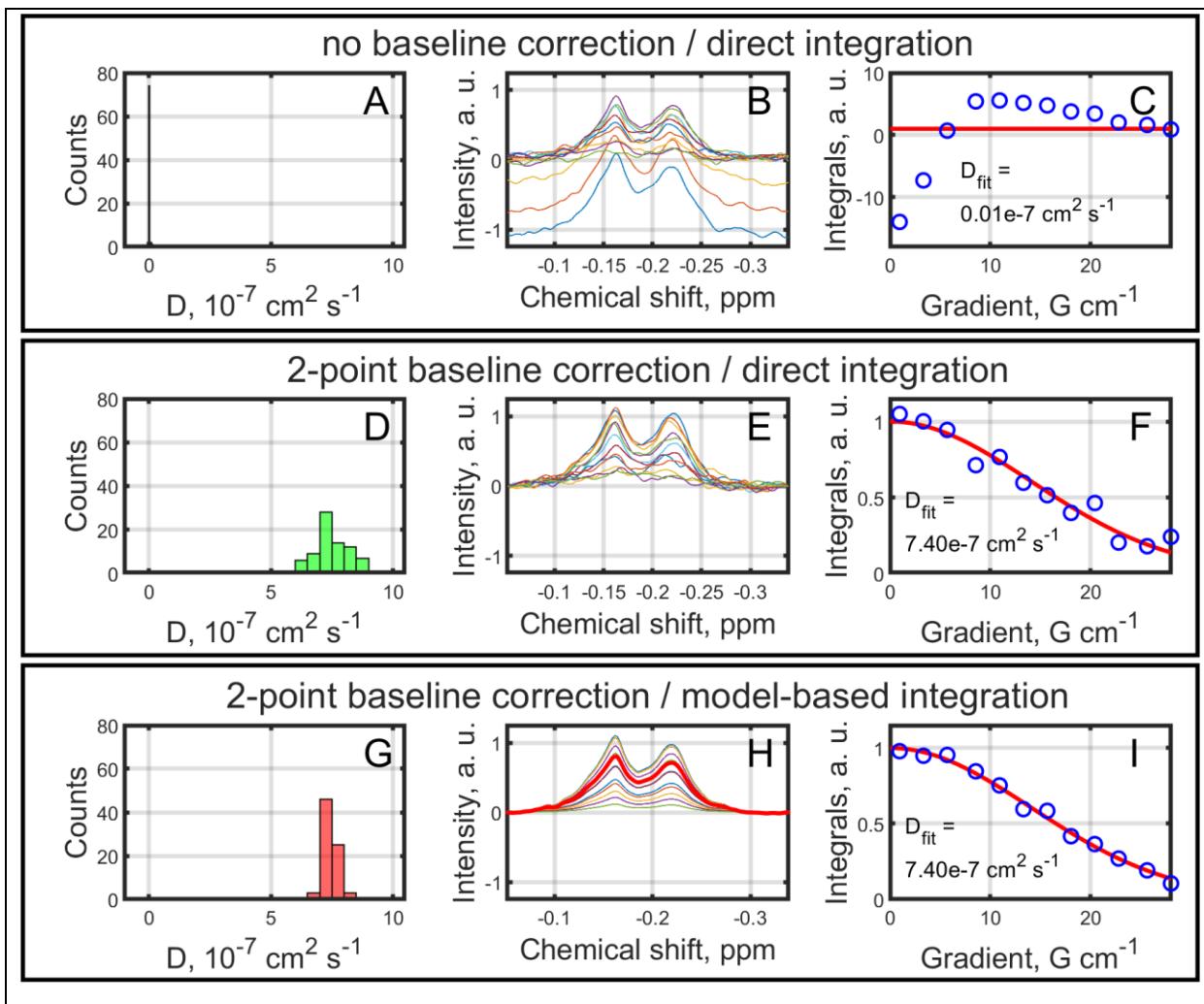
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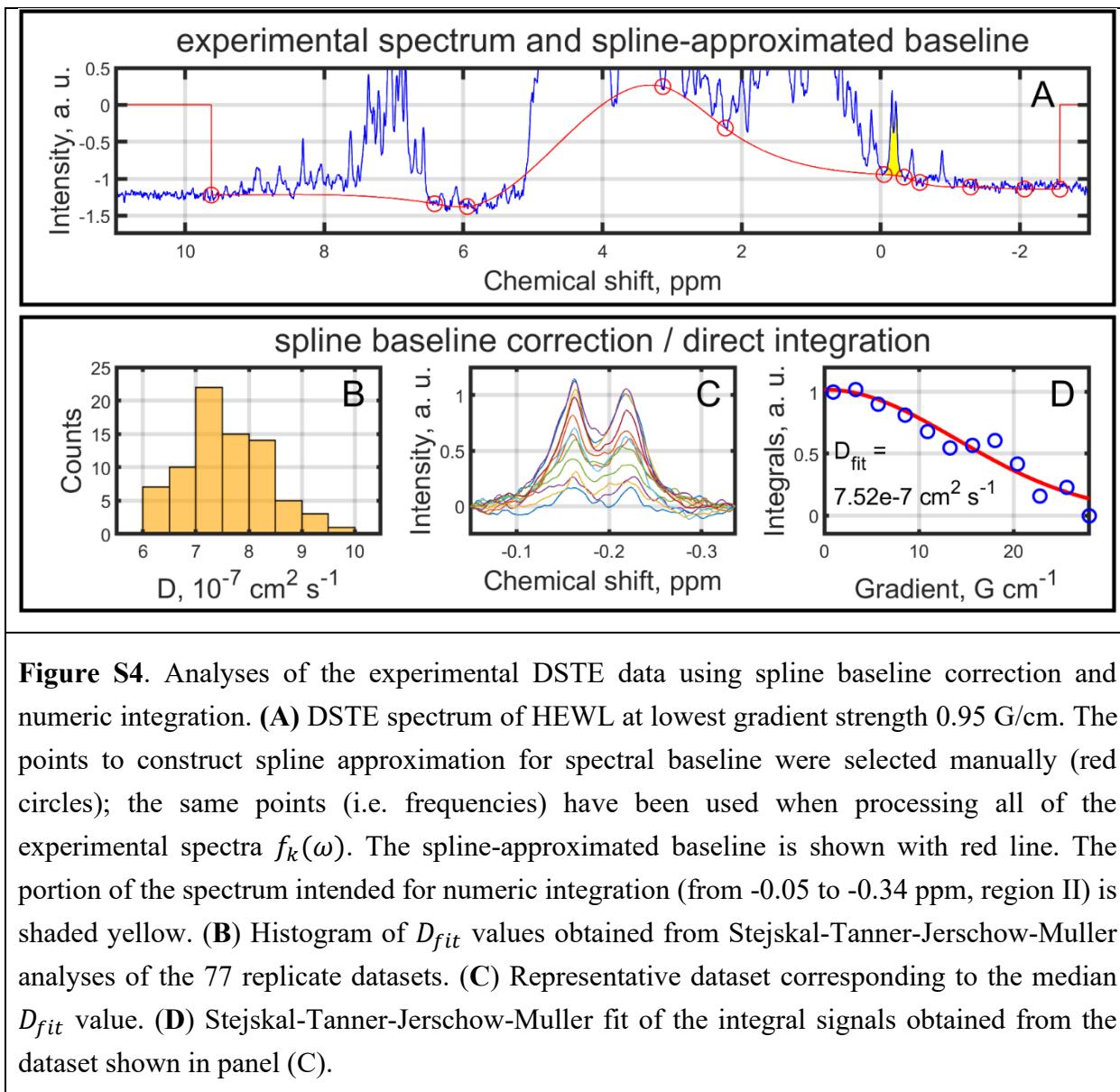
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**Figure S3.** Analyses of the experimental DSTE data from the sample of HEWL using different processing schemes. **Top row:** no baseline correction, direct numeric integration of the spectra in the specified region, from -0.05 to -0.34 ppm (region II). **(A)** Histogram of  $D_{fit}$  values obtained from Stejskal-Tanner-Jerschow-Muller analyses of 77 replicate datasets. **(B)** Representative dataset corresponding to the median  $D_{fit}$  value. **(C)** Stejskal-Tanner-Jerschow-Muller fit of the integral signals from the dataset shown in panel (B). The analysis is compromised by the baseline shift in some of the spectra (those recorded with weaker gradients), leading to dramatic underestimation of diffusion coefficients. **Middle row:** 2-point baseline correction, direct numeric integration of the spectra. **(D)** Histogram of  $D_{fit}$  values. **(E)** Representative dataset corresponding to the median  $D_{fit}$  value. **(F)** Stejskal-Tanner-Jerschow-Muller fit of the integral signals  $I_k^{(0)}$  from the dataset (E). **Bottom row:** 2-point baseline correction, model-based integration as implemented in DDfit. **(G)** Histogram of  $D_{fit}$  values. **(H)** Representative dataset corresponding to the median  $D_{fit}$  value, after DDfit treatment. **(I)** Stejskal-Tanner-Jerschow-Muller fit of the integral signals  $I_k^{(1)}$  obtained from the dataset (H). The plots similar to panels (H) and (I) are generated as a part of the demo by the DDfit server; these results are from the first DSTE dataset out of seventy seven experimental datasets, yielding  $D_{fit} = 7.59 \times 10^{-8} \text{ cm}^2 \text{ s}^{-1}$ .



**Figure S4.** Analyses of the experimental DSTE data using spline baseline correction and numeric integration. **(A)** DSTE spectrum of HEWL at lowest gradient strength 0.95 G/cm. The points to construct spline approximation for spectral baseline were selected manually (red circles); the same points (i.e. frequencies) have been used when processing all of the experimental spectra  $f_k(\omega)$ . The spline-approximated baseline is shown with red line. The portion of the spectrum intended for numeric integration (from -0.05 to -0.34 ppm, region II) is shaded yellow. **(B)** Histogram of  $D_{fit}$  values obtained from Stejskal-Tanner-Jerschow-Muller analyses of the 77 replicate datasets. **(C)** Representative dataset corresponding to the median  $D_{fit}$  value. **(D)** Stejskal-Tanner-Jerschow-Muller fit of the integral signals obtained from the dataset shown in panel (C).