## **Supporting Information for**

"Significant <sup>13</sup>C NMR Signal Enhancements in Amino Acids via Adiabatic Demagnetization and Remagnetization Cross Polarization"

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Parameter	Hartmann-Hahn CP	ADRF/ARRF CP
External field, B <sub>0</sub> (T)	7.05	
Spectrometer	Bruker Avance Neo	
Probe and sample holder	Bruker 3.2 mm HX MAS probe	
Pre-scan delay (s)	5	
Number of transients	128	
<sup>1</sup> H $\pi/2$ rf power (kHz)	62.5	
<sup>1</sup> H decoupling during acquisition	62.5 kHz	
<sup>1</sup> H transfer pulse:		
Shape	Linear Ramp	Downward tanH
Power (kHz)	55 to 70	50 at shape maximum
Length (ms)	6	5
<sup>13</sup> C transfer pulse:		
Shape	Constant	Upward tanH
Power (kHz)	62.5	55 at shape maximum
Length (ms)	6	40

**Table S1**: Experimental Parameters for NMR Experiments on Stationary Glycine Sample

**Table S2**: *Experimental Parameters for NMR Experiments on Stationary*<sup> $\dagger$ </sup> *and Spinning*<sup> $\ddagger$ </sup> *Samples of Tyrosine HCl* 

Parameter	Hartmann-Hahn CP	ADRF/ARRF CP	
External field, B <sub>0</sub> (T)	7.0	7.05	
	Bruker Ava	Bruker Avance Neo	
Probe and sample holder	Bruker 3.2 mm HX MAS probe		
Pre-scan delay (s)	5		
<sup>1</sup> H $\pi/2$ rf power (kHz)	62.	62.5	
<sup>1</sup> H decoupling during acquisition	62.5 kHz		
<sup>1</sup> H transfer pulse:			
Shape	Linear Ramp	Downward tanH	
Power (kHz)	55 to 70	40 at shape maximum	
Length (ms)	6	1	
<sup>13</sup> C transfer pulse:			
Shape	Constant	Upward tanH	
Power (kHz)	62.5	50 at shape maximum	
Length (ms)	6	40	

<sup>†</sup>Stationary-sample experiments used 128 transients for both HH CP and ADRF/ARRF CP <sup>‡</sup>Spinning-sample experiments used 1 transient, a 62.5 kHz rf power for the <sup>13</sup>C  $\pi/2$  flip-back and readout pulses, and 10 kHz sample rotation initiated during the 60 s z-storage period

## Bruker Pulse Sequence for ADRF/ARRF CP:

```
;CP using ADRF/ARRF
;Make AD/AR shapes with Bruker's waveform generator
1 ze
2 d1 do:f2
  (p3 pl12 ph1):f2
 (p20:sp0 ph20):f2
  (p21:sp1 ph21):f1
  1u cpds2:f2
                   ;decoupling sequence
  go=2 ph31
  1m do:f2
  wr #0
HaltAcqu, 1m
exit
ph0= 0
ph1= 1 3
ph20= 2 0
ph21= 0 2 1 3 ; cyclops receiver phase cycling
ph31= 1 2 1 3
```