

Supplementary material to:

Relaxation optimized double acquisition (RODA) as an alternative for virtual decoupling of NMR spectra

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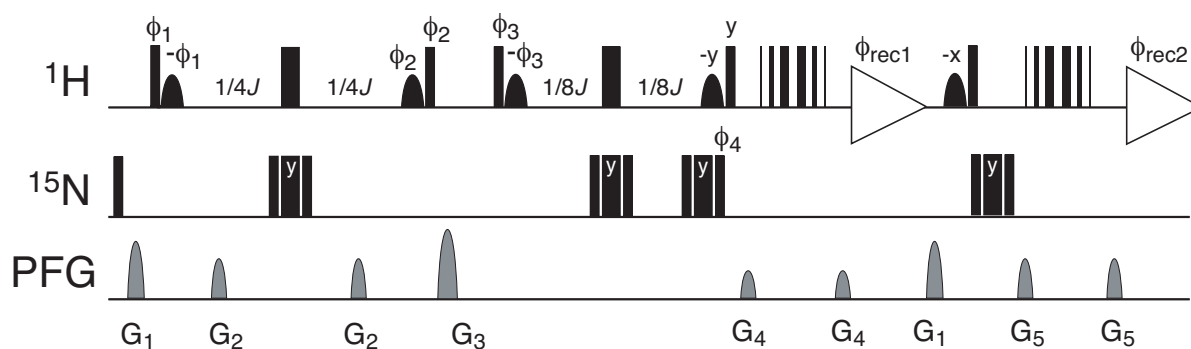


Figure S1. Pulse scheme used for (^{15}N , ^1H) isotope filtering and RODA spin state selection. On the lines labelled ^1H and ^{15}N , representing the respective radio-frequency (rf) channels, narrow and wide bars stand for non-selective 90° and 180° rf-pulses, respectively. Phases are x, unless indicated otherwise above the pulse bar. The ^1H selective pulses (curved shapes) are used to maintain the water magnetization along the +z axis. The relevant J coupling is $^1J_{\text{NH}} = 93$ Hz. For inversion of ^{15}N spins a 90_x - 240_y - 90_x composite pulse was chosen because of its wide inversion profile. To suppress the strong detergent signals in the region between 1 to 5 ppm in the measurements with the sample containing mixed OmpX/DHPC micelles, the refocusing 180° pulse on ^1H was replaced by a REBURP pulse of a duration of 1 ms centred at 8.5 ppm to selectively invert the amide proton spins. Isotope filtering is accomplished using $\phi_1 = y$, $-y$ and $\phi_2 = x$, x , $-x$, $-x$ and spin-state selection is achieved with $\phi_3 = 45^\circ$, 135° , 225° , 315° ; $\phi_4 = x$, $-x$ and $\phi_{\text{rec1}} = x$, x , $-x$, $-x$ and RODA $\phi_{\text{rec2}} = x$, $-x$. Pulsed field gradients (PFGs): $G_1 = 40$ G/cm, 1 ms; $G_2 = 9$ G/cm, 0.8 ms; $G_3 = 35$ G/cm, 0.3 ms; $G_4 = 40$ G/cm, 0.3 ms; $G_5 = 21$ G/cm, 0.3 ms. Water suppression was achieved by a WATERGATE sequence [1].

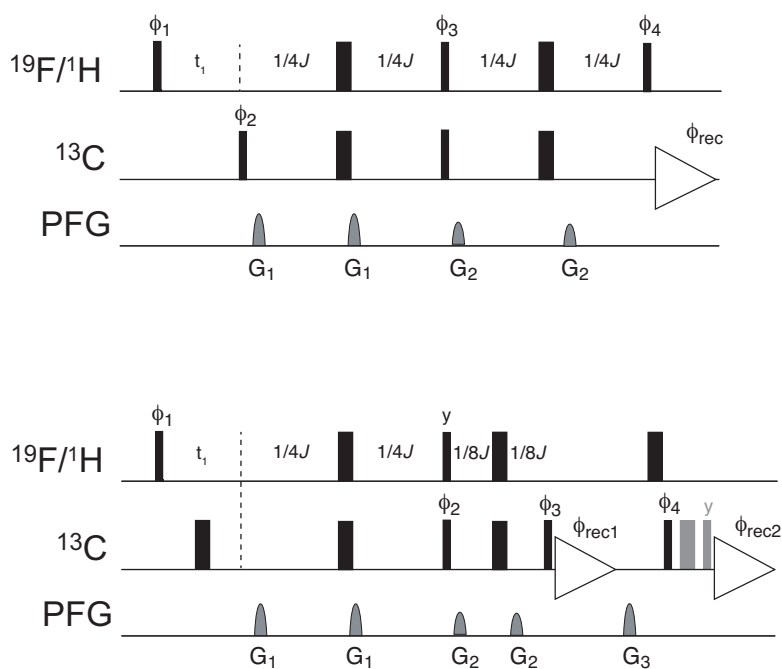


Figure S2. Pulse schemes used for 2D [^{19}F , ^{13}C] and [^1H , ^{13}C] ST2-PT (top) and RODA (bottom) experiments [2–4]. Same schematic representation as in supplementary figure 1.

The relevant J -coupling is $^1J_{\text{CH}}$ (≈ 160 Hz) or $^1J_{\text{FC}}$ (≈ 240 Hz). For ST2-PT the phase cycle is the following: $\phi_1 = y, -y, x, -x$ and $\phi_1' = -y, y, x, -x$ (for quadrature detection, see below); $\phi_2^{\text{D}} = -y$; $\phi_2^{\text{U}} = y$; $\phi_3^{\text{D}} = y$; $\phi_3^{\text{U}} = -y$, where the superscript D and U denote selection of the downfield or the upfield component, respectively, where ϕ_2 acts on the δ_1 and ϕ_3 on δ_2 dimension. $\phi_4 = x$ and $\phi_{\text{rec}} = x, -x, -y, y$. Quadrature detection in the F_1 dimension is performed using the echo-anti echo method. For each t_1 increment ϕ_1 and ϕ_{rec} are inverted, and two scans are recorded with alternating ϕ_1 and ϕ_1' phases and simultaneous inversion of ϕ_2 and ϕ_4 . Pulsed field gradients (PFGs) along the z -axis: $G_1 = 5.5$ G/cm, 0.6 ms; $G_2 = 15.5$ G/cm.

For the corresponding RODA pulse sequence (bottom) the phase cycle is: $\phi_1 = 4x, 4-x$; $\phi_2^{\text{D}} = 315^\circ, 135^\circ$; $\phi_2^{\text{U}} = 45^\circ, 225^\circ$; $\phi_3^{\text{D}} = -y, -y, y, y$; $\phi_3^{\text{U}} = y, y, -y, -y$; $\phi_4 = x, -x, -x, x$ and $\phi_{\text{rec1}} = \phi_{\text{rec2}} = x, -x, x, -x, -x, x, -x, x$. Quadrature detection in the F_1 dimension is performed using the States-TPPI method. For each t_1 increment two scans are recorded with ϕ_1 incremented by 90° . PFGs along the z -axis: $G_1 = 30.0$ G/cm, 0.6 ms; $G_2 = 22.5$ G/cm; $G_2 = 18.5$ G/cm. The grey pulses at the end of the sequence can be inserted for obtaining a similar baseline in both scans if distortions are observed.

Pulse program 1: 2D [¹H,¹HN]-NOESY with ¹⁵N editing in 2nd dimension, and TROSY component selection with (¹⁵N)-¹H RODA

(Pulse program for spectrum shown in Fig. 3 of main text)

```
; 2D [1H,1HN]-NOESY with 15N editing in F2, and TROSY component selection with (15N)-1H RODA
```

```
; A.D. Gossert, G. Wider, J. Magn. Res. (2022)
```

```
;optimization of water flip back:
```

```
;- optimize water flipdown (sp1,ph11,ph18)
```

```
;- optimize water flipup (sp2,ph26)
```

```
;Attention: Pulse sequence is not prosol compatible
```

```
;p11 : power for 1H
```

```
;p12 : power for 13C
```

```
;p13 : power for 15N
```

```
;sp1 : water flipup power
```

```
;sp2 : water flipdown power
```

```
;spnam1: gauss128_5
```

```
;spnam2: gauss128_5
```

```
;p1 : 90 degree hard pulse 1H
```

```
;p3 : 90 degree hard pulse 13C
```

```
;p4 : 180 degree hard pulse 13C (225d for 5/600)
```

```
;p5 : 90 degree hard pulse 15N
```

```
;p11 : water flipback pulse (1.5m)
```

```
;p21 : 900u (Gradient in first INEPT)
```

```
;p22 : 900u (Gradient in second INEPT)
```

```
;p23 : 400u (Gradient in 1st WG)
```

```
;p24 : 900u (z-filter Gradient)
```

```
;p25 : 400u (Gradient in 2nd WG)
```

```
;gpz1 : 80%
```

```
;gpz2 : 35%
```

```
;gpz3 : 70%
```

```
;gpz4 : 80%
```

```
;gpz5 : 70%
```

```
;d1 : relaxation delay (1 s)
```

```
;d2 : 1/2J delay (5.4 ms)
```

```
;d4 : 1/2J delay (5.4 ms)
```

```
;d5 : delay 3-9-19=1/(Hz between nulls) (170-250 us)
```

```
;in0 : 1/(2 SW) (Hz)
```

```
define delay DEL11
```

```
define delay DEL12
```

```
define delay DEL21
```

```
define delay DEL22
```

```
define delay DEL23
```

```
define delay DEL2
```

```
#include <Avance.incl>
```

```
#define GRADIENT1 10u p21:gp1 200u
```

```
#define GRADIENT2 10u p22:gp2 200u
```

```
#define GRADIENT3 10u p23:gp3 200u
```

```
#define GRADIENT4 10u p24:gp4 200u
```

```

#define GRADIENT5 10u p25:gp5 200u

#define WG3919 p1*0.2308 ph20 d5 p1*0.6923 ph20 d5 p1*1.4615 ph20 d5 p1*1.4615 ph22 d5 p1*0.6923 ph22
d5 p1*0.2308 ph22

"p2=2*p1"
"p4=2*p3"
"p8=2*p5"
"p6=8*(p5/3)"

"DEL11=(d2/2)-(p22+210u)-p5-p11-10u"
"DEL12=(d2/2)-(p22+210u)-p5-p11-10u"
"DEL21=(d2/4)-(p11+10u)-p6/2-p5+80u"
"DEL22=(d2/4)-(p11+10u)-p6/2-p5+80u"
"DEL23=6u+p5+p5+p6"

"d0=in0/2-(p1*2/3.1415)"

"l2 = 1"

1 10u ze
2 1m
5 10u
  d1
  20u p11:f1
  20u p12:f2
  20u p13:f3
  20u LOCKH_ON
;----- 1H evolution
  (refalign (p1 ph8 d0 d0 p1 ph25):f1 center (p8 ph20):f3 center (p3 ph20 2u p4 ph21 2u p3 ph20):f2)
;----- NOE mixing time
  d7
  (p5 ph20):f3
  GRADIENT1
  10u
;----- 15N-editing element
  (p1 ph1):f1
  10u
  (p11:sp1 ph11:r):f1 (p5 ph20 3u p6 ph21 3u p5 ph24):f3
  GRADIENT2
  DEL11
  (center (p2 ph20):f1 (p5 ph20 3u p6 ph21 3u p5 ph20):f3)
  GRADIENT2
  DEL12
  (refalign (p11:sp1 ph12:r):f1 ralign (p5 ph20 3u p6 ph21 3u p5 ph24):f3)
  10u p11:f1
  (p1 ph2):f1
;----- z-filter gradient against remaining H
  GRADIENT4
  (p1 ph3):f1
;----- 1/4J separate I13 I24
  (lalign (p5 ph20 3u p6 ph21 3u p5 ph26):f3 (10u p11:sp1 ph13:r):f1)
  DEL21 p11:f1
  (center (p2 ph20):f1 (p5 ph20 3u p6 ph21 3u p5 ph20):f3)
  DEL22 p11:f1
  (ralign (p5 ph20 3u p6 ph21 3u p5 ph26):f3 (p11:sp1 ph14:r 10u p11):f1)
  (p1 ph4:r):f1
;----- Watergate
  GRADIENT3
  (WG3919):f1
  GRADIENT3
;----- 1st acquisition I13

```

```

    goscpn ph31
;----- Clean up
    GRADIENT5
;----- Record second component
    (p11:sp2 ph15:r):f1
    10u p11:f1
    (p1 ph20):f1
    (p5 ph20 3u p6 ph21 3u p5 ph20):f3
    GRADIENT5
    (WG3919):f1
    GRADIENT5
    DEL23
;----- 2nd acquisition I13
    go=2 ph30
    1m mc #0 to 2
        F1PH(ip8, id0)

10u do:f1
10u do:f2
10u do:f3
10u do:f4
10u LOCKH_OFF
exit

;----- Phases

ph1 =    1 1 3 3
ph2 =    0 2 0 2 2 0 2 0
ph3 =(8) 1 3 5 7
ph4 =    1
ph8 =    0 0 0 0 0 0 0 0 2 2 2 2 2 2 2 2

ph11=    3 3 1 1
ph12=    2 0 2 0 0 2 0 2
ph13=(8) 5 7 1 3
ph14=    3
ph15=    2

ph20=    0
ph21=    1
ph22=    2
ph23=    3

ph24=    0 0 0 0 2 2 2 2
ph25=(8) 1
ph26=    2 0

ph30=3 1 3 1 3 1 3 1 1 3 1 3 1 3 1 3
ph31=0 0 0 0 2 2 2 2 2 2 2 2 0 0 0 0

```

Pulse program 2:

1D (¹⁹F)-¹³C RODA

(Pulse program for spectrum shown in Fig. 4 of main text)

```
; 1D (19F)-13C RODA

; A.D. Gossert, G. Wider, J. Magn. Res. (2022)

;- NBL : 2
;- ZGOPTNS: -DDOWNFIELD: select downfield component, default selects upfield component

;Channels for (19F)-13C-RODA
;p11 : power for 13C
;p12 : power for 19F
;p13 : power for 1H

;p1 : 90 degree hard pulse 13C
;p3 : 90 degree hard pulse 19F

;p16 : 1000u (long gradient)
;p19 : 300u (short gradient)
;gpz1 : 60%
;gpz2 : 35%
;gpz3 : 47%

;d1 : relaxation delay (1s)
;d2 : 1/2J delay

;cnst2 : J-coupling in Hz (240 Hz for 1J(13C,19F))

define delay DEL11
define delay DEL12
define delay DEL21
define delay DEL22
define delay DEL23

#include <Avance.incl>

#define GRADIENT1 10u p16:gp1 500u
#define GRADIENT2 10u p19:gp2 150u
#define GRADIENT3 10u p16:gp3 200u

"p2=2*p1"
"p4=2*p3"

"d2=1/2*cnst2"

"DEL21=(d2/4)-(p19+160u)-(larger(p2,p4))/2"
"DEL22=(d2/4)-(p19+160u)-(larger(p2,p4))/2"
"DEL23=p4"

1 10u ze
2 30m LOCKH_OFF
  10u do:f3
  10u do:f1
  d1
  10u st0
  20u p11:f1
  20u p12:f2
  20u p113:f3
  20u LOCKH_ON
;----- Purge z magnetisation on F
5 10u
  (p3 ph20):f2
  GRADIENT1
;----- 1/4J to separate I13 I24
  (p1 ph1):f1
```

```

GRADIENT2
DEL21
(center (p2 ph20):f1 (p4 ph20):f2)
GRADIENT2
DEL22
(p1 ph2:r):f1
;----- 1st acquisition I13
goscnp ph30 cpd3:f3
;----- Clean up
10u do:f3
GRADIENT3
10u st
20u pl13:f3
;----- Read I24 and convert to I13
(p1 ph3):f1
(p4 ph20):f2
(p2 ph4):f1
DEL23
(p1 ph21):f1
;----- 2nd acquisition I13
go=2 ph31 cpd3:f3
10u do:f3
10m wr #0

10u LOCKH_OFF
10u do:f1
10u do:f2
10u do:f3
10u LOCKH_OFF
exit

;----- Phases

#ifdef DOWNFIELD
    ph1 =(8) 7 3      ; Selection of downfield component
    ph2 =   3 3 1 1
#else
    ph1 =(8) 1 5      ; Selection of upfield component
    ph2 =   1 1 3 3
#endif

ph3 = 0 2 2 0
ph4 = 1 0

ph20= 0
ph21= 1
ph22= 2
ph23= 3

ph30= 0 2

ph31= 0 2

```


Pulse program 3:
2D [¹⁹F, ¹³C]-HSQC-RODA and 2D [¹H, ¹³C^{aro}]-HSQC-RODA
(Pulse program for spectrum shown in Fig. 5B of main text)

```
; 2D [19F,13C]-HSQC-RODA and 2D [1H,13Caro]-HSQC-RODA
; A.D. Gossert, G. Wider, J. Magn. Res. (2022)
;- Switch channels 2 and 3 for 2D [1H,13Caro]-HSQC-RODA
;- ZGOPTNS: -DDOWNFIELD: select downfield component, default selects upfield component

;p11 : power for 13C
;p12 : power for 19F
;p13 : power for 1H

;p1 : 90 degree hard pulse 13C
;p3 : 90 degree hard pulse 19F

;p16 : 1000u (long gradient pulse)
;p19 : 300u (short gradient pulse)
;gpz1 : 60%
;gpz2 : 35%
;gpz3 : 47%

;d1 : relaxation delay (1s)
;d2 : 1/2J delay

;cnst2 : J-coupling in Hz

define delay DEL11
define delay DEL12
define delay DEL21
define delay DEL22
define delay DEL23
define delay DELTA

#include <Avance.incl>

#define GRADIENT0 10u p16:gp0 500u
#define GRADIENT1 10u p19:gp1 200u
#define GRADIENT2 10u p19:gp2 150u
#define GRADIENT3 10u p16:gp3 200u

"p2=2*p1"
"p4=2*p3"

"d2=1/2*cnst2"

"d0=3u"
"in0=inf1/2"
"DELTA=d0*2+p2"

"DEL11=(d2/2)-(p19+210u)-(larger(p2,p4))/2"
"DEL12=(d2/2)+2*d0+p2-(p19+210u)-(larger(p2,p4))/2"
"DEL21=(d2/4)-(p19+160u)-(larger(p2,p4))/2" ;consider adding baseopt echo delay
"DEL22=(d2/4)-(p19+160u)-(larger(p2,p4))/2"
"DEL23=p4"

1 10u ze
2 30m LOCKH_OFF
10u do:f3
10u do:f1
d1
20u p11:f1
20u p12:f2
20u p113:f3
20u LOCKH_ON
;----- Purge z magnetisation on 13C
5 10u
```

```

    (p1 ph20):f1
    GRADIEN0
;----- First 1/2J INEPT from 19F to 13C
    (p3 ph10):f2
    d0
    (p2 ph20):f1
    d0
    GRADIEN1
    DEL11
    (center (p4 ph20):f2 (p2 ph20):f1 )
    GRADIEN1
    DEL12
    (p3 ph21):f2
;----- Refocussing 1/4J INEPT to separate I13 I24
#ifdef DOWNFIELD
    (p1 ph11):f1
#else
    (p1 ph1):f1
#endif
    GRADIEN2
    DEL21
    (center (p2 ph20):f1 (p4 ph20):f2)
    GRADIEN2
    DEL22
#ifdef DOWNFIELD
    (p1 ph12:r):f1
#else
    (p1 ph2:r):f1
#endif
;----- 1st acquisition I13
    goscpn ph30 cpd3:f3
;----- Clean up
    10u do:f3
    GRADIEN3
    20u p13:f3
;----- Read I24 and convert to I13
    (p1 ph3):f1
    (p4 ph20):f2
    (p2 ph4):f1
    DEL23
    (p1 ph21):f1
;----- 2nd acquisition I13
    go=2 ph31 cpd3:f3
    10u do:f3
    30m mc #0 to 2
                                F1PH(calph(ph10, +90), caldel(d0, +in0))

10u LOCKH_OFF
10u do:f1
10u do:f2
10u do:f3
10u LOCKH_OFF
exit

;----- Phases
ph10=   0 0 0 0 2 2 2 2
ph1 =(8) 1 5                ; Selection of upfield component
ph11=(8) 7 3                ; Selection of downfield component
ph2 =   1 1 3 3
ph12=   3 3 1 1                ; Inverted phase for correct phase in downfield component case

ph3 =   0 2 2 0
ph4 =   0 1

ph20=0
ph21=1
ph22=2
ph23=3

ph30=0 2 0 2 2 0 2 0
ph31=0 2 0 2 2 0 2 0

```

Pulse program 4:
2D [¹⁹F, ¹³C]-TROSY, 2D [¹H, ¹³C]-TROSY and 2D [¹H, ¹⁵N]-TROSY
(Pulse program for spectrum shown in Fig. 5 of main text)

```

; 2D [I,X]-TROSY with component selection by means of ST2PT, X-detected
; I = 19F or 1H, X = 13C or 15N.

; A.D. Gossert, G. Wider, J. Magn. Res. (2022)
; K. Takeuchi, H. Arthanari, I. Shimada, G. Wagner, J Biomol NMR. 63 (2015) 323-331

;-ZGPTNS:
;
;           Component selection: -DF1U -DF2D
;           F1 = Dimension 1, Up- or Downfield component
;
;           FC-TROSY: F1U, F2U
;           HN-TROSY: F1D, F2U
;           HC-TROSY: F1D, F2D
;           default FC-TROSY F1U, F2U
;
;           Component selection depends on several factors, test on your spectrometer, if correct
;           component is selected

;for FC TROSY: f1:13C, f2:19F, f3:1H, f4:2H
;p11 : power for 13C
;p12 : power for 19F
;p13 : power for 1H

;p1 : 90 degree hard pulse f1 (13C)
;p3 : 90 degree hard pulse f2 (19F)

;p16 : 1000u (long gradient pulse)
;p19 : 300u (short gradient pulse)
;gpz0: 3%
;gpz1: 30%
;gpz2: 80%
;gpz3: 45%
;gpz4: 50%
;gpz5: 16.2%
;gpnam0: SINE.100
;gpnam1: SINE.100
;gpnam2: SINE.50
;gpnam3: SINE.100
;gpnam4: SINE.100
;gpnam5: SINE.50

;d0 : incremented delay (2D) [6 usec]
;d1 : relaxation delay; 1-5 * T1
;d11: delay for disk I/O [30 msec]
;d16: delay for homospoil/gradient recovery

;d26 : 1/(4J)YH

;cnst4: = J(YH)
;inf1: 1/SW(X) = 2 * DW(X)
;in0: 1/(2 * SW(X)) = DW(X)
;nd0: 2
;FnMODE: echo-antiecho

#include <Avance.incl>
#include <Grad.incl>
#include <Delay.incl>

"p2=p1*2"
"p4=p3*2"

```

```

"d11=30m"
"d12=20u"

"d21=1s/(cnst4*4)"

"d0=3u"

"DELTA1=d21-p16-d16-4u"
"DELTA2=d21-p1-p16-d16-4u"

"i0=inf1/2"

"l0=1"

1 d11 ze
  d11 LOCKDEC_ON
  50u LOCKH_ON
  d11 H2_PULSE
  d11 p117:f4

2 d11 do:f4 do:f3 do:f2
3 d11 H2_LOCK
  9m LOCKH_OFF
  d1 p11:f1
  50u UNBLKGRAD
  d12 H2_PULSE

4 d11 p11:f1 p12:f2 p13:f3

;----- Start on 19F, evolution without decoupling
  if "l0 %2 == 1"
    {
      (p3 ph4):f2
    }
  else
    {
      (p3 ph5):f2
    }

  d0
  d0

;----- First INEPT to 13C
  (p1 ph6)
  4u
  p16:gp3
  d16
  DELTA2 p11:f1
  (center (p2 ph1) (p4 ph1):f2 )
  4u
  DELTA1 p116:f3
  p16:gp3
  d16

;----- Refocussing INEPT on 13C
  (center (p1 ph1) (p3 ph8):f2 )
  4u
  p16:gp4
  d16
  DELTA1
  (center (p2 ph1) (p4 ph1):f2 )
  DELTA1 p116:f3
  p16:gp4
  d16
  4u
  (p3 ph7):f2
  4u BLKGRAMP

;----- Acquisition on 13C
go=2 ph31 cpd3:f3 cpd4:f4
d11 do:f2 do:f3 do:f4 mc #0 to 2

```

```
F1EA(calph(ph6, +180) & calph(ph7, +180) & calclc(10, 1), caldel(d0, +in0) & calph(ph4, +180) & calph(ph5, +180) & calph(ph31, +180))
```

```
d11 do:f2 do:f3 do:f4
```

```
d11 H2_LOCK  
d11 LOCKH_OFF  
d11 LOCKDEC_OFF
```

```
exit
```

```
;----- Phases
```

```
ph1=0  
ph2=1  
ph3=3  
ph4=1 3 0 2  
ph5=3 1 0 2
```

```
#ifdef F1D  
    ph6=3  
#else  
    ph6=1  
#endif
```

```
#ifdef F2D  
    ph8=1  
#else  
    ph8=3  
#endif
```

```
ph7=0
```

```
ph31=0 2 3 1
```

Pulse program 5: 1D (¹³C')-¹³C^{ali} RODA

(Pulse program for spectrum shown in Fig. 6 of main text)

```
; 1D CaCO RODA

; A.D. Gossert, G. Wider, J. Magn. Res. (2022)

; W. Bermel, I. Bertini, L. Duma, I.C. Felli, L. Emsley, R. Pierattelli,
; P.R. Vasos, Angew. Chem. Int. Ed. 44, 3089-3092 (2005)

;p11 : f1 channel - power level for pulse (default)
;p112: f2 channel - power level for CPD/BB decoupling

;sp23: f1 channel - shaped pulse 90 degree (on resonance)
;sp24: f1 channel - shaped pulse 180 degree (on resonance)
;sp25: f1 channel - shaped pulse 90 degree (on resonance)
;
; for time reversed pulse
;sp27: f1 channel - shaped pulse 180 degree (Ca off resonance)

;p11: f1 channel - 90 degree shaped pulse
;p12: f1 channel - 180 degree shaped pulse
;cpd2: decoupling according to sequence defined by cpdprg2
;pcpd2: f2 channel - 90 degree pulse for decoupling sequence

;d1 : relaxation delay; 1-5 * T1
;d2 : 1/(8J(COCa)) [4.5 msec]
;cnst2: CaCO J-coupling (in Hz)
;cnst21: CO chemical shift (offset, in ppm)
;cnst22: Calpha chemical shift (offset, in ppm)
;o1p: CO chemical shift (cnst21)

;l0: flag to switch between inphase and antiphase
;inf1: 1/SW(Ca) = 2 * DW(Ca)
;in0: 1/(2 * SW(Ca)) = DW(Ca)
;nd0: 2

;p16: homospoil/gradient pulse [1 msec]
;d16: delay for homospoil/gradient recovery
;gpz1: 50%
;gpnam1: SMSQ10.100

prosol relations=<triple_c>

#include <Avance.incl>
#include <Delay.incl>
#include <Grad.incl>

"d11=30m"
"d12=20u"

"d2=1/(8*cnst2)"

"d0=3u"
"d20=d2-p12-4u-p12/2"
"in0=inf1/2"

"spoff23=0"
"spoff24=0"
"spoff25=0"
"spoff26=bf1*((cnst21-cnst22)/1000000)"
"spoff27=bf1*((cnst22-cnst21)/1000000)"
"spoff28=0"
```

```

1 ze
  d11 p112:f2
2 d11 do:f2
3 d1
  d12 cpd2:f2
  50u UNBLKGRAD
  p16:gp1
  d16 fq=cnst21(bf ppm):f1
;----- 1/4J INEPT to separate I13 I24
  (p11:sp23 ph1)
  4u
  (p12:sp27 ph20) ;Bloch-Siegert compensation
  d20
  (p12:sp24 ph20)
  4u
  (p12:sp27 ph20)
  d20
  (p11:sp25 ph2)
;----- 1st acquisition I13
  goscnp ph30
;----- Clean up
  p16:gp2
  d16
  10u st
  10u BLKGRAD
;----- Read I24 and convert to I13
  (p12:sp27 ph20):f1
  (p11:sp23 ph3):f1
;----- 2nd acquisition I13
  go=2 ph31
  10m do:f2
  10m wr #0

10u do:f1
10u do:f2
exit

;----- Phases

#ifdef DOWNFIELD
  ph1 =(8) 7 3      ; Selection of downfield component
; ph2 =  3 1      ; phase for -z
  ph2 =  1 3      ; phase for +z
#else
  ph1 =(8) 1 5      ; Selection of upfield component
; ph2 =  1 3      ; phase for -z
  ph2 =  3 1      ; phase for +z
#endif

ph3 =0
ph4 =1

ph20=0
ph21=1
ph22=2
ph23=3

ph30=2 0

ph31=0

```

References

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- [3] K. Takeuchi, H. Arthanari, I. Shimada, G. Wagner, Nitrogen detected TROSY at high field yields high resolution and sensitivity for protein NMR, *J Biomol NMR*. 63 (2015) 323–331. <https://doi.org/10.1007/s10858-015-9991-y>.
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