

Highly Conserved Histidine Plays a Dual Catalytic Role in Protein Splicing: a pK_a Shift Mechanism

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Supporting information

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- 1. Various mutants used to study the role of conserved histidines (Table S1);**
- 2. Histidine pK_a estimation from limited number of titration points (Table S2);**
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- 4. Complete Ref.48;**
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1. Table S1. Various mutants used to study the role of conserved histidines.

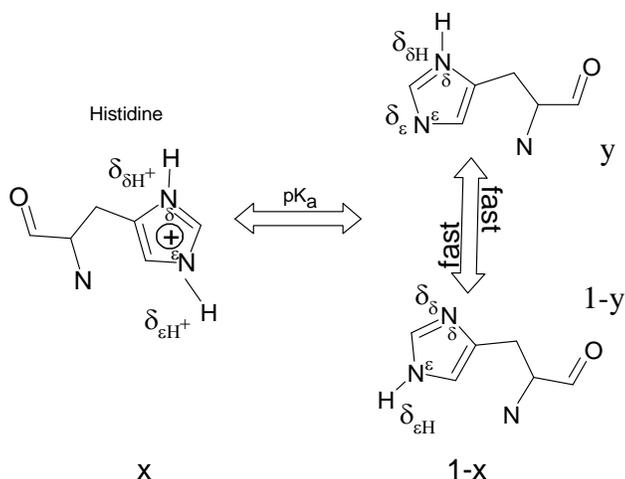
Name	N-extein	Changes relative to $\Delta\Delta\text{Ihh-SM}$ in intein	C-extein	Usage in the study	Relevance to the steps in splicing
$\Delta\Delta\text{Ihh-SM}$ (splicing mutant)	None	None	None	NMR pK_a determination in intein product	All four steps
$\Delta\Delta\text{Ihh-CM}$ (cleavage mutant)		D121G			C-terminal cleavage
NI	17 AAs	C1A		NMR pK_a determination in a precursor with N-extein	N-S acyl shift
MIC	Maltose binding protein (MBP)	None	C-terminal domain of I-TEV1	<i>In vivo</i> splicing and cleavage assay with SDS-PAGE and Western blot	All four steps

2. Histidine pK_a estimation based on average chemical shift of ring nitrogens

The average chemical shift of histidine ring nitrogens is a good indication of the ionization state of the histidine ring¹⁻³. When obtaining the full titration curve is not practical, the pK_a of histidines in NI was determined by the following equation:

$$pK_a = pH + \log \frac{(\tilde{A}_H - \tilde{A}_{obs})}{(\tilde{A}_{obs} - \tilde{A}_{H^+})} \quad (1),$$

where $\tilde{A}_{obs} = 1/2(\delta_{\delta(obs)} + \delta_{\epsilon(obs)})$, $\delta_{\delta(obs)}$ and $\delta_{\epsilon(obs)}$ corresponds to the ¹⁵N chemical shifts of histidine N^{δ1} and N^{ε2} atoms measured by HMQC NMR spectra at individual pHs; $\tilde{A}_H = 1/2(\delta_{\delta} + \delta_{\epsilon H}) = 1/2(\delta_{\epsilon} + \delta_{\delta H})$, δ_{δ} , δ_{ϵ} , $\delta_{\delta H}$, and $\delta_{\epsilon H}$ being the ¹⁵N chemical shifts of histidine (with either δ or ϵ position protonated) when the imidazole ring is neutral (Fig. below); $\tilde{A}_{H^+} = 1/2(\delta_{\delta H^+} + \delta_{\epsilon H^+})$, $\delta_{\delta H^+}$ and $\delta_{\epsilon H^+}$ is the ¹⁵N chemical shifts of histidine with both δ and ϵ position protonated in a positively charged the imidazole ring.



Let x be the fractional population of the charged histidine ring and y be the fractional population with protonated $N\delta$ in tautomer exchange. Based on the scheme in the above figure, we have:

$$\delta_{\delta}(obs) = \delta_{\delta H^+} x + (1-x)y\delta_{\delta H} + (1-x)(1-y)\delta_{\delta} \quad (2)$$

$$\delta_{\epsilon}(obs) = \delta_{\epsilon H^+} x + (1-x)y\delta_{\epsilon} + (1-x)(1-y)\delta_{\epsilon H} \quad (3)$$

Assuming $\delta_{\epsilon} = \delta_{\delta}$, $\delta_{\delta H} = \delta_{\epsilon H}$, and $\delta_{\delta H^+} = \delta_{\epsilon H^+} = \tilde{A}_{H^+}$, (2)+(3) divided by 2 yields,

$$\begin{aligned} & \frac{1}{2}[\delta_{\delta}(obs) + \delta_{\epsilon}(obs)] \\ &= \tilde{A}_{obs} = \delta_{\delta H^+} x + \frac{1}{2}(\delta_{\delta} + \delta_{\epsilon H})(1-x) = \frac{1}{2}(\delta_{\delta} + \delta_{\epsilon H}) - \left[\frac{1}{2}(\delta_{\delta} + \delta_{\epsilon H}) - \delta_{\delta H^+} \right] x \end{aligned} \quad (4)$$

Or

$$x = \frac{\frac{1}{2}(\delta_{\delta} + \delta_{\epsilon H}) - \tilde{A}_{obs}}{\frac{1}{2}(\delta_{\delta} + \delta_{\epsilon H}) - \delta_{\delta H^+}} = \frac{\tilde{A}_H - \tilde{A}_{obs}}{\tilde{A}_H - \tilde{A}_{H^+}} \quad (5)$$

And

$$1-x = \frac{\tilde{A}_{obs} - \tilde{A}_{H^+}}{\tilde{A}_H - \tilde{A}_{H^+}} \quad (6)$$

By definition,

$$pK_a = pH + \log \frac{x}{1-x} \quad (7)$$

Thus, equation (1) can be derived by substituting (5) and (6) into (7).

The canonical values of δ_{δ} or δ_{ϵ} , $\delta_{\delta H}$ or $\delta_{\epsilon H}$, and $\delta_{\delta H^+}$ or $\delta_{\epsilon H^+}$ are 249, 168, and 176 ppm respectively.¹ Incorporating those numbers can simplify equation 1 to,

$$pK_a = pH + \log \frac{(208.5 - \tilde{A}_{obs})}{(\tilde{A}_{obs} - 176)} \quad (8)$$

For His73 in the intein spliced product $\Delta\Delta_{\text{hhI-SM}}$, we have observed $\delta_{\delta} = 243.1$ ppm and $\delta_{\text{eH}} = 165.5$ ppm, resulting in $\tilde{A} = 204.3$. So a slightly modified version of equation 7 gives

$$pK_a = pH + \log \frac{(204.3 - \tilde{A}_{\text{obs}})}{(\tilde{A}_{\text{obs}} - 176)} \quad (8')$$

To eliminate the potentially big error in calculation pH when either the protonated form or the neutral form dominate, only data where pH is within one unit of pK_a , where $\frac{(208.5 - \tilde{A}_{\text{obs}})}{(\tilde{A}_{\text{obs}} - 176)}$ falls between 0.1 and 10, and $179.0 \text{ ppm} < \tilde{A}_{\text{obs}} < 205.5 \text{ ppm}$ should be used.

In table S2, we validate the method by comparing pK_a values determined from equation (8), (8') and from curve-fitting when the full range of pH titration data are available. As seen from the table, pK_a values determined from average chemical shifts of ring nitrogens are consistently within 0.6 of the value determined from full curve-fitting, as long as only chemical shift values at pH within 1 unit of pK_a are used. Equation 8 and 8' yield very similar results, indicating that small errors in the estimation of \tilde{A}_H and \tilde{A}_{H+} won't significantly affect the calculated pK_a values. To ensure that \tilde{A}_H and \tilde{A}_{H+} do not have large variations, we performed statistical analysis based on histidine chemical shifts we obtained and those published in the literature.^{4,5} The average and standard deviation of \tilde{A}_H in 48 neutral histidine rings are 208.2 ± 3.4 ppm. The average and standard deviation of \tilde{A}_{H+} from 18 imidazolium rings are 178 ± 2 ppm.

Table S2

$\Delta\Delta_{\text{hhI-SM}}$	pH	\tilde{A}_{obs} (ppm)	pK_a (Eqn. 8)	pK_a (Eqn. 8')	pK_a (Titration)
His 17	5.19	198.5	4.84	4.60	4.5±0.3
	4.67	191.4	4.71	4.59	
	4.02	182.5	4.62	4.54	
His 30	7.28	200.9	6.77	6.42	7.1±0.2
	6.83	193.9	6.74	6.60	
	6.37	184.1	6.85	6.77	
	5.79	179.9	6.66	6.59	
His 41	7.28	198.2	6.95	6.72	7.1±0.2
	6.83	193.1	6.78	6.65	
	6.37	184.9	6.79	6.71	
	5.79	179.9	6.65	6.58	
His 429	8.2	192.5	8.19	8.06	8.9±0.2
	7.28	181.6	7.96	7.89	

3. Supporting figures

Figure S1: Protein splicing and its side reactions: N-terminal and C-terminal cleavage. N-extein, intein, and C-extein are colored in green, red, and blue.

Figure S2. Assignment of histidine resonances in $\Delta\Delta\text{Ihh-SM}$. A), 600 MHz (HB)CB(CGCD)HD spectrum and B), 600 MHz HMQC spectrum for $\Delta\Delta\text{Ihh-SM}$ at pH 7.0. The assignments of the histidines in HMQC were based on chemical shift of $\text{H}^{\delta 2}$. Histidine $\text{H}^{\delta 2}$ was assigned in (HB)CB(CGCD)HD spectrum based on C^{β} derived from backbone sequential assignments.

Figure S3. Assignment of histidine resonances in $\Delta\Delta\text{Ihh-CM}$. A), 600 MHz (HB)CB(CGCD)HD spectrum and B), 600 MHz HMQC spectrum for $\Delta\Delta\text{Ihh-CM}$ at pH 7.0. The assignments of the histidines in HMQC were based on chemical shift of $\text{H}^{\delta 2}$. Histidine $\text{H}^{\delta 2}$ was assigned in (HB)CB(CGCD)HD spectrum based on C^{β} derived from backbone sequential assignments.

Figure S4. Assignment of histidine resonances in NI. A), 600 MHz (HB)CB(CGCD)HD spectrum and B), 600 MHz HMQC spectrum for NI at pH 7.0. The assignments of the histidines in HMQC were based on chemical shift of $\text{H}^{\delta 2}$. Histidine $\text{H}^{\delta 2}$ was assigned in (HB)CB(CGCD)HD spectrum based on C^{β} derived from backbone sequential assignments.

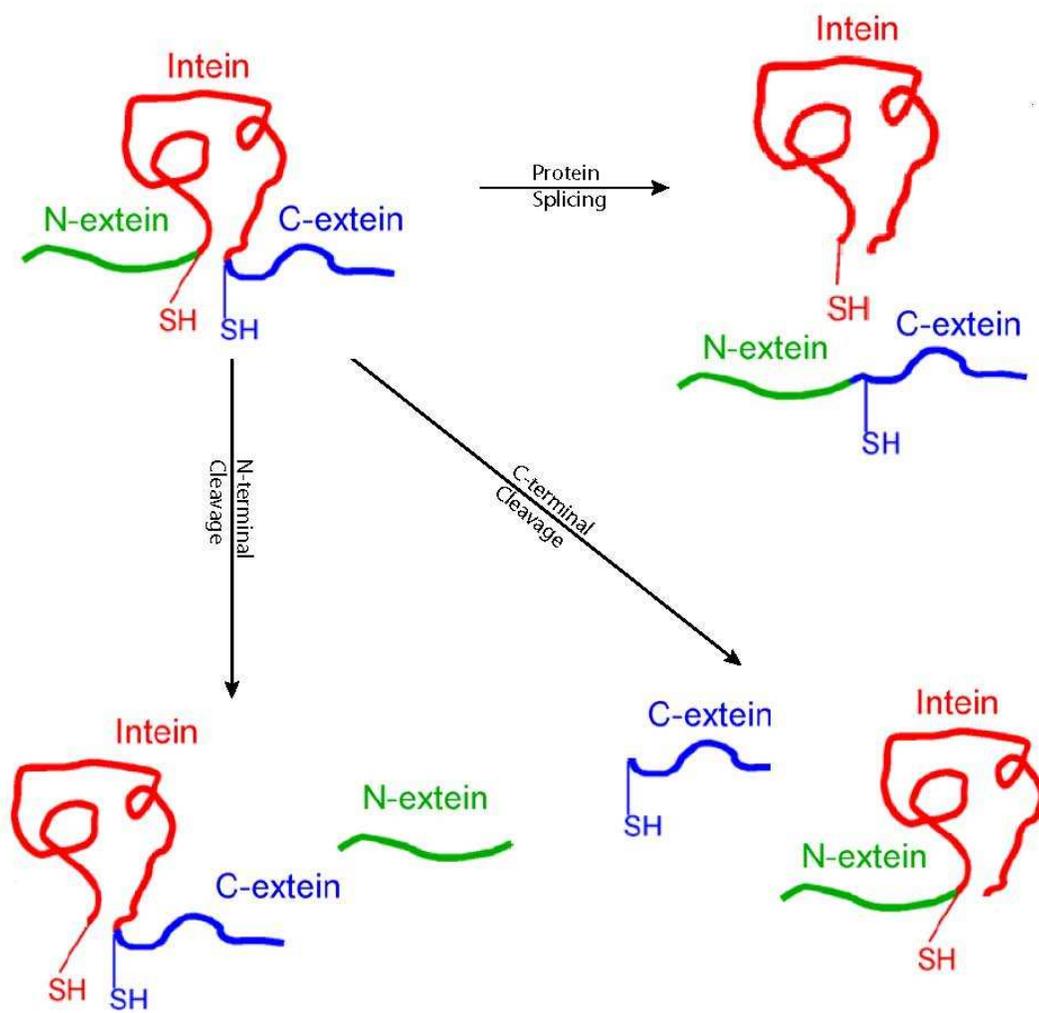


Figure. S1

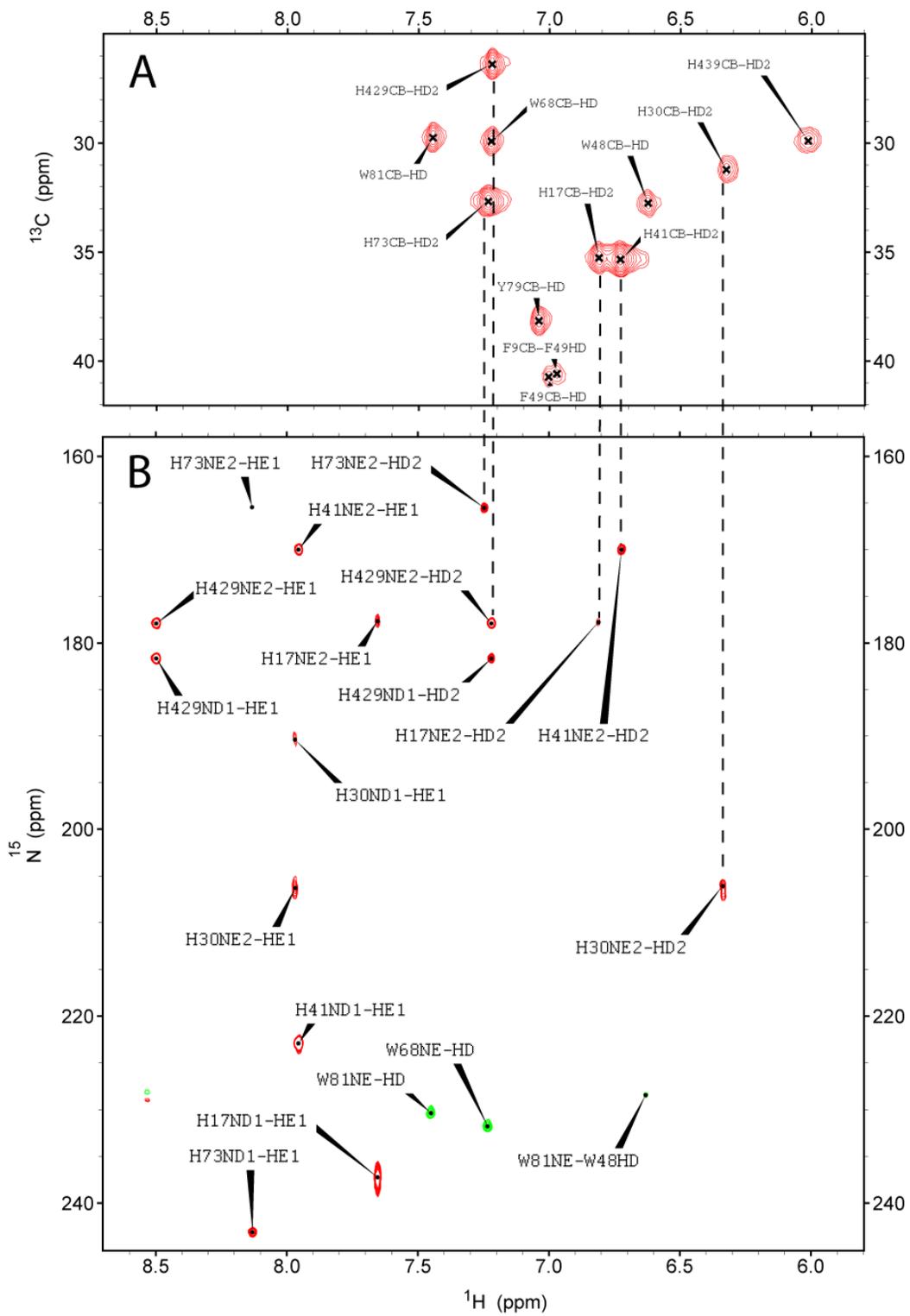


Figure.S2

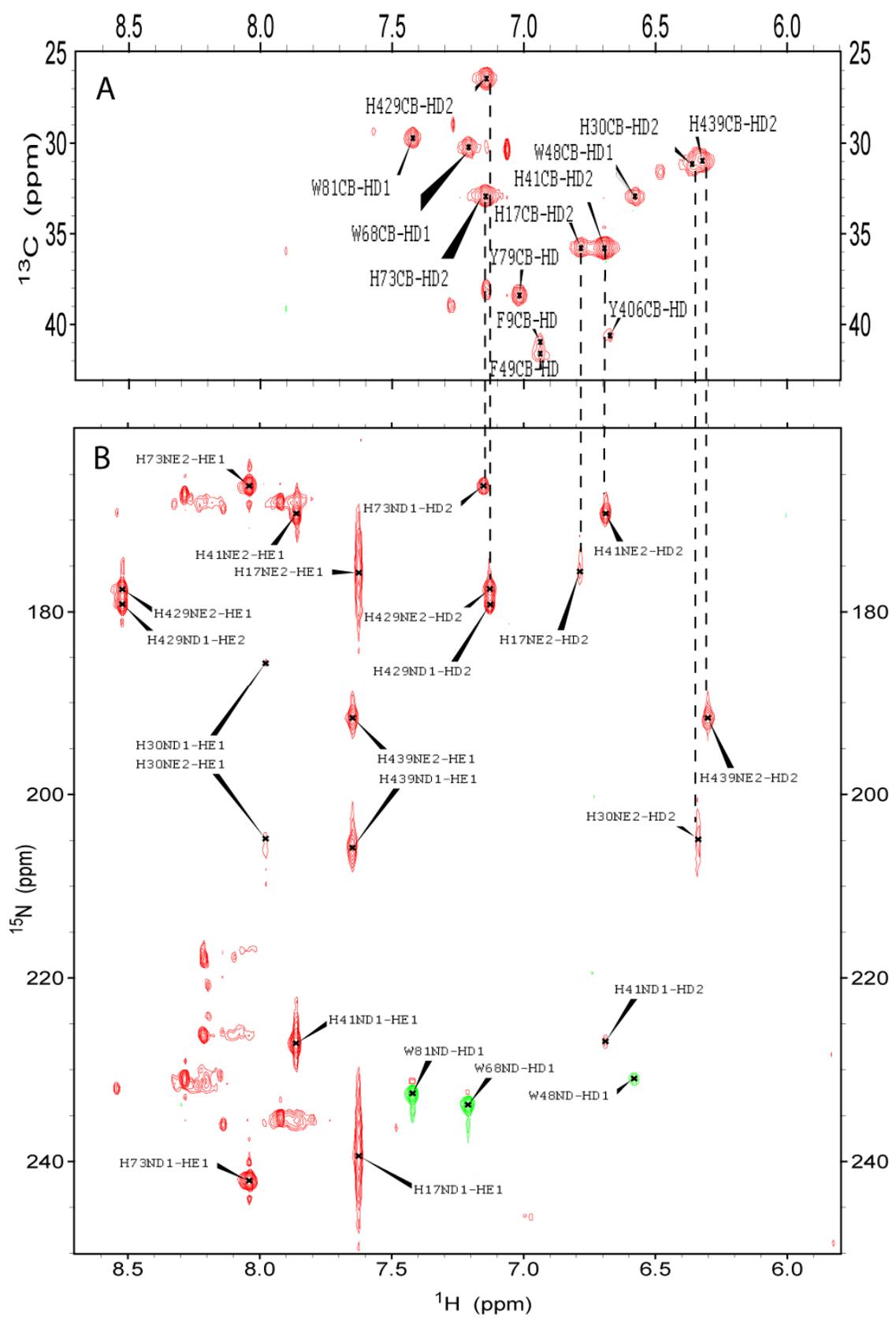


Figure.S3

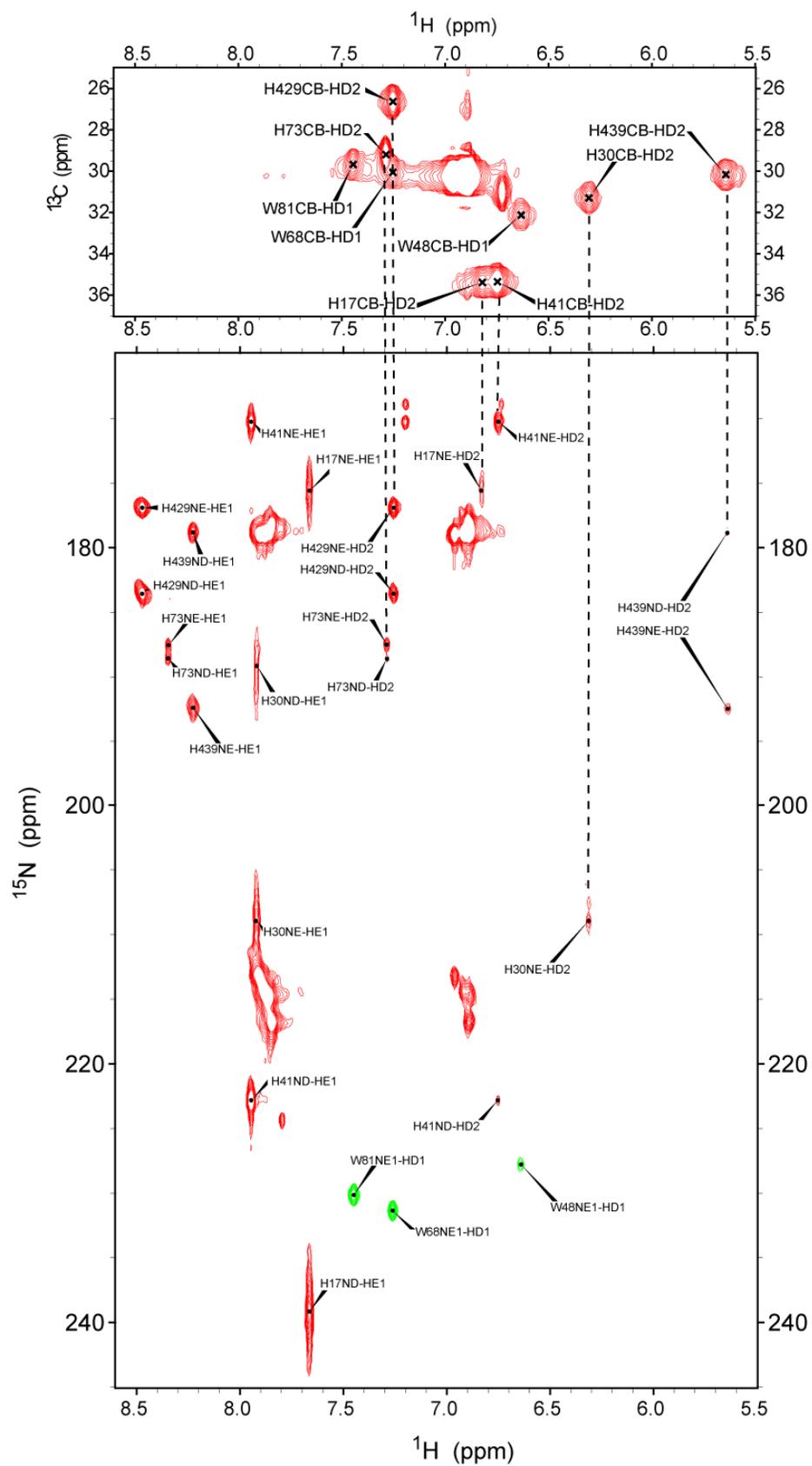


Figure.S4

4. Complete Ref. 48

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