

**Supplementary information for
*Spatially encoded diffusion-ordered NMR spectroscopy of reaction mixtures in
organic solvents***

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I- Supplementary figures

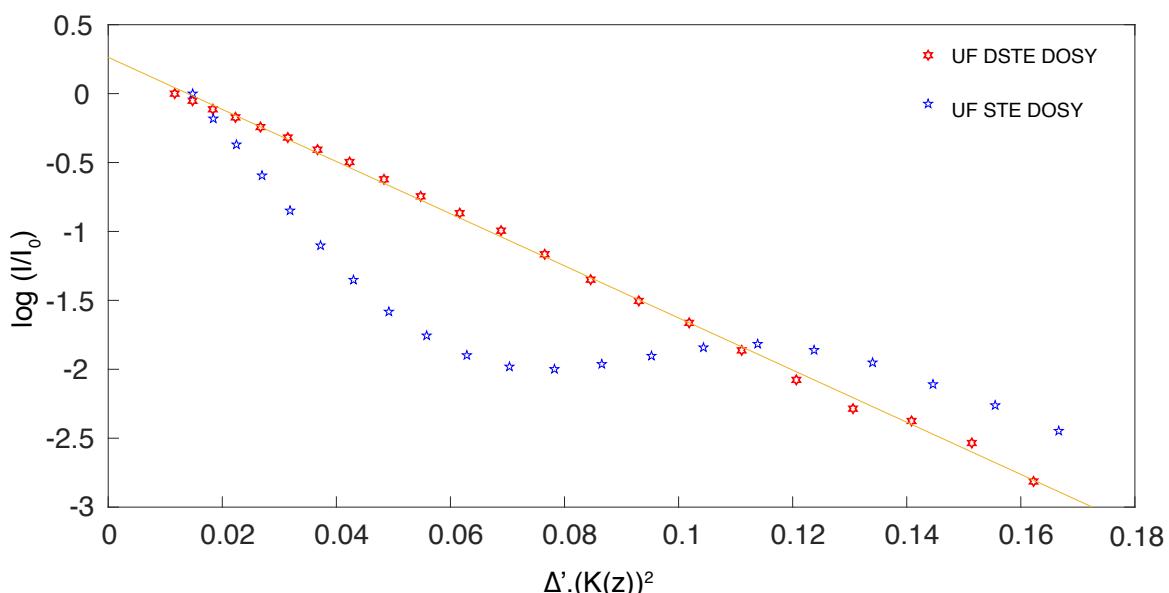


Figure S1 Signal decay as a function of $\Delta' \cdot (K(z))^2$, where Δ' is the effective diffusion delay and $K(z)$ is the effective gradient area (see Ref. ¹), for the cyclododecane resonance for the SPEN STE DOSY (blue) and SPEN DSTE DOSY (red) experiments carried out in CD_2Cl_2 . The diffusion delay was 140 ms for both SPEN STE and DSTE experiment.

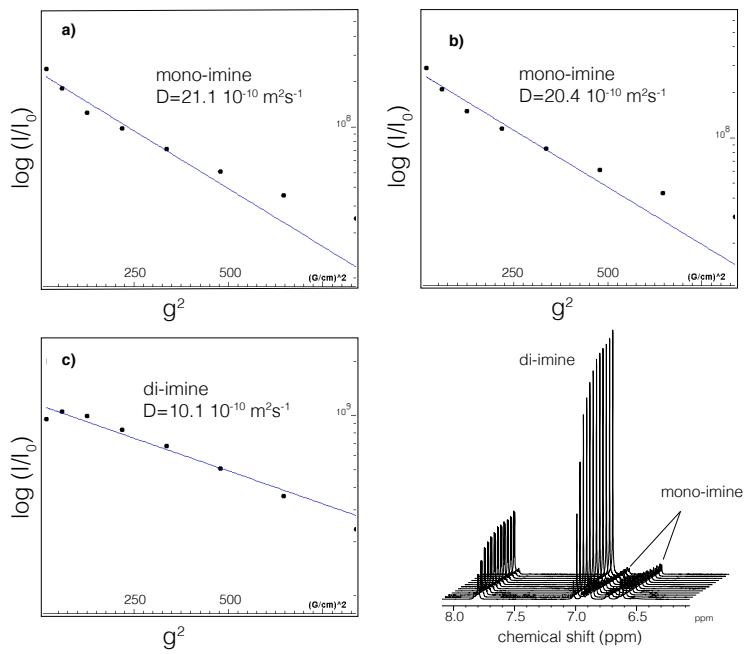


Figure S2. Signal attenuation as a function of the squared diffusion gradient amplitude for selected resonances of the mono- and di-imine compounds, obtained with a conventional DOSY sequence, in CD_3CN , during the course of a di-amination reaction.

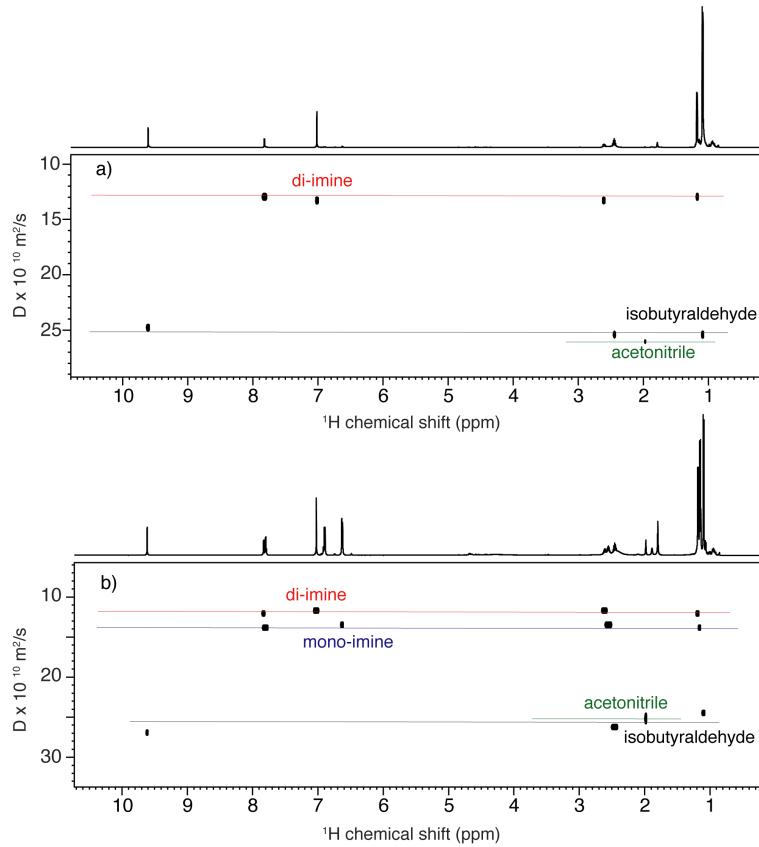


Figure S3 DOSY spectra of the reaction mixture in CD_3CN at 283 K, at equilibrium, obtained with (a) the conventional DSTE DOSY pulse sequence using stoichiometry i/; (b) the conventional DSTE DOSY experiment using stoichiometry ii/ (see sec. B for the stoichiometry).

II- Pulse sequence for spectrally selective SPEN DOSY (ssSPEN DSTE)

```

;ssSPENDSTE
#include <Avance.incl>
#include <Grad.incl>
#include <De.incl>
#include <Delay.incl>

"d20=(td*dw/(2*I3))-d6"
"DELTA1=d30/2-p11-2*d4-d11-2*p20-3*d16-p22-2*p1"
"p2=p1*2"

1 ze
20u st0
30m zd
2 30m
d1
50u UNBLKGRAD
20u pl1:f1
;-----excitation
p1 ph1
d5
;-----first spin echo
;-----spatial encoding
p20:gp20
d16 pl0:f1
d4 gron0
p11:sp1:f1 ph2
d11
d4 groff
p20:gp21
d16 pl1:f1
p1 ph1
p22:gp14
d16
DELTA1
p1 ph1
;-----decoding period
p20:gp22
d16 pl0:f1
d4 gron0
p11:sp1:f1 ph2
d11
d4 groff
p20:gp23
d16 pl1:f1
;-----Middle 180 pulse
p2 ph3
;-----second spin echo
;-----spatial encoding
p20:gp26
d16 pl0:f1
d4 gron0
p11:sp1:f1 ph2
d11
d4 groff
p20:gp27
d16 pl1:f1
p1 ph1
p22:gp15
d16
DELTA1
p1 ph1
;-----decoding period
p20:gp28
d16 pl0:f1
d4 gron0
p11:sp1:f1 ph2
d11

```

```

d4 groff
p20:gp29
d16
;-----region selection
p16:gp30
d16 pl1:f1
p12:sp2:f1 ph4
p16:gp31
d16
;-----pre-phasing
p25:gp25
d16
;-----acquisition
ACQ_START(ph30,ph31)
1u DWELL_GEN:f1
3 d20 gron2
d6 groff
d20 gron3
d6 groff
l0 to 3 times l3
rcyc=2
30m mc #0 to 2 F1QF(id2)
100u BLKGRAD
d17

;-----phase cycling
ph1=0
ph2=0
ph3=0
ph4=0
ph29=0
ph30=0
ph31=0

;pl1 : f1 channel - power level for pulse (default)
;sp1: shaped pulse power level for selective detection
;pl1 : f1 channel - 90 degree high power pulse
;p11: duration of the encoding chirp
;p20 :1ms coherence-selection delay
;p22 :20ms coherence-selection delay
;spnam1 : shaped pulse for spatial encoding
;d1: relaxation delay; 1-5 * T1
;d2: ramping period for gradient pulse (25us)
;d3: delay 10us
;d4: delay 25us
;d5: delay 5us
;d17: delay 10us
;d16: gradient recovery delay (200us)
;d30: diffusion time (big DELTA)
;d6: gradient ramp off during acquisition (25us)
;d20 + d6 : acquisition gradient duration
;GPZ0 : strength for excitation gradient
;GPZ2 and GPZ3 : strength for reversed acquisition gradient GPZ3 = -GPZ2
;GP20 and GP26 : coherence-selection gradients encoding (a)
;GP21 and GP27 : coherence-selection gradients encoding (a+c)
;GP22 and GP28 : coherence-selection gradients encoding (b)
;GP23 and GP29 : coherence-selection gradients encoding (b+c)
;GP30 and GP31 : coherence-selection gradients GP30 = GP31
;GP14 : spoiler Gradient Pulse
;GPZ15 : pre-phasing gradient GPZ15= -GPZ2/2
;GPNAM14= GPNAM15= GPNAM20= GPNAM21= GPNAM22= GPNAM23= GPNAM26= GPNAM27= SINE.100
;GPNAM28= GPNAM30= GPNAM31= SINE.100
;GPNAM25= RECT.1
;l3=number of loops for acquisition
;IMPORTANT: set d20 + d6 = DW x TD(f3)/(2xL3)

```

1. L. Guduff, I. Kuprov, C. van Heijenoort and J.-N. Dumez, *Chem. Commun.*, 2017, **53**, 701-704.