

Supporting Information:

Transferred-Rotational-Echo DOuble Resonance

(TREDOR)

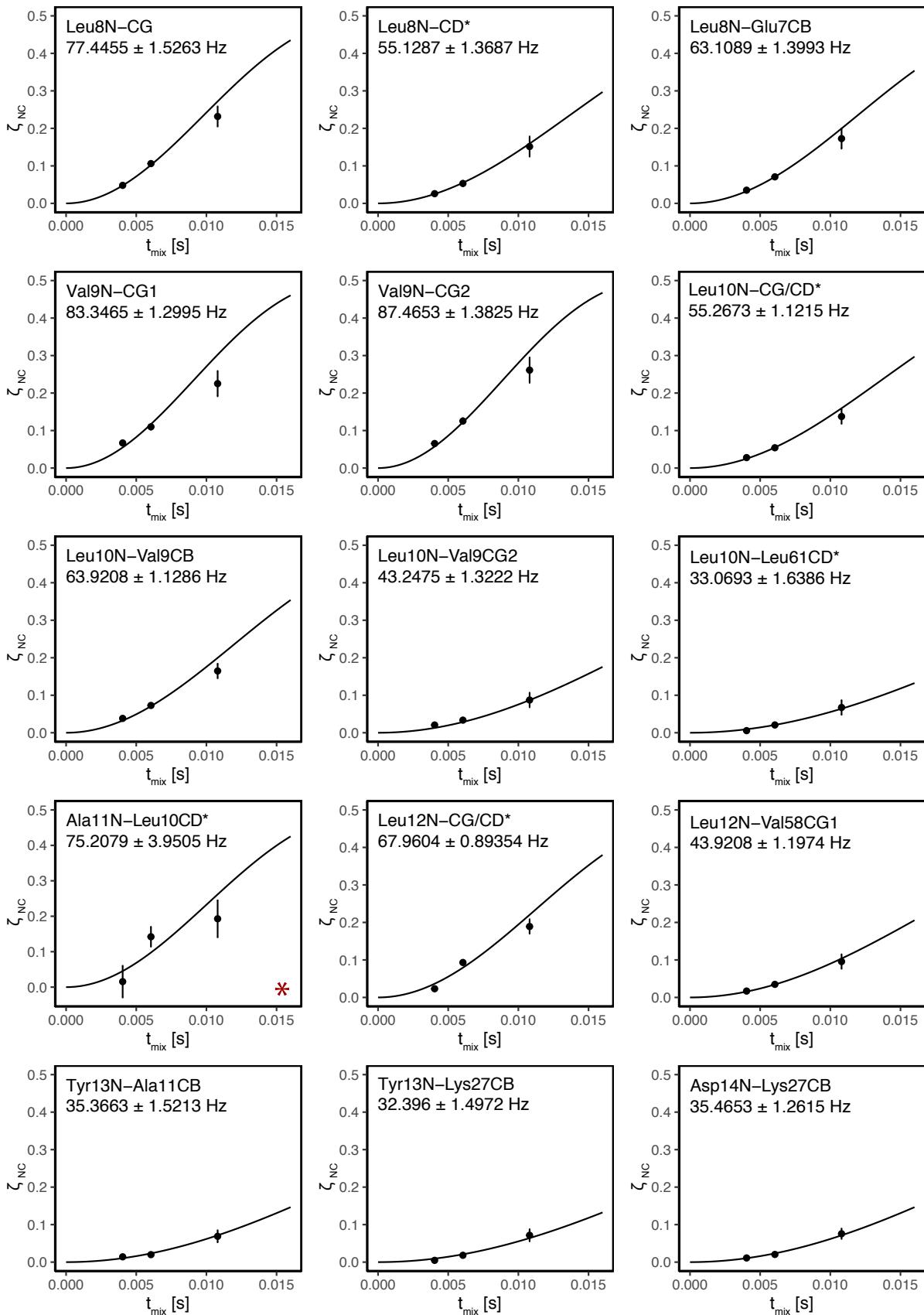
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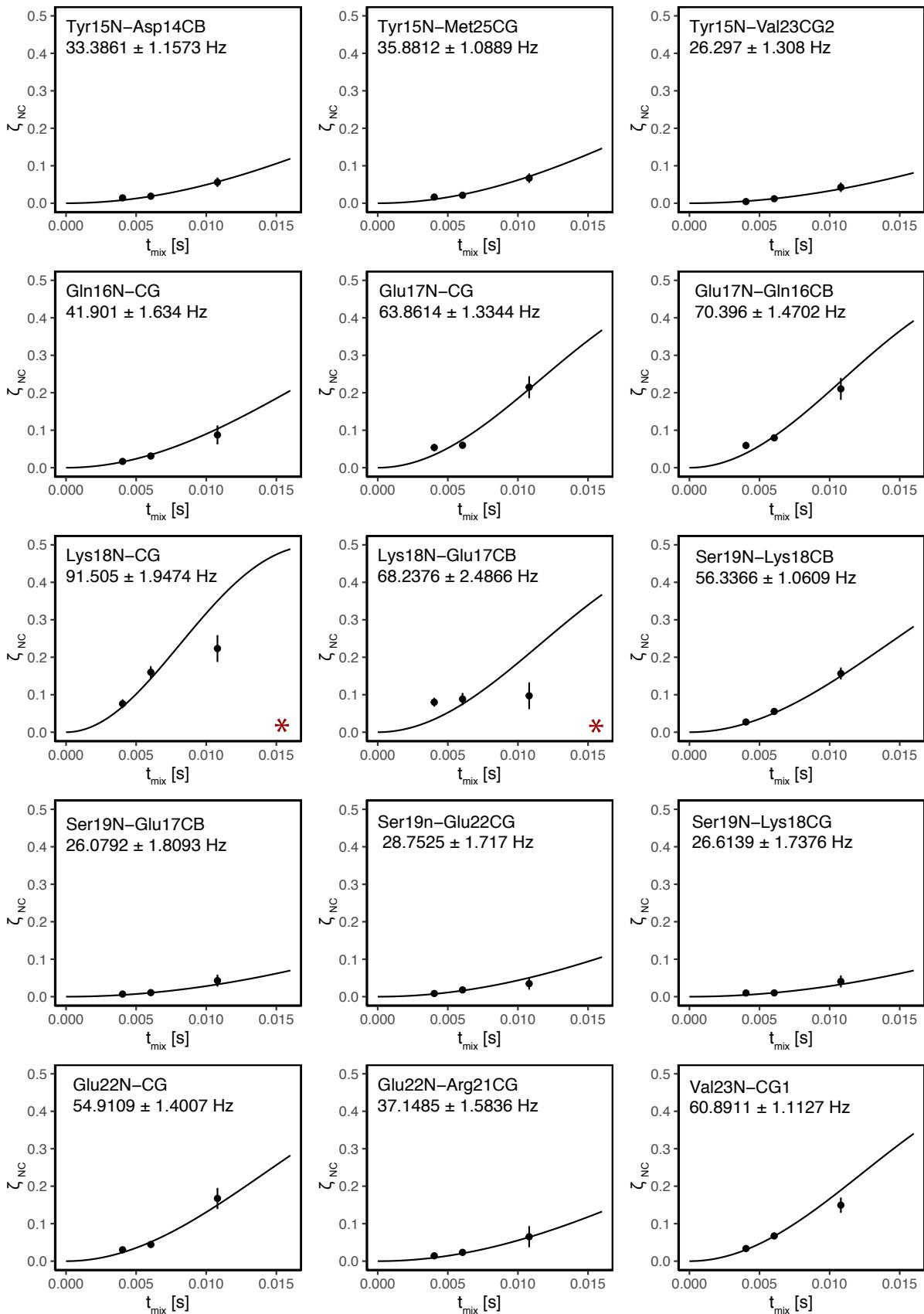
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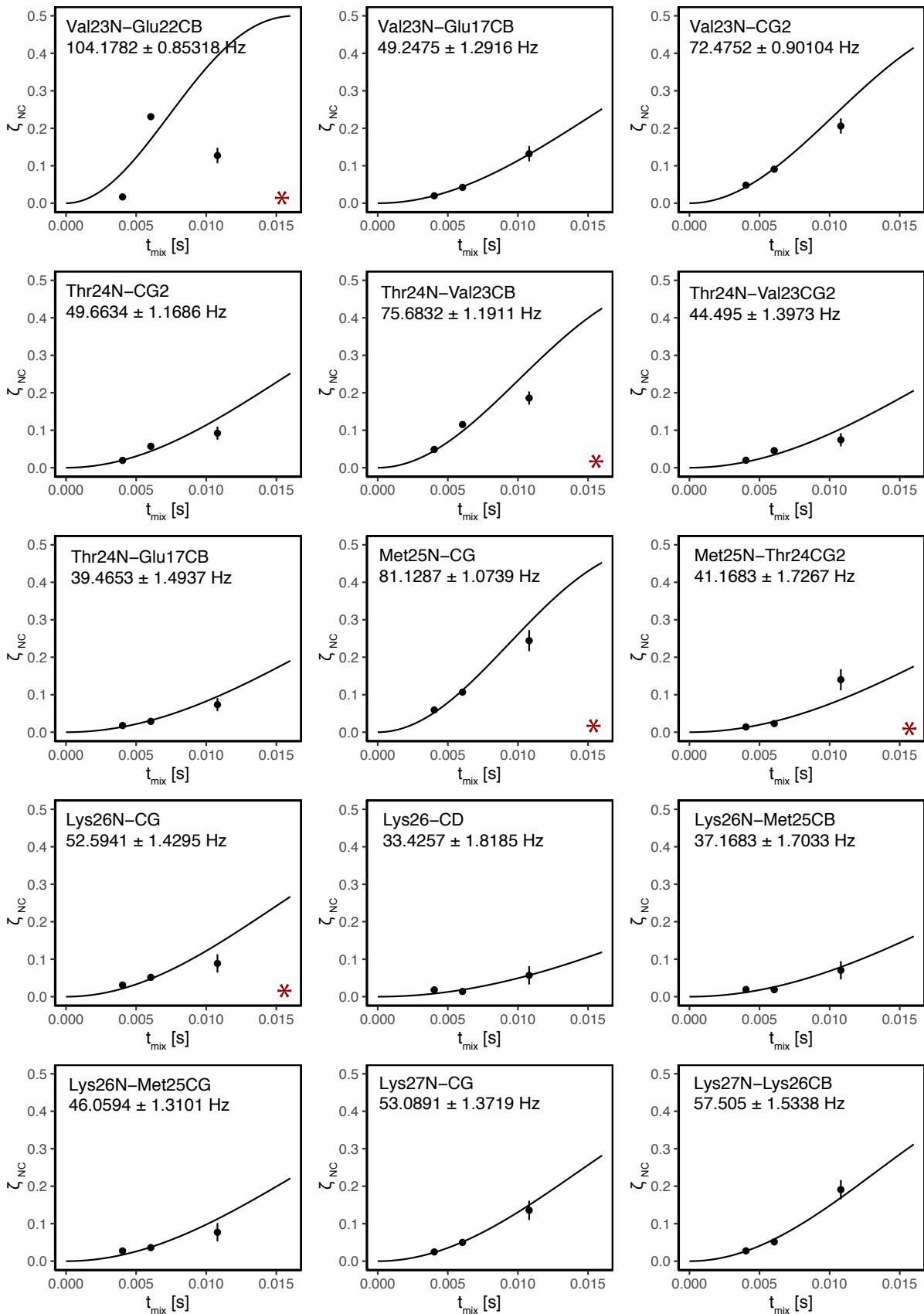
[‡]*Equal Contribution*

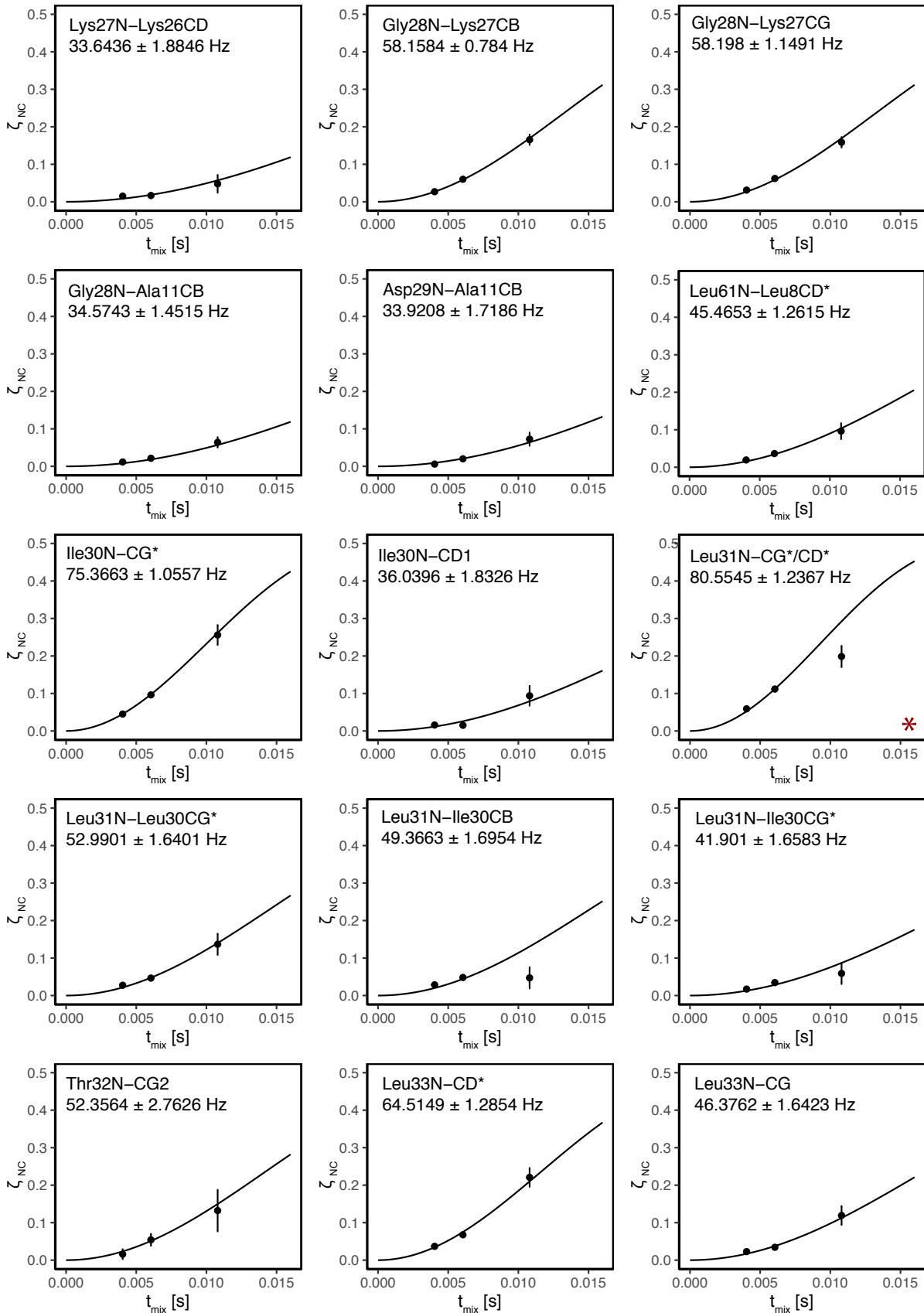
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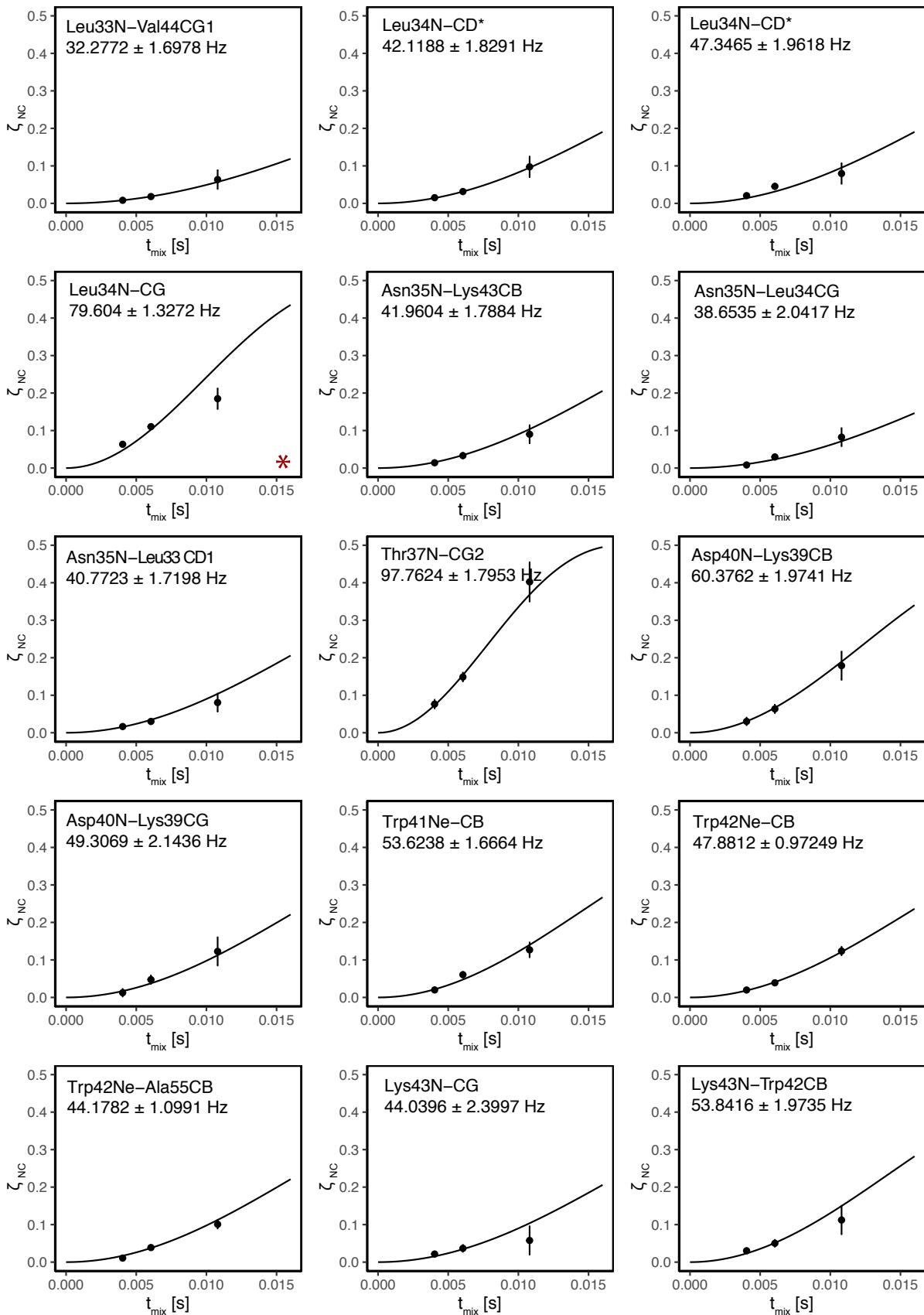
Figure S1. Fitting of all contacts observed in SH3 by pseudo-4D TREDOR

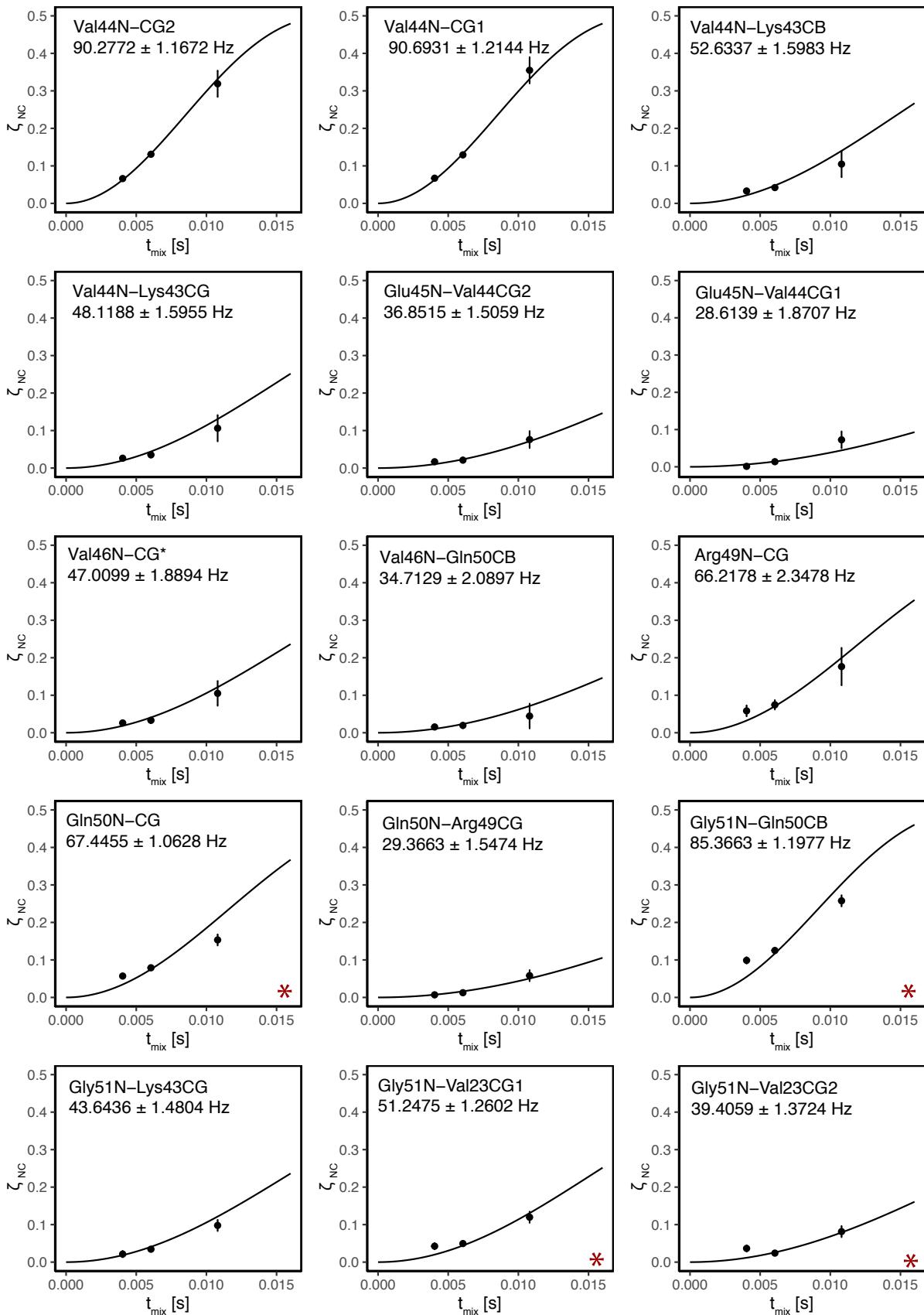


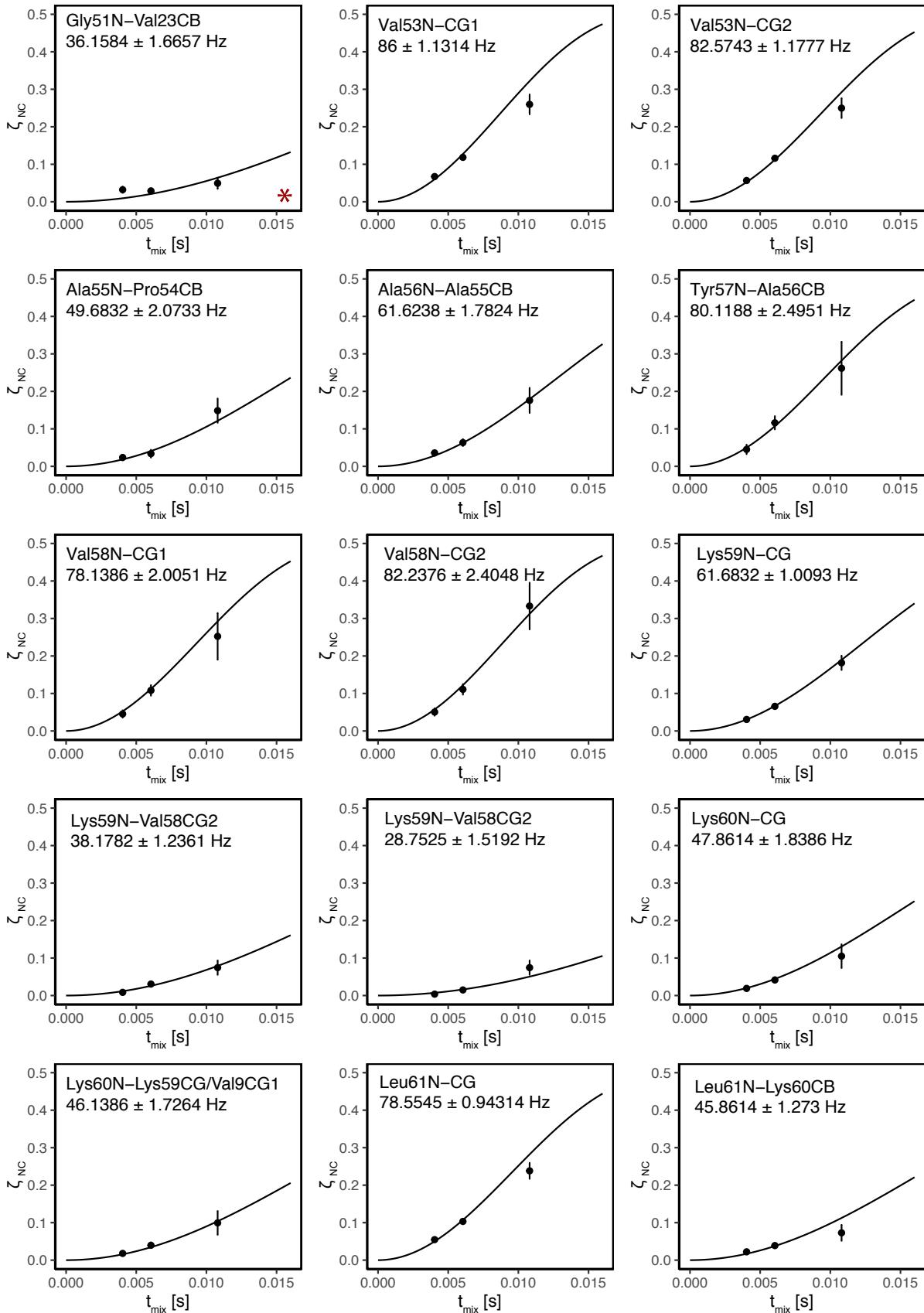


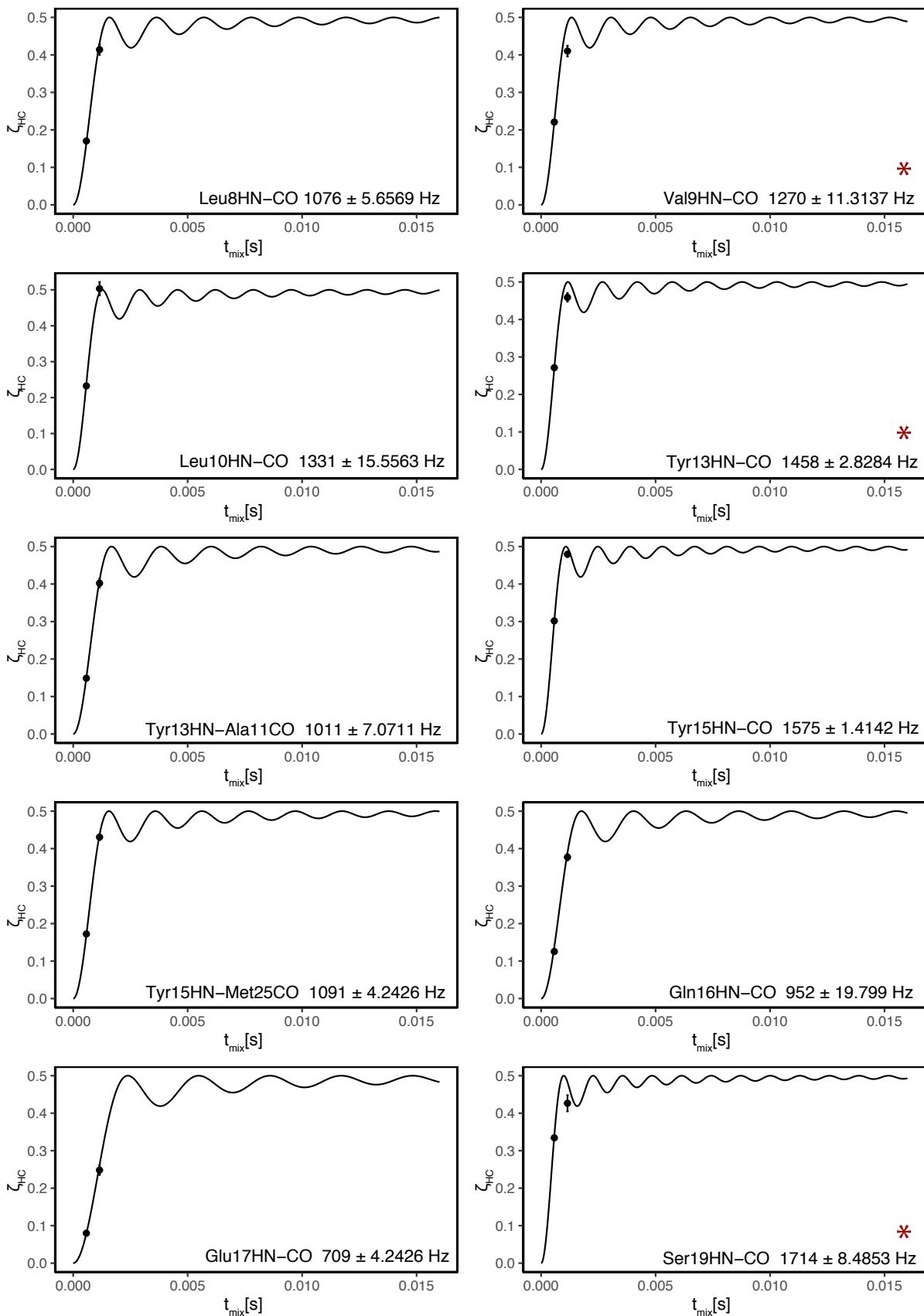


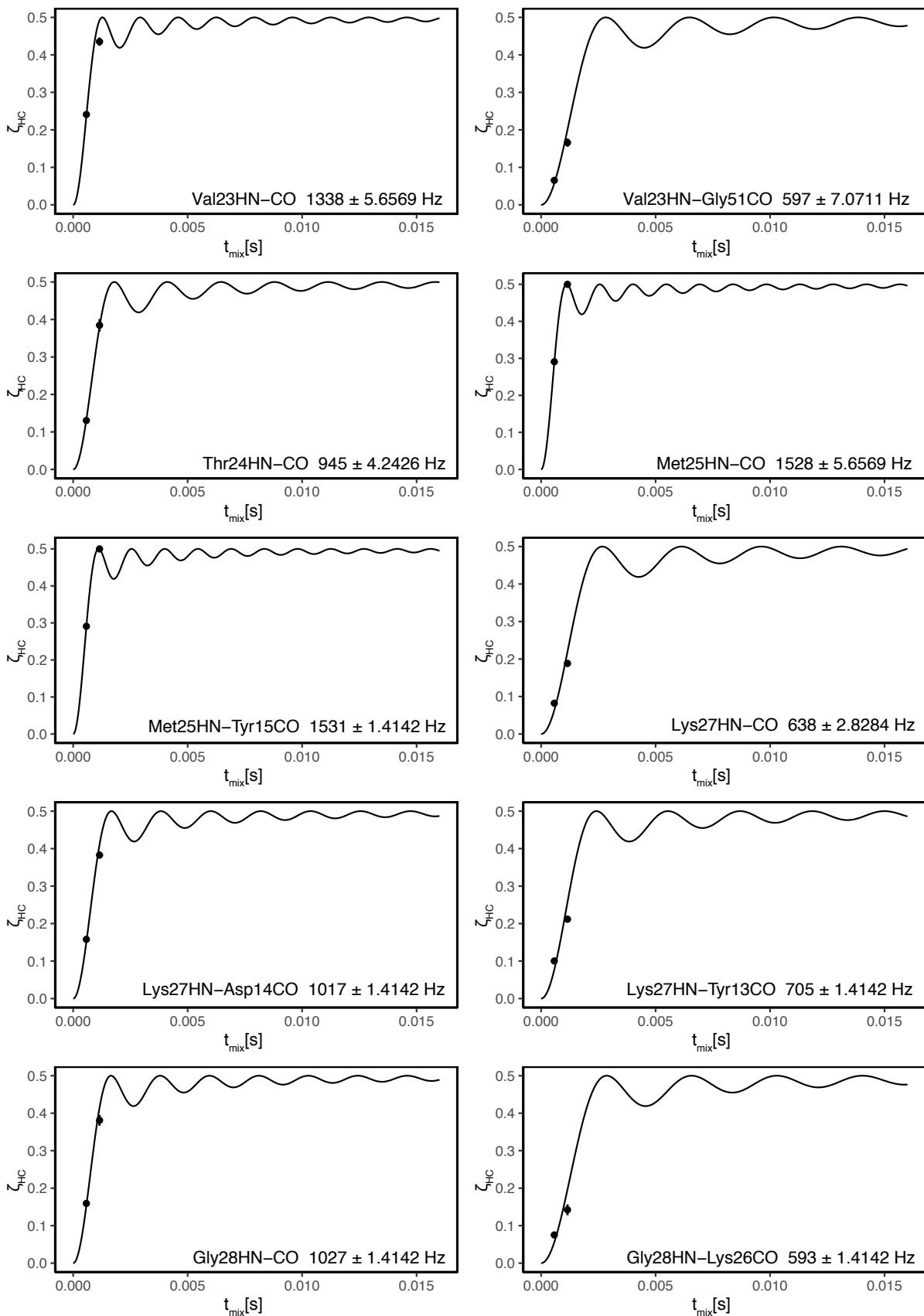


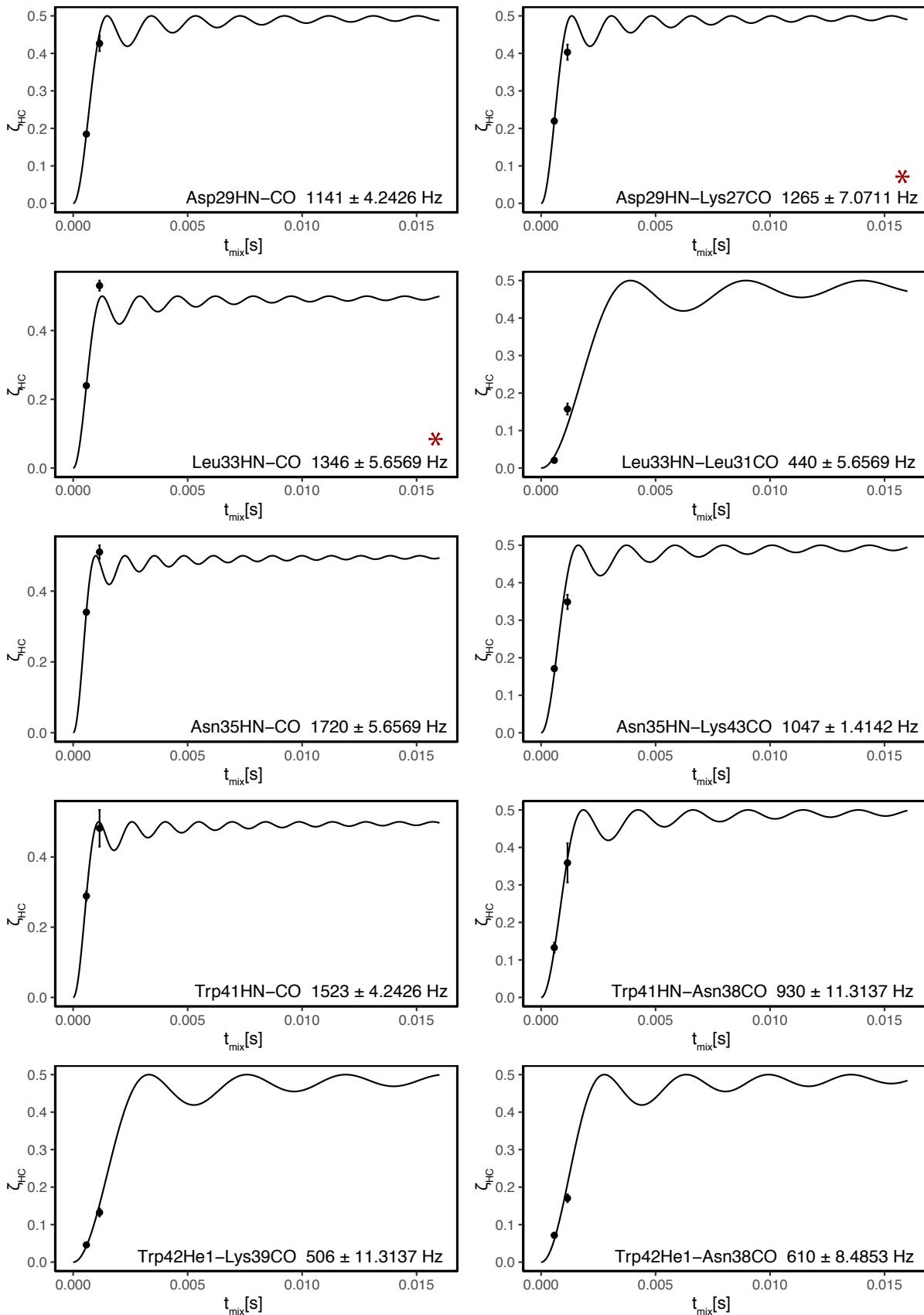


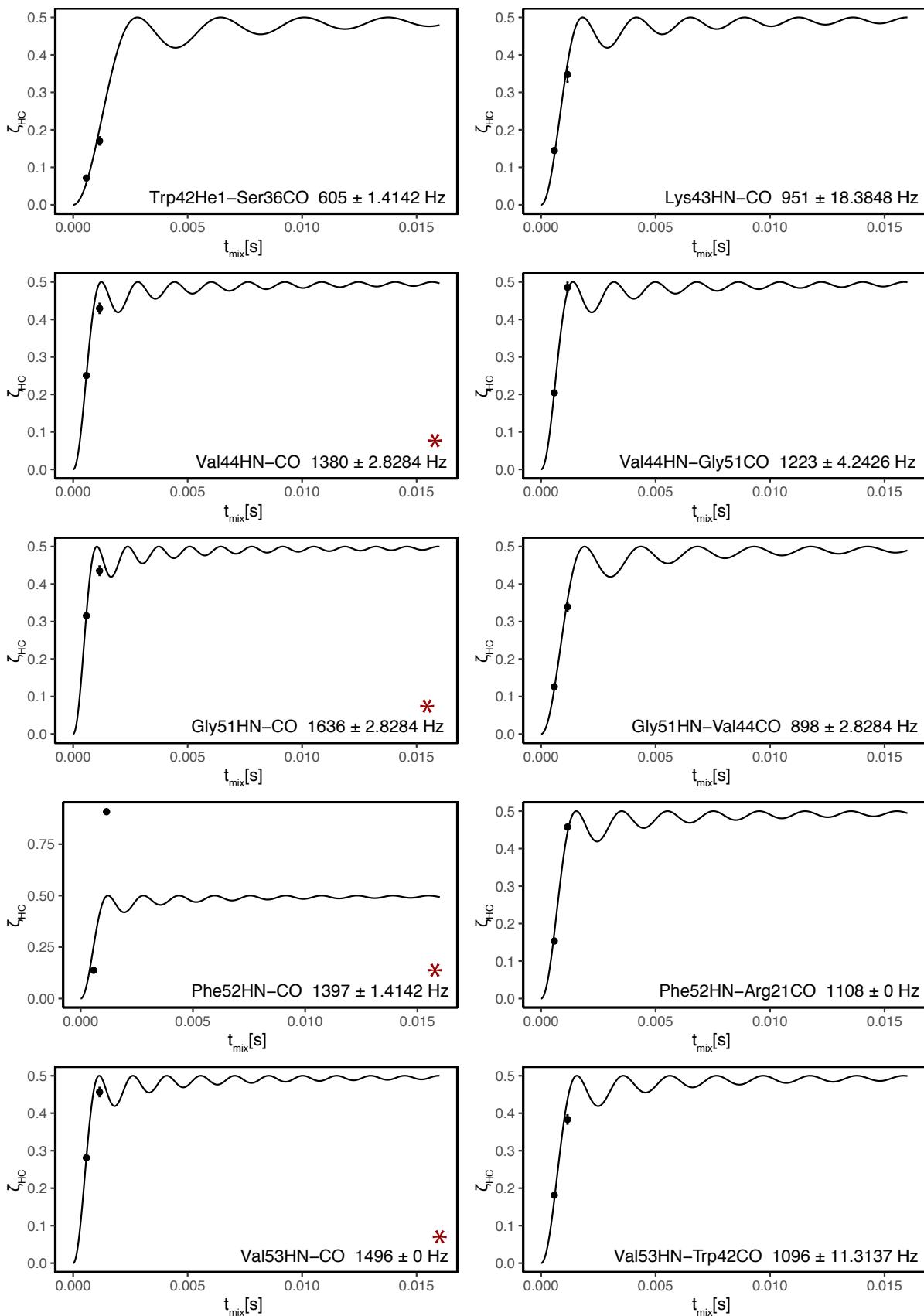


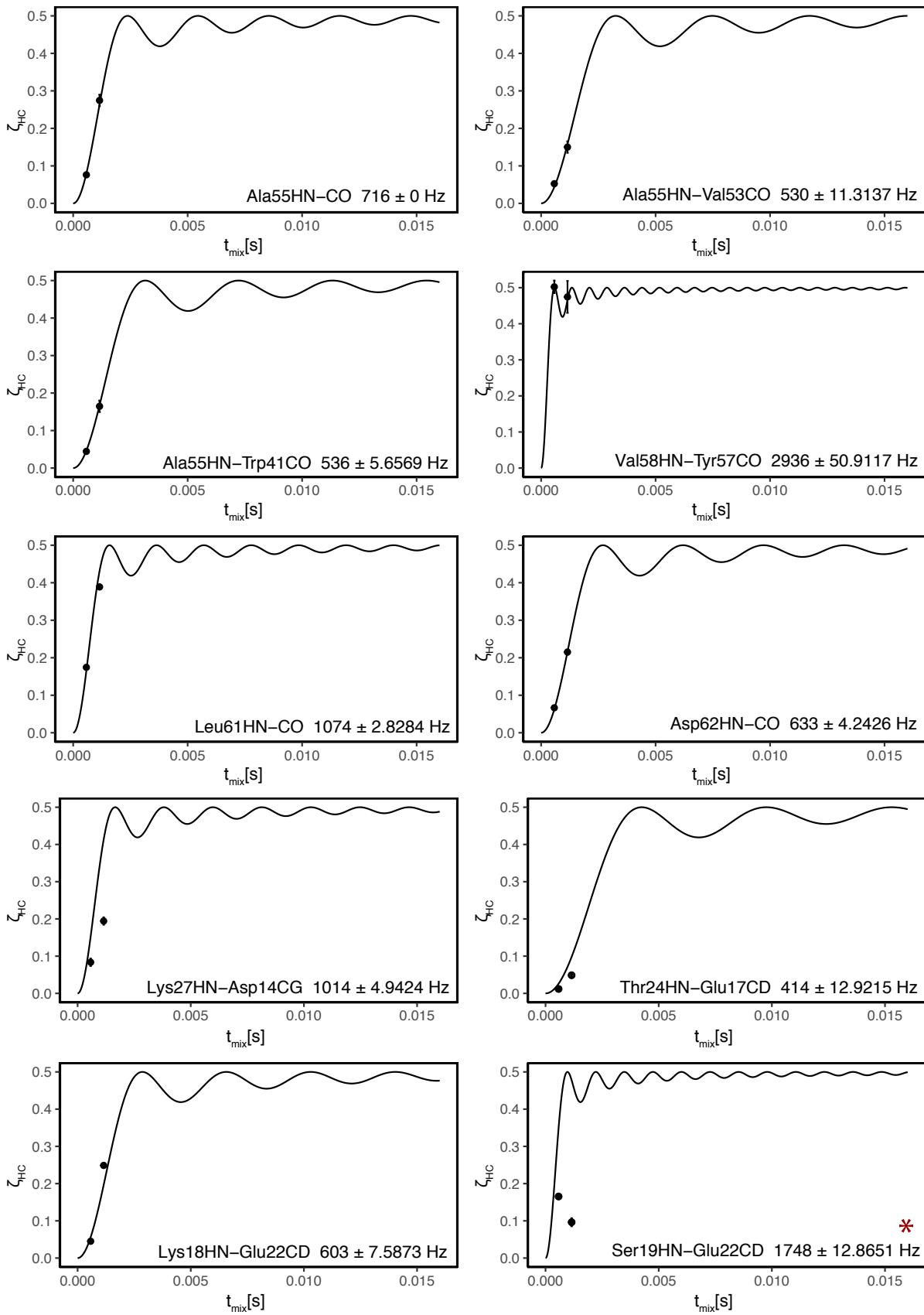


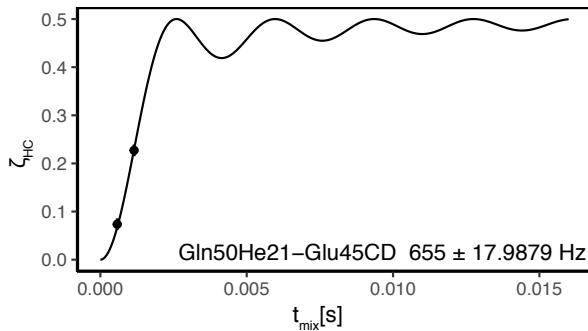
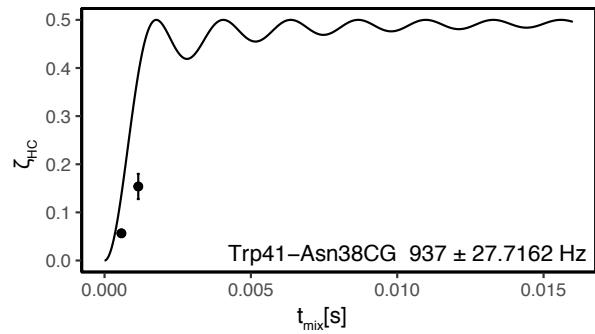
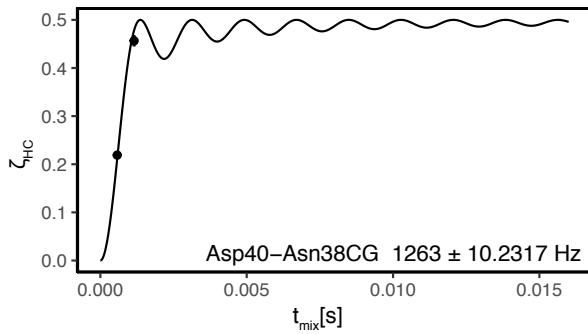












* indicates potentially compromised fit quality due to spectral overlap

Table S1. Relaxation rate Nitrogen T2 from hNH vs TREDOR relaxation rate

Residue	hNH t2 [ms]	TREDOR T2[ms]	Residue	hNH t2[ms]	TREDOR T2 [ms]
7GluN	19.55	3.87	34LeuN	64.94	4.34
8LeuN	68.45	4.51	35AsnN	56.15	4.02
9ValN	67.61	3.52	37ThrN	70.08	5.82
10LeuN	64.02	4.68	38AsnN	69.74	4.46
11Alan	103.95	0.91	39LysN	92.85	4.16
12LeuN	78.37	3.98	40AspN	49.85	5.12
13TyrN	84.67	4.93	41TrpN	64.31	3.27
14AspN	109.89	4.79	41TrpNe1	53.30	4.73
15TyrN	74.18	5.17	42TrpN	72.10	9.68
16GlnN	136.61	5.30	42TrpNe1	162.07	5.81
17GluN	88.34	5.72	43LysN	86.43	4.08
18LysN	83.61	4.42	44ValN	76.05	3.05
19SerN	65.40	5.72	45GluN	79.11	5.30
22GluN	74.35	3.57	49ArgN	52.88	6.50
23ValN	101.01	4.03	50GlnN	75.24	5.29
24ThrN	118.62	4.86	51GlyN	69.54	9.48
25MetN	79.74	3.90	52PheN	104.60	4.98
26LysN	108.70	4.21	53ValN	77.04	3.50
27LysN	106.27	3.73	55AlaN	93.37	4.64
28GlyN	70.32	6.40	56AlaN	73.42	4.73
29AspN	75.13	5.23	57TyrN	85.84	3.94
30IleN	91.49	3.59	58ValN	86.13	3.35
31LeuN	70.18	3.87	59LysN	60.28	3.14
32ThrN	90.33	4.92	60LysN	119.19	3.64
33LeuN	98.91	4.46	61LeuN	85.76	3.88

Table S2. Upper and lower distance restraints used in the structure calculation.
Ambiguous restraints are given in the CYANA format as putting them with 0 Å.

Contact		Upper limit [Å]	Lower limit [Å]
8 LEU N	8 LEU CG	3.44	2.82
8 LEU N	8 LEU CD1	3.91	3.2
8 LEU N	8 LEU CD2	0	0
8 LEU N	7 GLU CB	3.78	3.09
9 VAL N	9 VAL CG1	3.44	2.81
9 VAL N	9 VAL CG2	3.31	2.71
10 LEU N	10 LEU CG	3.92	3.21
10 LEU N	10 LEU CD1	0	0
10 LEU N	10 LEU CD2	0	0
10 LEU N	9 VAL CB	3.77	3.09
10 LEU N	9 VAL CG2	4.3	3.52
10 LEU N	61 LEU CD1	4.73	3.87
10 LEU N	61 LEU CD2	0	0
11 ALA N	10 LEU CD1	3.51	2.87
11 ALA N	10 LEU CD2	0	0
12 LEU N	12 LEU CG	3.6	2.94
12 LEU N	12 LEU CD1	0	0
12 LEU N	12 LEU CD2	0	0
12 LEU N	58 VAL CG1	4.28	3.51
13 TYR N	11 ALA CB	4.54	3.71
13 TYR N	27 LYS CB	4.77	3.9
14 ASP N	27 LYS CB	4.53	3.7
15 TYR N	14 ASP CB	4.77	3.24
15 TYR N	25 MET CG	4.45	3.64
15 TYR N	25 MET CE	5.05	4.01
16 GLN N	16 GLN CG	4.33	3.54
17 GLU N	17 GLU CG	3.76	3.08
17 GLU N	16 GLN CB	3.6	2.94
18 LYS N	18 LYS CG	3.3	2.7
18 LYS N	17 GLU CB	3.62	2.96
19 SER N	18 LYS CB	3.82	3.12
19 SER N	17 GLU CB	4.93	4.03
19 SER N	22 GLU CG	4.78	3.91
19 SER N	18 LYS CG	4.97	4.07
22 GLU N	22 GLU CG	3.91	3.2
22 GLU N	21 ARG CG	4.39	3.59
23 VAL N	23 VAL CG1	3.79	3.1
23 VAL N	22 GLU CB	3.18	2.6
23 VAL N	17 GLU CB	4.03	3.29
23 VAL N	23 VAL CG2	3.6	2.94
24 THR N	24 THR CG2	4.03	3.29
24 THR N	23 VAL CB	3.49	2.86
24 THR N	23 VAL CG2	4.23	3.46

24 THR N	17 GLU CB	4.34	3.55
25 MET N	25 MET CG	3.44	2.82
25 MET N	24 THR CG2	4.34	3.55
26 LYS N	26 LYS CG	4.02	3.29
26 LYS N	26 LYS CD	4.7	3.85
26 LYS N	25 MET CB	4.4	3.6
26 LYS N	25 MET CG	4.09	3.35
27 LYS N	27 LYS CG	4.02	3.29
27 LYS N	26 LYS CB	3.79	3.1
27 LYS N	26 LYS CD	4.69	3.83
28 GLY N	27 LYS CB	3.79	3.1
28 GLY N	27 LYS CG	3.79	3.1
28 GLY N	11 ALA CB	4.59	3.75
29 ASP N	11 ALA CB	4.7	3.85
30 ILE N	30 ILE CG1	3.49	2.86
30 ILE N	30 ILE CD1	4.5	3.68
31 LEU N	30 ILE CG1	4.03	3.29
31 LEU N	30 ILE CB	4.32	3.58
32 THR N	32 THR CG2	3.98	3.26
33 LEU N	33 LEU CD1	3.73	3.05
33 LEU N	33 LEU CD2	0	0
33 LEU N	33 LEU CG	4.07	3.33
33 LEU N	44 VAL CG1	4.75	3.88
34 LEU N	34 LEU CD1	4.31	3.52
34 LEU N	34 LEU CD2	0	0
34 LEU N	34 LEU CD1	4.06	3.32
34 LEU N	34 LEU CD2	0	0
34 LEU N	34 LEU CG	3.44	2.82
35 ASN N	43 LYS CB	4.32	3.53
35 ASN N	34 LEU CG	4.35	3.56
35 ASN N	33 LEU CD1	4.33	3.54
37 THR N	37 THR CG2	3.2	2.62
40 ASP N	39 LYS CB	3.78	3.1
40 ASP N	39 LYS CG	4.05	3.31
43 LYS N	43 LYS CG	4.2	3.44
43 LYS N	42 TRP CB	3.96	3.24
44 VAL N	44 VAL CG2	3.31	2.71
44 VAL N	44 VAL CG1	3.31	2.71
44 VAL N	43 LYS CB	4.02	3.29
44 VAL N	43 LYS CG	4.03	3.3
45 GLU N	44 VAL CG2	4.38	3.59
45 GLU N	44 VAL CG1	4.8	3.93
46 VAL N	46 VAL CG1	4.07	3.33
46 VAL N	50 GLN CB	4.56	3.73
49 ARG N	49 ARG CG	3.64	2.98
50 GLN N	50 GLN CG	3.6	2.95
50 GLN N	49 ARG CG	4.77	3.91

51 GLY N	50 GLN CB	3.36	2.75
51 GLY N	43 LYS CG	4.28	3.51
51 GLY N	23 VAL CG1	4.03	3.29
51 GLY N	23 VAL CG2	4.34	3.55
51 GLY N	23 VAL CB	4.46	3.65
53 VAL N	53 VAL CG1	3.33	2.73
53 VAL N	53 VAL CG2	3.44	2.81
55 ALA N	54 PRO CB	4.03	3.3
56 ALA N	55 ALA CB	3.79	3.1
57 TYR N	56 ALA CB	3.44	2.81
58 VAL N	58 VAL CG1	3.45	2.83
58 VAL N	58 VAL CG2	3.44	2.81
59 LYS N	59 LYS CG	3.79	3.1
59 LYS N	58 VAL CG2	4.34	3.55
59 LYS N	58 VAL CG1	4.78	3.91
60 LYS N	60 LYS CG	4.07	3.33
60 LYS N	59 LYS CG	4.11	3.36
60 LYS N	9 VAL CG1	4.11	3.36
61 LEU N	61 LEU CG	3.44	2.82
61 LEU N	60 LYS CB	4.09	3.34
61 LEU N	61 LEU CD1	4.16	3.4
61 LEU N	61 LEU CD2	0	0
41 TRP NE1	41 TRP CB	3.98	3.26
42 TRP NE1	42 TRP CB	4.03	3.29
42 TRP NE1	55 ALA CB	4.29	3.51
31 LEU N	31 LEU CD2	3.44	2.82
31 LEU N	31 LEU CG	4	3.27
31 LEU N	31 LEU CD1	0	0
31 LEU N	31 LEU CD2	0	0
8 LEU H	8 LEU C	3.1	2.54
9 VAL H	9 VAL C	2.95	2.41
10 LEU H	10 LEU C	2.89	2.37
13 TYR H	13 TYR C	2.81	2.3
13 TYR H	11 ALA C	3.17	2.59
15 TYR H	15 TYR C	2.73	2.23
15 TYR H	25 MET C	3.09	2.53
16 GLN H	16 GLN C	2.65	2.17
17 GLU H	17 GLU C	3.56	2.92
19 SER H	19 SER C	2.66	2.18
23 VAL H	23 VAL C	2.88	2.36
23 VAL H	51 GLY C	3.8	3.11
24 THR H	24 THR C	3.25	2.66
25 MET H	25 MET C	2.76	2.26
25 MET H	15 TYR C	2.76	2.26
27 LYS H	27 LYS C	3.69	3.02
27 LYS H	14 ASP C	3.17	3.17
27 LYS H	13 TYR C	3.56	2.92

28 GLY H	28 GLY C	3.15	2.57
28 GLY H	26 LYS C	3.78	3.1
29 ASP H	29 ASP C	3.04	2.48
29 ASP H	27 LYS C	2.95	2.41
33 LEU H	33 LEU C	2.88	2.36
33 LEU H	31 LEU C	4.2	3.44
35 ASN H	35 ASN C	2.65	2.17
35 ASN H	43 LYS C	3.14	2.57
41 TRP H	41 TRP C	2.75	2.25
41 TRP H	38 ASN C	3.25	2.66
42 TRP H	53 VAL C	3.05	2.49
42 TRP HE1	39 LYS C	4	3.28
42 TRP HE1	38 ASN C	3.76	3.08
42 TRP HE1	36 SER C	3.76	3.08
43 LYS H	43 LYS C	3.22	2.64
44 VAL H	44 VAL C	2.86	2.34
44 VAL H	51 GLY C	2.98	2.44
51 GLY H	51 GLY C	2.7	2.21
51 GLY H	44 VAL C	3.3	2.7
52 PHE H	52 PHE C	2.84	2.32
52 PHE H	21 ARG C	3.07	2.51
53 VAL H	53 VAL C	2.78	2.28
53 VAL H	42 TRP C	3.08	2.52
55 ALA H	55 ALA C	3.54	2.9
55 ALA H	53 VAL C	3.89	3.19
55 ALA H	41 TRP C	3.93	3.21
58 VAL H	58 VAL C	2.89	2.37
61 LEU H	61 LEU C	3.1	2.54
62 ASP H	62 ASP C	3.7	3.02
27 LYS H	14 ASP CG	3.17	2.59
24 THR H	17 GLU CD	4.27	3.49
18 LYS H	22 GLU CD	3.77	3.08
19 SER H	22 GLU CD	2.64	2.16
40 ASP H	38 ASN CG	2.94	2.41
41 TRP H	38 ASN CG	3.25	2.66
50 GLN HE21	45 GLU CD	3.45	2.82

Table S3. TALOS-N angle restraints.

Residue		Angle	Angle margin	
8	LEU	PHI	-155.7	-108
8	LEU	PSI	135.2	175.2
9	VAL	PHI	-148.2	-108.2
9	VAL	PSI	131.4	171.4
10	LEU	PHI	-145.1	-105.1
10	LEU	PSI	116.5	156.5
11	ALA	PHI	-94.3	-52.8
11	ALA	PSI	106.7	146.7
12	LEU	PHI	-103.9	-63.1
12	LEU	PSI	-60.8	-19.7
13	TYR	PHI	-182.5	-131.4
13	TYR	PSI	138	178
16	GLN	PHI	-149.5	-86.9
16	GLN	PSI	109.4	169.9
23	VAL	PHI	-175.6	-72.5
23	VAL	PSI	129.5	177.3
24	THR	PHI	-132.1	-83.3
24	THR	PSI	111.5	151.5
25	MET	PHI	-151.5	-89.7
25	MET	PSI	113.5	171.6
26	LYS	PHI	-118	-56.7
26	LYS	PSI	130.5	170.5
27	LYS	PHI	-76.5	-36.5
27	LYS	PSI	113.2	153.2
29	ASP	PHI	-84.6	-44.6
29	ASP	PSI	126.3	166.3
30	ILE	PHI	-129.2	-89.2
30	ILE	PSI	104.5	144.5
31	LEU	PHI	-130	-89.1
31	LEU	PSI	103.8	158.4
32	THR	PHI	-97.3	-52.6
32	THR	PSI	107.9	147.9
33	LEU	PHI	-117.8	-50.3
33	LEU	PSI	88.7	168.2
34	LEU	PHI	-79.4	-39.4
34	LEU	PSI	-51.9	-11.9
36	SER	PHI	-88.8	-48.8
36	SER	PSI	-55.9	-15.9
37	THR	PHI	-85.9	-45.9
37	THR	PSI	-59.5	-19.5
38	ASN	PHI	-81.9	-41.9
39	LYS	PSI	-38.9	1.1
40	ASP	PHI	-105.7	-65.7
40	ASP	PSI	-29.6	10.4
41	TRP	PHI	-123.6	-67.1
41	TRP	PSI	112.2	159.1

42	TRP	PHI	-144.3	-104.3
42	TRP	PSI	114.3	175.6
43	LYS	PHI	-138.7	-62.6
43	LYS	PSI	103.5	151
44	VAL	PHI	-146.1	-94.4
44	VAL	PSI	112.9	178.5
50	GLN	PHI	-162.6	-88.7
50	GLN	PSI	132.5	172.5
51	GLY	PHI	153.3	193.3
51	GLY	PSI	-192.9	-152.9
52	PHE	PHI	-119.7	-67
52	PHE	PSI	116.3	163.4
53	VAL	PHI	-157.6	-104
53	VAL	PSI	124.7	175.2
54	PRO	PSI	108.5	148.5
55	ALA	PHI	-82.5	-42.5
55	ALA	PSI	-51.7	-11.7
56	ALA	PHI	-87.5	-47.5
56	ALA	PSI	-39.1	3.8
59	LYS	PHI	-149.6	-94.6
59	LYS	PSI	107.6	172.9
60	LYS	PHI	-99.5	-50.9
60	LYS	PSI	108.3	148.3
61	LEU	PHI	-104	-50.2
61	LEU	PSI	115.2	155.2

- Simultaneous multi-curve TREDOR fitting starting with Equation 1. This result can be extended¹ to a set of n_i equations, which is used to simultaneously fit the buildup curves of the individual carbon spins coupled to N₁.

$$n_i \left\{ \begin{array}{l} V_{11}(t_{mix}) = \Lambda_1(1 - [J_0(\sqrt{2}D_{11}t_{mix})]^2) \prod_{k=2}^{n_i} (1 + [J_0\sqrt{2}D_{1k}t_{mix}]^2) \\ V_{12}(t_{mix}) = \Lambda_1(1 - [J_0(\sqrt{2}D_{12}t_{mix})]^2) \prod_{k=1 \neq 2}^{n_i} (1 + [J_0\sqrt{2}D_{1k}t_{mix}]^2) \\ \dots \\ V_{1i}(t_{mix}) = \Lambda_1(1 - [J_0(\sqrt{2}D_{1i}t_{mix})]^2) \prod_{k=1 \neq i}^{n_i} (1 + [J_0\sqrt{2}D_{1k}t_{mix}]^2) \end{array} \right. \quad (1)$$

Where Λ_1 is given as:

$$\Lambda_1 = \frac{1}{2^{n_i}} V_1(0) \lambda_1 \exp(-\Gamma_1 t_{mix}) \quad (2)$$

The Γ_1 gives the coherence decay rate of spin coherence, which can be modeled as a single exponential. λ_1 is an amplitude scaling factor, which is typically fit freely since the starting signal is not recorded in a TEDOR measurement. So in a standard TEDOR fitting, to extract the dipolar couplings, one needs to fit the overall decay rate and an arbitrary scaling factor which are irrelevant to the parameter of concern. This consequently puts more demand on the data density, something that might be unachievable especially for samples with low sensitivity or short coherence times. In the pseudo-4D TREDOR experiment, we address this problem by recording the intensity of the untransferred part of the magnetization, so that the ratio $\zeta_{1i}(t_{mix})$ can be formed between transferred and total magnetization

$$\zeta_{1i}(t_{mix}) = \frac{V_{1i}(t_{mix})}{V_1(t_{mix}) + \sum_{k=1}^{n_i} V_{1k}(t_{mix})} \quad (3)$$

$$= \frac{V_{1i}(t_{mix})}{V_{total}(t_{mix})} \quad (4)$$

Where $V_1(t_{mix})$ is the intensity of the untransferred nitrogen signal, and the $V_{total}(t_{mix})$ is the total signal at a certain mixing time. The untransferred signal $V_1(t_{mix})$ appears

with the same scaling and typically also the same decay parameter, and hence the total signal also follows a single exponential decay of the original signal intensity.

$$V_{total}(t_{mix}) = V_1(0) \exp(-\Gamma_1 t_{mix}) \quad (5)$$

Normalization by the total signal results in a coherence decay-free TEDOR curve, for which the parameters λ_1 and $V_1(0)$ no longer appear. Combining Equation 1 and Equation 3, it is straightforward to show that we can fit the simultaneous buildup curve of several ^{13}C spins coupled to the same N_1 as the following set of equations:

$$\left. \begin{array}{l} \zeta_{11}(t_{mix}) = \frac{1}{2^{n_i}} (1 - [J_0(\sqrt{2}D_{11}t_{mix})]^2) \prod_{k=2}^{n_i} (1 + [J_0\sqrt{2}D_{1k}t_{mix}]^2) \\ \zeta_{12}(t_{mix}) = \frac{1}{2^{n_i}} (1 - [J_0(\sqrt{2}D_{12}t_{mix})]^2) \prod_{k=1 \neq 2}^{n_i} (1 + [J_0\sqrt{2}D_{1k}t_{mix}]^2) \\ \dots \\ \zeta_{1i}(t_{mix}) = \frac{1}{2^{n_i}} (1 - [J_0(\sqrt{2}D_{1i}t_{mix})]^2) \prod_{k=1 \neq i}^{n_i} (1 + [J_0\sqrt{2}D_{1k}t_{mix}]^2) \end{array} \right\} \quad (6)$$

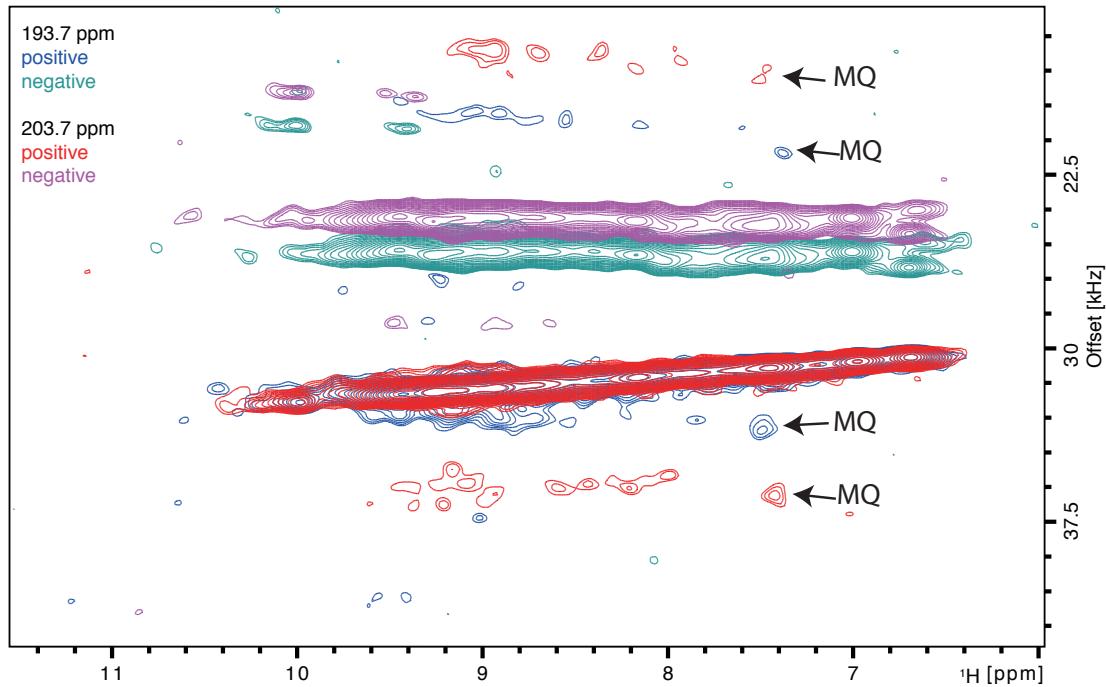


Figure S2. Comparison of positions of multiple quantum artifacts in pseudo-3D H-CO TREDOR. With a shift of the offset from 193.7 ppm to 203.7 ppm, the artifacts changed chemical shifts on the H-CO dimension each by 20 ppm, confirming the major artifacts left are only the leftover double quantum terms but not zero quantum terms. When the offset is placed at 203.7 ppm, the artifacts are placed away from the signal of interest enough for quantitative analysis.

Table S3. CO T2 rho in SH3 in ms with different hard pulse power in kHz and offsets in ppm

Power	Offset	53 ppm	100 ppm
85 kHz		11.1 ms	64.7 ms
100 kHz		10.9 ms	60.4 ms

Table S4. CA T2 rho in SH3 in ms with different hard pulse power in kHz and offsets in ppm

Power	Offset	53 ppm	100 ppm
85 kHz		41.0 ms	41.1 ms
100 kHz		47.5 ms	48.5 ms

References

- (1) Jaroniec, C. P.; Filip, C.; Griffin, R. G. 3D TEDOR NMR experiments for the simultaneous measurement of multiple carbon-nitrogen distances in uniformly ¹³C,¹⁵N-labeled solids. *J. Am. Chem. Soc.* **2002**, *124*, 10728–10742.