

Supplementary Information

Biomimetic non-heme iron-catalyzed epoxidation of challenging terminal alkenes using aqueous H₂O₂ as an environmentally friendly oxidant

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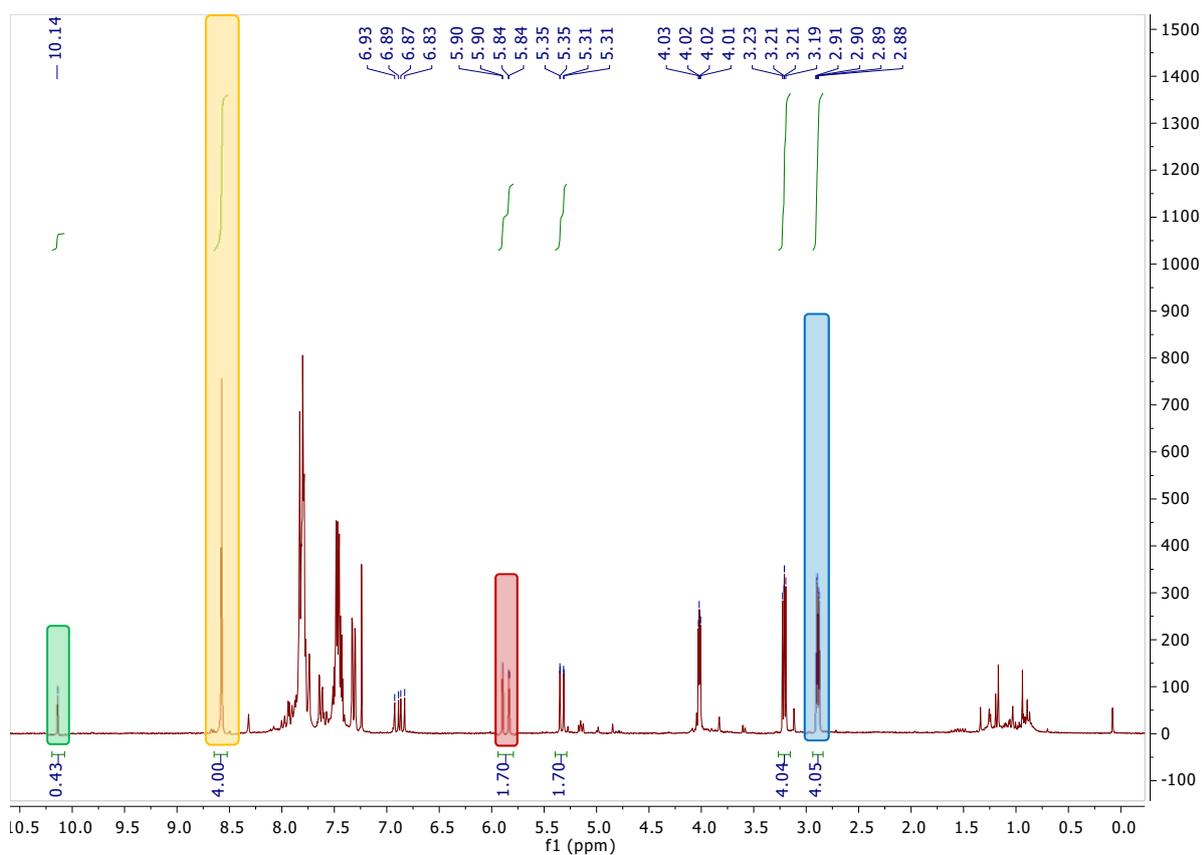
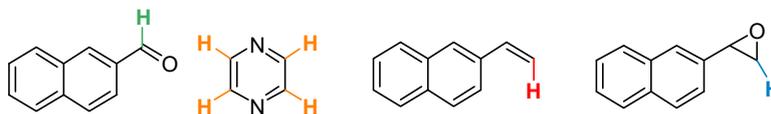
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1 Determination of yield via ^1H NMR using pyrazine as internal standard

Exemplary: Table 2, Entry 1:

^1H -NMR of reaction mixture after SiO_2 -plug containing pyrazine as internal standard (m = 4.32 mg, n = 53.9 μmol)



1H (pyrazine) \cong 53.9 μmol

1H (epoxide) \cong 53.9 $\mu\text{mol} \times 4.05 = 218 \mu\text{mol}$ (n (epoxide))

\rightarrow yield = n (epoxide) / n (alkene) = 218 μmol / 500 μmol = 0.44 \rightarrow 44%

2 Screening of reaction conditions for non-enantioselective epoxidation

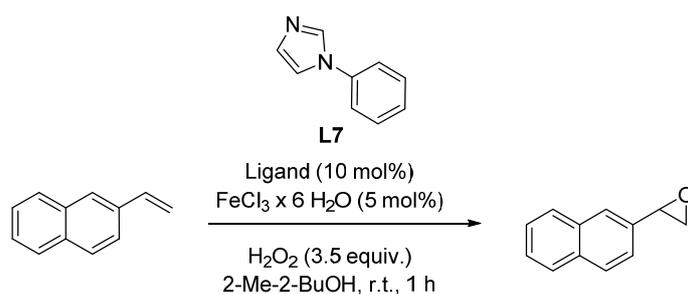


Table 1. Screening of reaction conditions.

Entry	Modification of reaction conditions	yield [%] ^{a)}
1	-	53
2	1 h <i>in situ</i> catalyst generation (under inert conditions)	traces
3	1 h <i>in situ</i> catalyst generation (not under inert conditions)	54
4	1 h <i>in situ</i> catalyst generation and 5 mol% $\text{CH}_3\text{CO}_2\text{H}$	44
5	2 h reaction time and 2 h H_2O_2 addition.	45
6	15°C	22
7	0.05 M substrate	42
8	0.2 M substrate	25
9	addition of H_2O_2 over 3 h period	52
10	10 mol% catalyst loading	55
11	0°C	13
12	50°C	27
13	1 h H_2O_2 addition plus 15 h stirring	50
14	1 h H_2O_2 addition + 1 h stirring	53
15	+ 5 mol% H_2Pydic	19
16	+ 5 mol% mandelic acid	26
17	irradiation with cold light	52

a) Determined via pyrazine as internal standard.

3 Screening of reaction conditions for enantioselective epoxidation

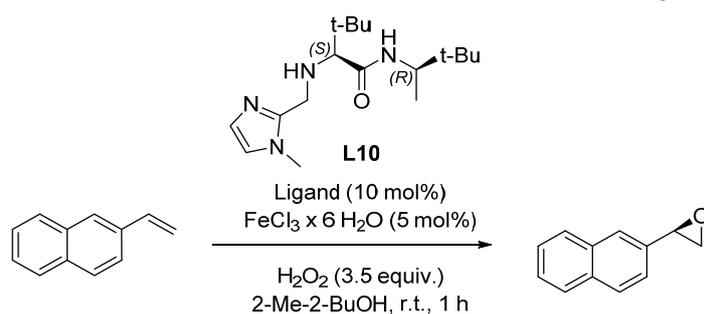


Table 2. Screening of reaction conditions.

Entry	Modification of reaction conditions	yield [%] ^{a)}	ee-value (<i>R</i>) [%]
1	-	44	26
2	1 h <i>in situ</i> catalyst generation with air-bubbling	42	24
3	at 0°C	23	36
4	+ 5 mol% mandelic acid	32	27
5	L10 (5 mol%)	31	26

a) Determined via pyrazine as internal standard.

4 Rearrangement of Epoxide

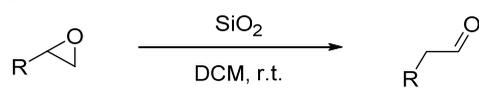


Table 3. First rearrangement attempts from epoxide towards corresponding aldehyde using SiO_2 .

Entry	Substrate	Time	Temperature	Yield [%] ^{c)}
1 ^{a), b)}		5 h	reflux	17
2		18 h	reflux	34
3 ^{a)}		18 h	r.t.	29
4 ^{a)}		2.5 h	reflux	32

a) 3.33 g/mmol SiO_2 ; b) CDCl_3 ; c) Determined via pyrazine as internal standard.

5 Comparative Table

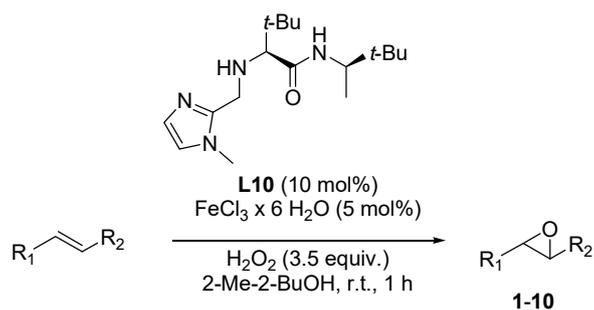


Table 4. Results obtained in this work contrasted with the results described in literature for non-heme iron-catalyzed epoxidation of the corresponding olefins.

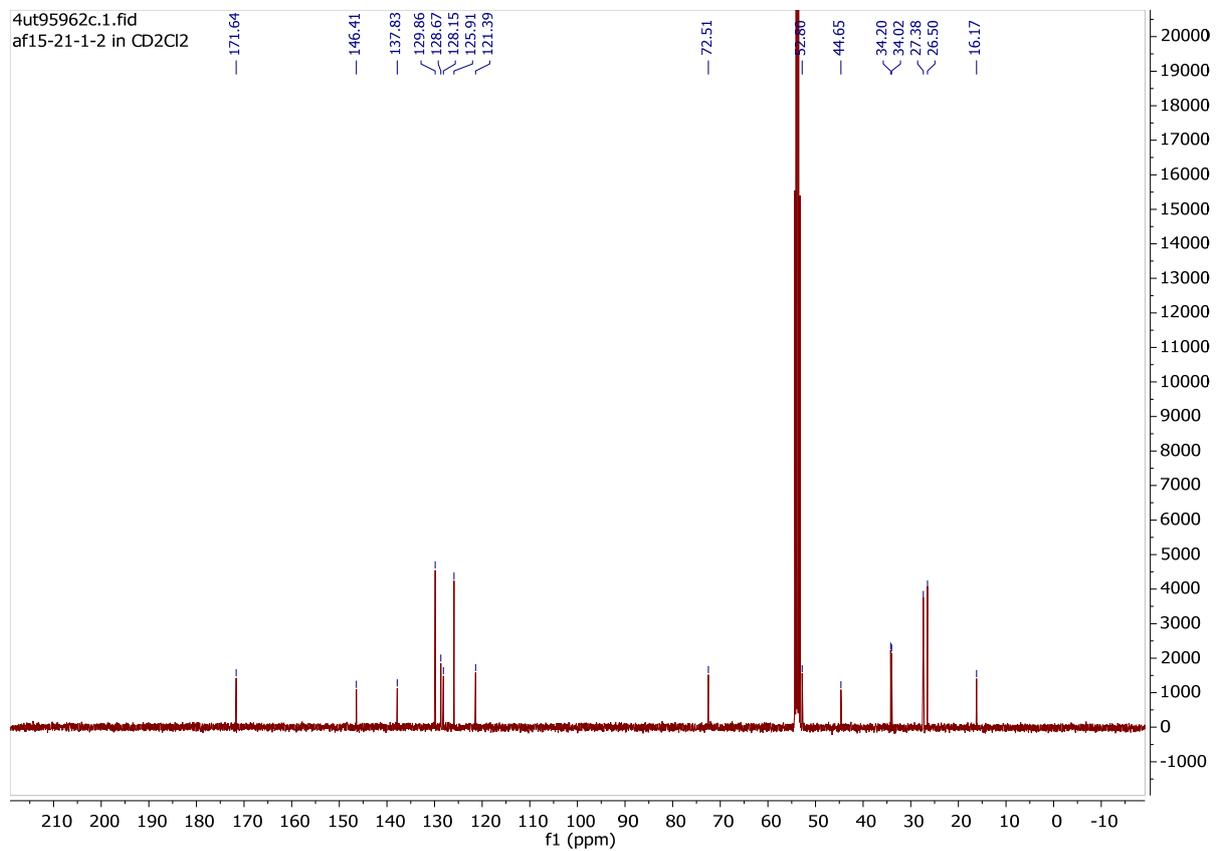
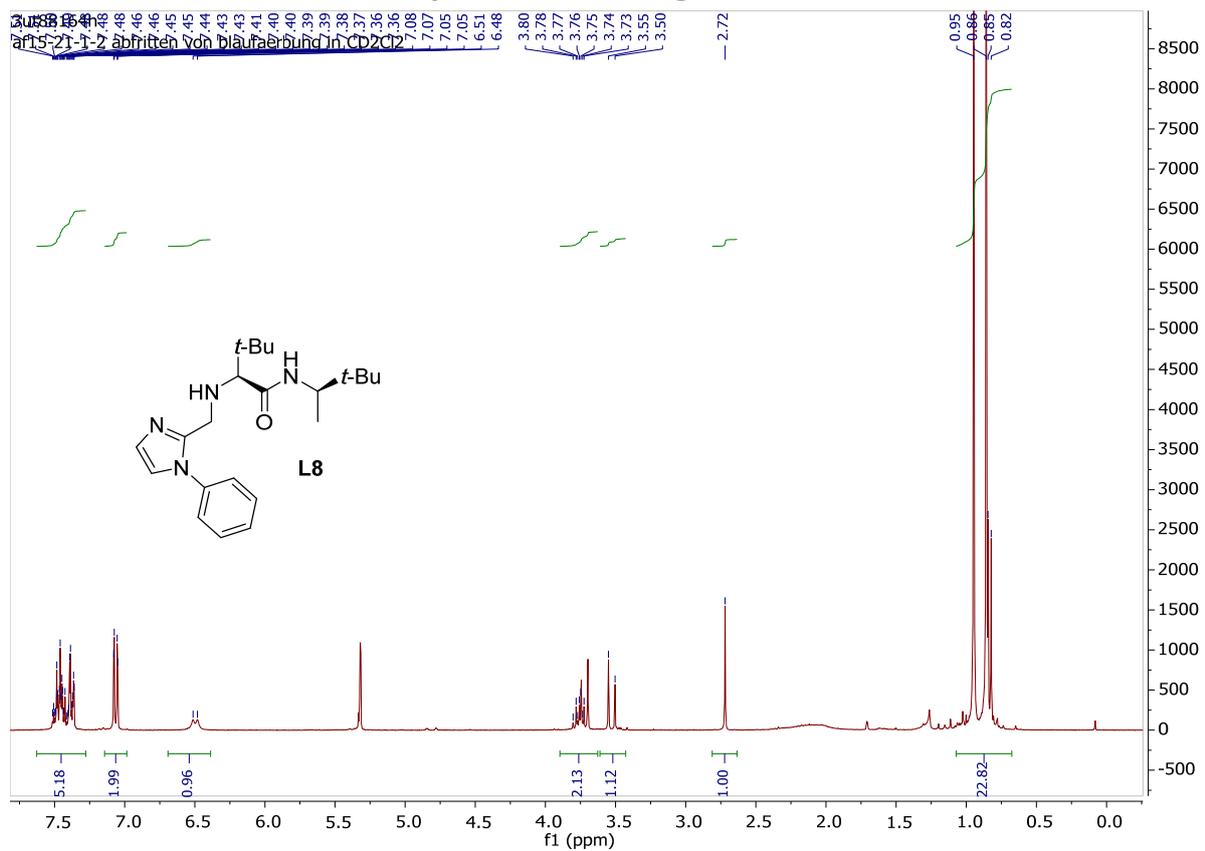
Entry	Product	Results of this work		Results described in literature		
		Yield ^{a),b)}	ee ^{c)}	Yield ^{d)}	ee ^{d)}	Ref.
1		44	26 (<i>R</i>)	40	–	[1]
2		30	27 (<i>R</i>)	84	8	[2]
				30	–	[3]
3		14	29 (<i>S</i>)	15	–	[4]
				66	–	[5]
4		25	25 (<i>R</i>) ^{e)}	90	–	[6]
				–	–	–
5		5 ^{e)}	14 (<i>R</i>) ^{e)}	82	–	[7]
				70	–	[8]
6		22	16 (<i>S</i>)	67	45	[9]
7		<5 ^{f)}	n.d. ^{g)}	60	–	[10]
8		32	50	32	19	[9]
				49	–	[11]
9		27	16 (<i>R</i>)	44	–	[3]
				72	–	[8]
10		<5	27 (2 <i>R</i> ,3 <i>S</i>)	38	90	[9]
11 ^{h)}		7	51 (2 <i>R</i> ,3 <i>S</i>)	–	–	–

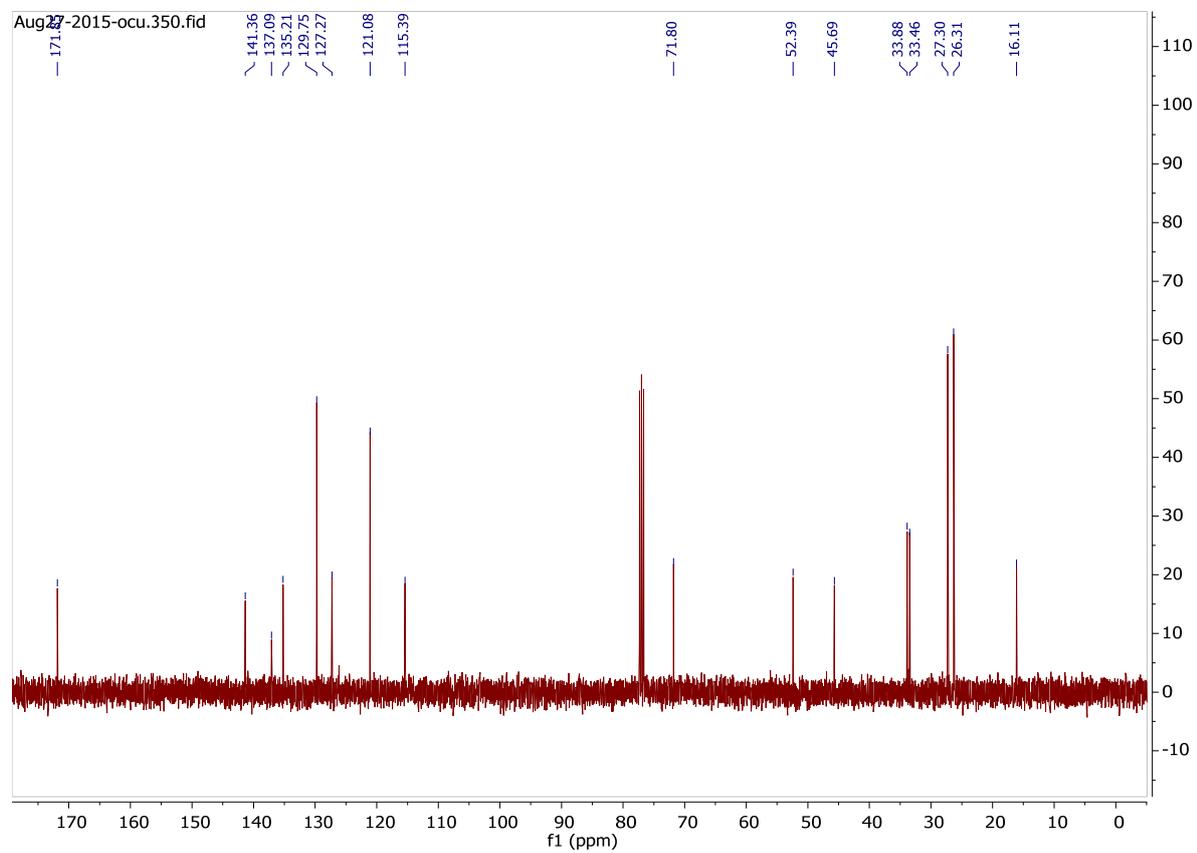
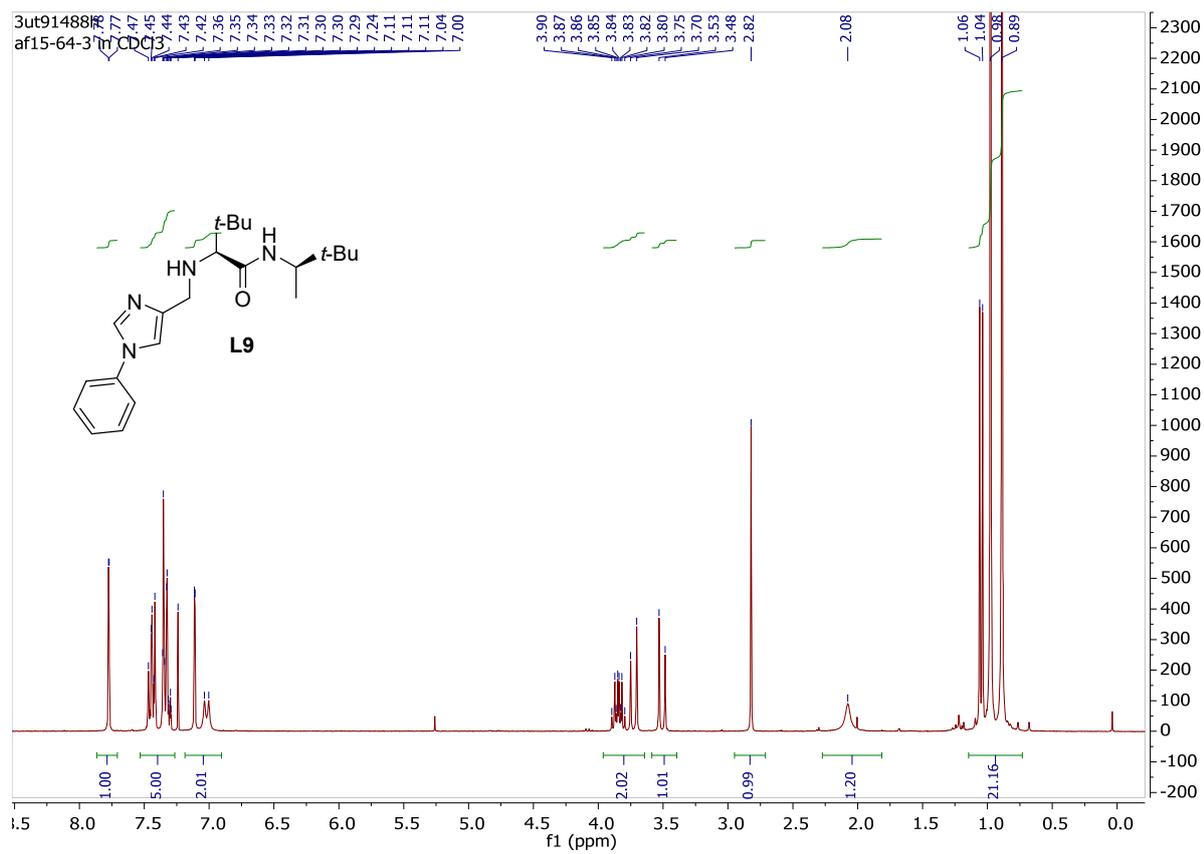
a) Yields determined via ¹H NMR with pyrazine as internal standard. b) Results described in this work. c) ee values determined via chiral HPLC measurement. d) Results described in literature for non-heme iron-catalyzed epoxidation employing H₂O₂ and miscellaneous ligands and/or additives (see references for further details). e) ee value determined after derivatization: aminolysis with isopropylamine towards corresponding β-aminoalcohol. f) Detected via ¹H-NMR and ESI-MS. g) n. d.: not determined. h) *in situ* catalyst generation with **L10** (5 mol%), FeCl₃·6H₂O (5 mol%), (*S*)-(+)-mandelic acid (15 mol%), 1.6 mL 2-Me-2-BuOH, followed by the addition of alkene (0.166 mmol) and 2 eq. H₂O₂ *via* syringe pump at r. t. (3 h reaction time).

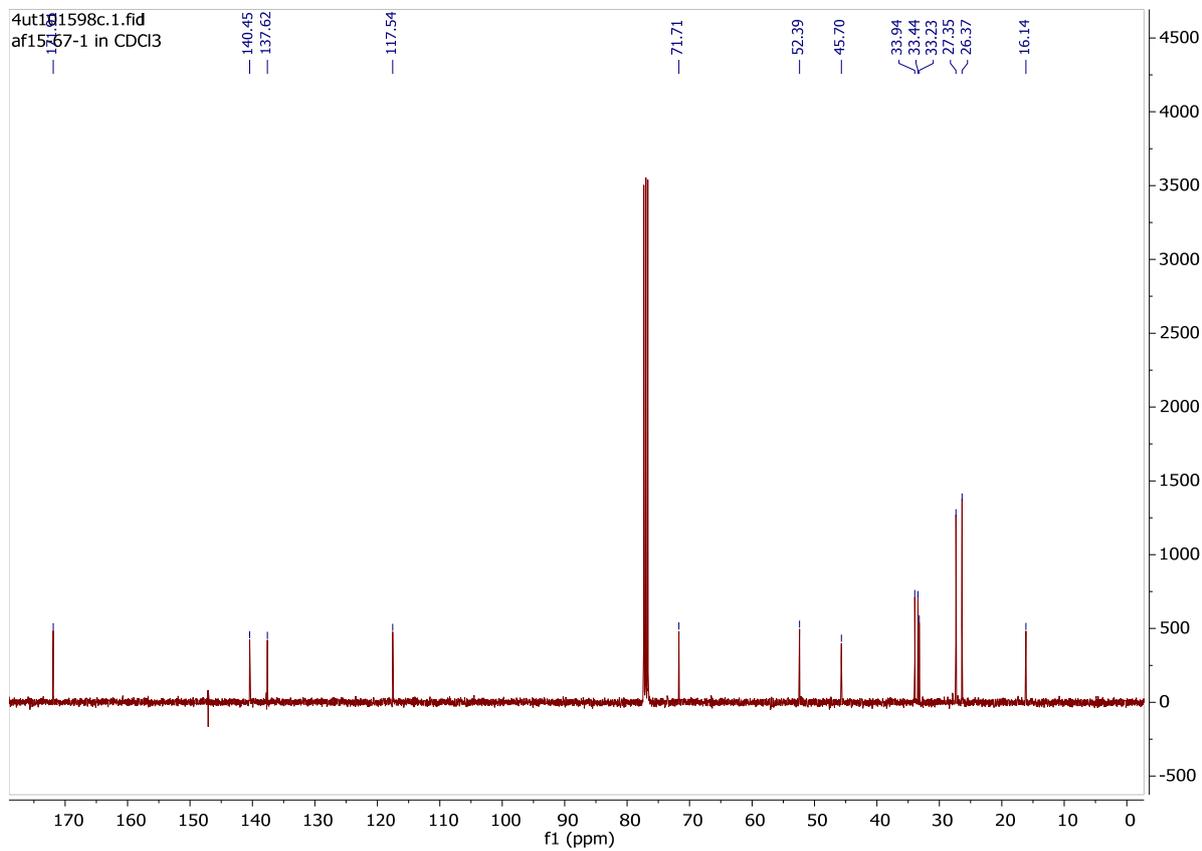
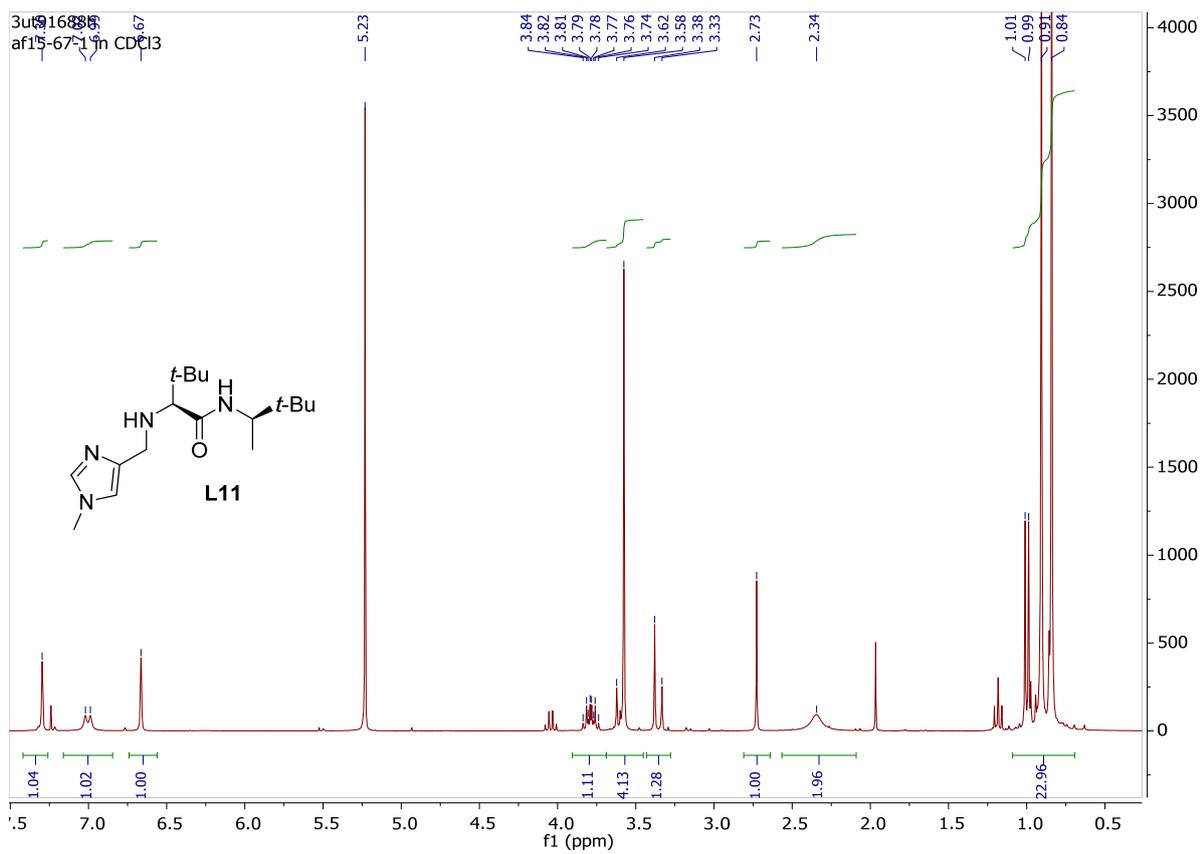
References (see Table 4)

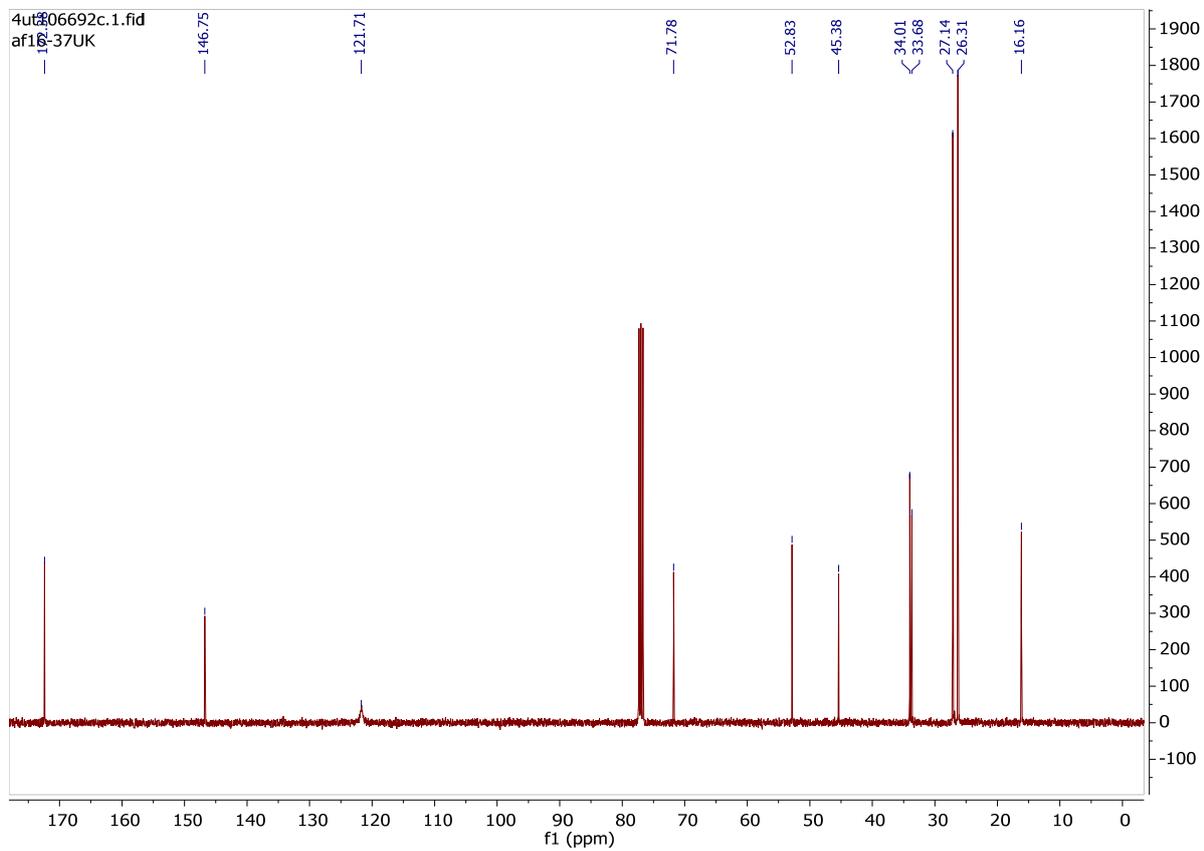
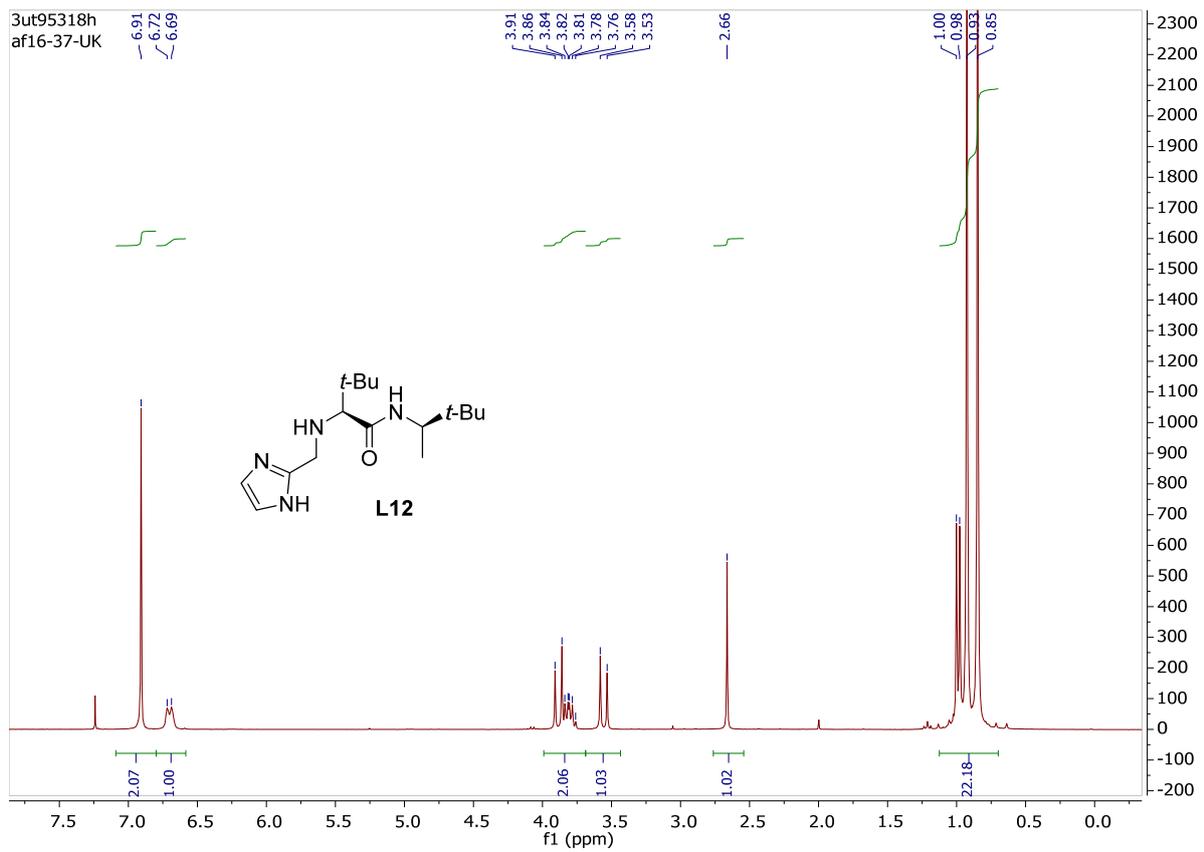
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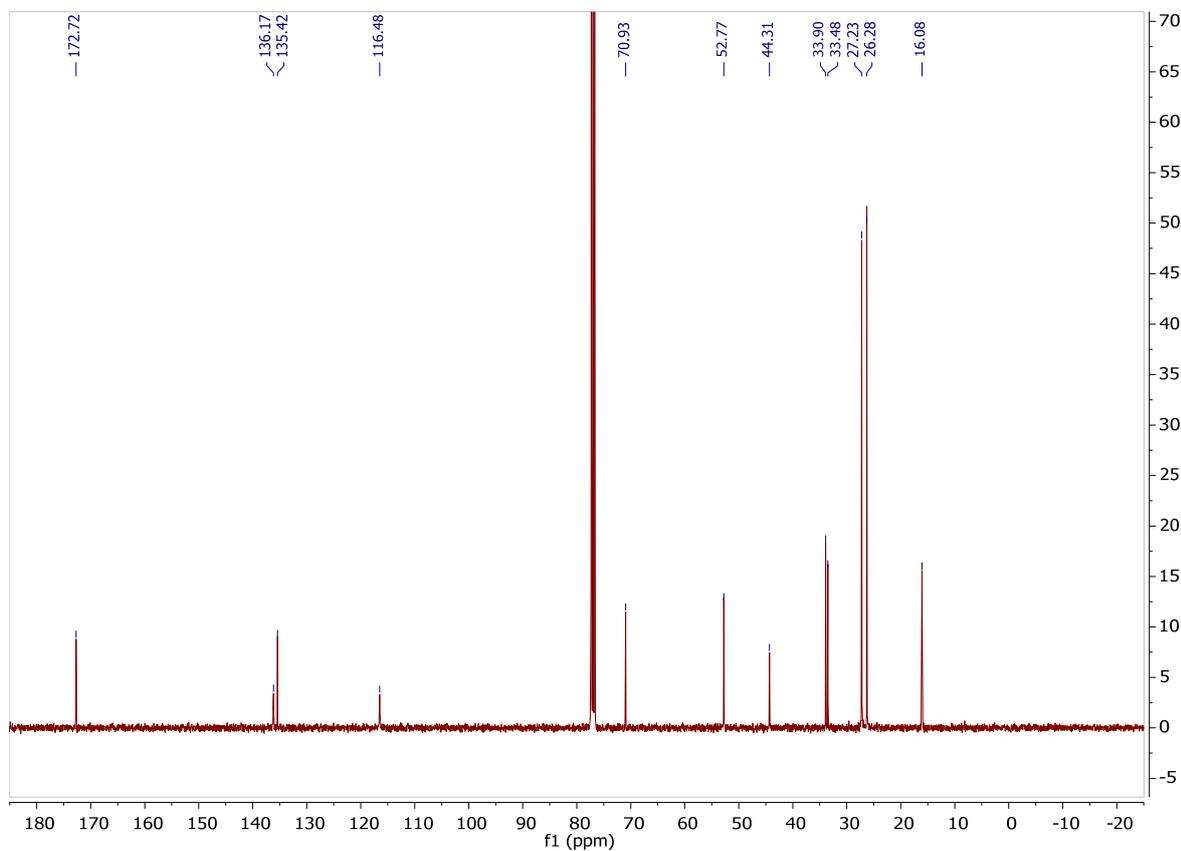
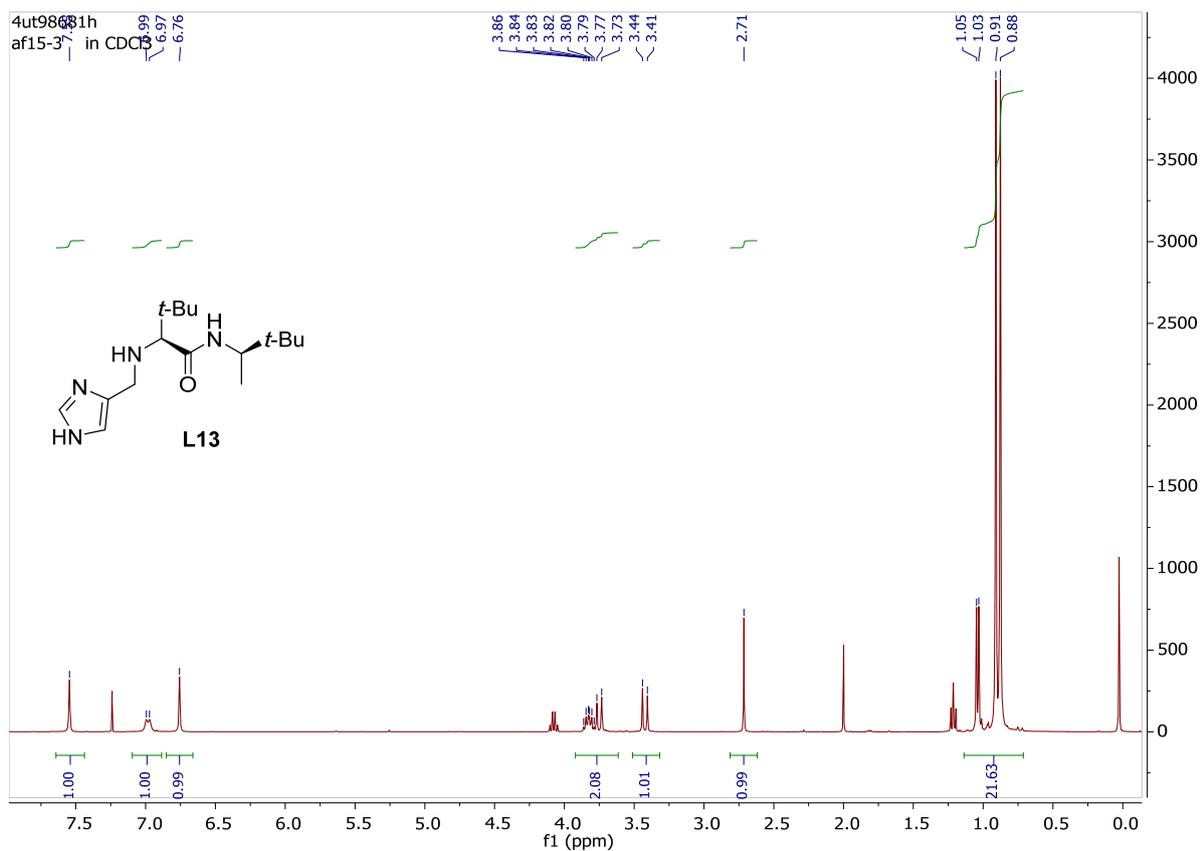
6 ¹H NMR and ¹³C NMR spectra of new ligands

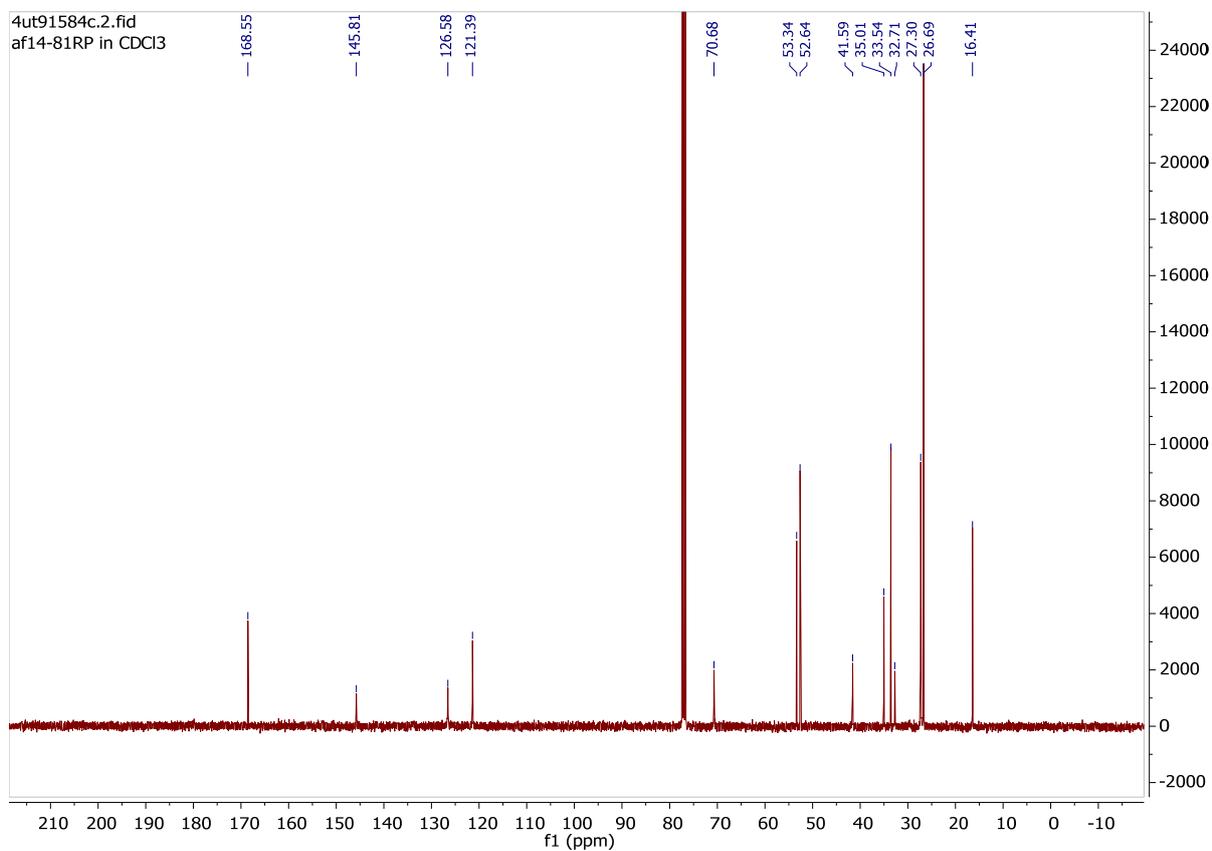
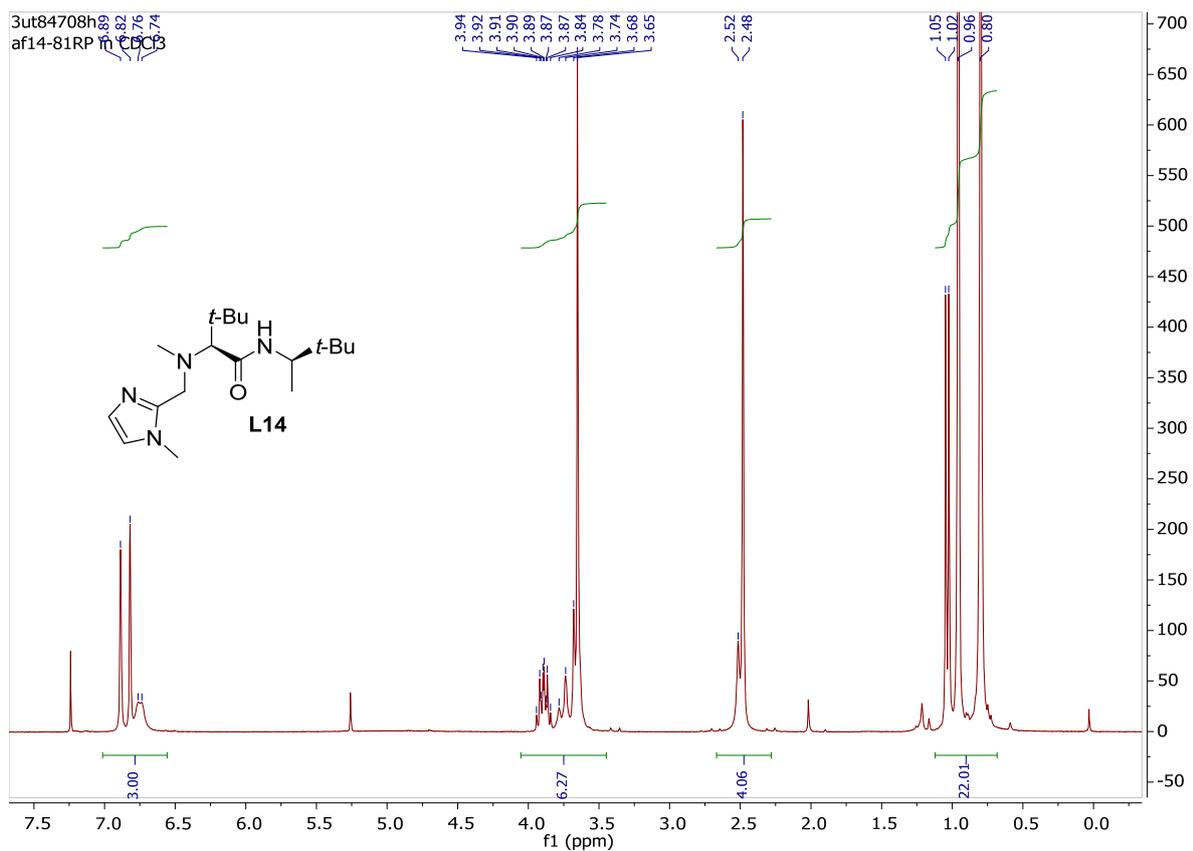


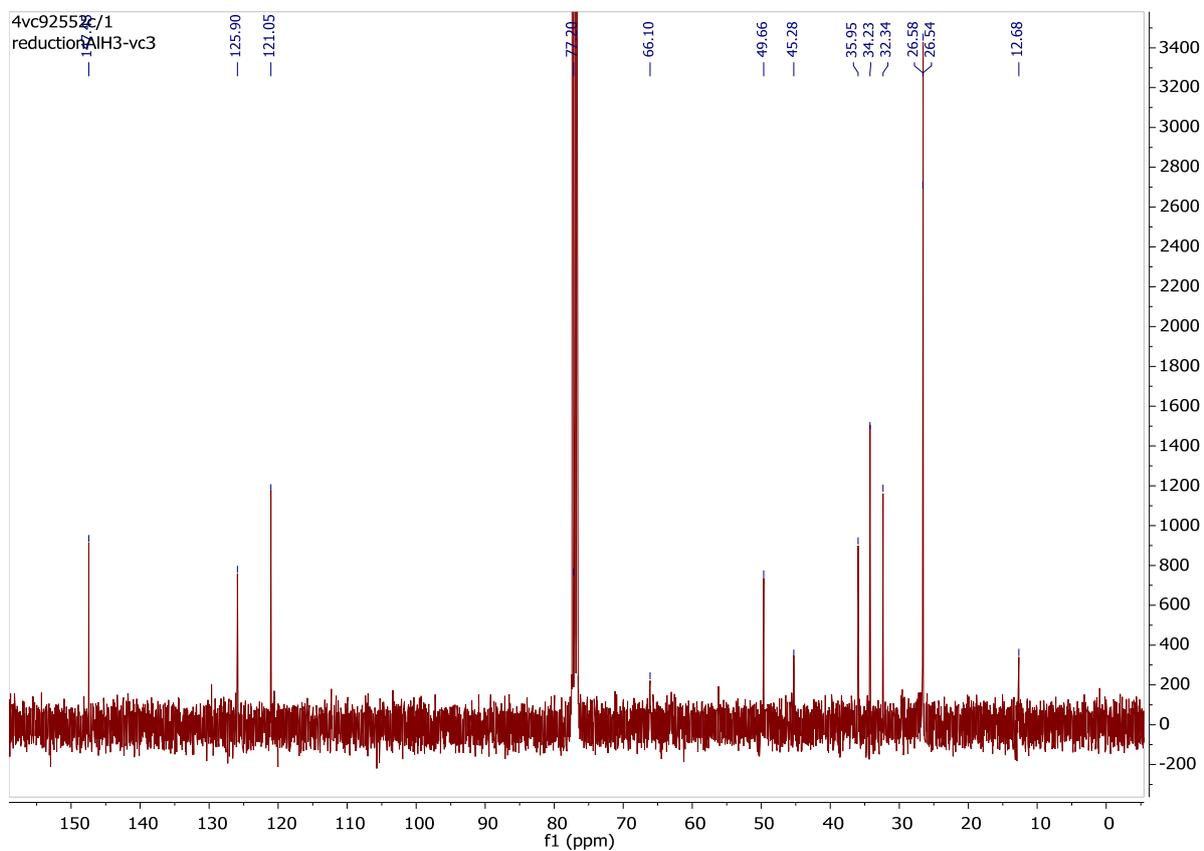
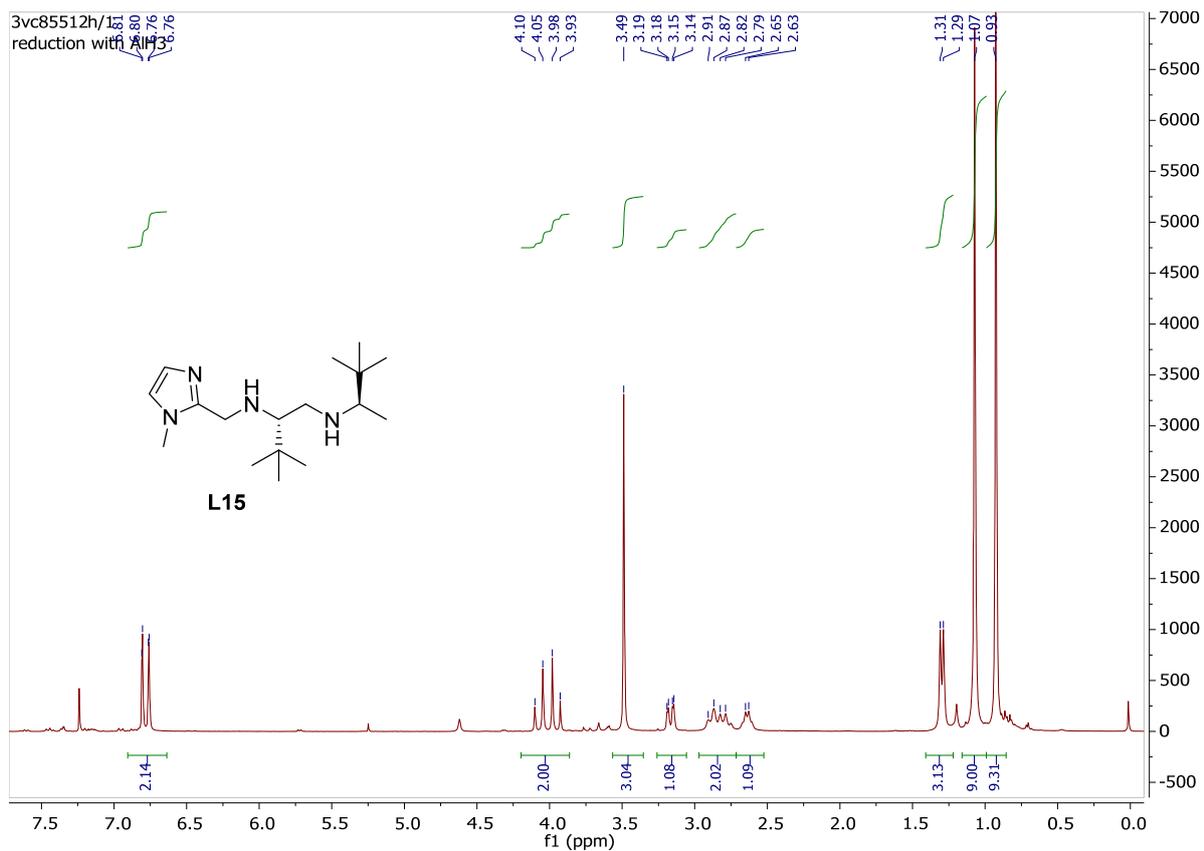


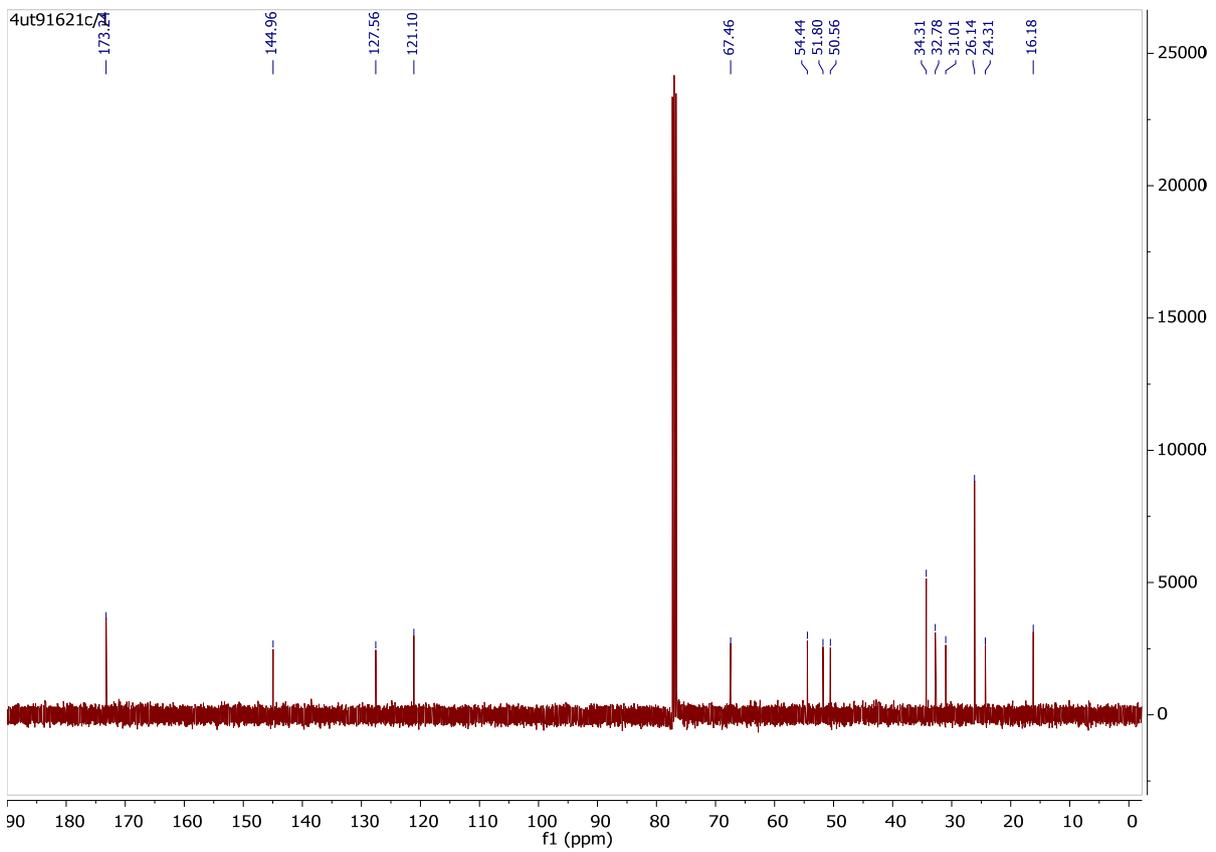
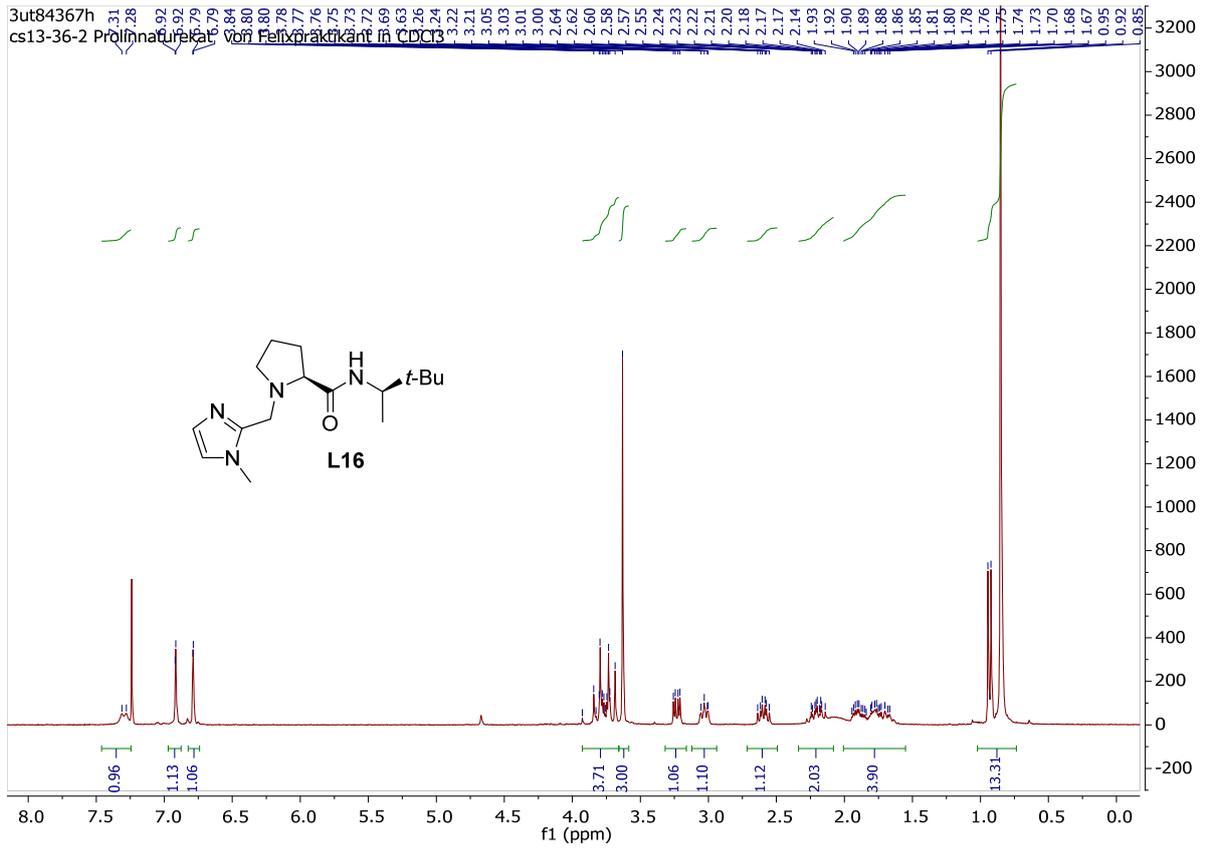








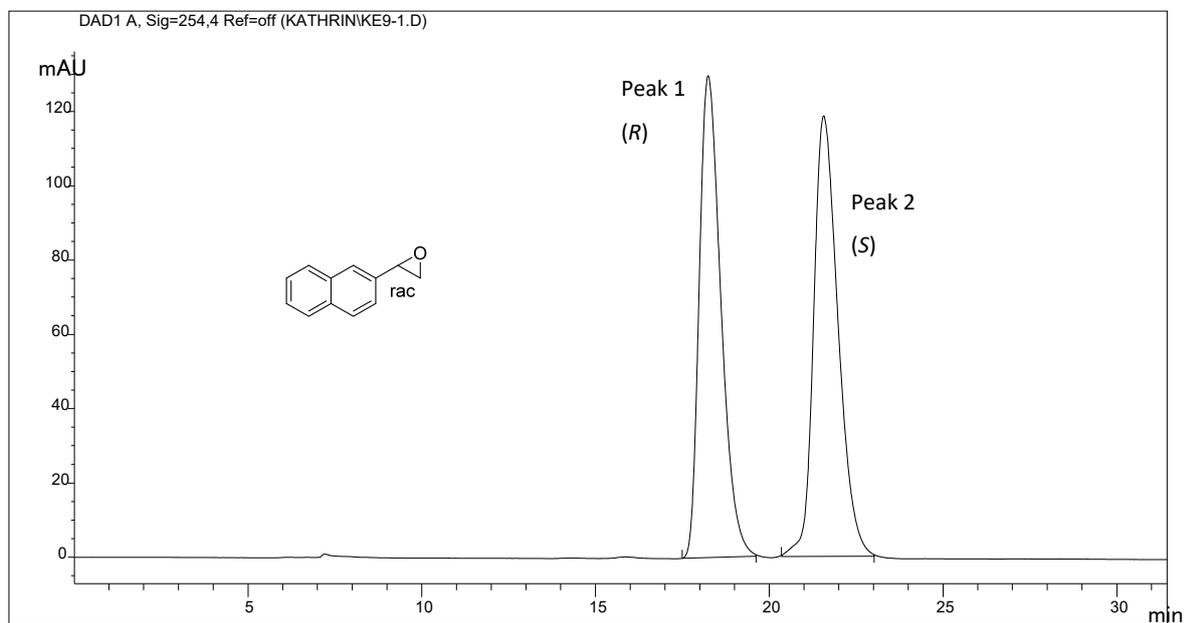




7 HPLC chromatograms

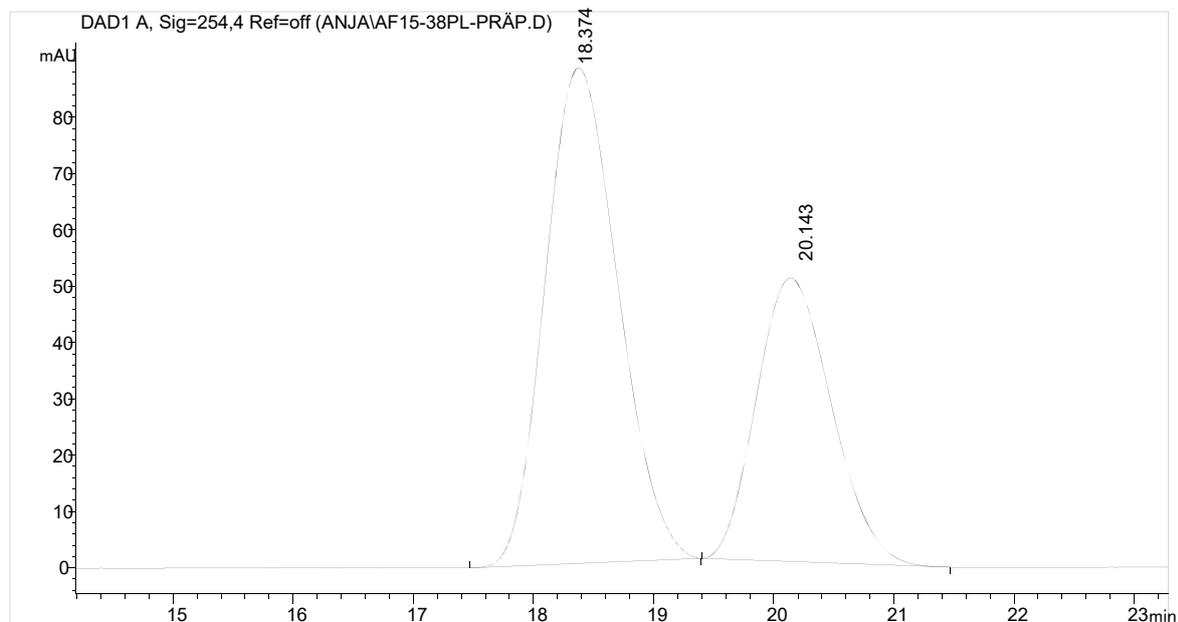
2-Naphthalen-2-yl-oxirane

Chiral HPLC (Daicel Chiralpak AS): *n*-hexane/*i*-PrOH 99:1, 0.5 ml/min flow rate, $\lambda = 254 \text{ nm}$;



Peak	t_R	Area	Area (%)
1	18.236	5621.8	49.24
2	21.563	5796	50.76

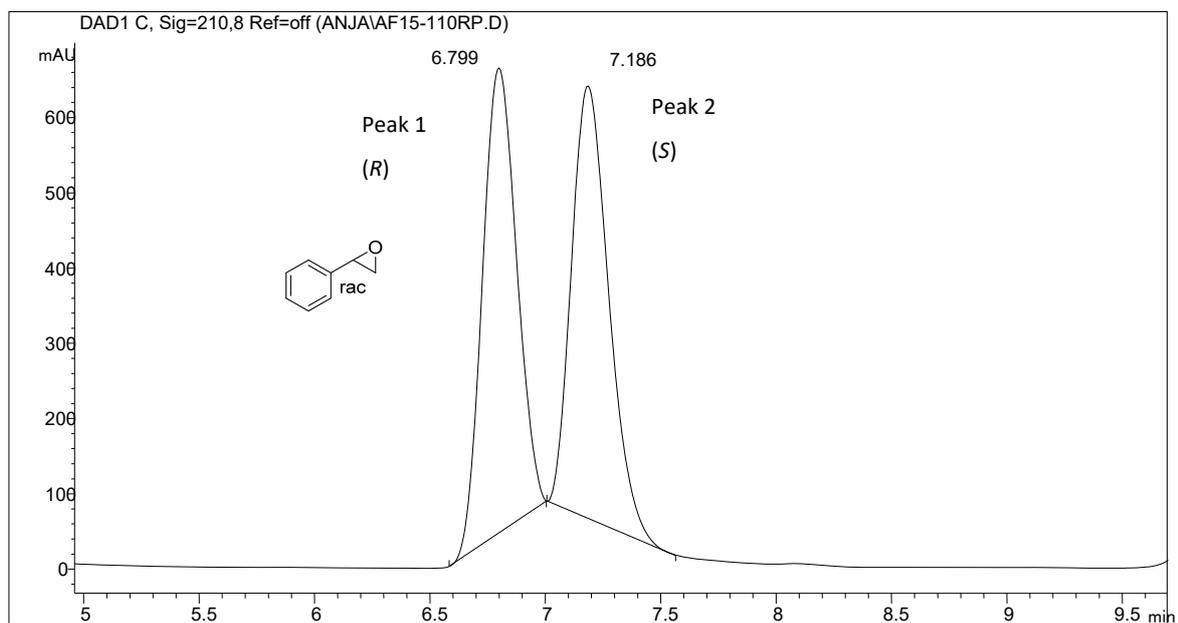
See Entry 1, Table 2



Peak	t_R	Area	Area (%)
1	18.374	3715.3	62.867
2	20.143	2194.5	37.133

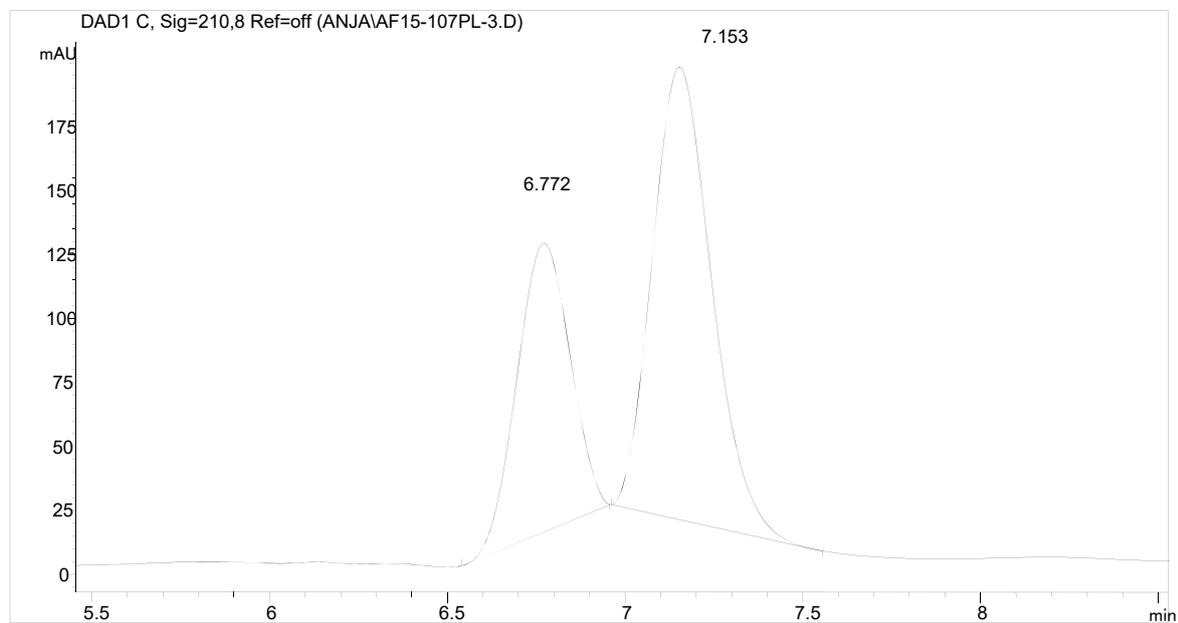
2-Phenyloxirane

Chiral HPLC (Macherey-Nagel OD): *n*-hexane/*i*-PrOH 99:1, 1.0 ml/min flow rate, $\lambda = 210$ nm;



Peak	t_R	Area	Area (%)
1	6.799	6357.5	49.627
2	7.186	6453	50.373

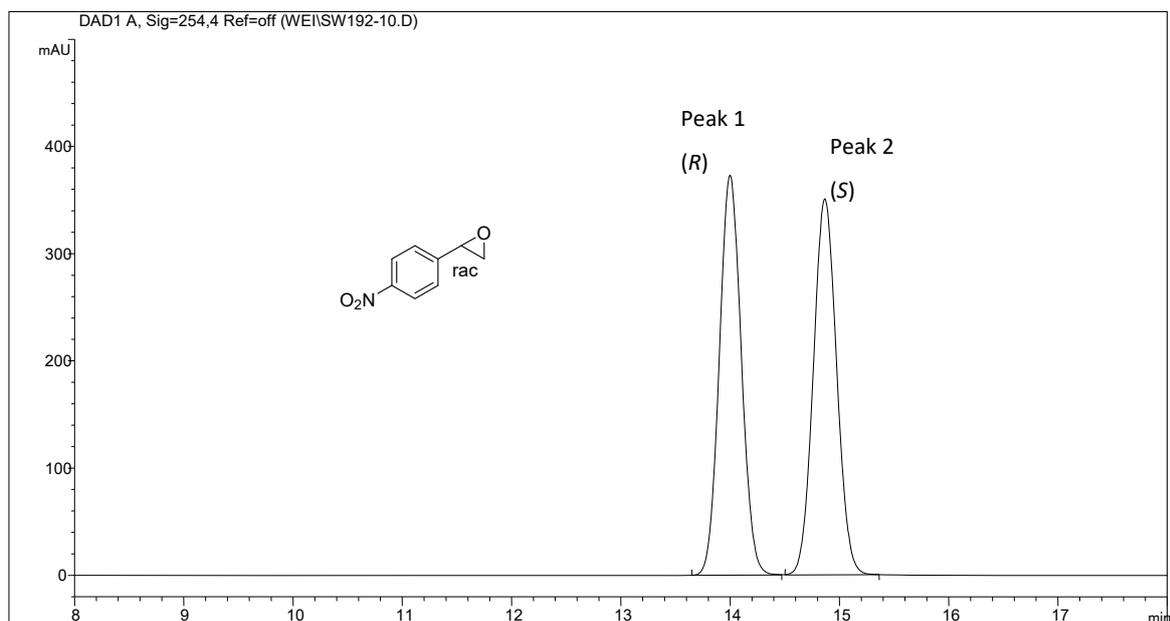
See Entry 2, Table 2



Peak	t_R	Area	Area (%)
1	6.772	1174.8	36.338
2	7.153	2058.1	63.662

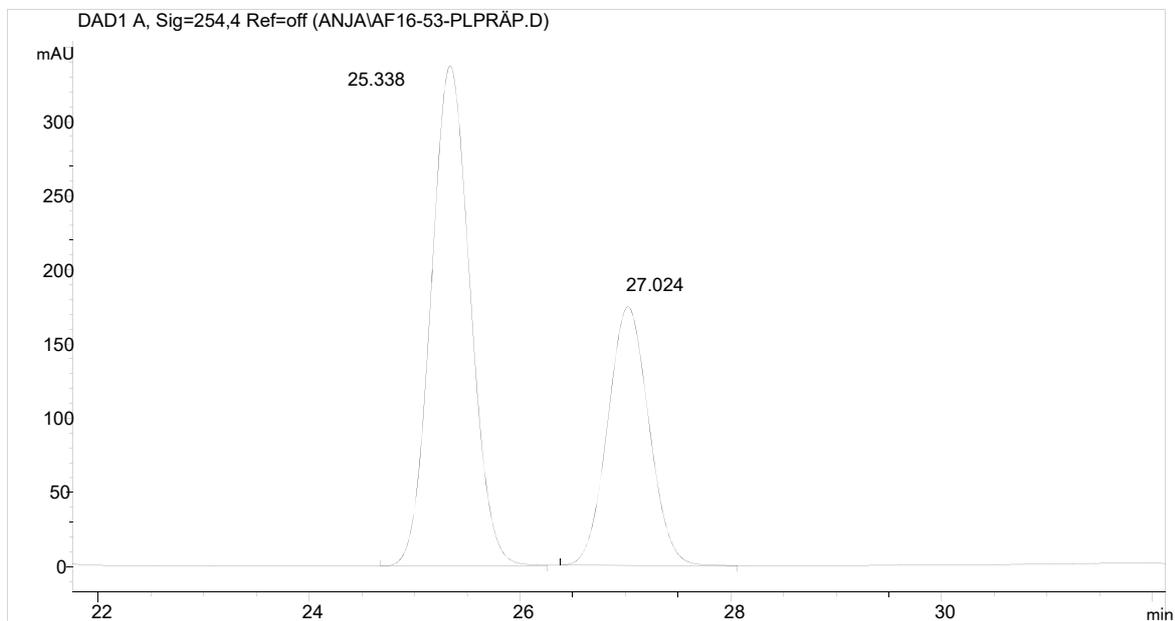
2-(4-Nitro-phenyl)-oxirane

Chiral HPLC (Daicel Chiralpak IC): *n*-hexane/*i*-PrOH 95:5, 1.0 ml/min flow rate, $\lambda = 254 \text{ nm}$;



Peak	t _R	Area	Area (%)
1	13.998	360.44	49.94
2	14.865	339.40	50.06

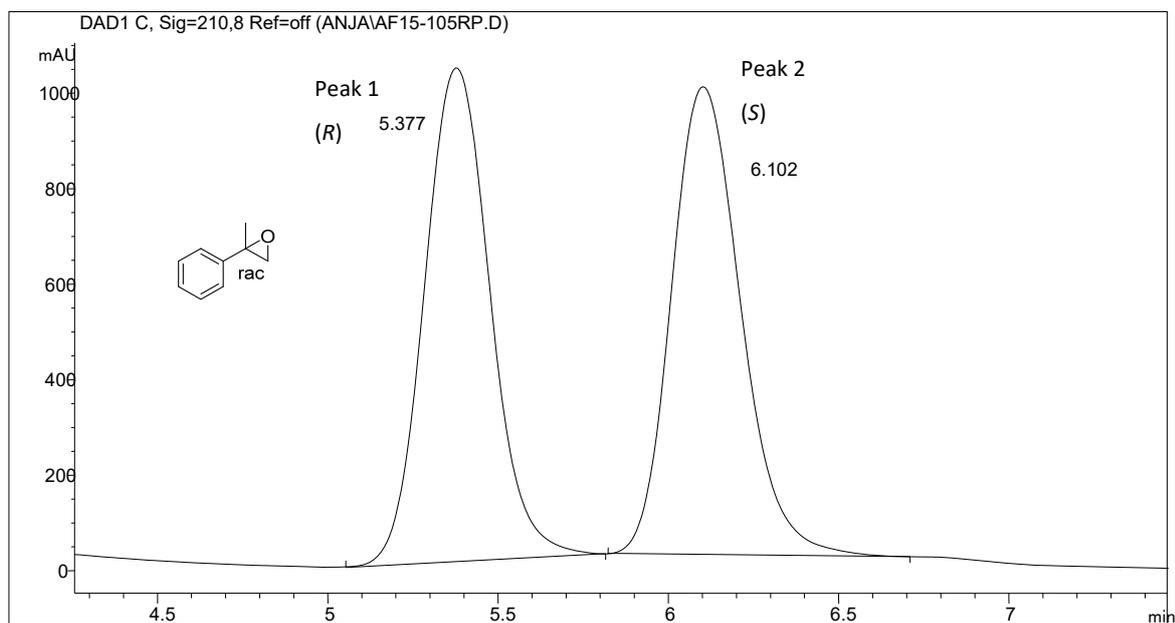
See Entry 3, Table 2 (different retention behaviour due to pressure problems in HPLC system)



Peak	t _R	Area	Area (%)
1	25.338	8679.2	64.625
2	27.024	4750.9	35.375

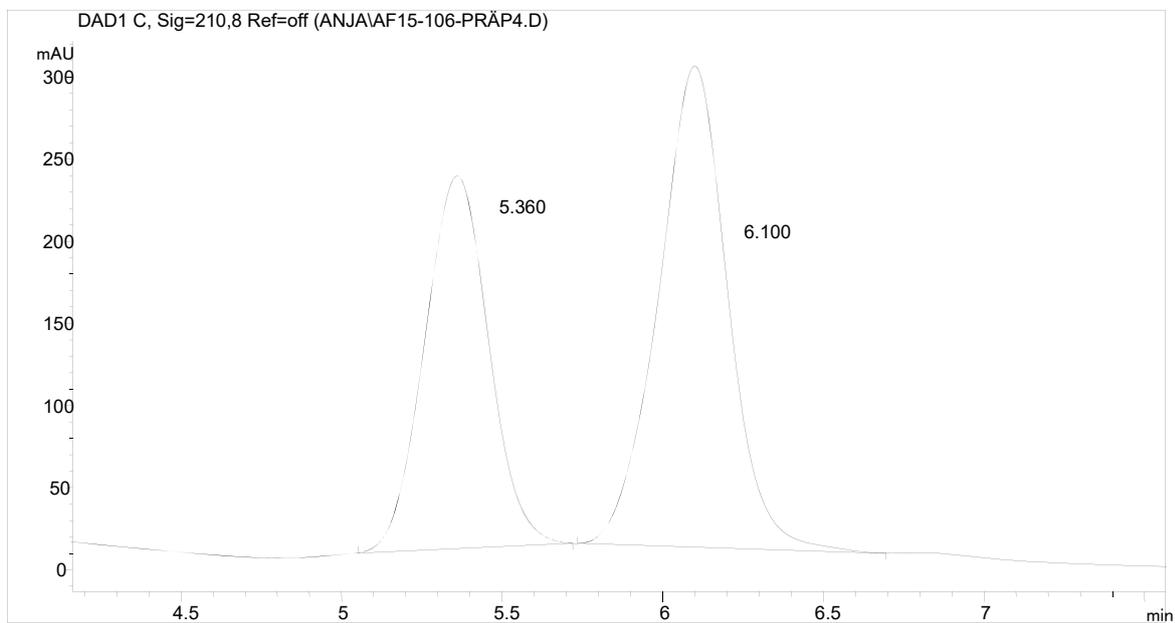
2-Methyl-2-phenyloxirane

Chiral HPLC (Macherey-Nagel OD): *n*-hexane/*i*-PrOH 99:1, 1.0 ml/min flow rate, $\lambda = 210$ nm;



Peak	t _R	Area	Area (%)
1	5.377	13850.2	50.000
2	6.102	13850	50.000

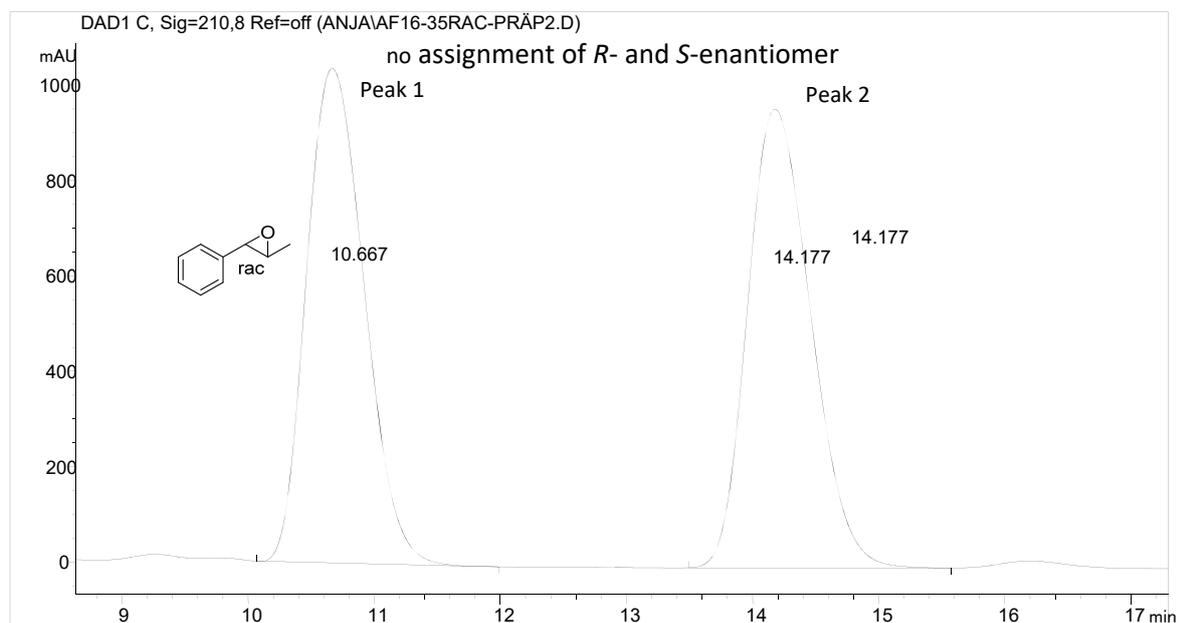
See Entry 6, Table 2



Peak	t _R	Area	Area (%)
1	5.36	3153	41.812
2	6.1	4387.9	58.188

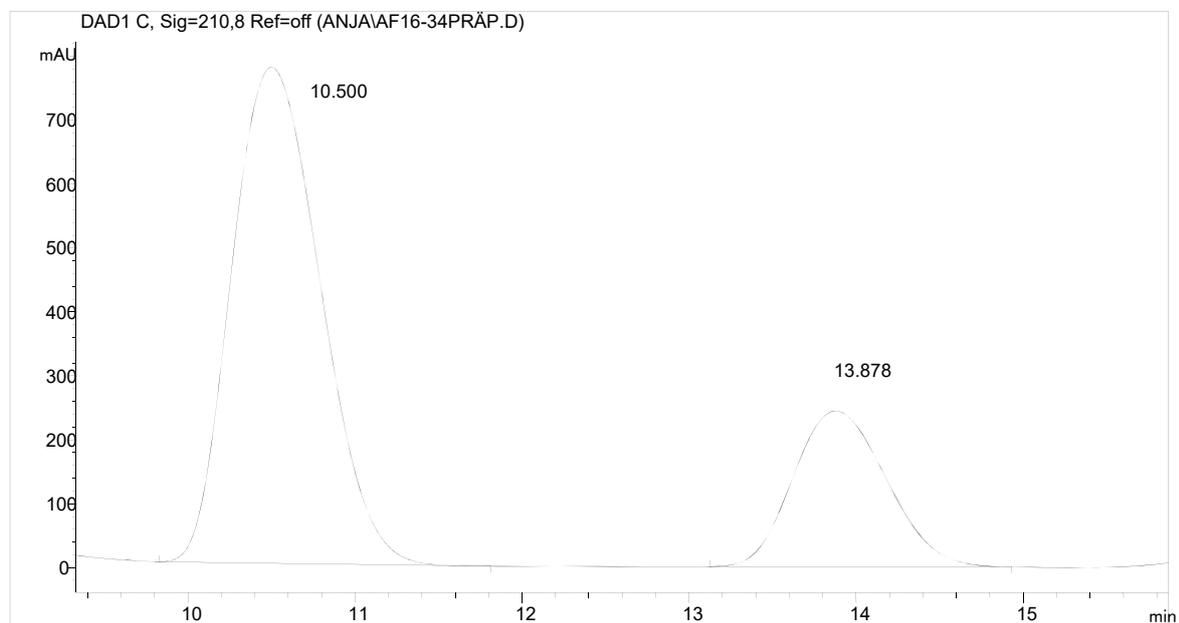
2-Methyl-3-phenyloxirane

Chiral HPLC (Daicel Chiralpak AS): *n*-hexane/*i*-PrOH 99:1, 0.5 ml/min flow rate, $\lambda = 210 \text{ nm}$;



Peak	t_R	Area	Area (%)
1	10.667	32920.4	49.569
2	14.177	33493	50.431

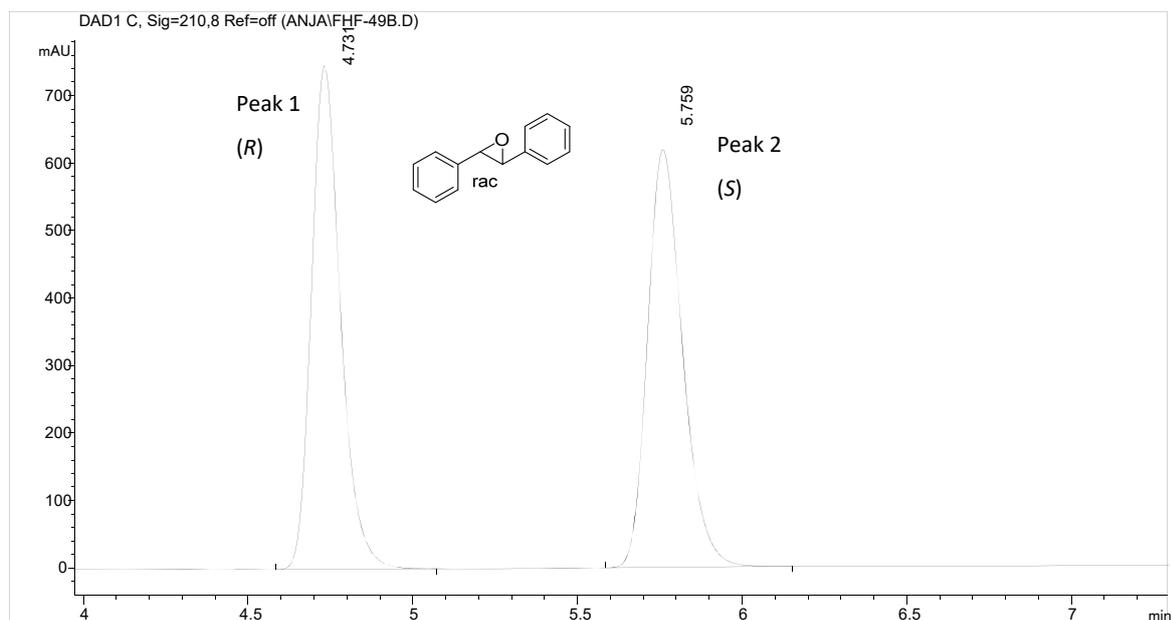
See Entry 8, Table 2



Peak	t_R	Area	Area (%)
1	10.5	28623.3	75.183
2	13.878	9448.3	24.817

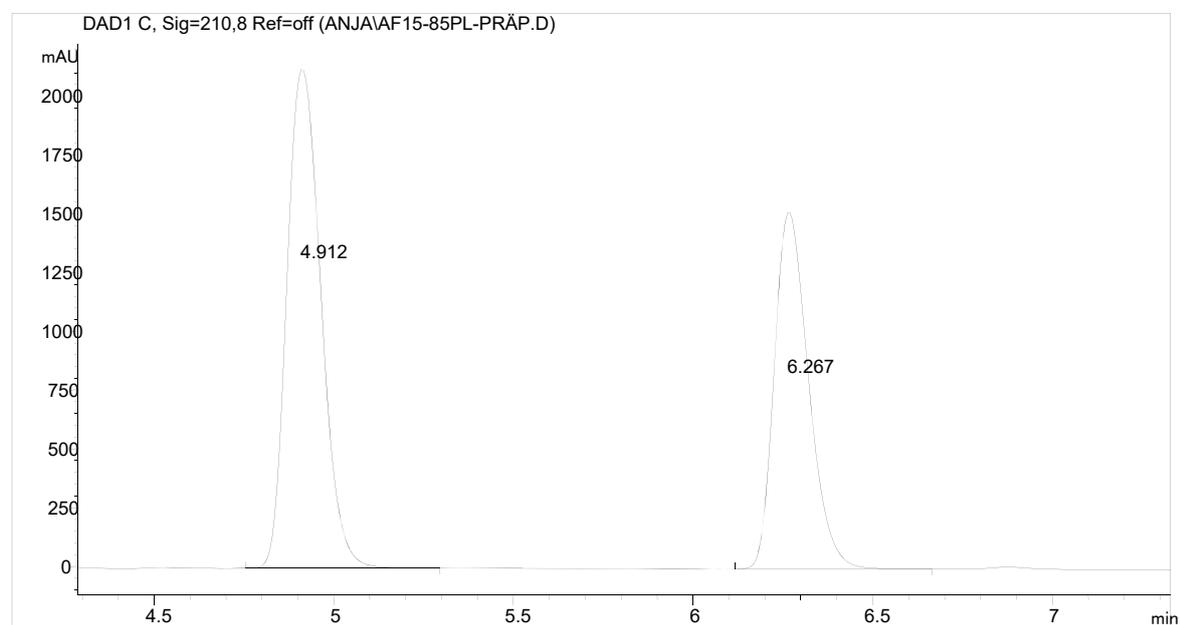
2,3-Diphenyloxirane

Chiral HPLC (Daicel Chiralpak IB): *n*-hexane/*i*-PrOH 92:8, 1.0 ml/min flow rate, $\lambda = 210$ nm;



Peak	t_R	Area	Area (%)
1	4.731	4576.1	49.973
2	5.759	4581.1	50.027

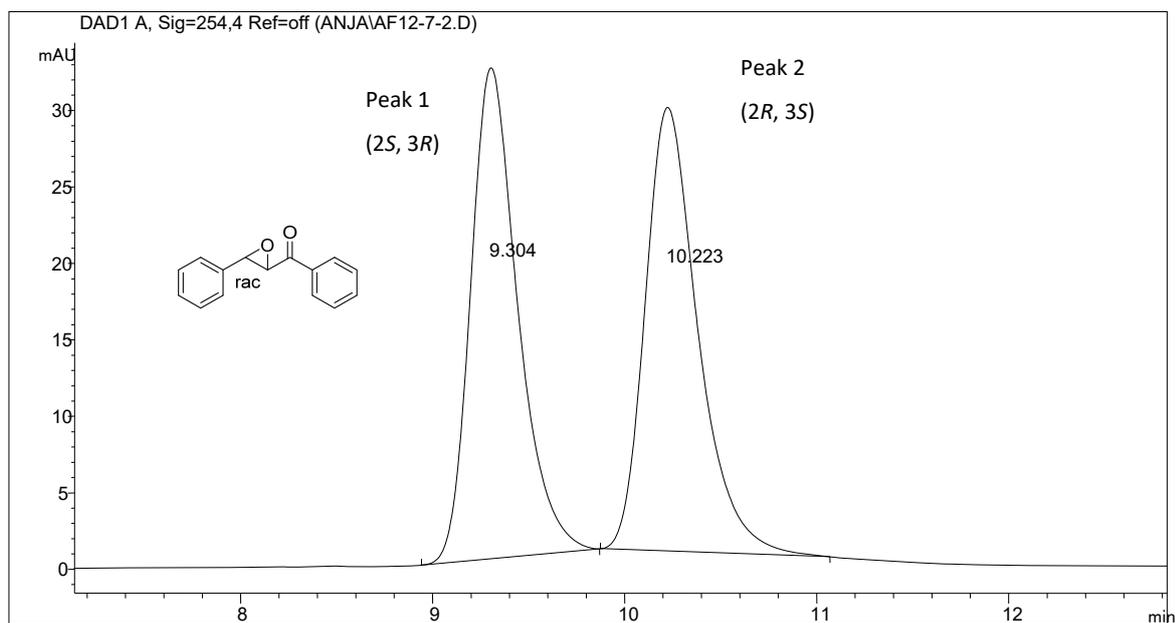
See Entry 9, Table 2



Peak	t_R	Area	Area (%)
1	4.912	13954.8	58.083
2	6.267	10070.9	41.917

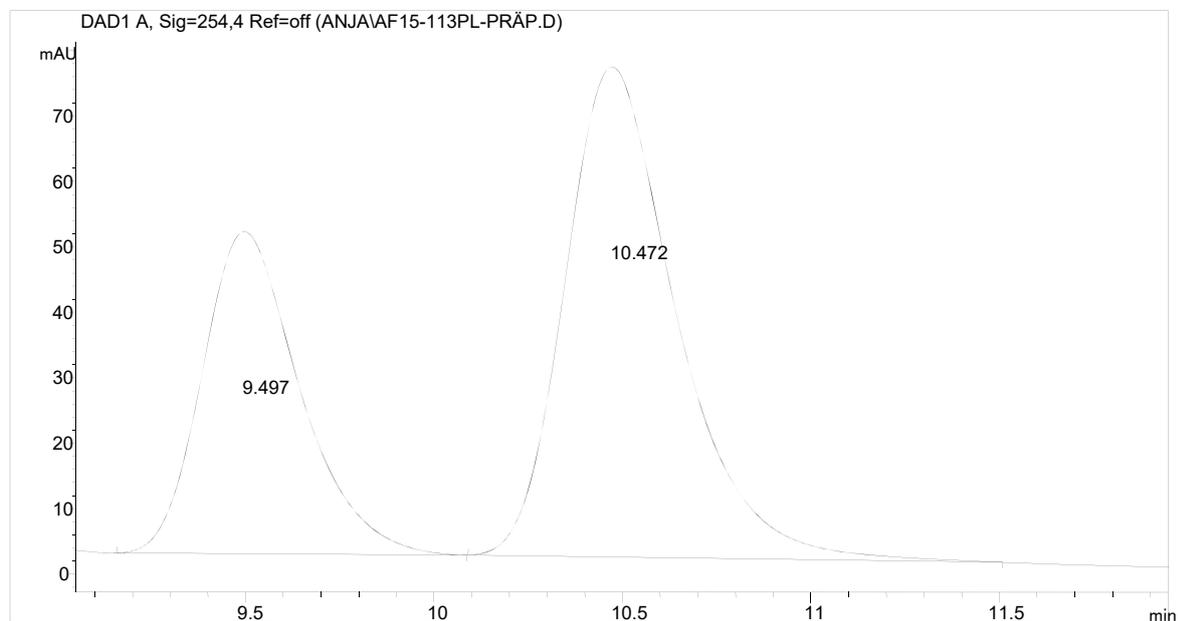
Phenyl(3-phenyloxiran-2-yl)methanone

Chiral HPLC (Macherey-Nagel OD): *n*-hexane/*i*-PrOH 90:10, 1.0 ml/min flow rate, $\lambda = 245$ nm;



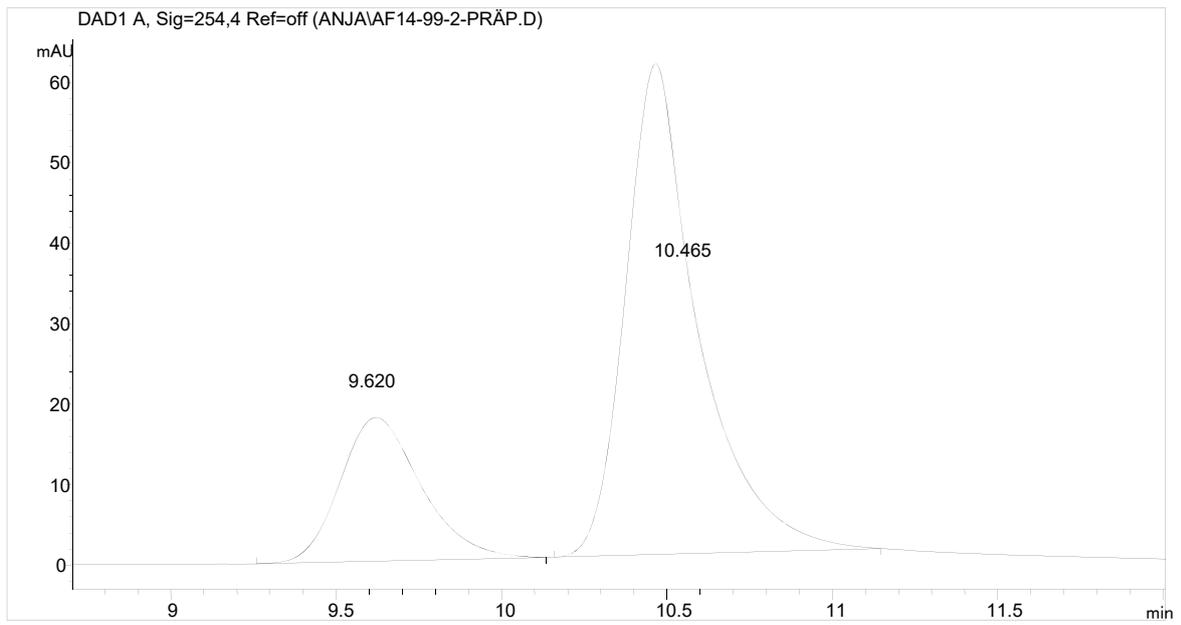
Peak	t _R	Area	Area (%)
1	9.304	547.6	49.702
2	10.223	554.1	50.298

See Entry 10, Table 2



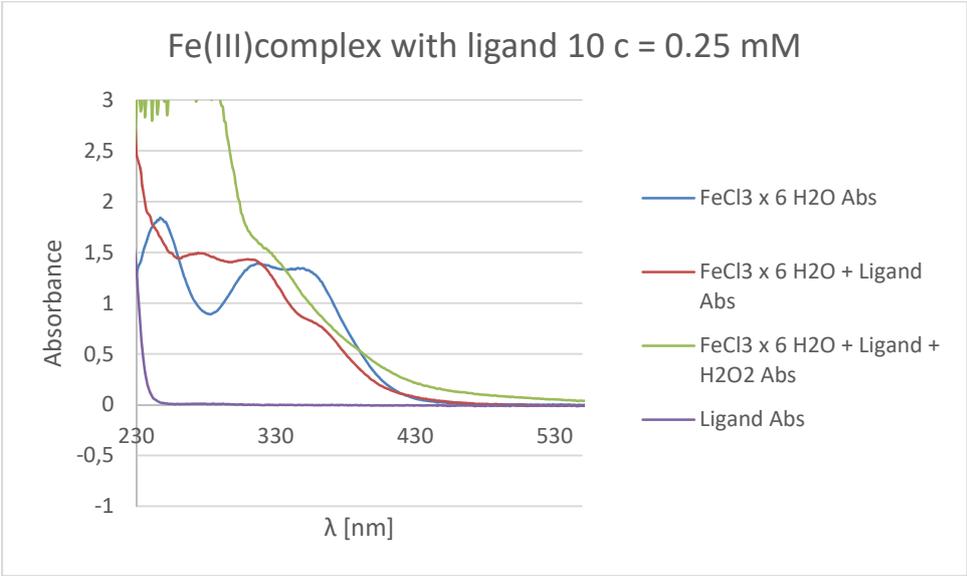
Peak	t _R	Area	Area (%)
1	9.497	884.6	36.474
2	10.472	1540.7	63.526

See Entry 11, Table 2

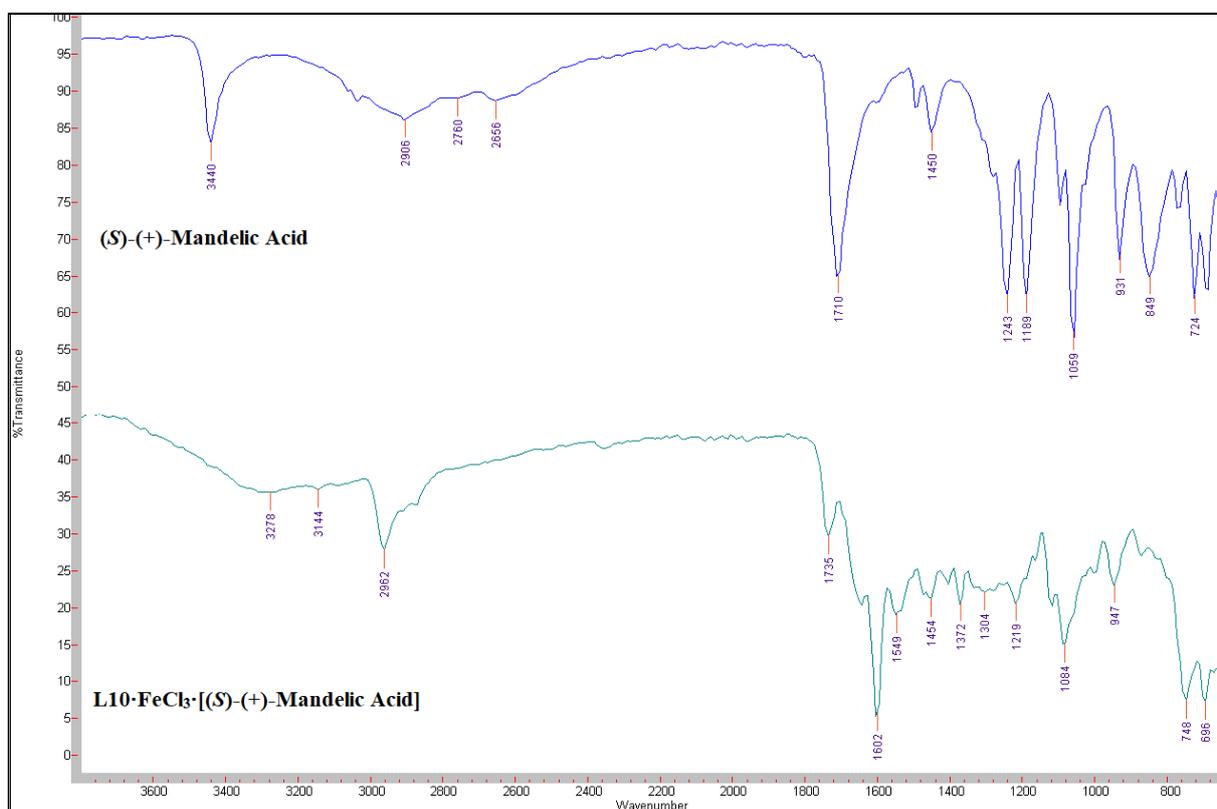
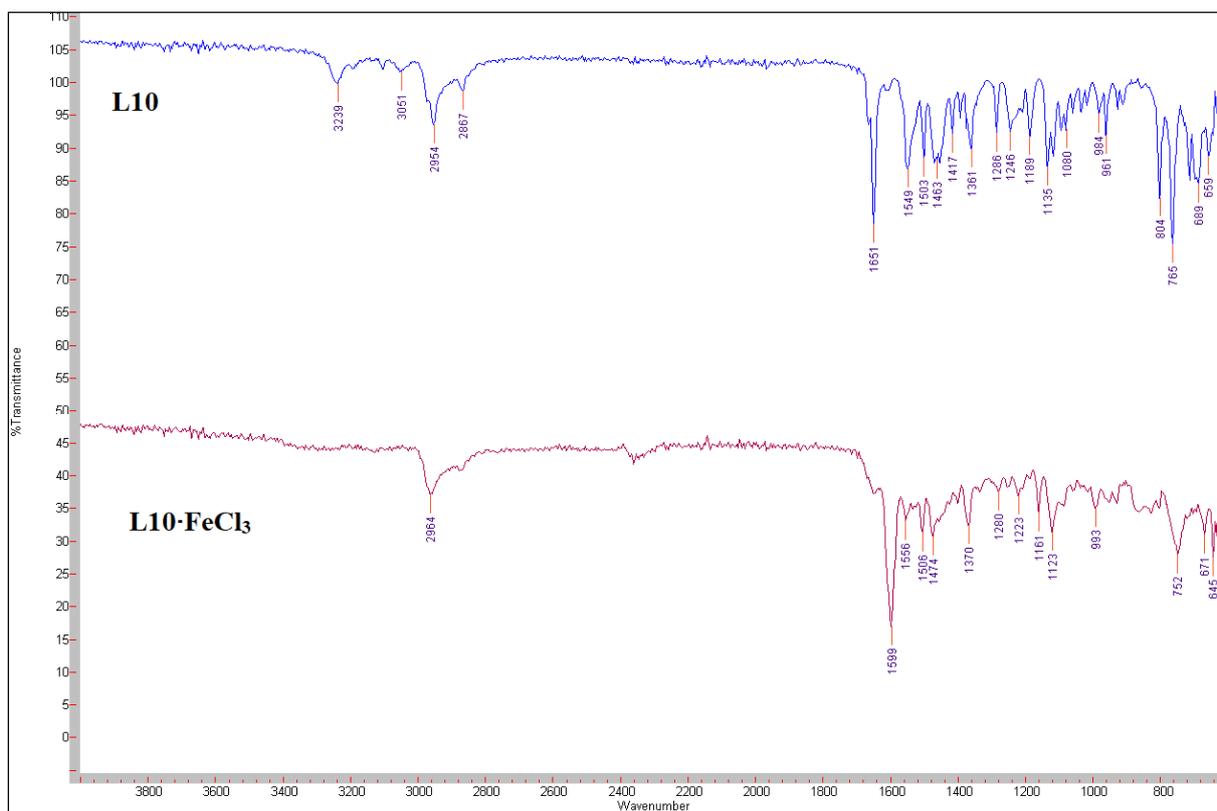


Peak	t _R	Area	Area (%)
1	9.62	304.5	24.589
2	10.465	934	75.411

8 Investigation of Fe(III) complex via UV-Vis spectroscopy

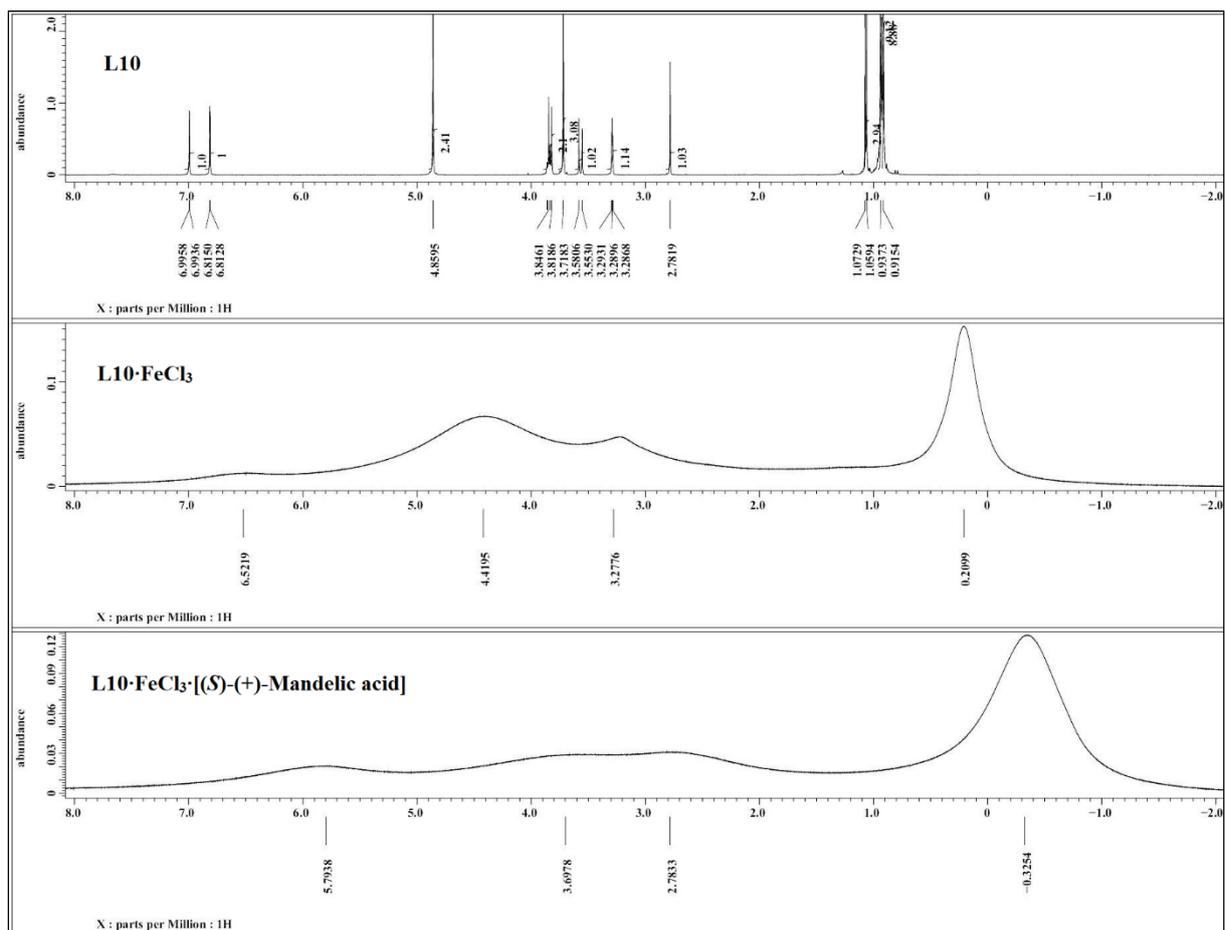


9 Infrared Experiments of Fe(III) complex

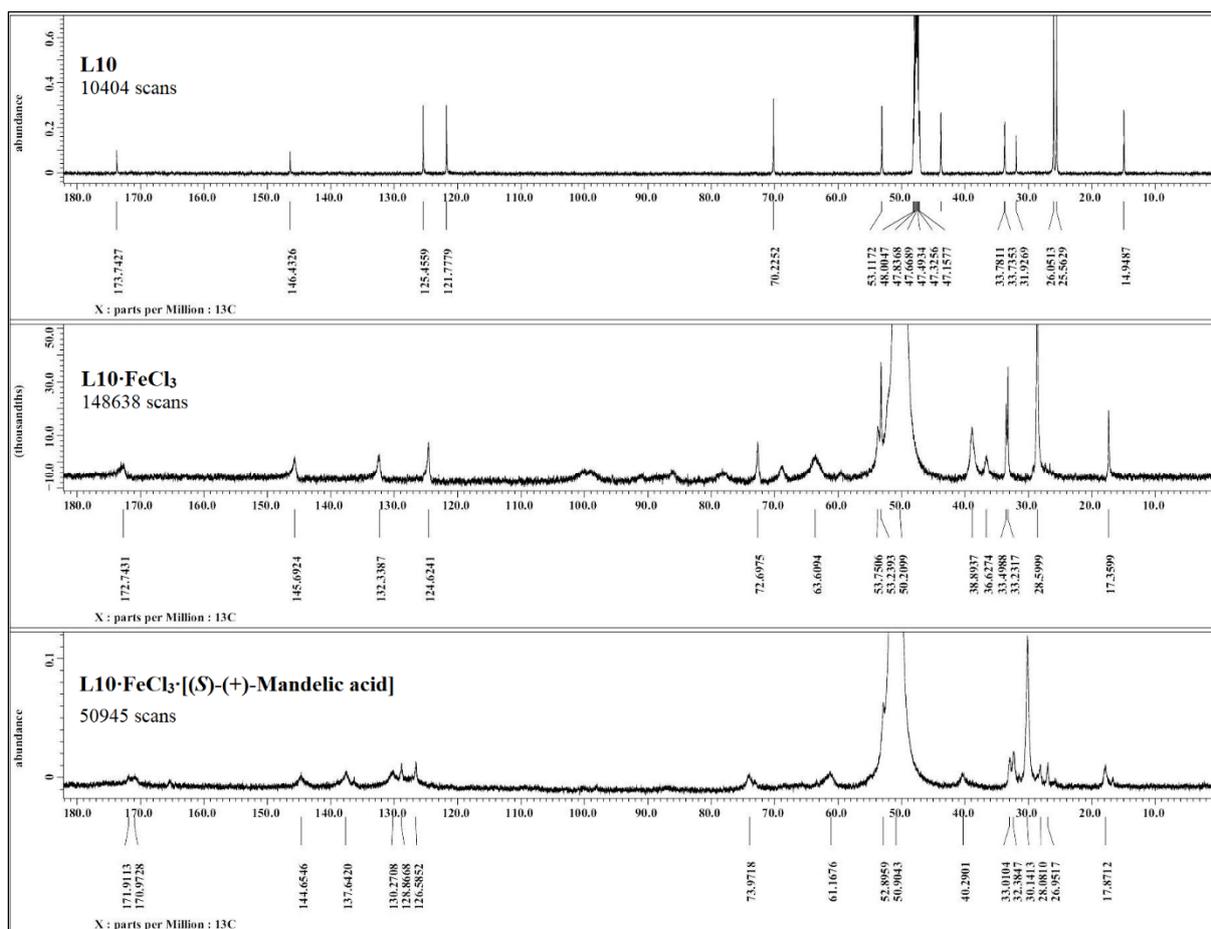


9 NMR Experiments of Fe(III) complex (Jeol ECA 500)

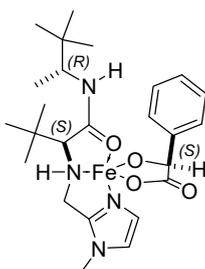
^1H , 500 MHz, methanol- d_3



^{13}C , 125.765 MHz, methanol- d_3



10 MS-ESI Experiment of Fe(III) complex



$C_{25}H_{38}FeN_4O_4$
Mol. Wt.: 514.4386



Table A
Experimental details

Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.038, 0.096, 1.09
No. of reflections	3632
No. of parameters	216
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\max}, \Delta\rho_{\min}$ (e Å ⁻³)	0.23, -0.18
Absolute structure	Refined as an inversion twin.
Absolute structure parameter	0.1 (4)
Computer programs: Bruker <i>APEX3</i> software, <i>SAINT</i> V8.38A integration software, <i>SORTAV</i> (Blessing, 1995), <i>SHELXS2013</i> (Sheldrick, 2008), <i>SHELXL2018/3</i> (Sheldrick, 2018), <i>ORTEP-3 for Windows</i> (Farrugia, 2012), <i>WinGX</i> publication routines (Farrugia, 2012).	

Table B
Hydrogen-bond geometry (Å, °)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C4—H4 \cdots O11 ⁱ	0.95	2.45	3.356 (2)	159
C6—H6A \cdots N8	0.98	2.55	3.027 (2)	110
C6—H6C \cdots O11 ⁱⁱ	0.98	2.54	3.482 (2)	162
N12—H12 \cdots N8	0.83 (3)	2.12 (2)	2.597 (2)	117 (2)

Symmetry codes: (i) $-x+1/2, -y+1, z+1/2$; (ii) $-x+3/2, -y+1, z+1/2$.

Computing details

Data collection: Bruker *APEX3* software; cell refinement: *SAINT* V8.38A integration software; data reduction: *SORTAV* (Blessing, 1995); program(s) used to solve structure: *SHELXS2013* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2018); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *WinGX* publication routines (Farrugia, 2012).

(shelx)

Crystal data

$C_{17}H_{30}N_4O$
 $M_r = 306.45$
 Orthorhombic, $P2_12_12_1$
 Hall symbol: P 2ac 2ab
 $a = 6.9534$ (3) Å
 $b = 14.0923$ (5) Å
 $c = 18.9394$ (7) Å
 $V = 1855.86$ (12) Å³
 $Z = 4$

$F(000) = 672$
 $D_x = 1.097$ Mg m⁻³
 Cu $K\alpha$ radiation, $\lambda = 1.54178$ Å
 Cell parameters from 9103 reflections
 $\theta = 3.9\text{--}72.3^\circ$
 $\mu = 0.55$ mm⁻¹
 $T = 141$ K
 Block, colourless
 $0.45 \times 0.39 \times 0.26$ mm

Data collection

Bruker D8 VENTURE
 diffractometer
 Radiation source: sealed x-ray microsource
 Multilayer mirrors monochromator
 φ or ω oscillation scans
 Absorption correction: multi-scan
 SADABS2016/2 - Bruker AXS area detector scaling
 and absorption correcti
 $T_{\min} = 0.677, T_{\max} = 0.754$

12734 measured reflections
 3632 independent reflections
 3452 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.035$
 $\theta_{\max} = 72.3^\circ, \theta_{\min} = 3.9^\circ$
 $h = -8 \rightarrow 8$
 $k = -17 \rightarrow 16$
 $l = -22 \rightarrow 23$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.096$
 $S = 1.09$
3632 reflections
216 parameters
0 restraints
0 constraints
Primary atom site location: dual

Hydrogen site location: mixed
H atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F^2) + (0.0536P)^2 + 0.1847P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.23 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.18 \text{ e } \text{\AA}^{-3}$
Absolute structure: Refined as an inversion twin.
Absolute structure parameter: 0.1 (4)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refined as a 2-component inversion twin.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C2	0.3729 (3)	0.51502 (13)	0.81217 (10)	0.0274 (4)
C4	0.2396 (3)	0.48785 (15)	0.91227 (11)	0.0371 (4)
H4	0.144996	0.478742	0.947808	0.045*
C5	0.4333 (3)	0.48187 (15)	0.92274 (10)	0.0338 (4)
H5	0.496783	0.46861	0.96604	0.041*
C6	0.7258 (3)	0.49392 (16)	0.84670 (10)	0.0352 (4)
H6A	0.763954	0.544904	0.814422	0.053*
H6B	0.75851	0.432403	0.825763	0.053*
H6C	0.793877	0.501301	0.891665	0.053*
C7	0.3917 (3)	0.53921 (13)	0.73808 (10)	0.0272 (4)
C9	0.5455 (3)	0.56919 (12)	0.63061 (9)	0.0247 (4)
H9	0.408052	0.573869	0.61549	0.03*
C10	0.6423 (2)	0.48977 (13)	0.58749 (9)	0.0248 (3)
C13	0.8078 (3)	0.33434 (13)	0.59699 (10)	0.0300 (4)
H13	0.818462	0.342356	0.544672	0.036*
C14	1.0118 (3)	0.31556 (14)	0.62544 (11)	0.0331 (4)
C15	1.1452 (3)	0.39353 (17)	0.59928 (16)	0.0516 (6)
H15A	1.272017	0.386047	0.621168	0.077*
H15B	1.091777	0.455574	0.611966	0.077*
H15C	1.157823	0.389187	0.547843	0.077*
C16	1.0829 (3)	0.22042 (17)	0.59710 (18)	0.0551 (7)
H16A	1.069924	0.219389	0.545578	0.083*
H16B	1.00607	0.169023	0.61759	0.083*
H16C	1.21828	0.211756	0.609928	0.083*
C17	1.0160 (5)	0.3147 (3)	0.70620 (14)	0.0720 (9)
H17A	1.145193	0.297563	0.722449	0.108*
H17B	0.922929	0.268148	0.723832	0.108*
H17C	0.98247	0.377843	0.72406	0.108*
C18	0.6634 (3)	0.25497 (17)	0.61048 (17)	0.0544 (7)
H18A	0.711664	0.195721	0.589983	0.082*
H18B	0.539995	0.271339	0.588671	0.082*
H18C	0.645806	0.246831	0.66146	0.082*
C19	0.6426 (3)	0.66769 (13)	0.61868 (10)	0.0303 (4)
C20	0.8590 (3)	0.66265 (16)	0.63368 (12)	0.0403 (5)

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{iso} */ <i>U</i> _{eq}
H20A	0.880126	0.634832	0.680459	0.06*
H20B	0.913638	0.726724	0.632361	0.06*
H20C	0.921503	0.623174	0.597792	0.06*
C21	0.6082 (4)	0.70108 (16)	0.54278 (12)	0.0447 (5)
H21A	0.672692	0.657792	0.509924	0.067*
H21B	0.660107	0.765247	0.536865	0.067*
H21C	0.469829	0.701575	0.532976	0.067*
C22	0.5494 (3)	0.73969 (15)	0.66897 (12)	0.0417 (5)
H22A	0.576565	0.721556	0.717924	0.062*
H22B	0.41006	0.740513	0.661321	0.062*
H22C	0.602216	0.802981	0.659783	0.062*
N1	0.5181 (2)	0.49860 (11)	0.85900 (8)	0.0280 (3)
N3	0.2014 (2)	0.50889 (13)	0.84311 (9)	0.0339 (4)
N8	0.5499 (2)	0.54364 (11)	0.70524 (8)	0.0257 (3)
N12	0.7308 (2)	0.42302 (12)	0.62562 (8)	0.0286 (3)
O11	0.6345 (2)	0.48868 (10)	0.52280 (7)	0.0374 (3)
H7	0.267 (3)	0.5536 (16)	0.7145 (12)	0.033 (6)*
H12	0.721 (3)	0.4291 (17)	0.6690 (13)	0.035 (6)*

Atomic displacement parameters (Å²)

	<i>U</i> ¹¹	<i>U</i> ²²	<i>U</i> ³³	<i>U</i> ¹²	<i>U</i> ¹³	<i>U</i> ²³
C2	0.0268 (8)	0.0257 (8)	0.0299 (8)	0.0000 (7)	0.0033 (7)	-0.0030 (7)
C4	0.0392 (10)	0.0389 (10)	0.0331 (9)	0.0029 (9)	0.0128 (8)	0.0010 (9)
C5	0.0422 (10)	0.0336 (9)	0.0257 (9)	0.0024 (9)	0.0059 (8)	-0.0005 (8)
C6	0.0261 (9)	0.0481 (12)	0.0315 (9)	0.0003 (8)	-0.0005 (7)	-0.0014 (9)
C7	0.0244 (9)	0.0268 (8)	0.0303 (9)	0.0020 (7)	-0.0002 (7)	-0.0024 (7)
C9	0.0223 (8)	0.0281 (8)	0.0236 (8)	0.0020 (7)	-0.0017 (6)	0.0016 (7)
C10	0.0229 (7)	0.0267 (8)	0.0250 (8)	-0.0016 (7)	-0.0014 (6)	-0.0008 (7)
C13	0.0301 (10)	0.0266 (9)	0.0333 (9)	0.0043 (7)	0.0007 (7)	-0.0022 (8)
C14	0.0295 (9)	0.0306 (9)	0.0393 (10)	0.0049 (8)	0.0009 (8)	0.0036 (8)
C15	0.0342 (11)	0.0348 (11)	0.0858 (18)	-0.0030 (10)	0.0005 (12)	0.0015 (12)
C16	0.0311 (11)	0.0326 (11)	0.102 (2)	0.0059 (9)	0.0098 (12)	-0.0033 (12)
C17	0.0647 (18)	0.105 (3)	0.0463 (14)	0.0288 (18)	-0.0145 (13)	0.0212 (16)
C18	0.0322 (11)	0.0338 (11)	0.097 (2)	0.0000 (9)	0.0089 (12)	-0.0044 (12)
C19	0.0337 (10)	0.0264 (9)	0.0309 (9)	0.0000 (8)	-0.0005 (8)	0.0011 (7)
C20	0.0350 (11)	0.0392 (11)	0.0467 (11)	-0.0094 (9)	0.0005 (9)	-0.0035 (10)
C21	0.0614 (15)	0.0350 (10)	0.0377 (11)	-0.0012 (10)	0.0006 (10)	0.0097 (9)
C22	0.0489 (12)	0.0280 (9)	0.0481 (12)	0.0027 (9)	0.0043 (10)	-0.0035 (9)
N1	0.0289 (7)	0.0285 (8)	0.0267 (7)	0.0000 (6)	0.0027 (6)	-0.0030 (6)
N3	0.0307 (8)	0.0362 (9)	0.0348 (8)	0.0010 (7)	0.0082 (7)	-0.0013 (7)
N8	0.0268 (7)	0.0265 (7)	0.0237 (7)	0.0018 (6)	-0.0009 (6)	-0.0004 (6)
N12	0.0326 (8)	0.0302 (8)	0.0229 (7)	0.0076 (7)	-0.0012 (6)	-0.0013 (6)
O11	0.0494 (8)	0.0394 (8)	0.0235 (6)	0.0102 (7)	-0.0039 (6)	-0.0009 (6)

Geometric parameters

<i>Bond lengths (Å)</i>			
C2—N3	1.331 (2)	C14—C17	1.530 (3)
C2—N1	1.364 (2)	C15—H15A	0.98
C2—C7	1.450 (2)	C15—H15B	0.98
C4—C5	1.363 (3)	C15—H15C	0.98
C4—N3	1.369 (3)	C16—H16A	0.98

Bond lengths (Å)

C4—H4	0.95	C16—H16B	0.98
C5—N1	1.364 (2)	C16—H16C	0.98
C5—H5	0.95	C17—H17A	0.98
C6—N1	1.464 (2)	C17—H17B	0.98
C6—H6A	0.98	C17—H17C	0.98
C6—H6B	0.98	C18—H18A	0.98
C6—H6C	0.98	C18—H18B	0.98
C7—N8	1.265 (2)	C18—H18C	0.98
C7—H7	1.00 (2)	C19—C21	1.531 (3)
C9—N8	1.459 (2)	C19—C20	1.533 (3)
C9—C10	1.540 (2)	C19—C22	1.535 (3)
C9—C19	1.560 (3)	C20—H20A	0.98
C9—H9	1	C20—H20B	0.98
C10—O11	1.227 (2)	C20—H20C	0.98
C10—N12	1.336 (2)	C21—H21A	0.98
C13—N12	1.464 (2)	C21—H21B	0.98
C13—C18	1.525 (3)	C21—H21C	0.98
C13—C14	1.540 (3)	C22—H22A	0.98
C13—H13	1	C22—H22B	0.98
C14—C15	1.521 (3)	C22—H22C	0.98
C14—C16	1.527 (3)	N12—H12	0.83 (3)

Bond Angles (°)

N3—C2—N1	111.45 (16)	C14—C16—H16C	109.5
N3—C2—C7	121.50 (17)	H16A—C16—H16C	109.5
N1—C2—C7	127.03 (16)	H16B—C16—H16C	109.5
C5—C4—N3	110.12 (17)	C14—C17—H17A	109.5
C5—C4—H4	124.9	C14—C17—H17B	109.5
N3—C4—H4	124.9	H17A—C17—H17B	109.5
C4—C5—N1	106.73 (18)	C14—C17—H17C	109.5
C4—C5—H5	126.6	H17A—C17—H17C	109.5
N1—C5—H5	126.6	H17B—C17—H17C	109.5
N1—C6—H6A	109.5	C13—C18—H18A	109.5
N1—C6—H6B	109.5	C13—C18—H18B	109.5
H6A—C6—H6B	109.5	H18A—C18—H18B	109.5
N1—C6—H6C	109.5	C13—C18—H18C	109.5
H6A—C6—H6C	109.5	H18A—C18—H18C	109.5
H6B—C6—H6C	109.5	H18B—C18—H18C	109.5
N8—C7—C2	124.48 (17)	C21—C19—C20	109.96 (19)
N8—C7—H7	121.9 (13)	C21—C19—C22	108.27 (17)
C2—C7—H7	113.6 (13)	C20—C19—C22	109.26 (18)
N8—C9—C10	108.98 (14)	C21—C19—C9	109.99 (16)
N8—C9—C19	110.54 (14)	C20—C19—C9	110.90 (16)
C10—C9—C19	112.38 (15)	C22—C19—C9	108.40 (16)
N8—C9—H9	108.3	C19—C20—H20A	109.5
C10—C9—H9	108.3	C19—C20—H20B	109.5
C19—C9—H9	108.3	H20A—C20—H20B	109.5
O11—C10—N12	123.46 (17)	C19—C20—H20C	109.5
O11—C10—C9	121.30 (16)	H20A—C20—H20C	109.5
N12—C10—C9	115.23 (15)	H20B—C20—H20C	109.5
N12—C13—C18	108.88 (17)	C19—C21—H21A	109.5

Bond Angles (°)

N12—C13—C14	110.73 (16)	C19—C21—H21B	109.5
C18—C13—C14	114.94 (17)	H21A—C21—H21B	109.5
N12—C13—H13	107.3	C19—C21—H21C	109.5
C18—C13—H13	107.3	H21A—C21—H21C	109.5
C14—C13—H13	107.3	H21B—C21—H21C	109.5
C15—C14—C16	108.80 (18)	C19—C22—H22A	109.5
C15—C14—C17	108.6 (2)	C19—C22—H22B	109.5
C16—C14—C17	109.8 (2)	H22A—C22—H22B	109.5
C15—C14—C13	108.88 (17)	C19—C22—H22C	109.5
C16—C14—C13	109.04 (18)	H22A—C22—H22C	109.5
C17—C14—C13	111.64 (19)	H22B—C22—H22C	109.5
C14—C15—H15A	109.5	C2—N1—C5	106.52 (16)
C14—C15—H15B	109.5	C2—N1—C6	129.38 (15)
H15A—C15—H15B	109.5	C5—N1—C6	124.04 (16)
C14—C15—H15C	109.5	C2—N3—C4	105.17 (17)
H15A—C15—H15C	109.5	C7—N8—C9	118.07 (16)
H15B—C15—H15C	109.5	C10—N12—C13	124.71 (16)
C14—C16—H16A	109.5	C10—N12—H12	115.1 (17)
C14—C16—H16B	109.5	C13—N12—H12	119.1 (17)
H16A—C16—H16B	109.5		

Torsion Angles (°)

N3—C4—C5—N1	-0.5 (2)	C10—C9—C19—C22	-174.58 (15)
N3—C2—C7—N8	-178.00 (19)	N3—C2—N1—C5	-0.4 (2)
N1—C2—C7—N8	3.9 (3)	C7—C2—N1—C5	177.86 (18)
N8—C9—C10—O11	170.40 (17)	N3—C2—N1—C6	176.74 (19)
C19—C9—C10—O11	-66.7 (2)	C7—C2—N1—C6	-5.0 (3)
N8—C9—C10—N12	-8.7 (2)	C4—C5—N1—C2	0.5 (2)
C19—C9—C10—N12	114.21 (18)	C4—C5—N1—C6	-176.78 (19)
N12—C13—C14—C15	-64.0 (2)	N1—C2—N3—C4	0.1 (2)
C18—C13—C14—C15	172.1 (2)	C7—C2—N3—C4	-178.29 (18)
N12—C13—C14—C16	177.39 (18)	C5—C4—N3—C2	0.3 (2)
C18—C13—C14—C16	53.5 (3)	C2—C7—N8—C9	-179.87 (16)
N12—C13—C14—C17	55.9 (3)	C10—C9—N8—C7	-122.76 (17)
C18—C13—C14—C17	-68.0 (3)	C19—C9—N8—C7	113.26 (18)
N8—C9—C19—C21	-170.77 (17)	O11—C10—N12—C13	-8.3 (3)
C10—C9—C19—C21	67.2 (2)	C9—C10—N12—C13	170.72 (16)
N8—C9—C19—C20	67.4 (2)	C18—C13—N12—C10	-98.8 (2)
C10—C9—C19—C20	-54.6 (2)	C14—C13—N12—C10	133.88 (19)
N8—C9—C19—C22	-52.6 (2)		

L10

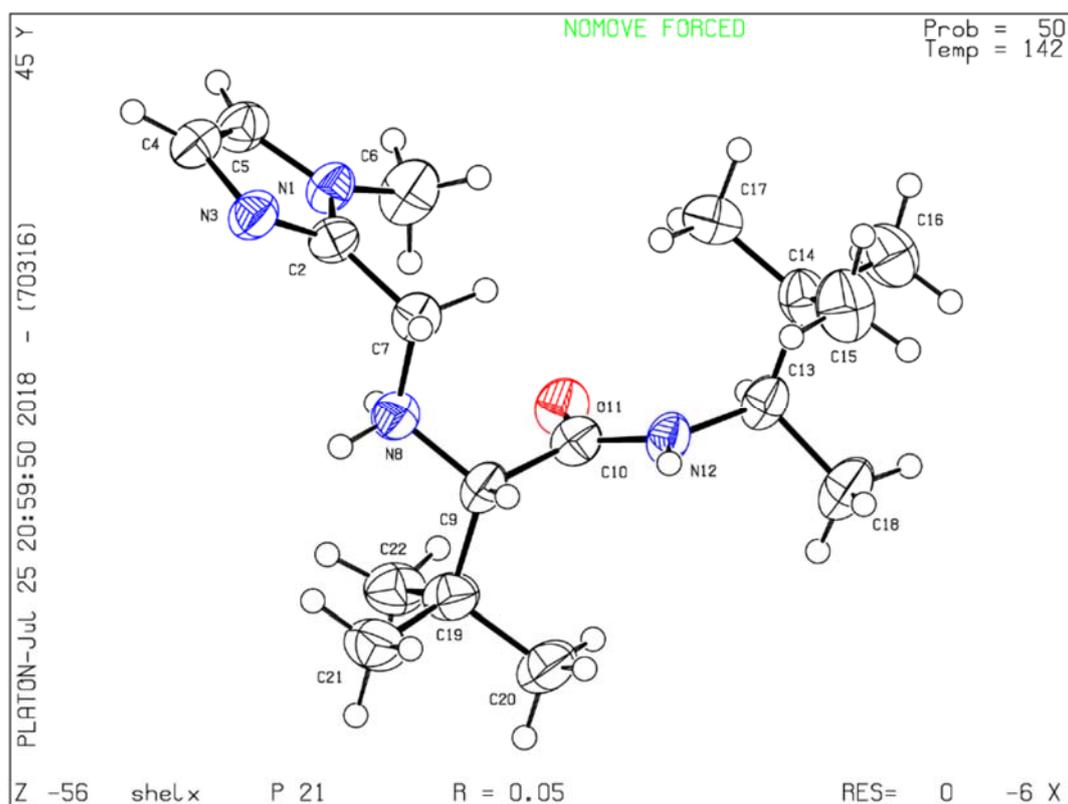


Table A

Experimental details

Crystal data	
Chemical formula	C ₁₇ H ₃₃ N ₄ O
<i>M</i> _r	309.47
Crystal system, space group	Monoclinic, <i>P</i> 2 ₁
Temperature (K)	142
<i>a</i> , <i>b</i> , <i>c</i> (Å)	8.0138 (5), 13.0776 (8), 9.4290 (6)
β (°)	108.996 (4)
<i>V</i> (Å ³)	934.36 (10)
<i>Z</i>	2
Radiation type	Cu <i>K</i> α
μ (mm ⁻¹)	0.55
Crystal size (mm)	0.32 × 0.17 × 0.11
Data collection	
Diffractometer	Bruker D8 VENTURE
Absorption correction	Multi-scan <i>SADABS2016/2</i> - Bruker AXS area detector scaling and absorption correction
<i>T</i> _{min} , <i>T</i> _{max}	0.575, 0.752
No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>) reflections	12600, 2823, 2188
<i>R</i> _{int}	0.085
θ _{max} (°)	61.4
(<i>sin</i> θ / λ) _{max} (Å ⁻¹)	0.570
Refinement	
<i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.053, 0.136, 1.05

Table A

Experimental details

No. of reflections	2823
No. of parameters	212
No. of restraints	1
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\max}, \Delta\rho_{\min}$ (e Å ⁻³)	0.14, -0.36
Absolute structure	Flack x determined using 788 quotients [(I+)-(I-)]/[(I+)+(I-)] (Parsons, Flack and Wagner, Acta Cryst. B69 (2013) 249-259).
Absolute structure parameter	0.2 (3)

Computer programs: Bruker *APEX3* software, *SAINTE* V8.38A integration software, *SORTAV* (Blessing, 1995), *SHELXS2013* (Sheldrick, 2008), *SHELXL2018/3* (Sheldrick, 2018), *ORTEP* for Windows (Farrugia, 2012), *WinGX* publication routines (Farrugia, 2012).

Table B

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C5—H5 \cdots O11 ⁱ	0.95	2.54	3.461 (6)	162
N12—H12 \cdots N3 ⁱⁱ	1.00 (6)	2.12 (6)	3.112 (6)	173 (5)

Symmetry codes: (i) $-x, y+1/2, -z$; (ii) $-x+1, y-1/2, -z+1$.

Computing details

Data collection: Bruker *APEX3* software; cell refinement: *SAINTE* V8.38A integration software; data reduction: *SORTAV* (Blessing, 1995); program(s) used to solve structure: *SHELXS2013* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2018); molecular graphics: *ORTEP* for Windows (Farrugia, 2012); software used to prepare material for publication: *WinGX* publication routines (Farrugia, 2012).

(shelx)

Crystal data

C₁₇H₃₃N₄O
M_r = 309.47
 Monoclinic, *P*2₁
 Hall symbol: P2yb
a = 8.0138 (5) Å
b = 13.0776 (8) Å
c = 9.4290 (6) Å
 β = 108.996 (4)°
V = 934.36 (10) Å³
Z = 2

F(000) = 342
D_x = 1.1 Mg m⁻³
 Cu *K* α radiation, λ = 1.54178 Å
 Cell parameters from 4408 reflections
 θ = 5.0–61.0°
 μ = 0.55 mm⁻¹
T = 142 K
 Needle, yellow
 0.32 × 0.17 × 0.11 mm

Data collection

Bruker D8 VENTURE
 diffractometer
 Radiation source: sealed x-ray microsource
 Graphite monochromator
 φ or ω oscillation scans
 Absorption correction: multi-scan
SADABS2016/2 - Bruker AXS area detector scaling
 and absorption correction
T_{min} = 0.575, *T_{max}* = 0.752

12600 measured reflections
 2823 independent
 reflections 2188
 reflections with $I > 2\sigma(I)$
R_{int} = 0.085
 θ_{\max} = 61.4°, θ_{\min} = 5.0°
h = -9→9
k = -14→14
l = -10→10

Refinement

Refinement on F^2	H atoms treated by a mixture of independent and constrained refinement
Least-squares matrix: full	
$R[F^2 > 2\sigma(F^2)] = 0.053$	$w = 1/[\sigma^2(F_o^2) + (0.0604P)^2 + 0.2044P]$
$wR(F^2) = 0.136$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.05$	$(\Delta/\sigma)_{\max} < 0.001$
2823 reflections	$\Delta\rho_{\max} = 0.14 \text{ e } \text{\AA}^{-3}$
212 parameters	$\Delta\rho_{\min} = -0.36 \text{ e } \text{\AA}^{-3}$
1 restraint	Extinction correction: <i>SHELXL2018/3</i> (Sheldrick 2018),
0 constraints	$F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Primary atom site location: dual	Extinction coefficient: 0.015 (2)
Secondary atom site location: difference Fourier map	Absolute structure: Flack x determined using 788 quotients
Hydrogen site location: mixed	$[(I^+)-(I^-)]/[(I^+)+(I^-)]$
	(Parsons, Flack and Wagner, Acta Cryst. B69 (2013) 249-259).
	Absolute structure parameter: 0.2 (3)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C2	0.3348 (6)	0.6980 (4)	0.3103 (5)	0.0382 (11)
C4	0.3312 (7)	0.8611 (4)	0.2862 (6)	0.0462 (13)
H4	0.35655	0.931443	0.307661	0.055*
C5	0.2177 (7)	0.8236 (4)	0.1544 (6)	0.0455 (13)
H5	0.150157	0.861998	0.069571	0.055*
C6	0.1311 (8)	0.6446 (4)	0.0569 (6)	0.0610 (16)
H6A	0.071904	0.679762	-0.038017	0.092*
H6B	0.043321	0.608073	0.089694	0.092*
H6C	0.216946	0.595744	0.04262	0.092*
C7	0.3643 (6)	0.5916 (4)	0.3689 (5)	0.0402 (12)
H7A	0.375645	0.545345	0.289401	0.048*
H7B	0.475741	0.588434	0.454325	0.048*
C9	0.2174 (7)	0.4479 (3)	0.4451 (5)	0.0399 (12)
H9	0.33545	0.428946	0.517268	0.048*
C10	0.1924 (7)	0.3884 (4)	0.2987 (5)	0.0411 (12)
C13	0.2989 (7)	0.2483 (3)	0.1769 (6)	0.0441 (13)
H13	0.182689	0.263213	0.099002	0.053*
C14	0.4412 (6)	0.2703 (4)	0.1042 (5)	0.0466 (12)
C15	0.6244 (8)	0.2388 (5)	0.2048 (7)	0.0696 (18)
H15A	0.712189	0.259567	0.158441	0.104*
H15B	0.628447	0.164366	0.217997	0.104*
H15C	0.650458	0.272049	0.30275	0.104*
C16	0.3938 (8)	0.2135 (5)	-0.0458 (6)	0.0645 (17)
H16A	0.274134	0.232295	-0.108318	0.097*
H16B	0.399434	0.139552	-0.027595	0.097*
H16C	0.477638	0.232147	-0.097413	0.097*
C17	0.4435 (8)	0.3847 (4)	0.0725 (6)	0.0594 (16)
H17A	0.489048	0.422025	0.167537	0.089*
H17B	0.323367	0.407958	0.01827	0.089*
H17C	0.519617	0.397686	0.011484	0.089*
C18	0.2916 (9)	0.1377 (4)	0.2241 (6)	0.0618 (16)

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{iso} */ <i>U</i> _{eq}
H18A	0.277223	0.092684	0.137794	0.093*
H18B	0.191415	0.128578	0.260932	0.093*
H18C	0.401311	0.120407	0.303971	0.093*
C19	0.0762 (7)	0.4195 (4)	0.5182 (5)	0.0473 (13)
C20	0.0639 (8)	0.3039 (4)	0.5296 (7)	0.0593 (15)
H20A	-0.015704	0.286892	0.586023	0.089*
H20B	0.181366	0.275884	0.581448	0.089*
H20C	0.018026	0.274557	0.428684	0.089*
C21	0.1366 (9)	0.4653 (5)	0.6773 (6)	0.0627 (17)
H21A	0.141411	0.540003	0.670733	0.094*
H21B	0.254068	0.439047	0.733583	0.094*
H21C	0.052905	0.44613	0.728686	0.094*
C22	-0.1051 (7)	0.4627 (5)	0.4306 (6)	0.0557 (15)
H22A	-0.193409	0.435466	0.472006	0.084*
H22B	-0.135983	0.442913	0.32489	0.084*
H22C	-0.102243	0.537427	0.438568	0.084*
N1	0.2217 (5)	0.7193 (3)	0.1702 (5)	0.0412 (10)
N3	0.4031 (5)	0.7818 (3)	0.3831 (4)	0.0442 (10)
N8	0.2189 (5)	0.5574 (3)	0.4178 (5)	0.0420 (11)
H8A	0.22442	0.591095	0.503666	0.05*
H8B	0.115367	0.57492	0.346681	0.05*
N12	0.3115 (6)	0.3156 (3)	0.3053 (5)	0.0415 (10)
O11	0.0704 (4)	0.4108 (3)	0.1833 (4)	0.0496 (10)
H12	0.410 (8)	0.308 (5)	0.402 (7)	0.080 (19)*

Atomic displacement parameters (\AA^2)

	<i>U</i> ¹¹	<i>U</i> ²²	<i>U</i> ³³	<i>U</i> ¹²	<i>U</i> ¹³	<i>U</i> ²³
C2	0.037 (3)	0.038 (3)	0.037 (3)	-0.002 (2)	0.009 (2)	0.003 (2)
C4	0.054 (3)	0.036 (3)	0.050 (3)	0.000 (2)	0.018 (3)	0.006 (2)
C5	0.047 (3)	0.043 (3)	0.044 (3)	0.007 (2)	0.011 (2)	0.011 (2)
C6	0.070 (4)	0.053 (4)	0.043 (3)	-0.002 (3)	-0.004 (3)	-0.002 (3)
C7	0.040 (3)	0.040 (3)	0.038 (3)	0.000 (2)	0.009 (2)	0.000 (2)
C9	0.048 (3)	0.030 (3)	0.036 (3)	0.000 (2)	0.005 (2)	0.001 (2)
C10	0.042 (3)	0.039 (3)	0.039 (3)	-0.003 (2)	0.008 (2)	0.001 (2)
C13	0.052 (3)	0.032 (3)	0.040 (3)	0.000 (2)	0.004 (2)	-0.001 (2)
C14	0.048 (3)	0.048 (3)	0.040 (3)	0.005 (3)	0.010 (2)	-0.005 (2)
C15	0.064 (4)	0.073 (5)	0.064 (4)	0.012 (3)	0.010 (3)	-0.008 (3)
C16	0.073 (4)	0.073 (5)	0.046 (3)	-0.001 (3)	0.018 (3)	-0.012 (3)
C17	0.068 (4)	0.061 (4)	0.059 (3)	-0.004 (3)	0.035 (3)	0.003 (3)
C18	0.090 (4)	0.036 (3)	0.048 (3)	-0.002 (3)	0.007 (3)	0.001 (3)
C19	0.052 (3)	0.047 (3)	0.043 (3)	-0.010 (3)	0.016 (2)	0.001 (2)
C20	0.073 (4)	0.043 (4)	0.061 (4)	-0.010 (3)	0.021 (3)	0.006 (3)
C21	0.077 (4)	0.068 (4)	0.050 (3)	-0.016 (3)	0.030 (3)	-0.007 (3)
C22	0.047 (3)	0.061 (4)	0.061 (4)	-0.004 (3)	0.021 (3)	0.006 (3)
N1	0.044 (2)	0.034 (3)	0.041 (2)	0.0030 (18)	0.0081 (19)	0.0042 (18)
N3	0.052 (2)	0.035 (2)	0.045 (2)	-0.005 (2)	0.0149 (19)	0.000 (2)
N8	0.046 (3)	0.037 (3)	0.043 (2)	-0.0019 (18)	0.015 (2)	-0.0006 (18)
N12	0.052 (3)	0.030 (2)	0.037 (2)	0.001 (2)	0.009 (2)	-0.0033 (17)
O11	0.051 (2)	0.048 (2)	0.041 (2)	0.0042 (17)	0.0036 (17)	0.0014 (17)

Geometric parameters

Bond lengths (\AA)			
C2—N3	1.314 (6)	C15—H15A	0.98
C2—N1	1.366 (6)	C15—H15B	0.98

Bond lengths (Å)

C2—C7	1.488 (7)	C15—H15C	0.98
C4—C5	1.369 (7)	C16—H16A	0.98
C4—N3	1.378 (6)	C16—H16B	0.98
C4—H4	0.95	C16—H16C	0.98
C5—N1	1.372 (6)	C17—H17A	0.98
C5—H5	0.95	C17—H17B	0.98
C6—N1	1.455 (6)	C17—H17C	0.98
C6—H6A	0.98	C18—H18A	0.98
C6—H6B	0.98	C18—H18B	0.98
C6—H6C	0.98	C18—H18C	0.98
C7—N8	1.456 (6)	C19—C20	1.520 (8)
C7—H7A	0.99	C19—C22	1.527 (7)
C7—H7B	0.99	C19—C21	1.540 (7)
C9—N8	1.456 (6)	C20—H20A	0.98
C9—C10	1.539 (7)	C20—H20B	0.98
C9—C19	1.550 (7)	C20—H20C	0.98
C9—H9	1	C21—H21A	0.98
C10—O11	1.238 (5)	C21—H21B	0.98
C10—N12	1.335 (6)	C21—H21C	0.98
C13—N12	1.473 (6)	C22—H22A	0.98
C13—C18	1.521 (7)	C22—H22B	0.98
C13—C14	1.537 (7)	C22—H22C	0.98
C13—H13	1	N8—H8A	0.91
C14—C15	1.522 (7)	N8—H8B	0.91
C14—C17	1.528 (8)	N12—H12	1.00 (6)
C14—C16	1.532 (7)		

Bond Angles (°)

N3—C2—N1	111.6 (4)	H16A—C16—H16C	109.5
N3—C2—C7	126.7 (4)	H16B—C16—H16C	109.5
N1—C2—C7	121.7 (4)	C14—C17—H17A	109.5
C5—C4—N3	110.1 (5)	C14—C17—H17B	109.5
C5—C4—H4	124.9	H17A—C17—H17B	109.5
N3—C4—H4	124.9	C14—C17—H17C	109.5
C4—C5—N1	105.8 (5)	H17A—C17—H17C	109.5
C4—C5—H5	127.1	H17B—C17—H17C	109.5
N1—C5—H5	127.1	C13—C18—H18A	109.5
N1—C6—H6A	109.5	C13—C18—H18B	109.5
N1—C6—H6B	109.5	H18A—C18—H18B	109.5
H6A—C6—H6B	109.5	C13—C18—H18C	109.5
N1—C6—H6C	109.5	H18A—C18—H18C	109.5
H6A—C6—H6C	109.5	H18B—C18—H18C	109.5
H6B—C6—H6C	109.5	C20—C19—C22	109.5 (5)
N8—C7—C2	111.0 (4)	C20—C19—C21	109.0 (5)
N8—C7—H7A	109.4	C22—C19—C21	109.0 (5)
C2—C7—H7A	109.4	C20—C19—C9	110.1 (4)
N8—C7—H7B	109.4	C22—C19—C9	112.0 (4)
C2—C7—H7B	109.4	C21—C19—C9	107.2 (4)
H7A—C7—H7B	108	C19—C20—H20A	109.5
N8—C9—C10	110.2 (4)	C19—C20—H20B	109.5
N8—C9—C19	111.3 (4)	H20A—C20—H20B	109.5
C10—C9—C19	112.1 (4)	C19—C20—H20C	109.5
N8—C9—H9	107.7	H20A—C20—H20C	109.5
C10—C9—H9	107.7	H20B—C20—H20C	109.5
C19—C9—H9	107.7	C19—C21—H21A	109.5
O11—C10—N12	123.8 (5)	C19—C21—H21B	109.5

Bond Angles (°)

O11—C10—C9	120.3 (5)	H21A—C21—H21B	109.5
N12—C10—C9	115.8 (4)	C19—C21—H21C	109.5
N12—C13—C18	109.1 (4)	H21A—C21—H21C	109.5
N12—C13—C14	113.1 (4)	H21B—C21—H21C	109.5
C18—C13—C14	114.4 (5)	C19—C22—H22A	109.5
N12—C13—H13	106.6	C19—C22—H22B	109.5
C18—C13—H13	106.6	H22A—C22—H22B	109.5
C14—C13—H13	106.6	C19—C22—H22C	109.5
C15—C14—C17	108.5 (5)	H22A—C22—H22C	109.5
C15—C14—C16	110.0 (5)	H22B—C22—H22C	109.5
C17—C14—C16	108.1 (4)	C2—N1—C5	107.0 (4)
C15—C14—C13	112.1 (5)	C2—N1—C6	126.1 (4)
C17—C14—C13	109.2 (4)	C5—N1—C6	126.8 (4)
C16—C14—C13	108.9 (4)	C2—N3—C4	105.6 (4)
C14—C15—H15A	109.5	C9—N8—C7	114.3 (4)
C14—C15—H15B	109.5	C9—N8—H8A	108.7
H15A—C15—H15B	109.5	C7—N8—H8A	108.7
C14—C15—H15C	109.5	C9—N8—H8B	108.7
H15A—C15—H15C	109.5	C7—N8—H8B	108.7
H15B—C15—H15C	109.5	H8A—N8—H8B	107.6
C14—C16—H16A	109.5	C10—N12—C13	122.1 (4)
C14—C16—H16B	109.5	C10—N12—H12	117 (4)
H16A—C16—H16B	109.5	C13—N12—H12	120 (4)
C14—C16—H16C	109.5		

Torsion Angles (°)

N3—C4—C5—N1	-0.7 (6)	C10—C9—C19—C21	168.3 (4)
N3—C2—C7—N8	99.0 (5)	N3—C2—N1—C5	-0.5 (6)
N1—C2—C7—N8	-77.9 (5)	C7—C2—N1—C5	176.8 (5)
N8—C9—C10—O11	-50.8 (6)	N3—C2—N1—C6	176.3 (5)
C19—C9—C10—O11	73.7 (6)	C7—C2—N1—C6	-6.3 (7)
N8—C9—C10—N12	127.4 (5)	C4—C5—N1—C2	0.7 (6)
C19—C9—C10—N12	-108.1 (5)	C4—C5—N1—C6	-176.1 (5)
N12—C13—C14—C15	70.0 (6)	N1—C2—N3—C4	0.1 (5)
C18—C13—C14—C15	-55.8 (6)	C7—C2—N3—C4	-177.1 (5)
N12—C13—C14—C17	-50.2 (5)	C5—C4—N3—C2	0.4 (6)
C18—C13—C14—C17	-176.0 (4)	C10—C9—N8—C7	-63.2 (5)
N12—C13—C14—C16	-168.0 (4)	C19—C9—N8—C7	171.8 (4)
C18—C13—C14—C16	66.2 (6)	C2—C7—N8—C9	167.0 (4)
N8—C9—C19—C20	173.8 (4)	O11—C10—N12—C13	-5.5 (8)
C10—C9—C19—C20	49.9 (6)	C9—C10—N12—C13	176.3 (4)
N8—C9—C19—C22	51.7 (5)	C18—C13—N12—C10	-121.9 (5)
C10—C9—C19—C22	-72.2 (5)	C14—C13—N12—C10	109.5 (5)
N8—C9—C19—C21	-67.8 (5)		
