

Supplemental Material

**Facile synthesis of new *N*-(aminocycloalkylene)amino acid compounds
using chiral triflate esters with *N*-Boc-aminopyrrolidines and *N*-Boc-
aminopiperidines**

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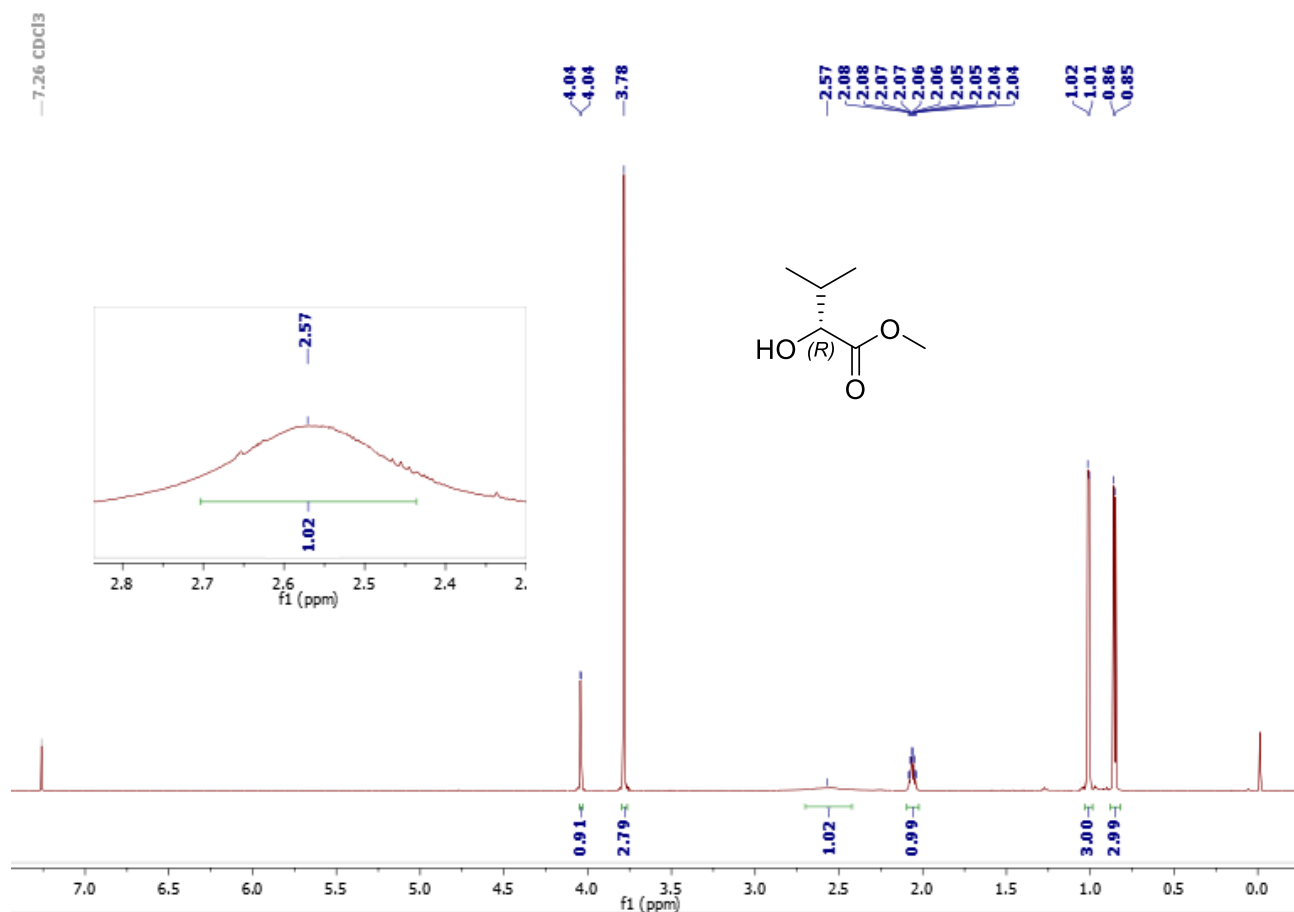


Figure S1. Methyl (2R)-2-hydroxy-3-methylbutanoate ((R)-1b). ¹H NMR spectrum (700 MHz, CDCl₃).

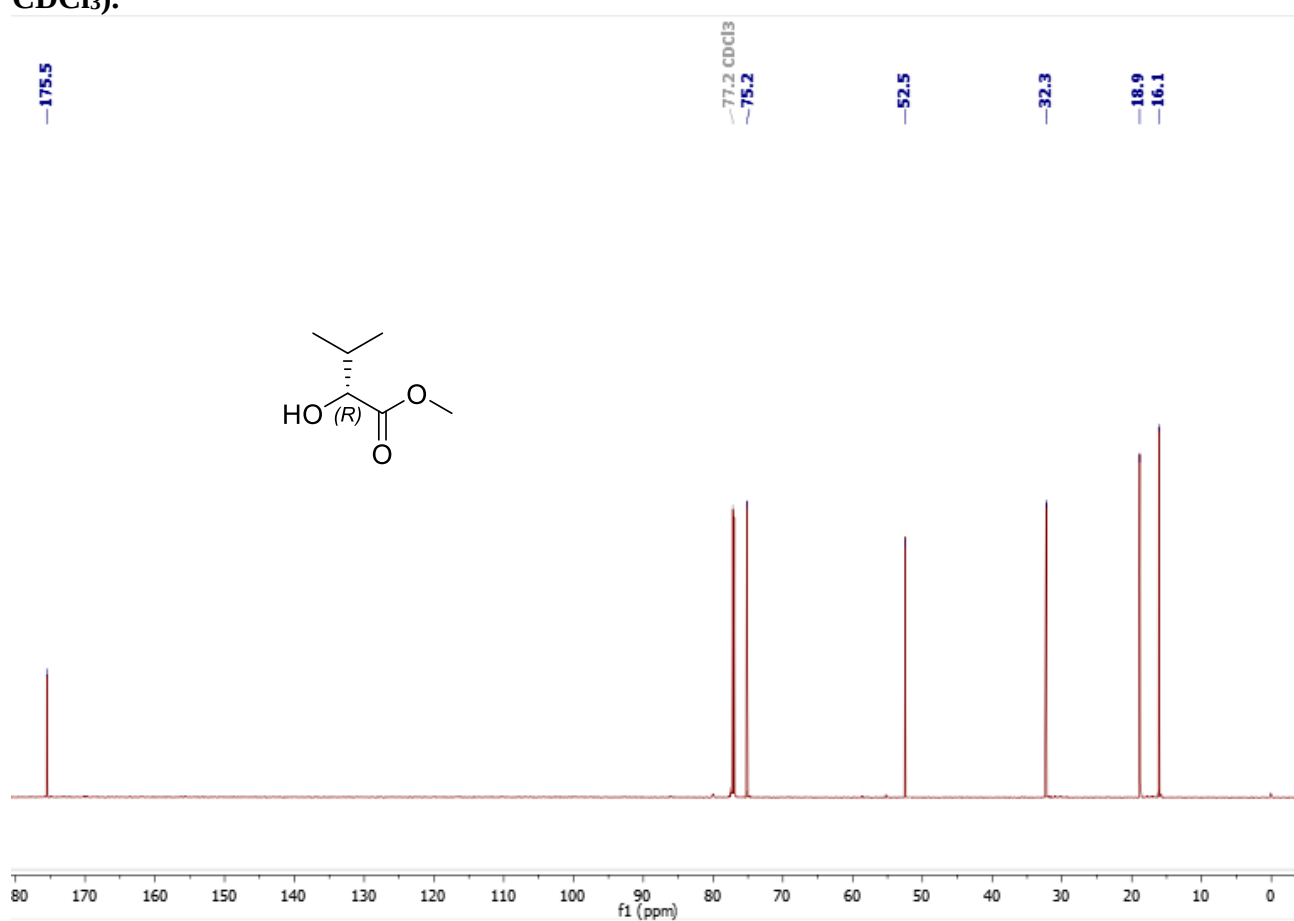


Figure S2. Methyl (2R)-2-hydroxy-3-methylbutanoate ((R)-1b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Compound Spectrum SmartFormula Report

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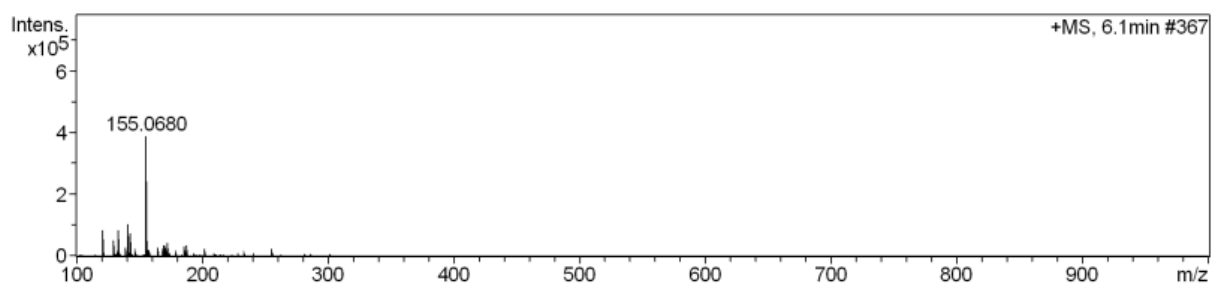
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Instrument micrOTOF-Q III 8228888.20448

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#	RT [min]	Area	Int. Type	I	S/N	Chromatogram	Max. m/z	FWHM [min]
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+MS, 6.1min #367



Meas. m/z	#	Ion Formula	m/z	err [ppm]	mSigma	# Sigma	Score	rdb	e ⁻ Conf	N-Rule
155.0680	1	C6H12NaO3	155.0679	-0.8	25.6	1	100.00	0.5	even	ok

Figure S3. Methyl (2*R*)-2-hydroxy-3-methylbutanoate ((*R*)-1b). HRMS (ESI-TOF).

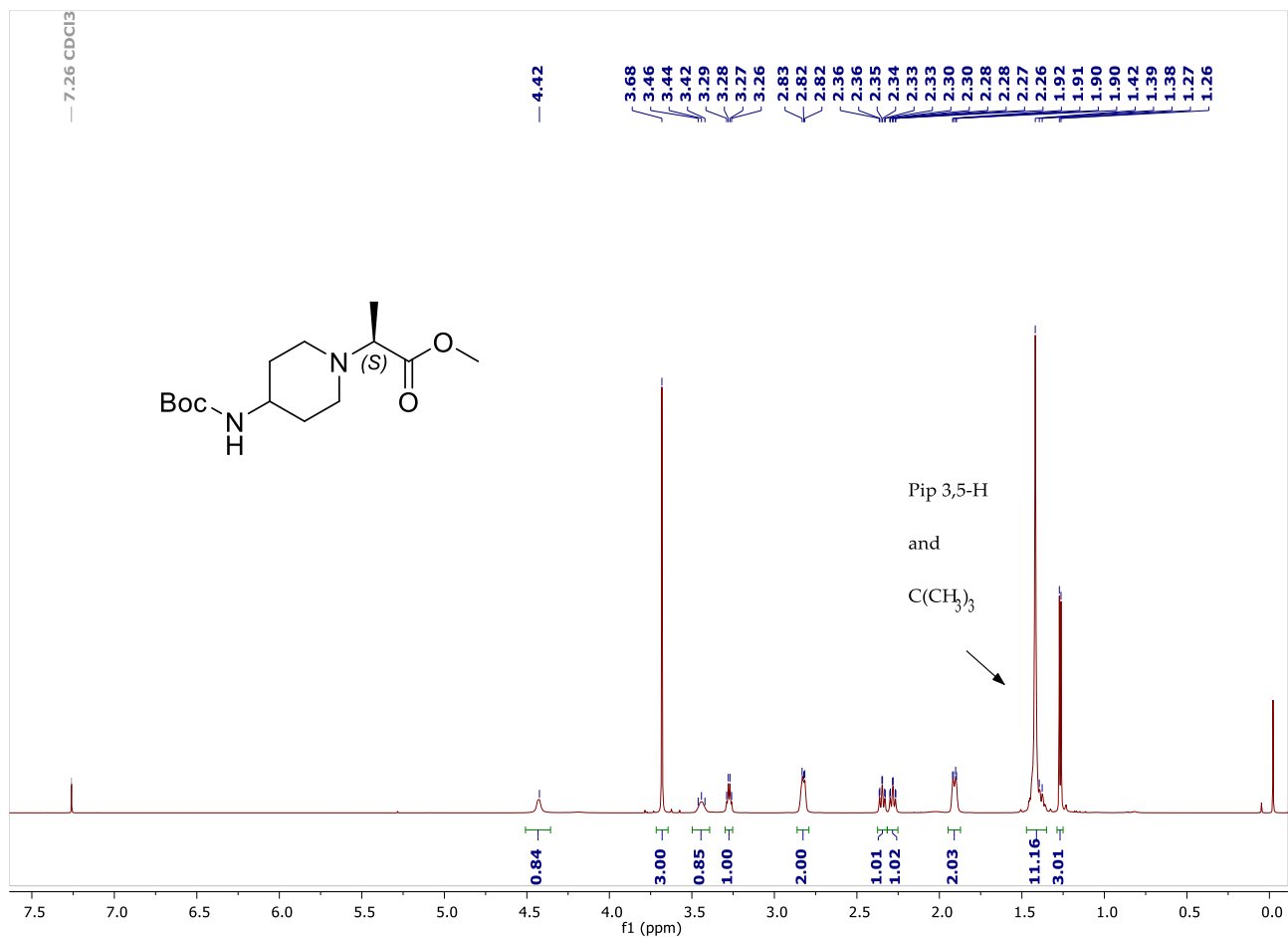


Figure S4. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((S)-3a). ¹H NMR spectrum (700 MHz, CDCl₃).

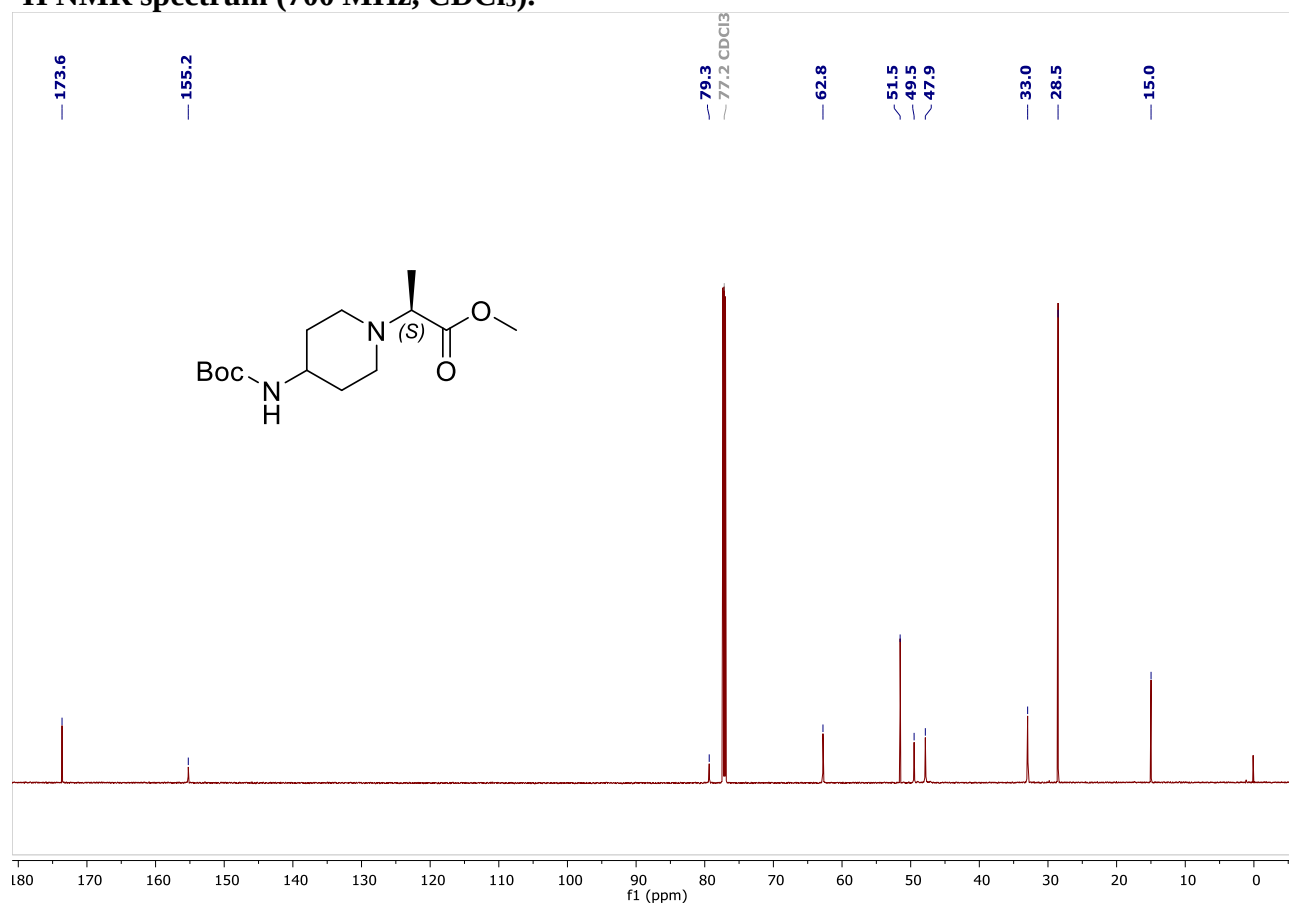


Figure S5. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((S)-3a). ¹³C NMR spectrum (176 MHz, CDCl₃).

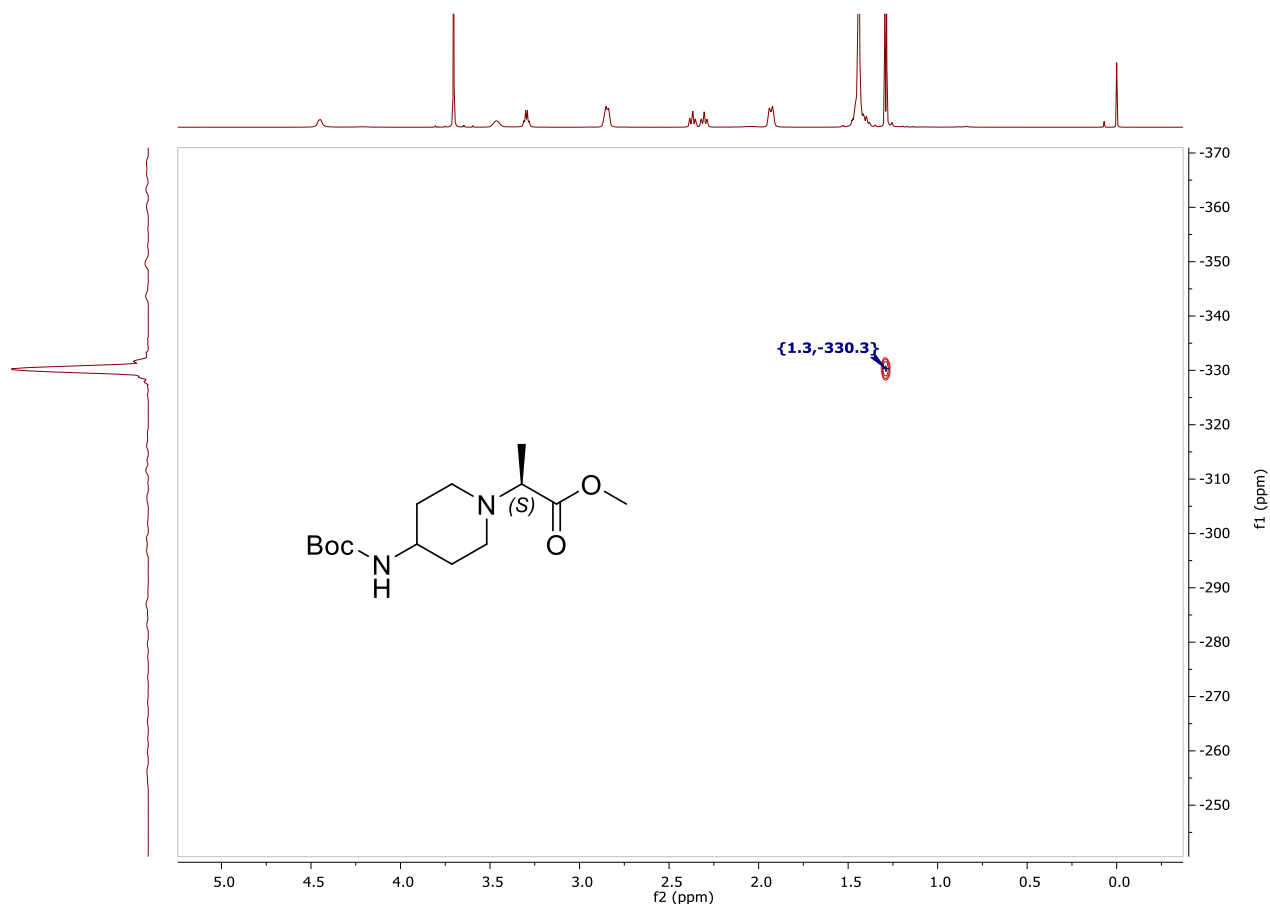


Figure S6. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((S)-3a). ^1H - ^{15}N HMBC spectrum (71 MHz, CDCl_3).

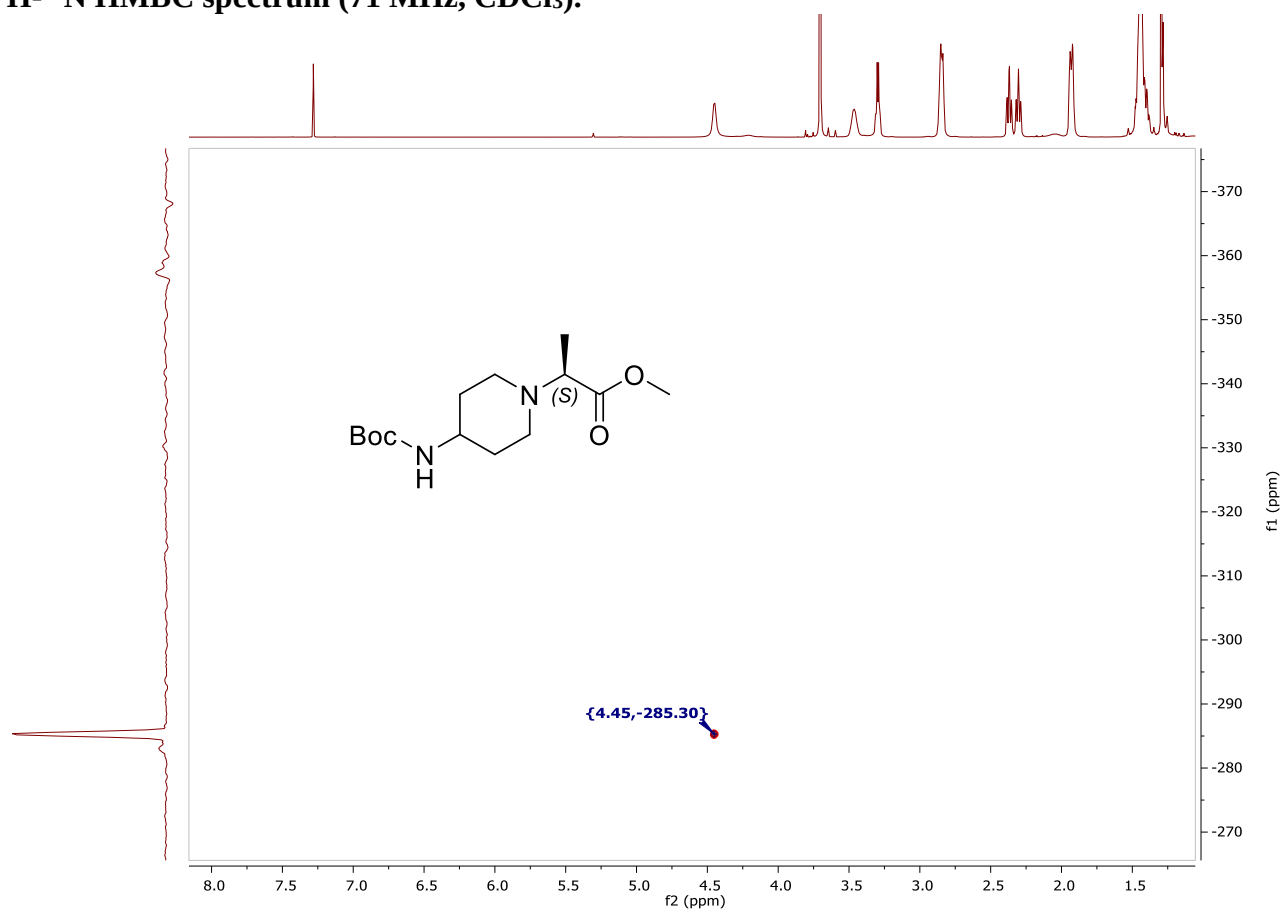


Figure S7. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((S)-3a). ^1H - ^{15}N HSQC spectrum (71 MHz, CDCl_3).

Mass Spectrum SmartFormula Report

Analysis Info

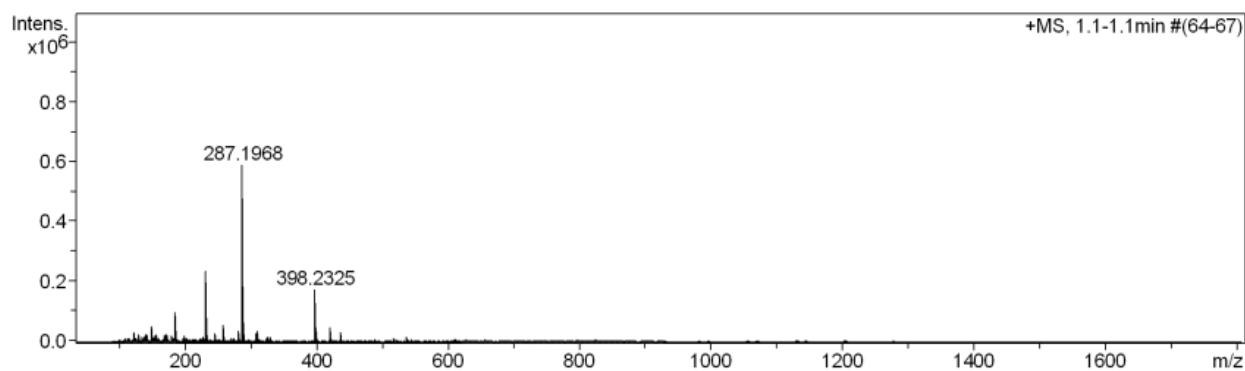
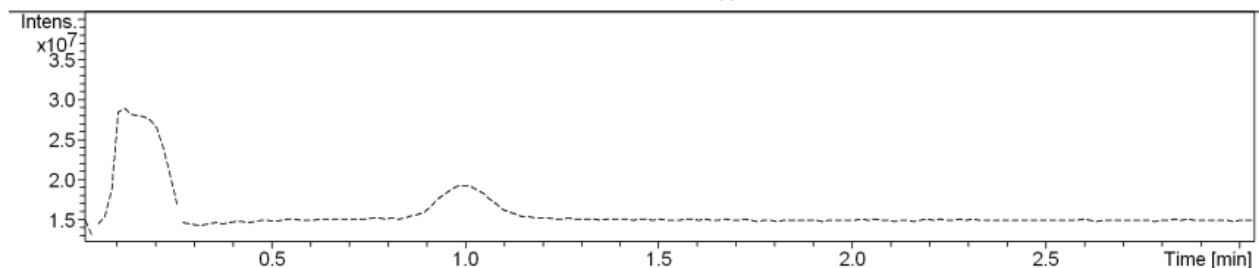
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Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

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Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



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287.1968	1	C 14 H 27 N 2 O 4	100.00	287.1965	-0.8	-0.5	15.5	2.5	even	ok

Figure S8. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((S)-3a). HRMS (ESI-TOF).

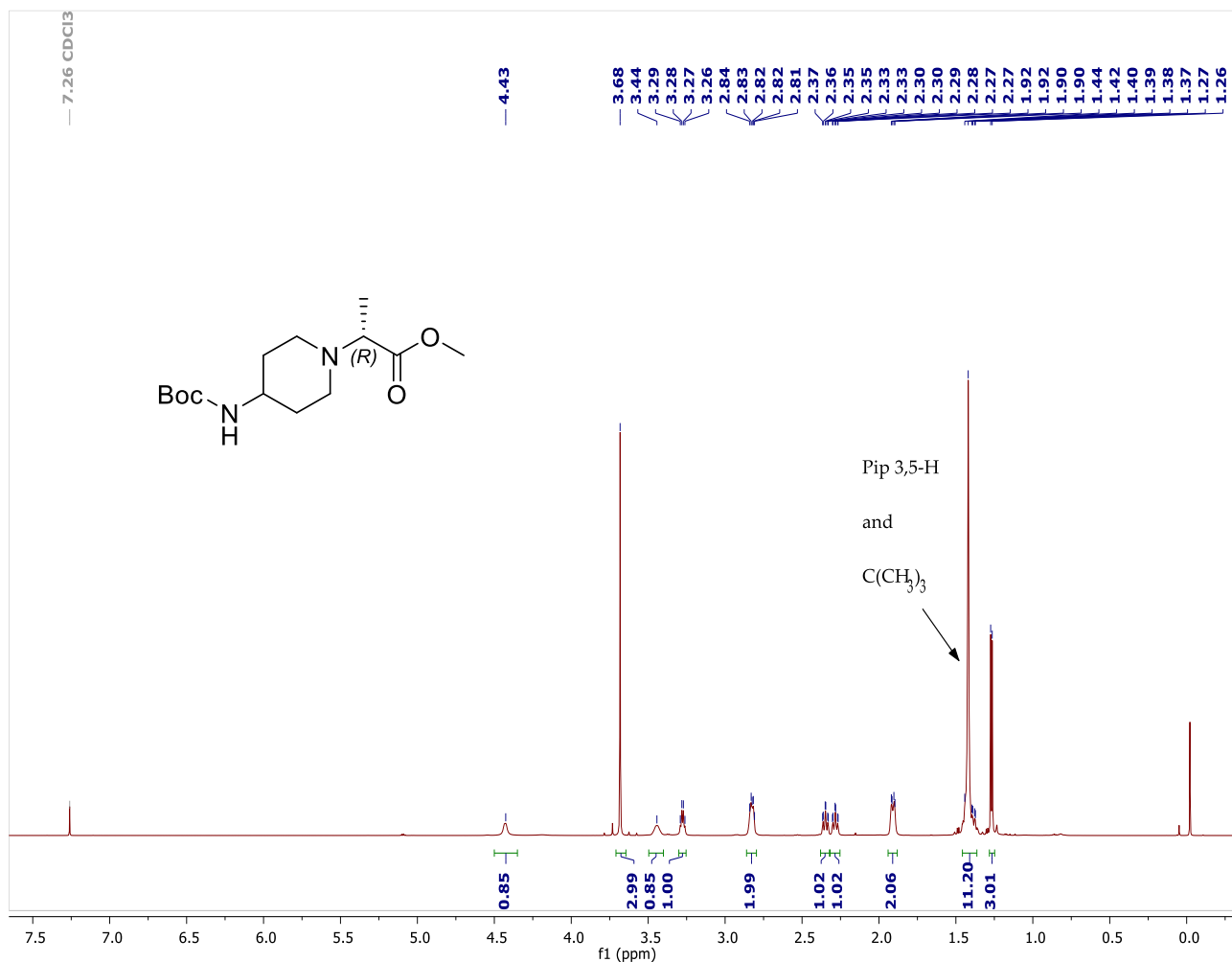


Figure S9. Methyl (2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((*R*)-3a). ¹H NMR spectrum (700 MHz, CDCl₃).

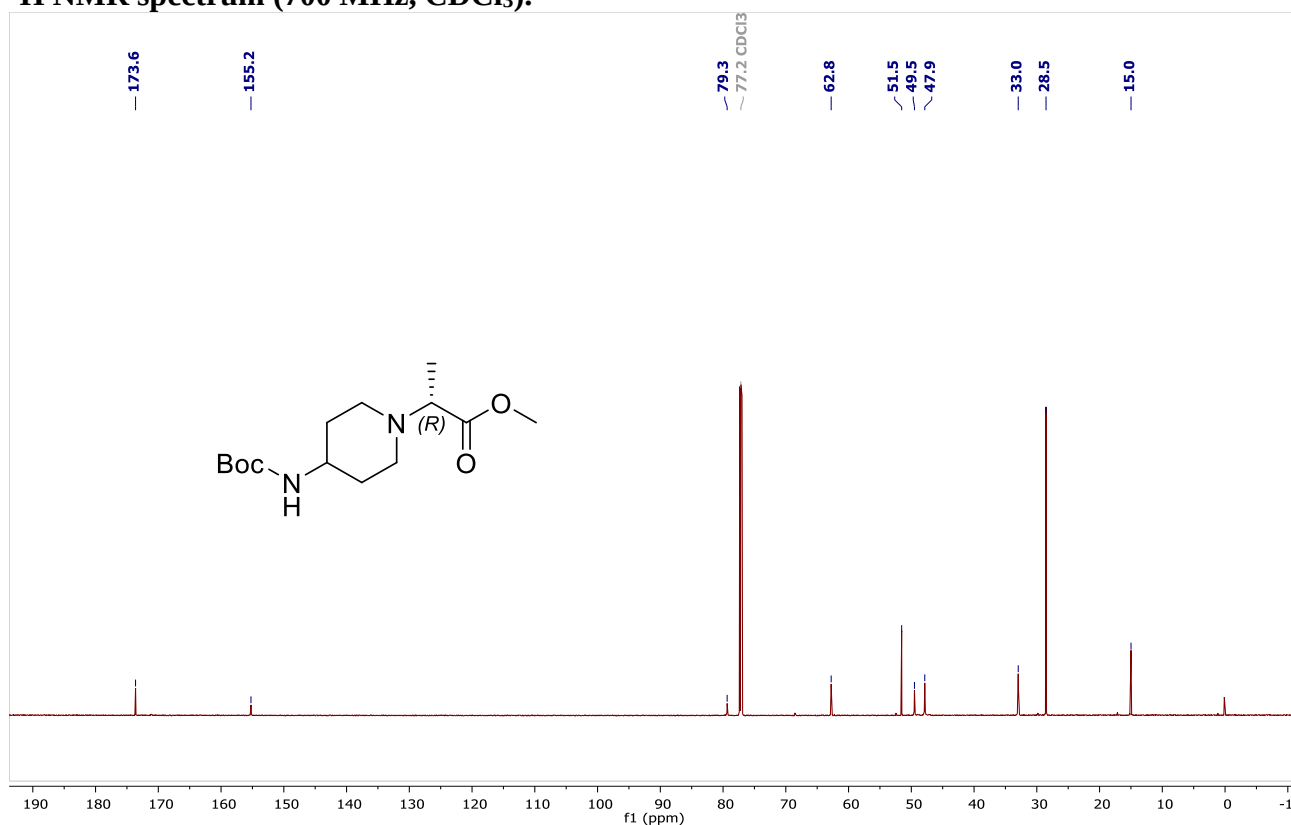


Figure S10. Methyl (2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((*R*)-3a). ¹³C NMR spectrum (176 MHz, CDCl₃).

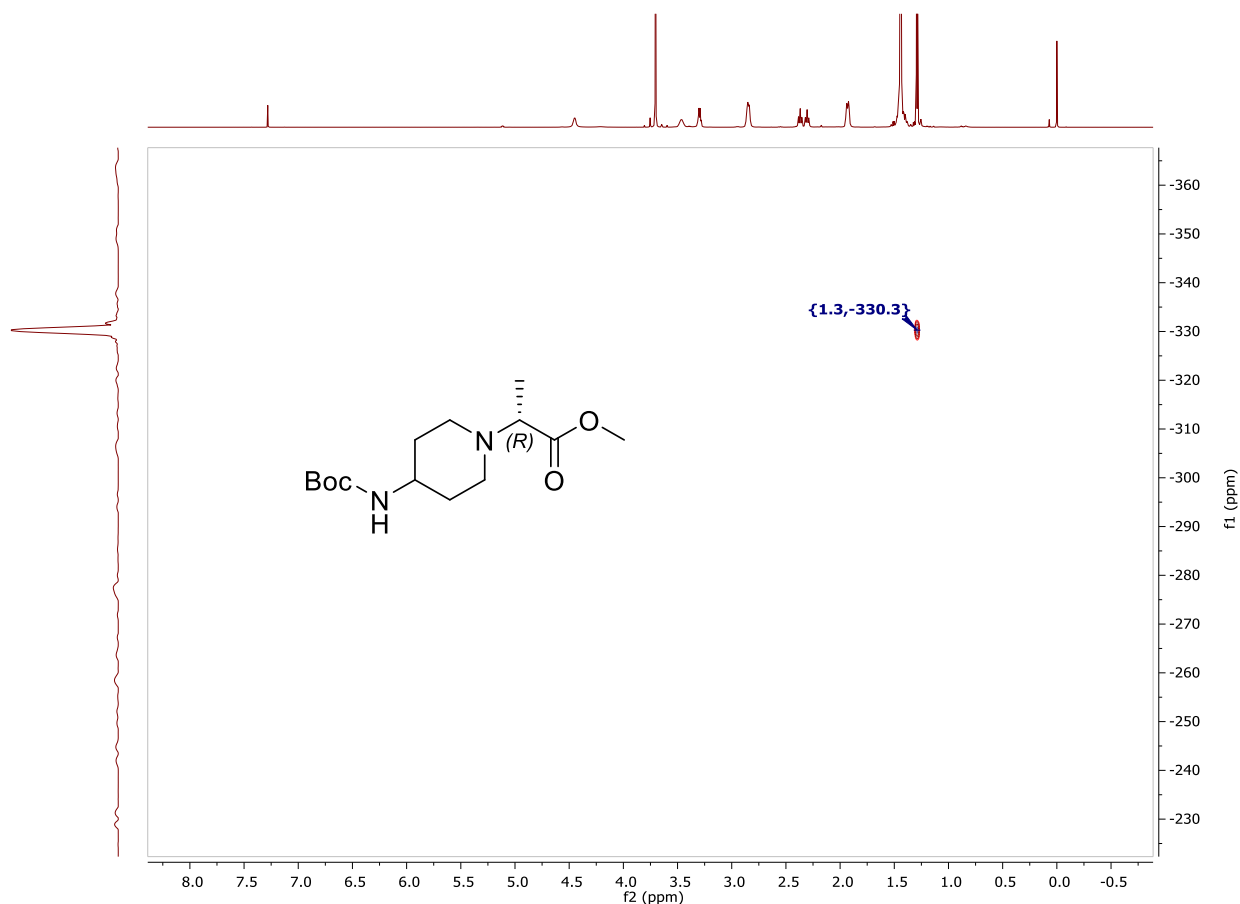


Figure S11. Methyl (2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((*R*)-3a). ^1H - ^{15}N HMBC spectrum (71 MHz, CDCl_3).

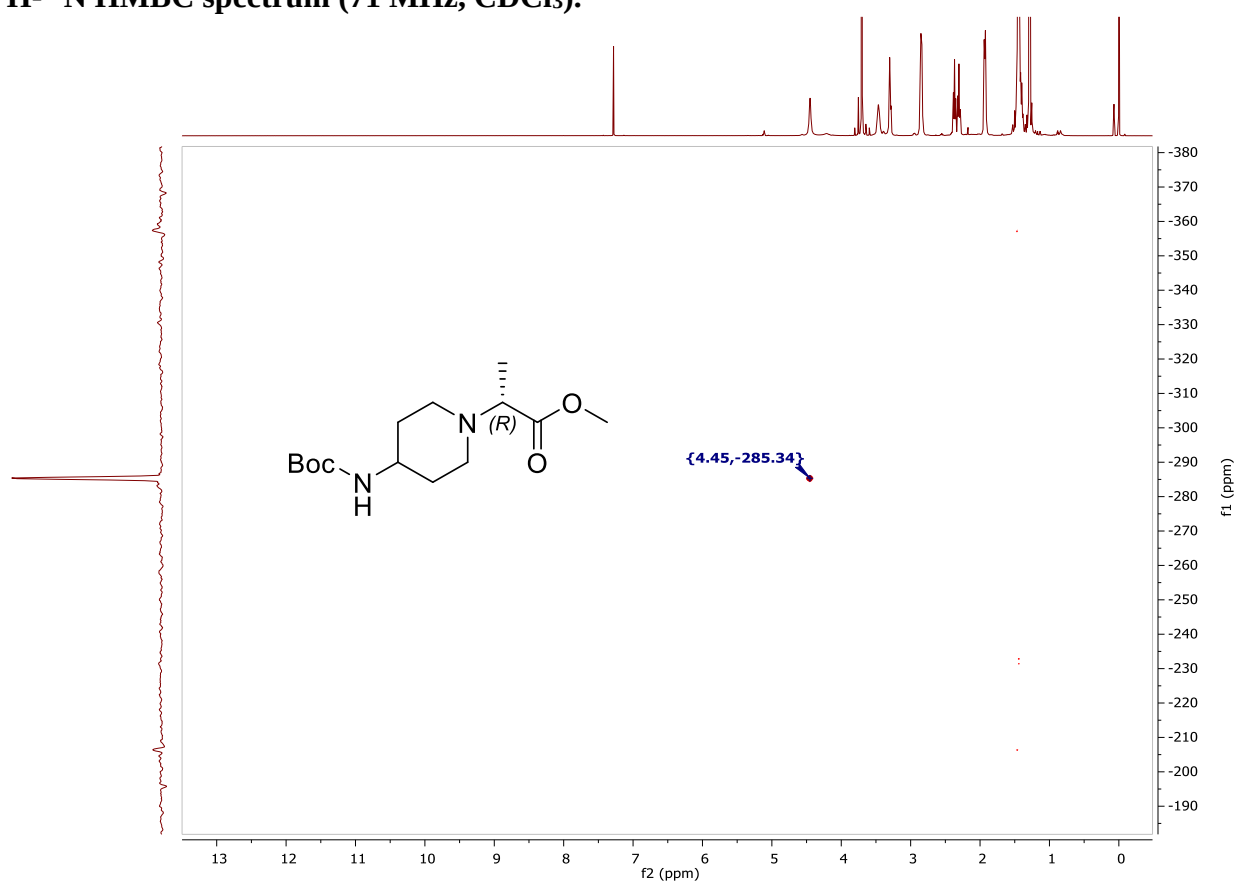


Figure S12. Methyl (2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((*R*)-3a). ^1H - ^{15}N HSQC spectrum (71 MHz, CDCl_3).

Mass Spectrum SmartFormula Report

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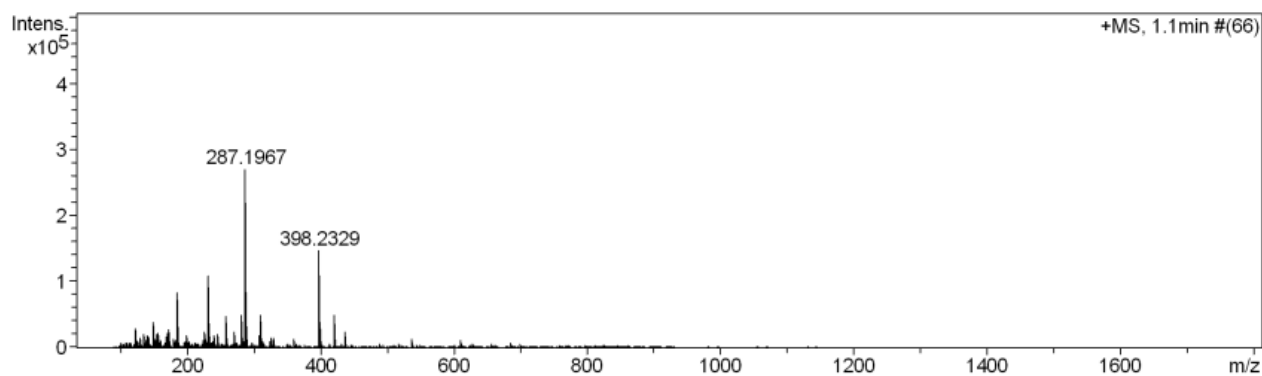
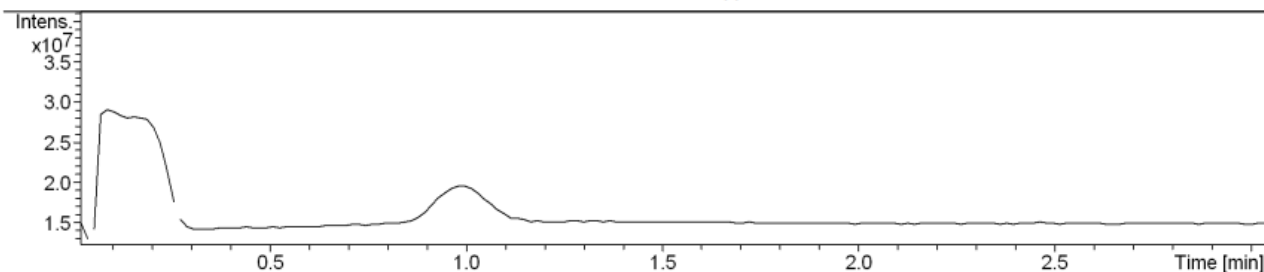
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Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

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Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
287.1967	1	C 14 H 27 N 2 O 4	100.00	287.1965	-0.6	-0.4	12.2	2.5	even	ok

Figure S13. Methyl (2R)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoate ((*R*)-3a). HRMS (ESI-TOF).

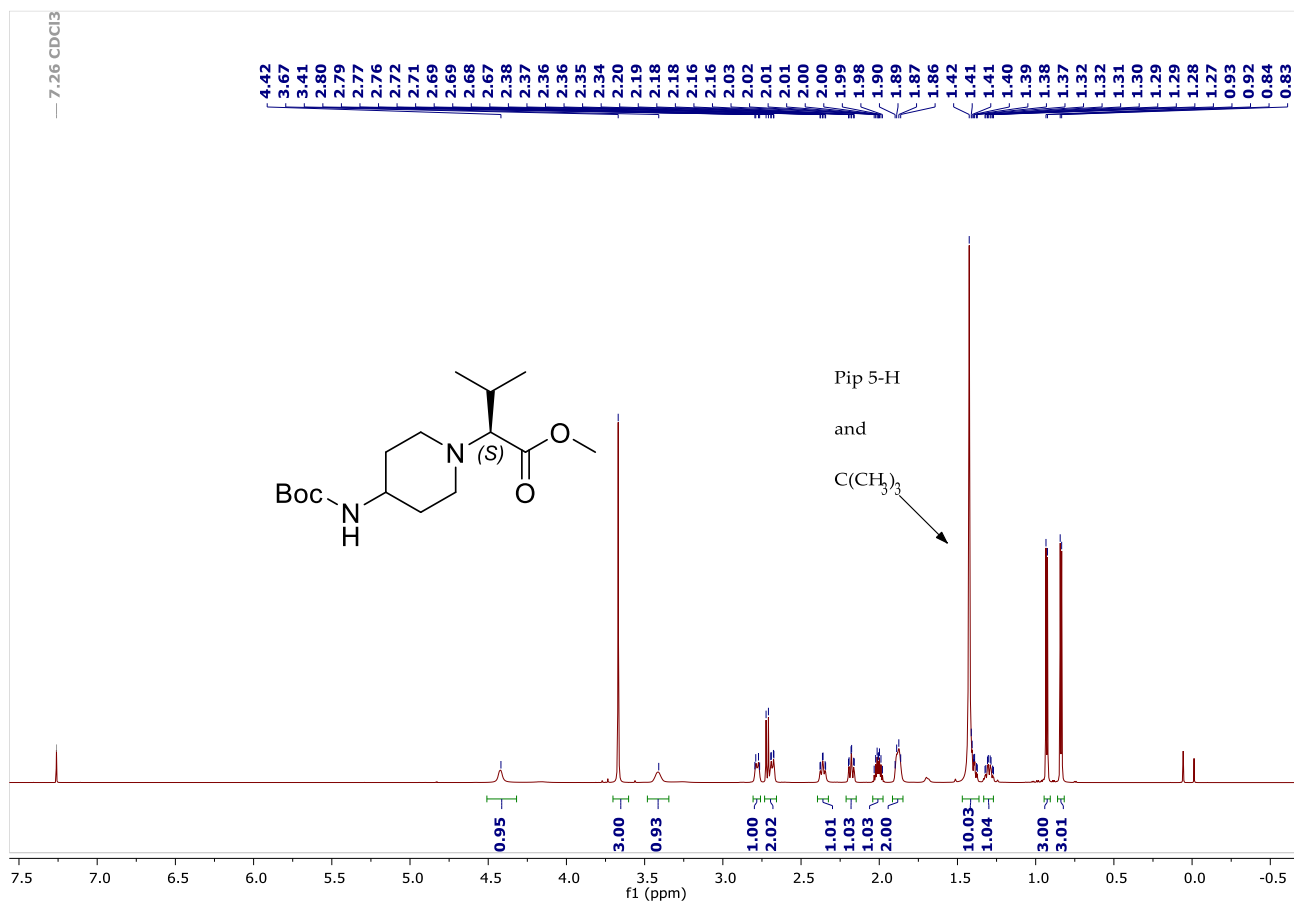


Figure S14. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-3-methylbutanoate ((S)-3b). ¹H NMR spectrum (700 MHz, CDCl₃).

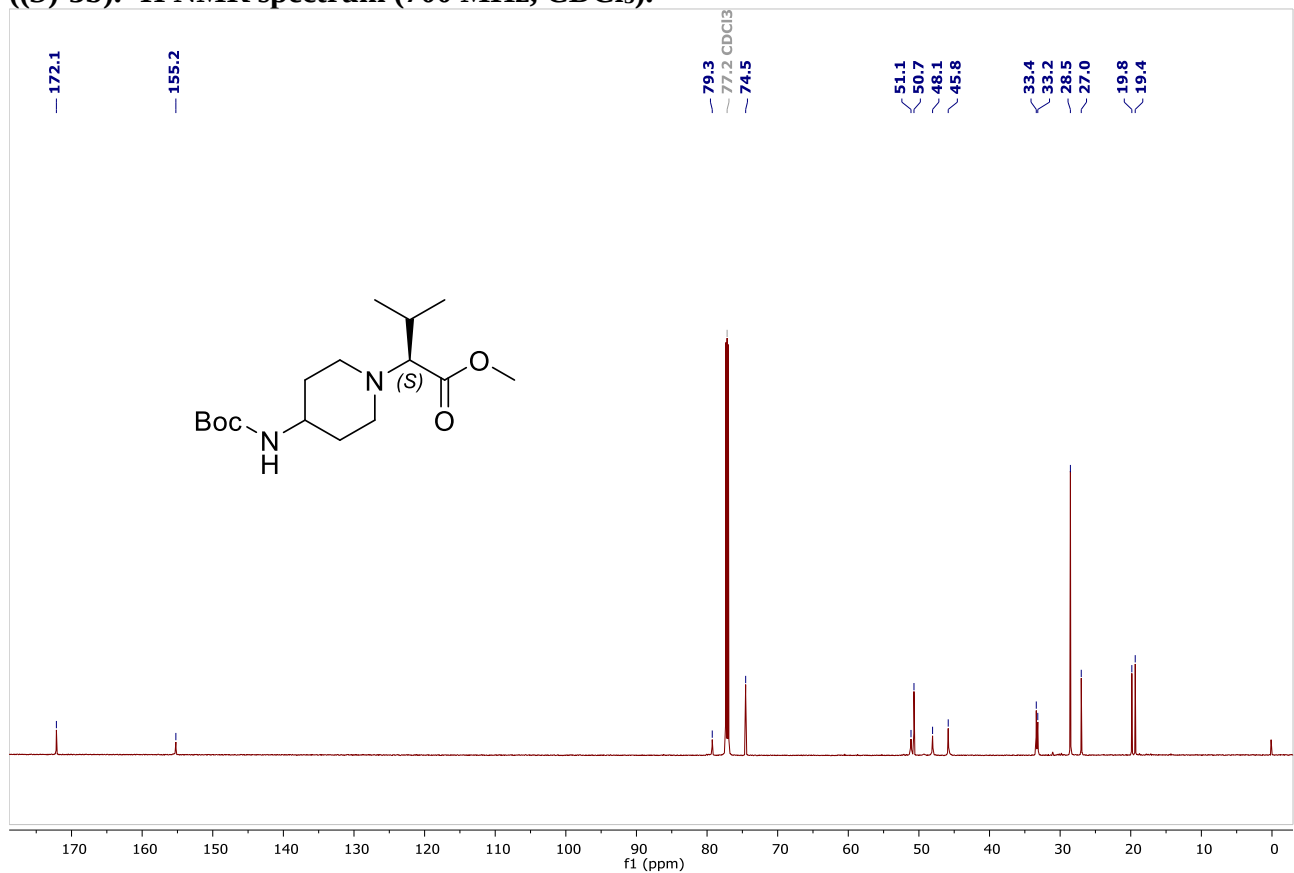


Figure S15. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-3-methylbutanoate ((S)-3b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

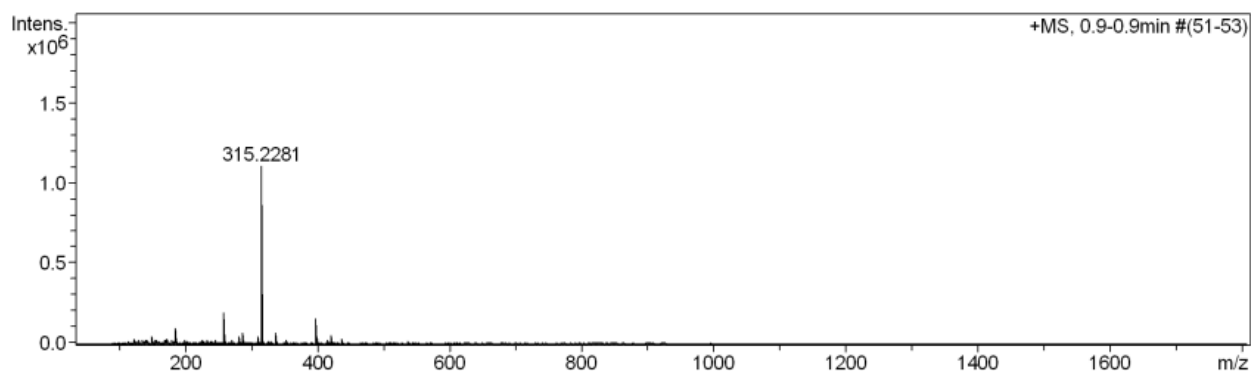
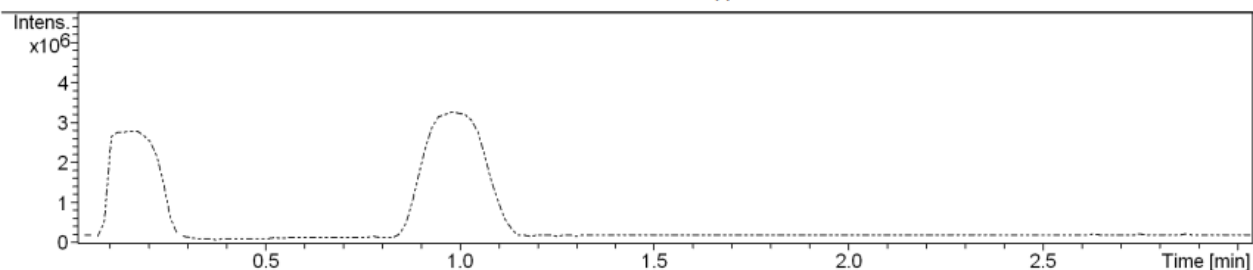
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Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

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Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
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	2	C ₁₇ H ₂₇ N ₆	51.82	315.2292	3.3	3.5	29.2	7.5	even	ok

Figure S16. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-3-methylbutanoate ((S)-3b). HRMS (ESI-TOF).

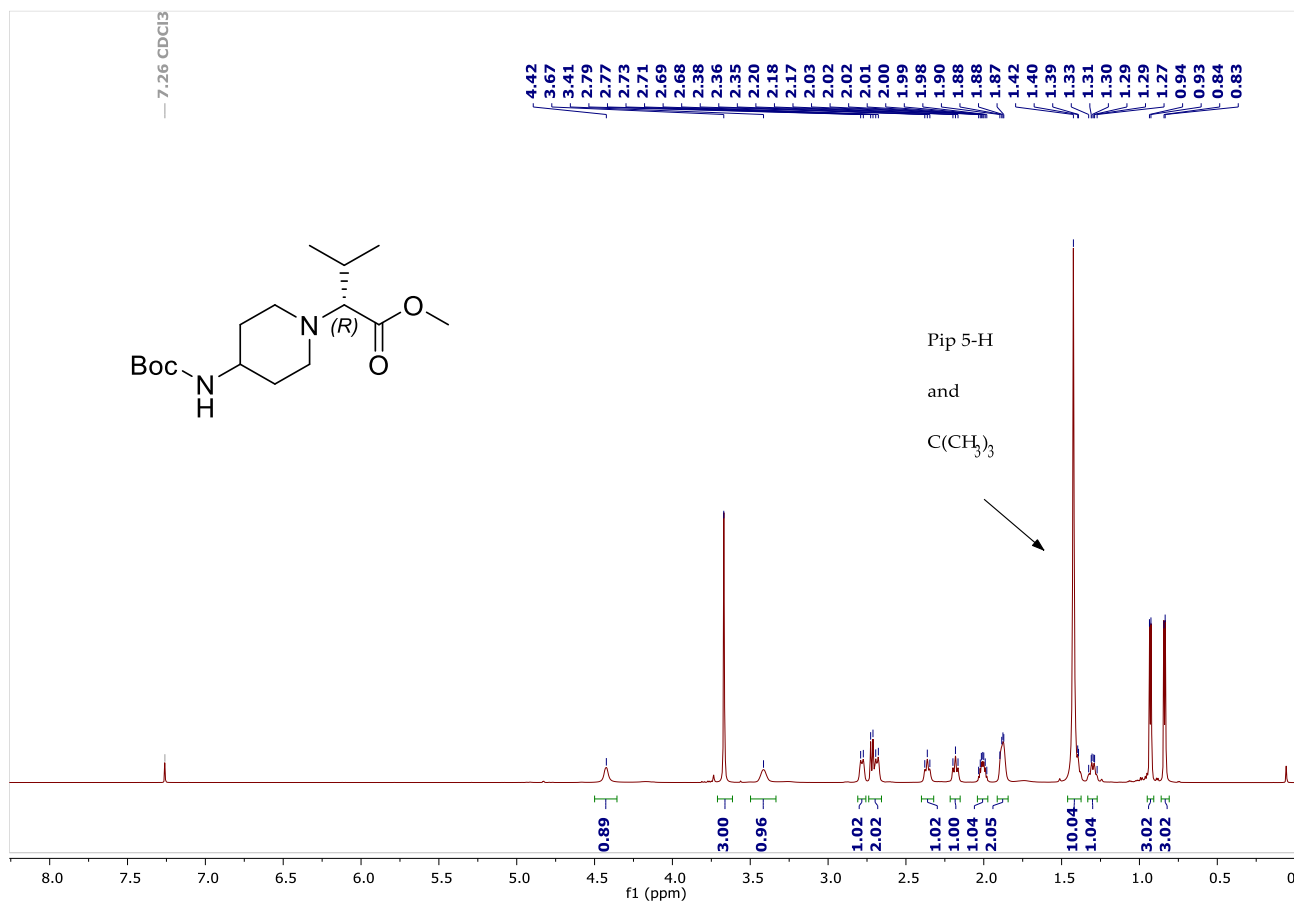


Figure S17. Methyl (2R)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-3-methylbutanoate ((R)-3b). ¹H NMR spectrum (700 MHz, CDCl₃).

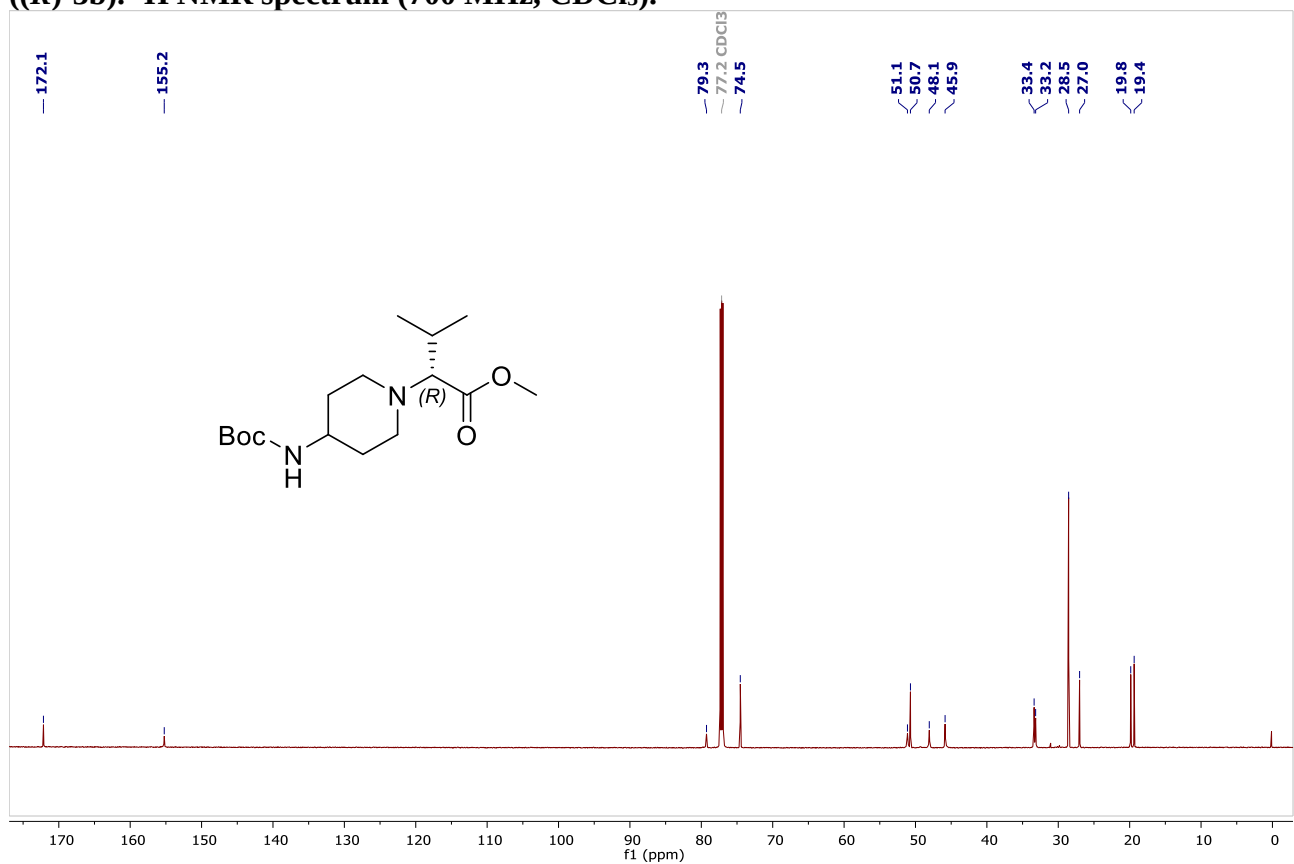


Figure S18. Methyl (2R)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-3-methylbutanoate ((R)-3b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

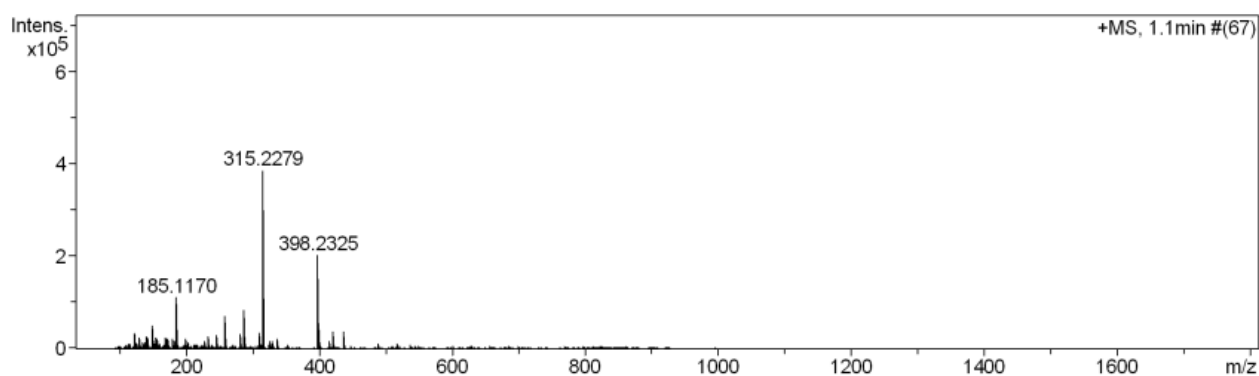
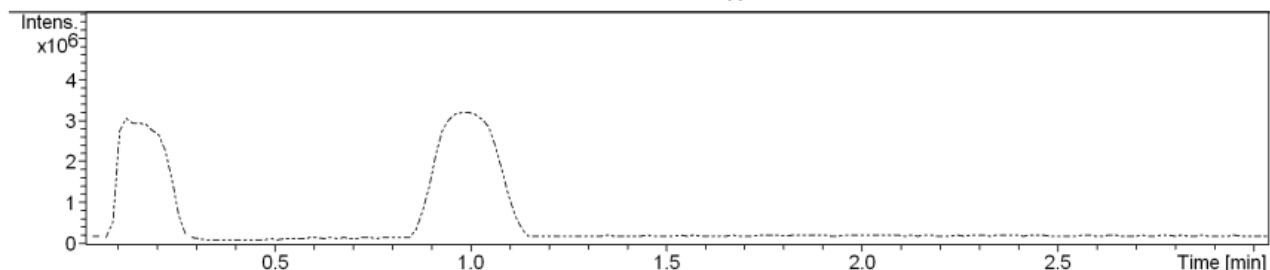
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Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

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Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



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	2	C 17 H 27 N 6	45.53	315.2292	3.9	4.0	27.9	7.5	even	ok

Figure S19. Methyl (2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-3-methylbutanoate ((*R*)-3b). HRMS (ESI-TOF).

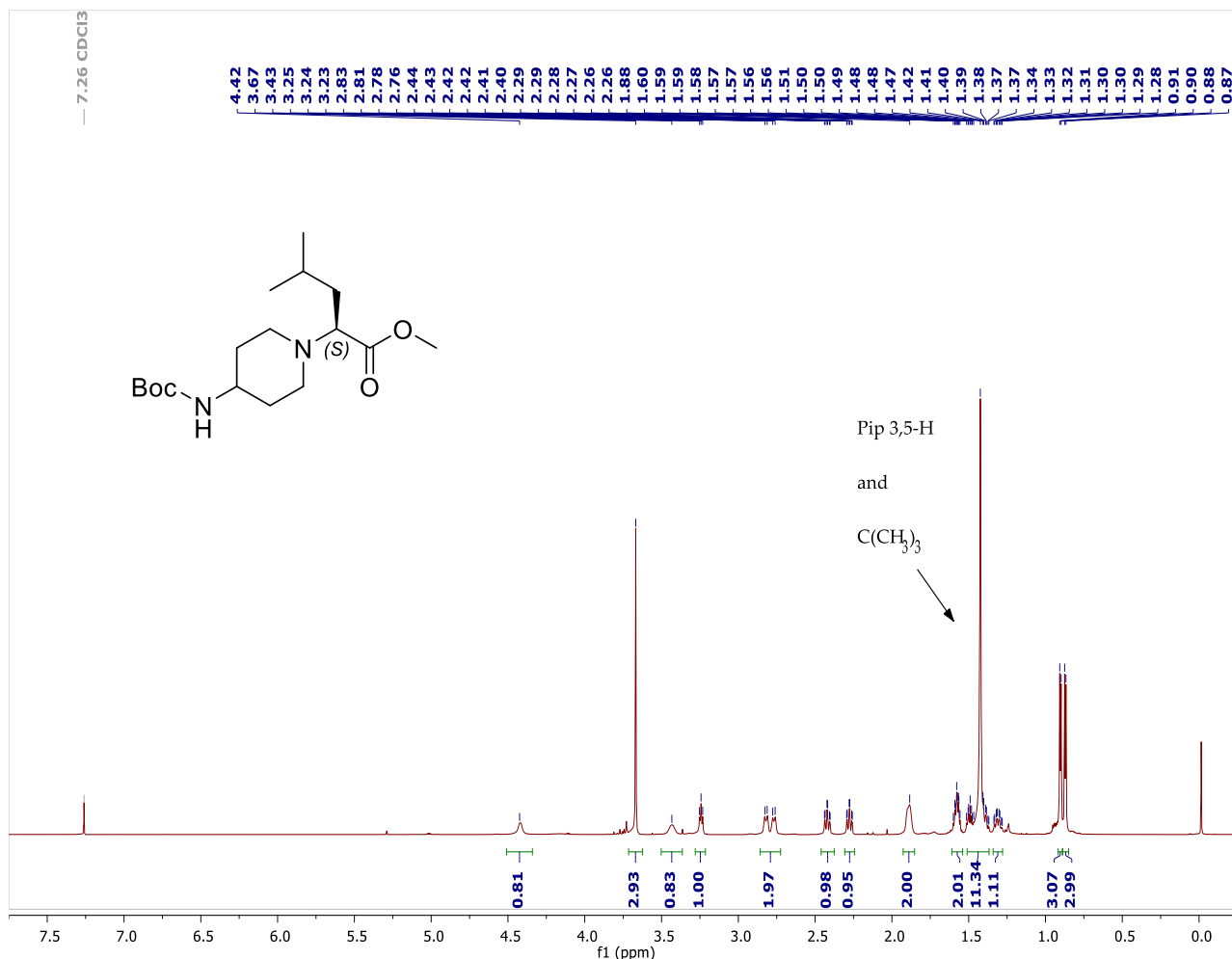


Figure S20. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-4-methylpentanoate ((S)-3c). ¹H NMR spectrum (700 MHz, CDCl₃).

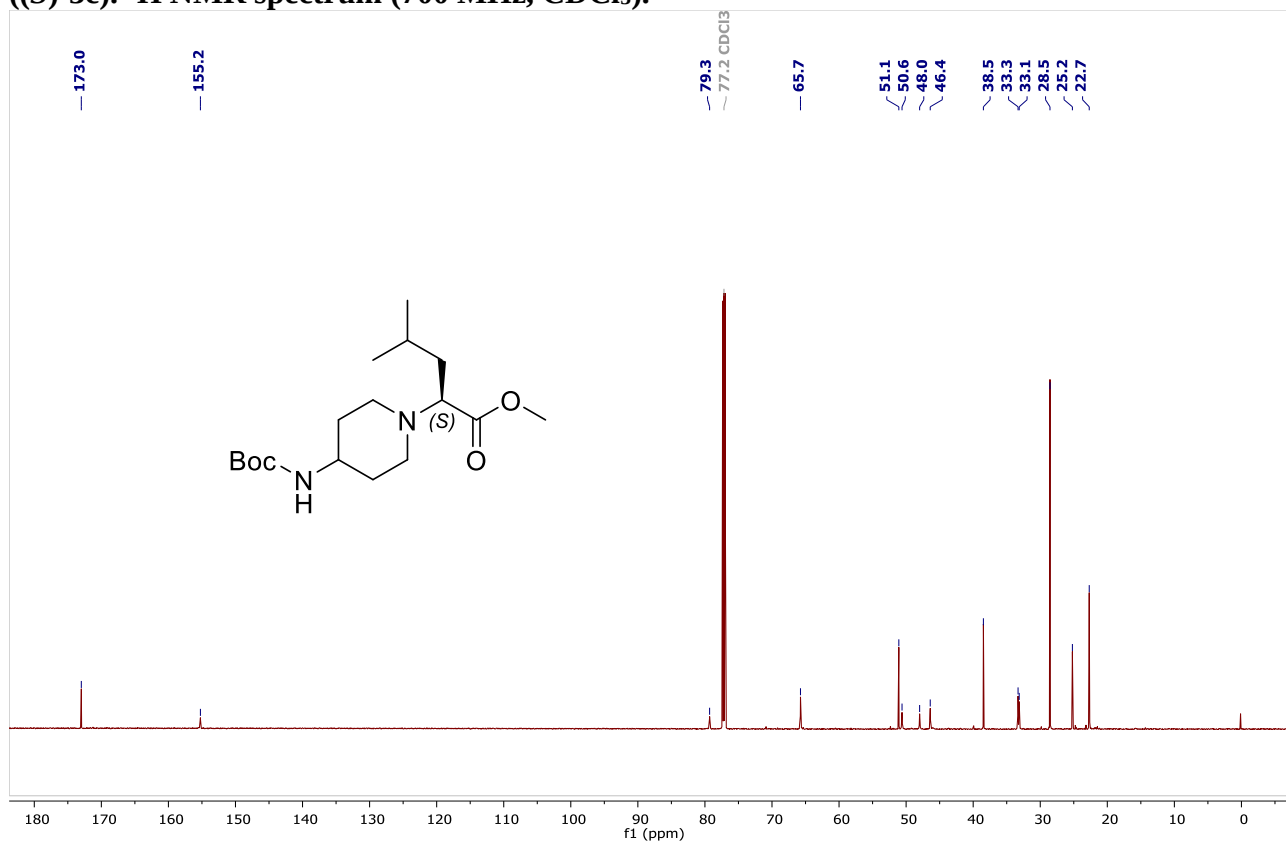


Figure S21. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-4-methylpentanoate ((S)-3c). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

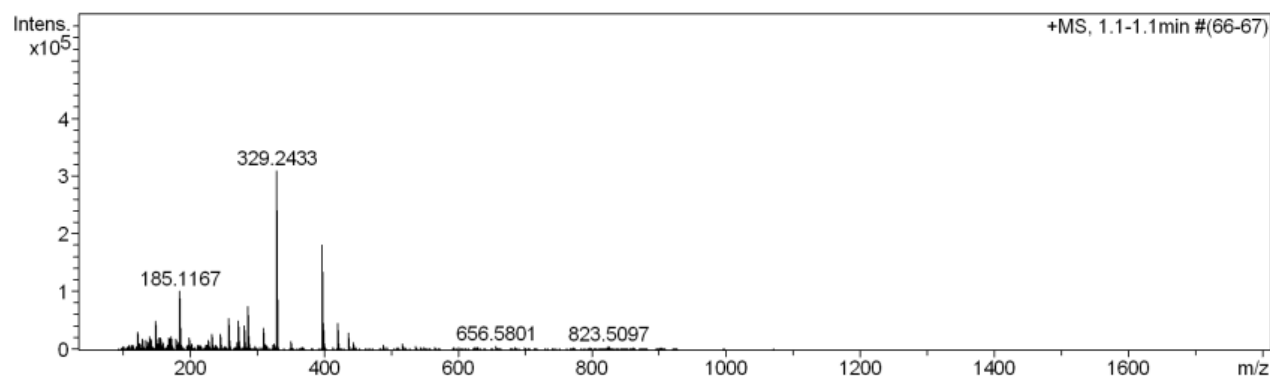
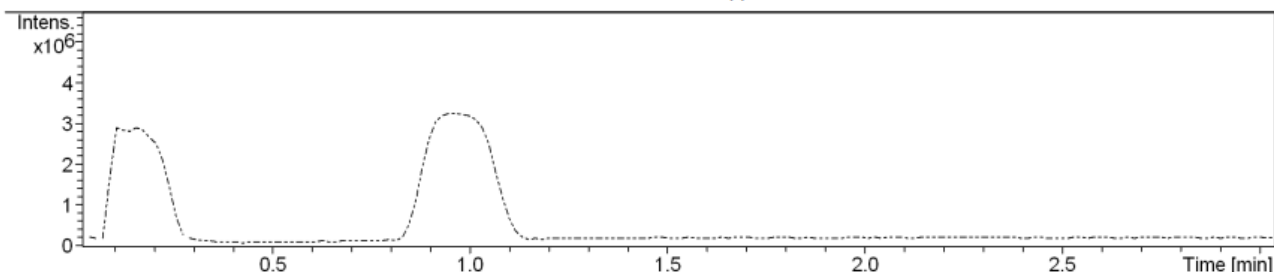
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 Instrument / Ser# maXis 4G 20218

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329.2433	1	C 17 H 33 N 2 O 4	100.00	329.2435	0.6	0.9	16.7	2.5	even	ok
	2	C 18 H 29 N 6	36.31	329.2448	4.7	4.9	29.3	7.5	even	ok

Figure S22. Methyl (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-4-methylpentanoate ((S)-3c). HRMS (ESI-TOF).

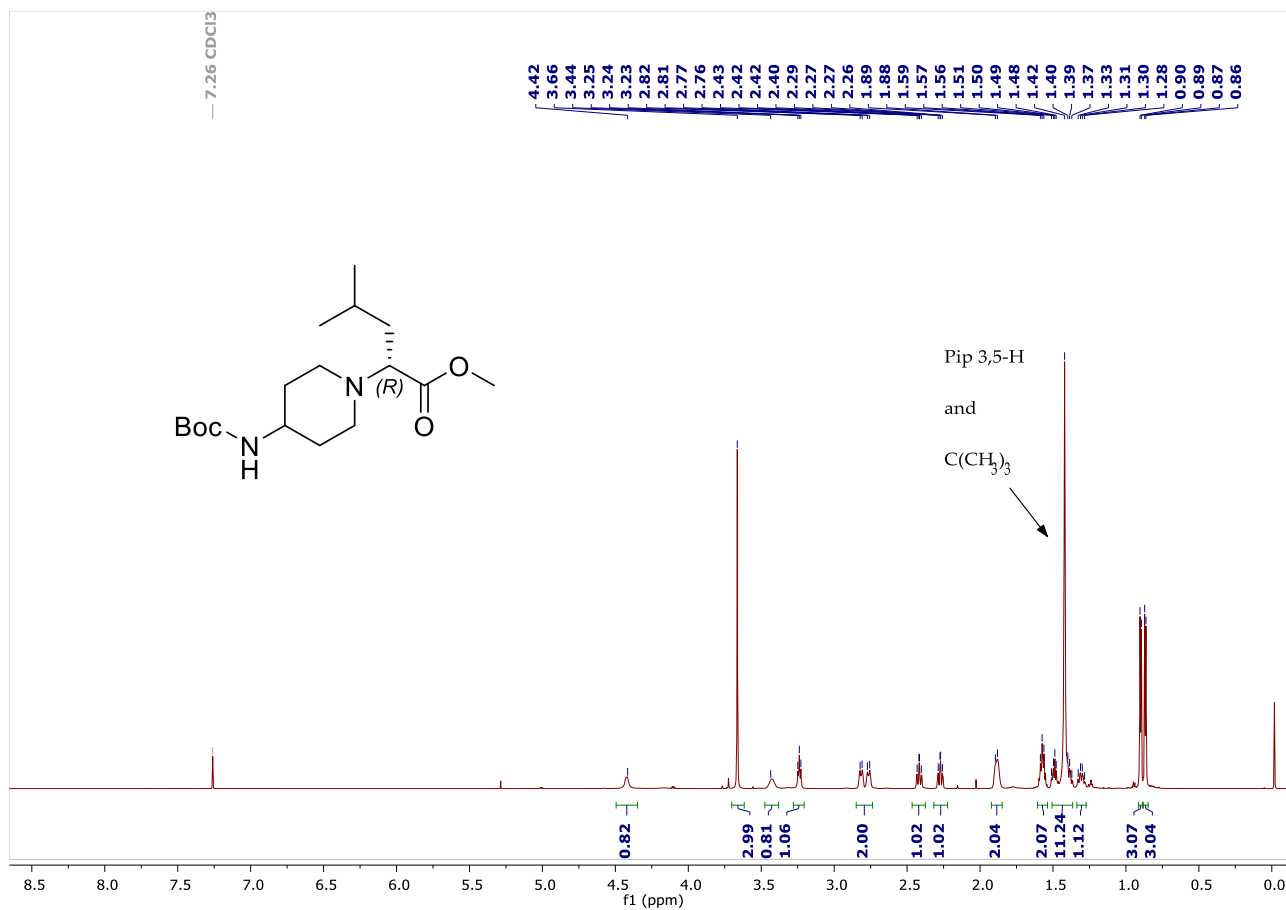


Figure S23. Methyl (2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-4-methylpentanoate ((*R*)-3c). ¹H NMR spectrum (700 MHz, CDCl₃).

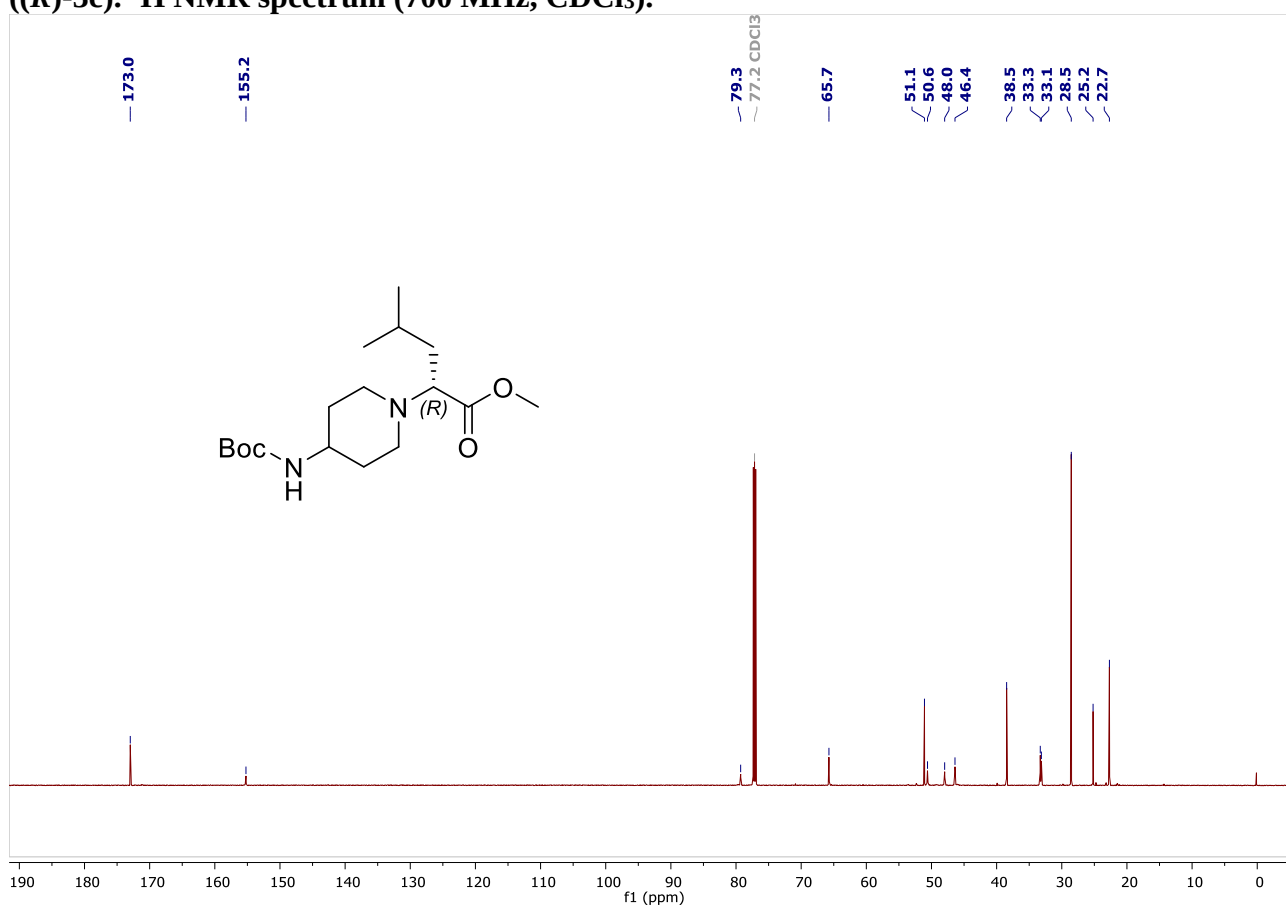


Figure S24. Methyl (2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-4-methylpentanoate ((*R*)-3c). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

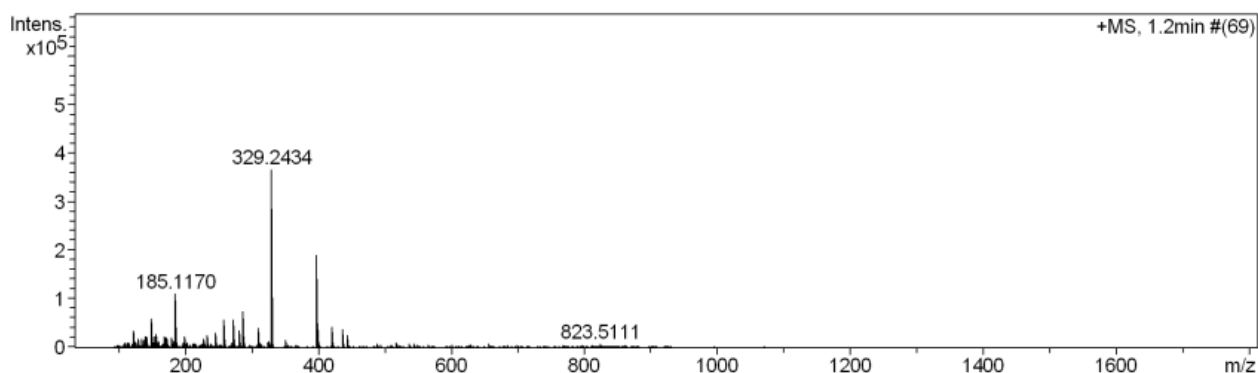
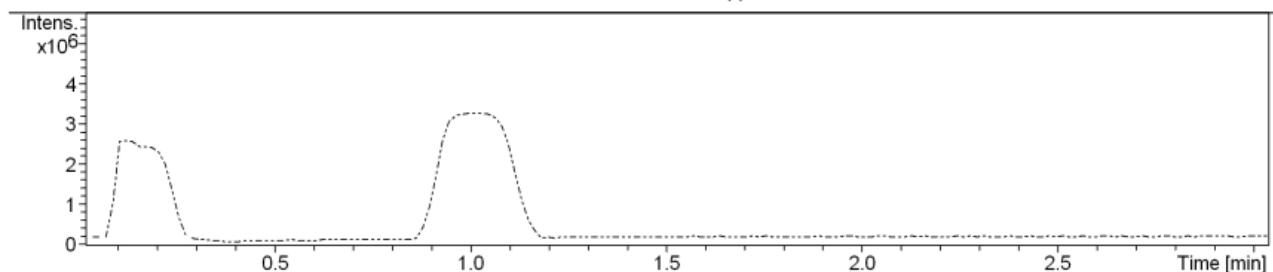
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 Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

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Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
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	2	C 18 H 29 N 6	37.00	329.2448	4.5	4.7	27.8	7.5	even	ok

Figure S25. Methyl (2R)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}-4-methylpentanoate ((R)-3c). HRMS (ESI-TOF).

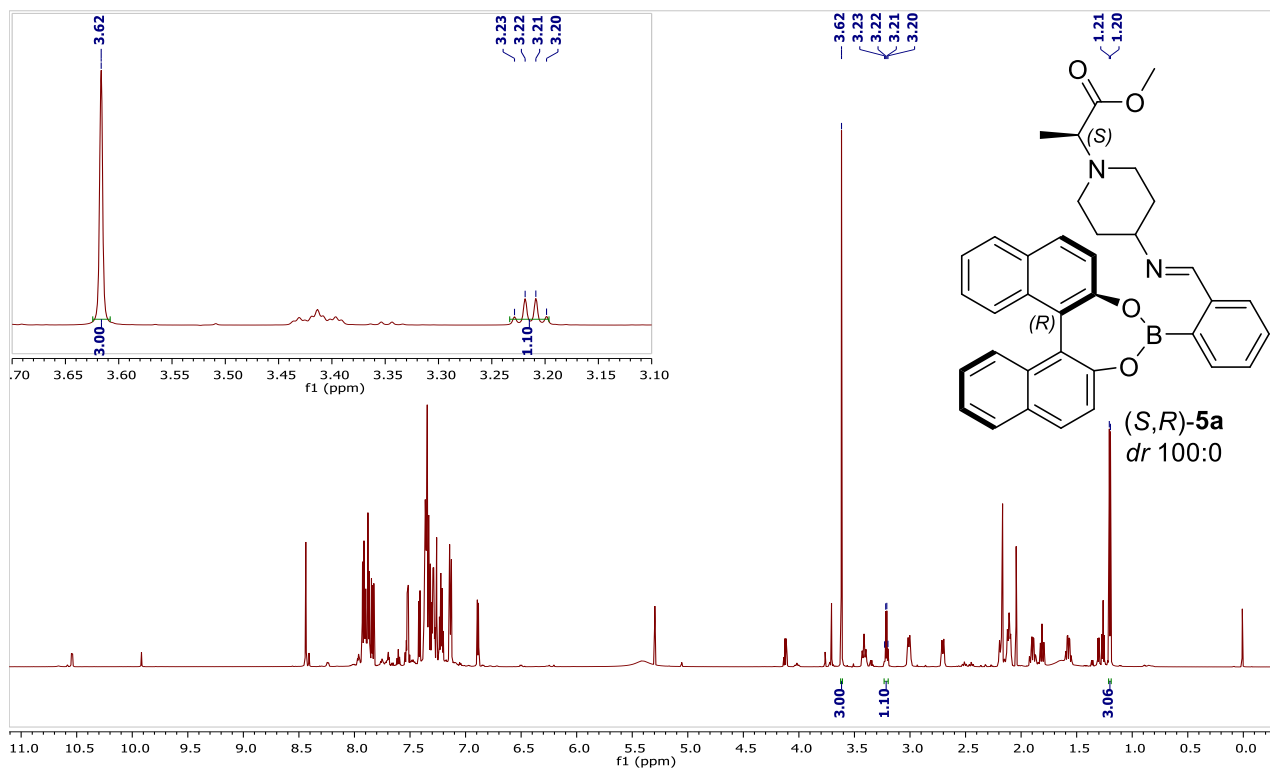


Figure S26. Crude sample from reaction mixture of iminoboronate ester complex (*(S,R)*-5a). ^1H NMR spectrum (700 MHz, CDCl_3).

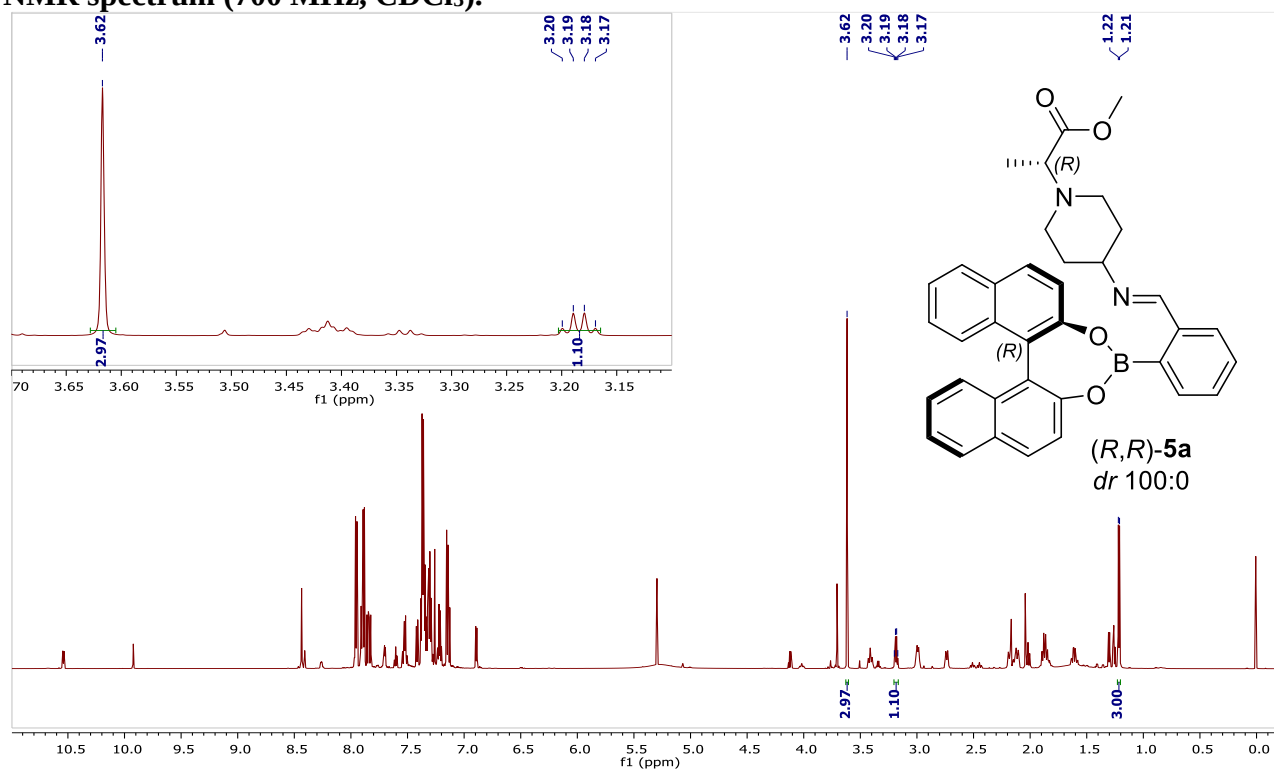


Figure S27. Crude sample from reaction mixture of iminoboronate ester complex (*(R,R)*-5a). ^1H NMR spectrum (700 MHz, CDCl_3).

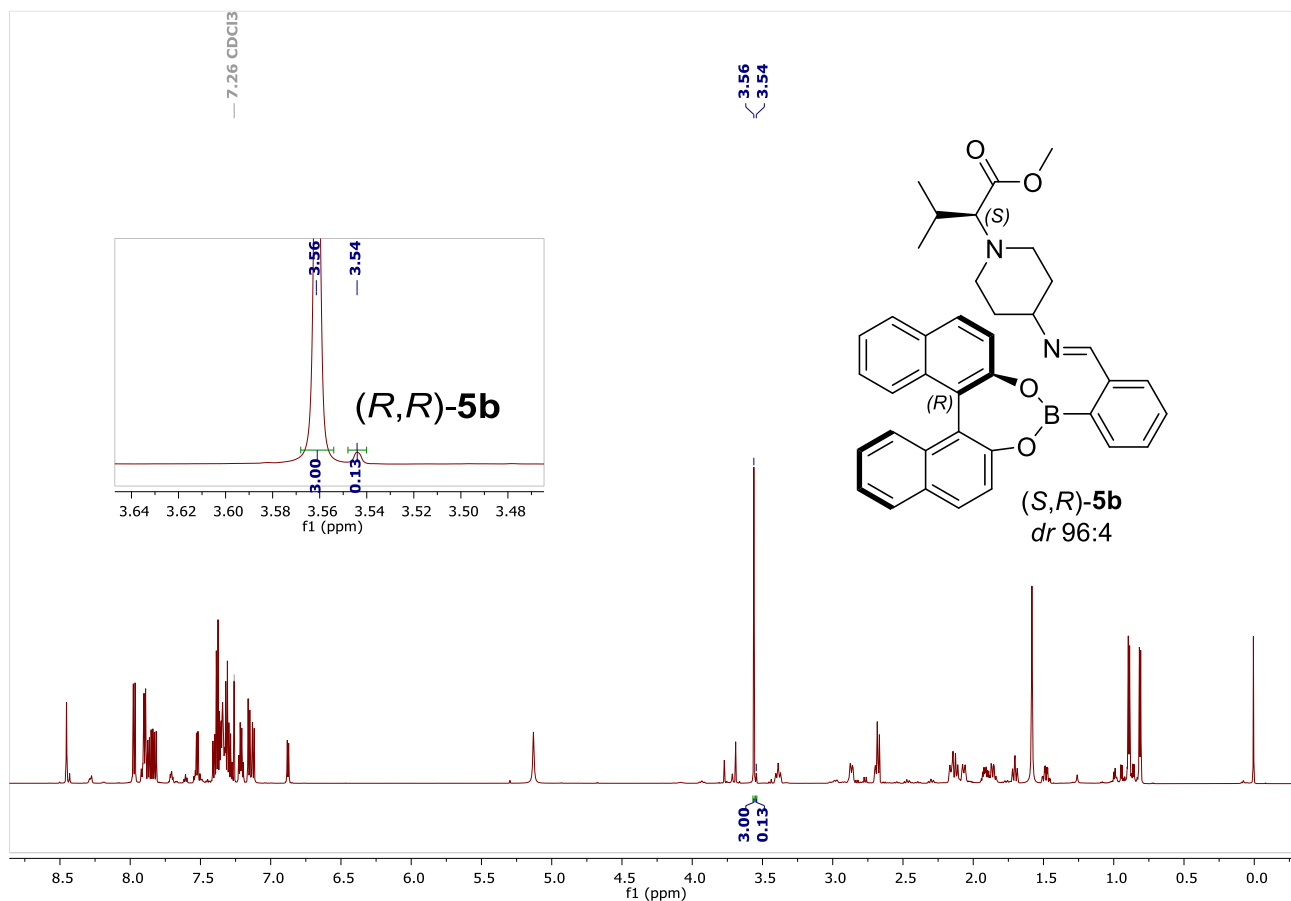


Figure S28. Crude sample from reaction mixture of iminoboronate ester complex ((*S,R*)-5b). ^1H NMR spectrum (700 MHz, CDCl_3).

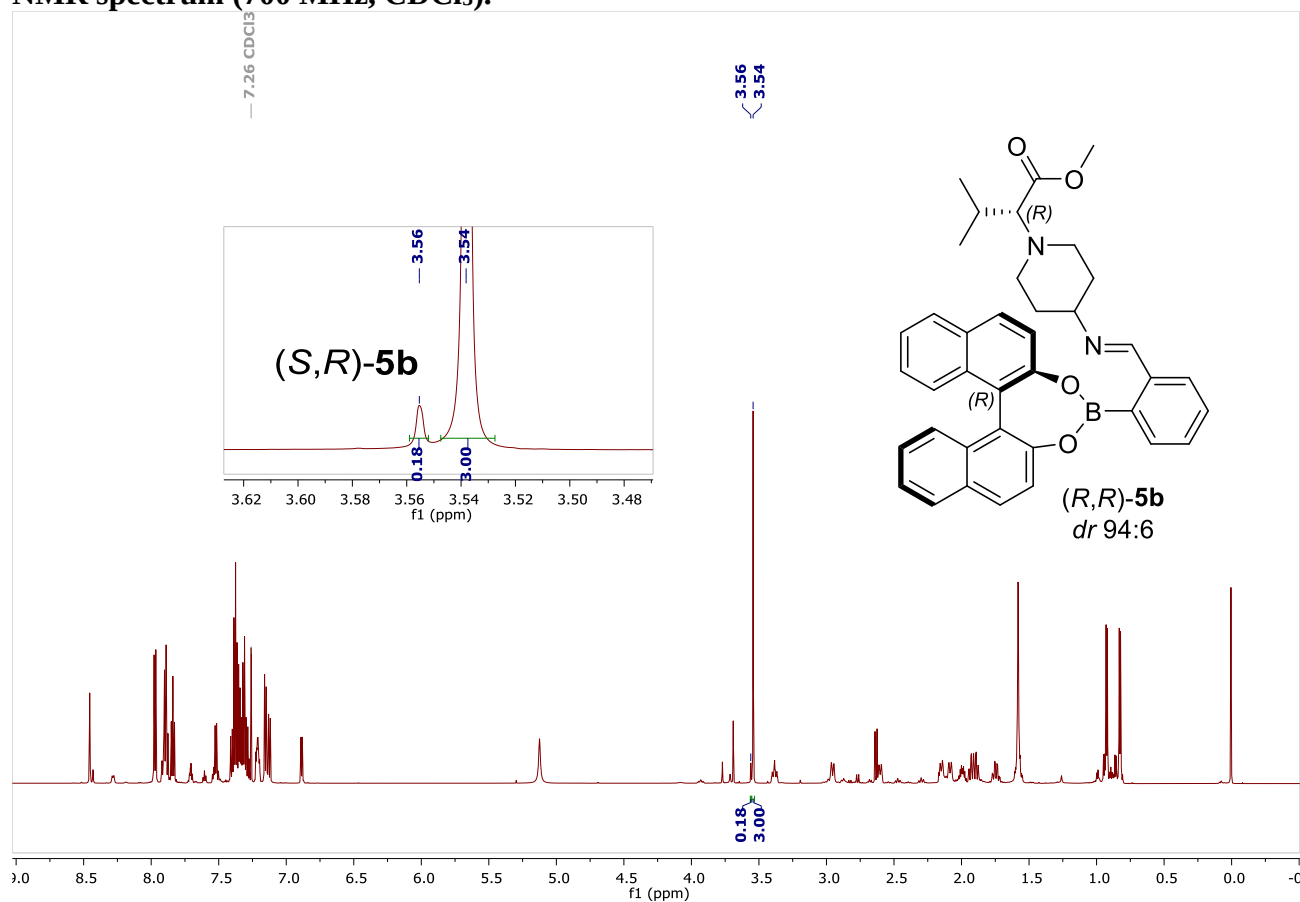


Figure S29. Crude sample from reaction mixture of iminoboronate ester complex ((*R,R*)-5b). ^1H NMR spectrum (700 MHz, CDCl_3).

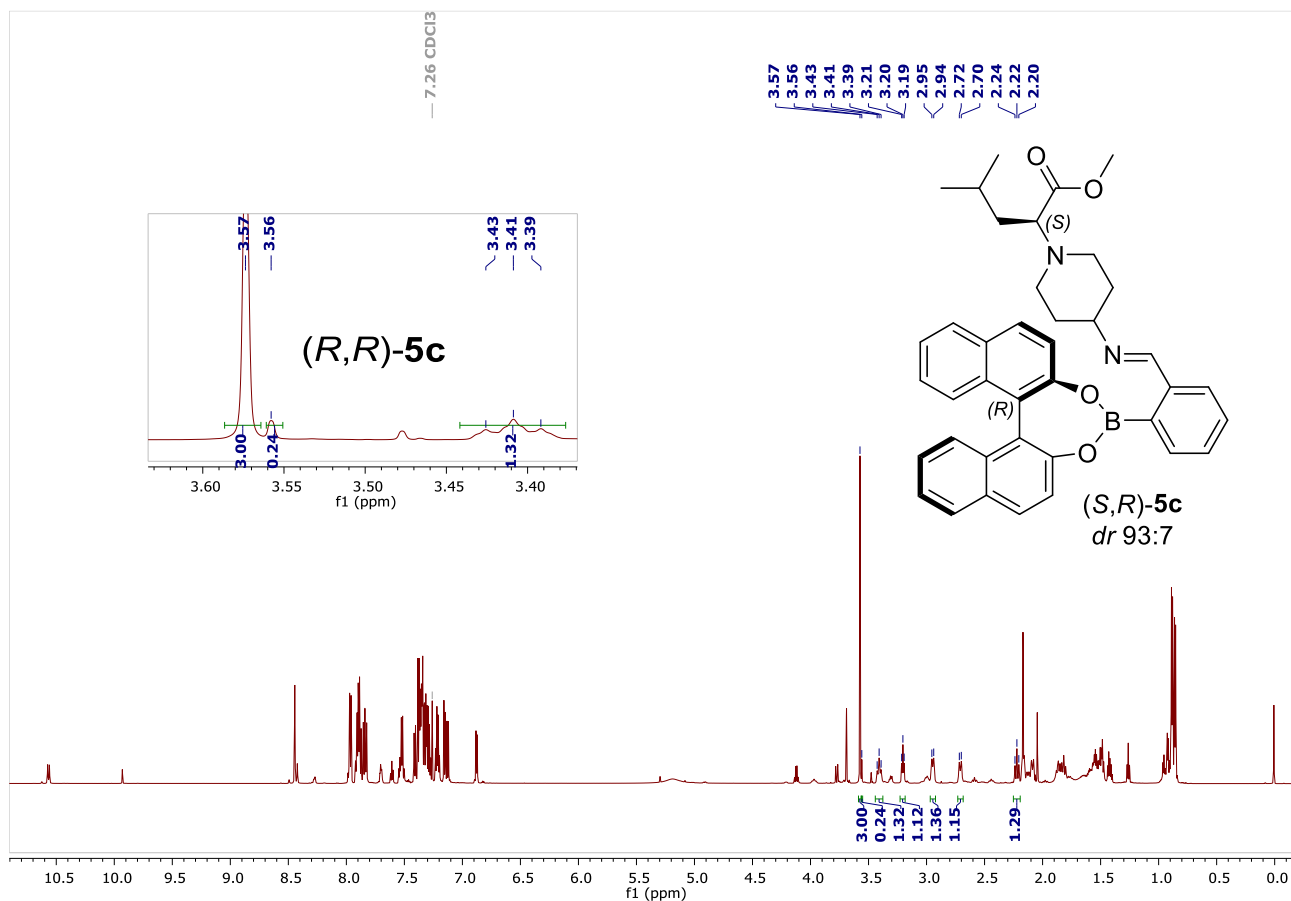


Figure S30. Crude sample from reaction mixture of iminoboronate ester complex ((*S,R*)-5c). ^1H NMR spectrum (700 MHz, CDCl_3).

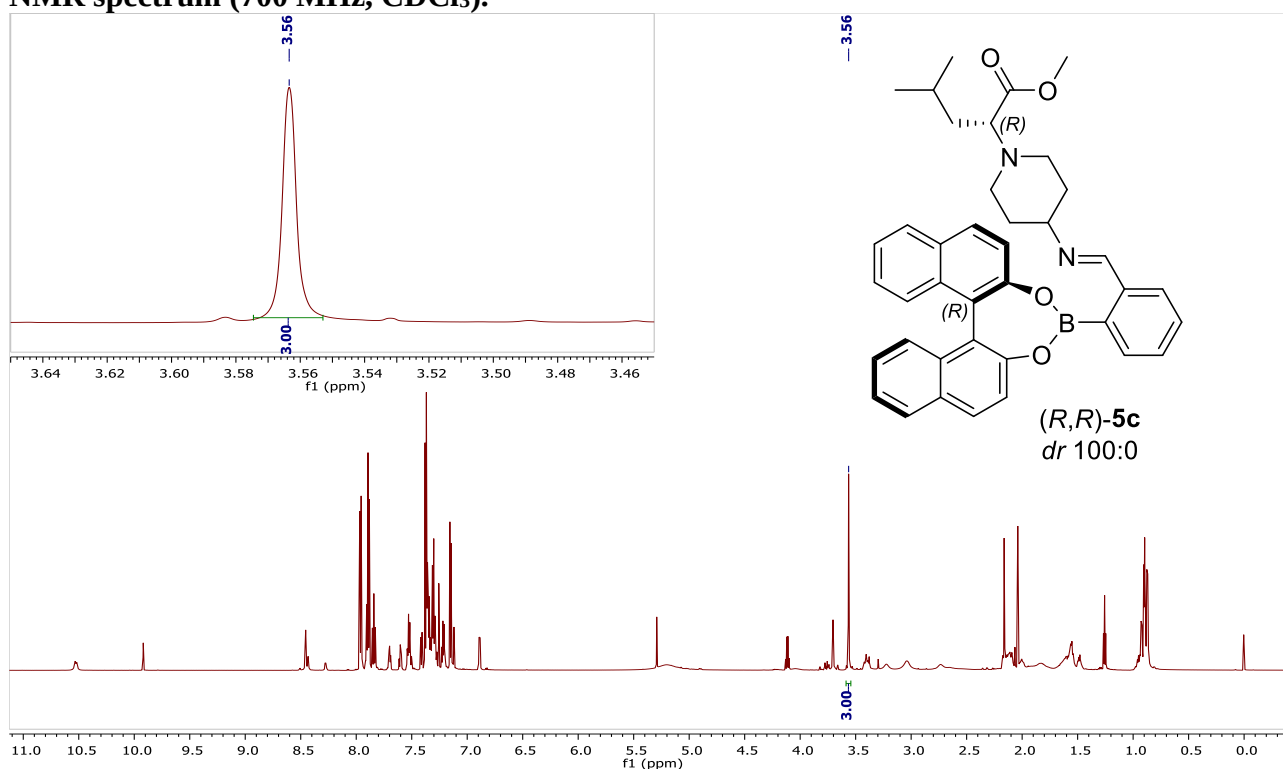


Figure S31. Crude sample from reaction mixture of iminoboronate ester complex ((*R,R*)-5c). ^1H NMR spectrum (700 MHz, CDCl_3).

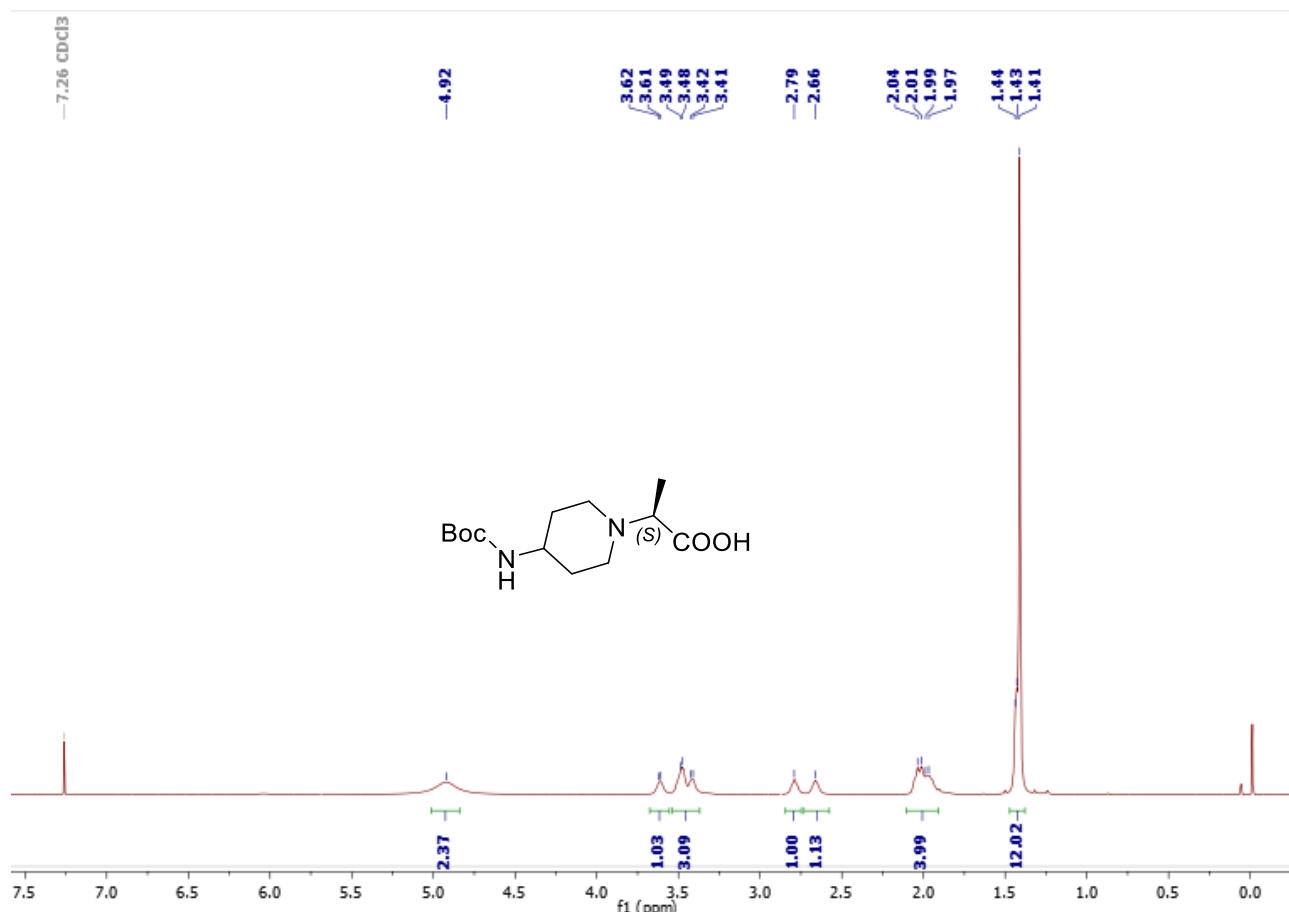


Figure S32. (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoic acid ((S)-6). ¹H NMR spectrum (700 MHz, CDCl₃).

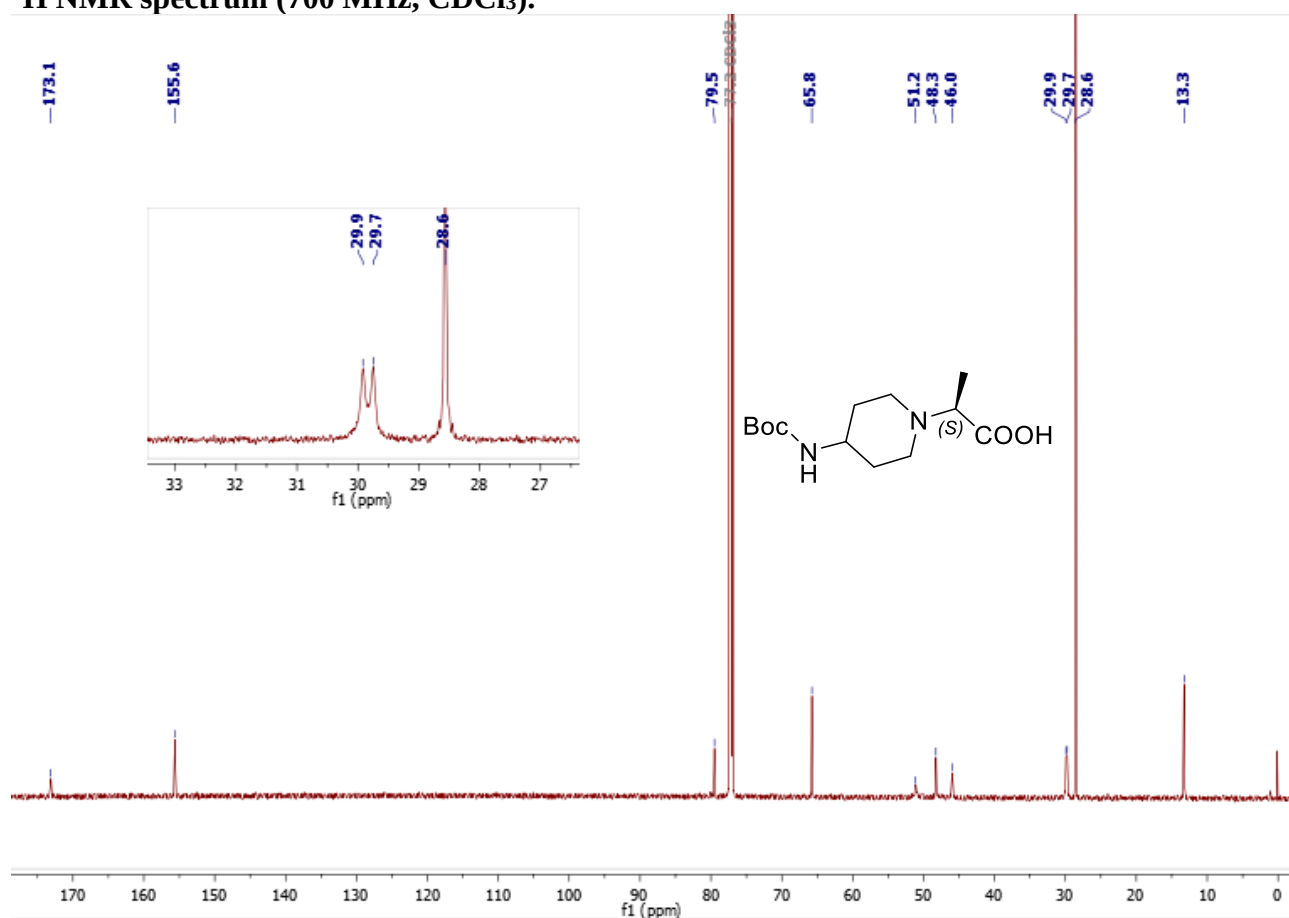


Figure S33. (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoic acid ((S)-6). ¹³C NMR spectrum (176 MHz, CDCl₃).

Compound Spectrum SmartFormula Report

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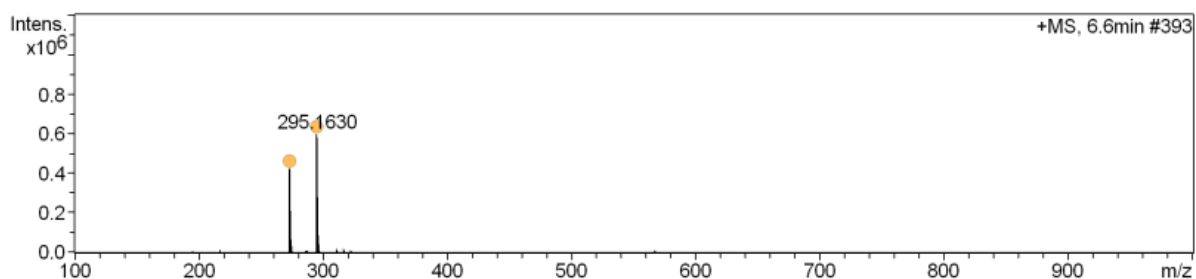
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Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	0.4 Bar
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Scan Begin	50 m/z	Set End Plate Offset	-500 V	Set Dry Gas	4.0 l/min
Scan End	1000 m/z	Set Collision Cell RF	140.0 Vpp	Set Divert Valve	Waste

#	RT [min]	Area	Int. Type	I	S/N	Chromatogram	Max. m/z	FWHM [min]
n.a.	6.6	n.a.	Single spectrum	n.a.	n.a.	n.a.	295.1630	n.a.

+MS, 6.6min #393



Meas. m/z	#	Ion Formula	m/z	err [ppm]	mSigma	# Sigma	Score	rdb	e ⁻ Conf	N-Rule
273.1805	1	C13H25N2O4	273.1809	-1.4	6.9	1	100.00	2.5	even	ok
295.1630	1	C13H24N2NaO4	295.1628	-0.6	2.2	1	100.00	2.5	even	ok

Figure S34. (2S)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoic acid ((S)-6). HRMS (ESI-TOF).

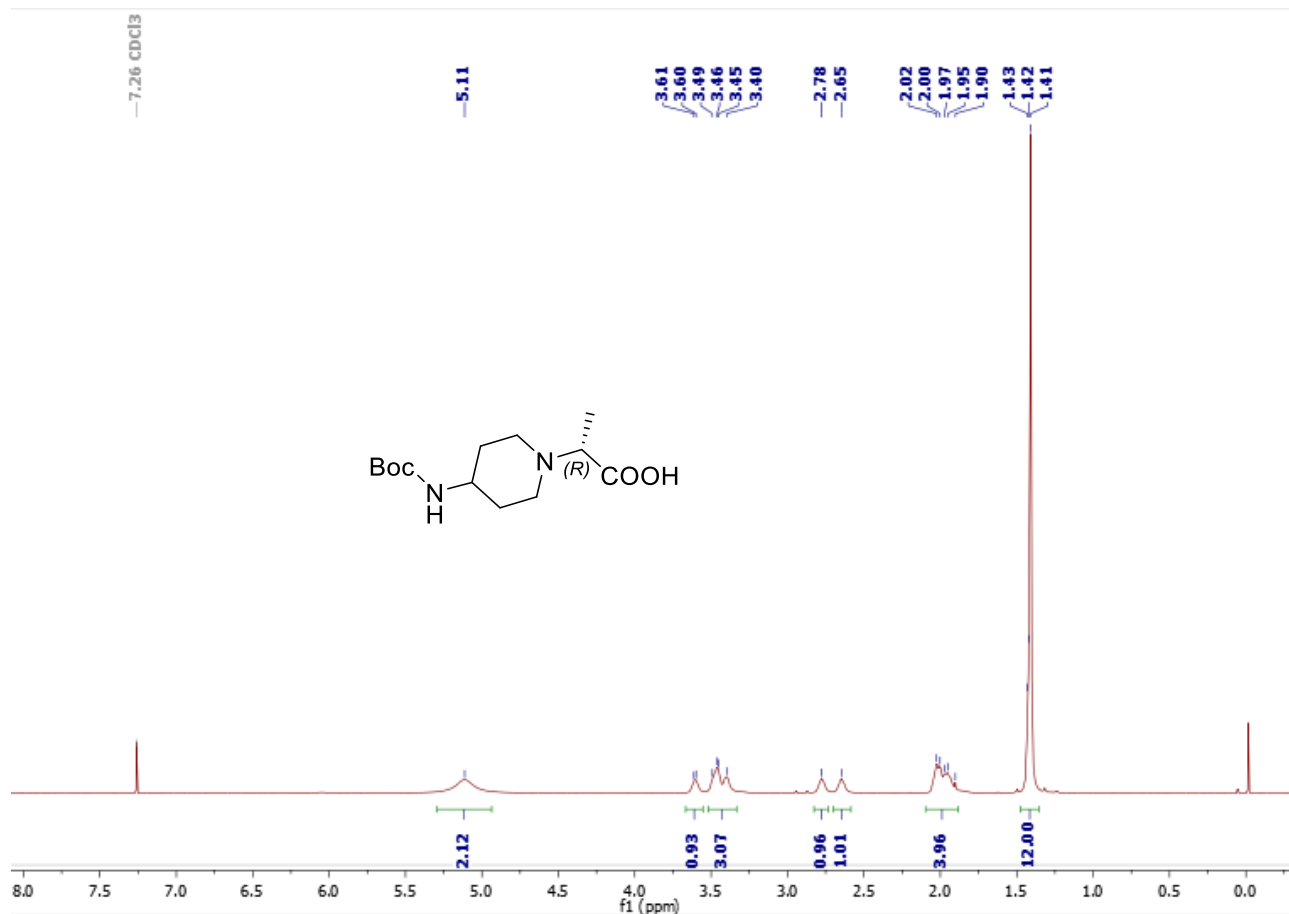


Figure S35. (2R)-2-{4-[(tert-butoxycarbonyl)amino]piperidin-1-yl}propanoic acid ((R)-6). ¹H NMR spectrum (700 MHz, CDCl₃).

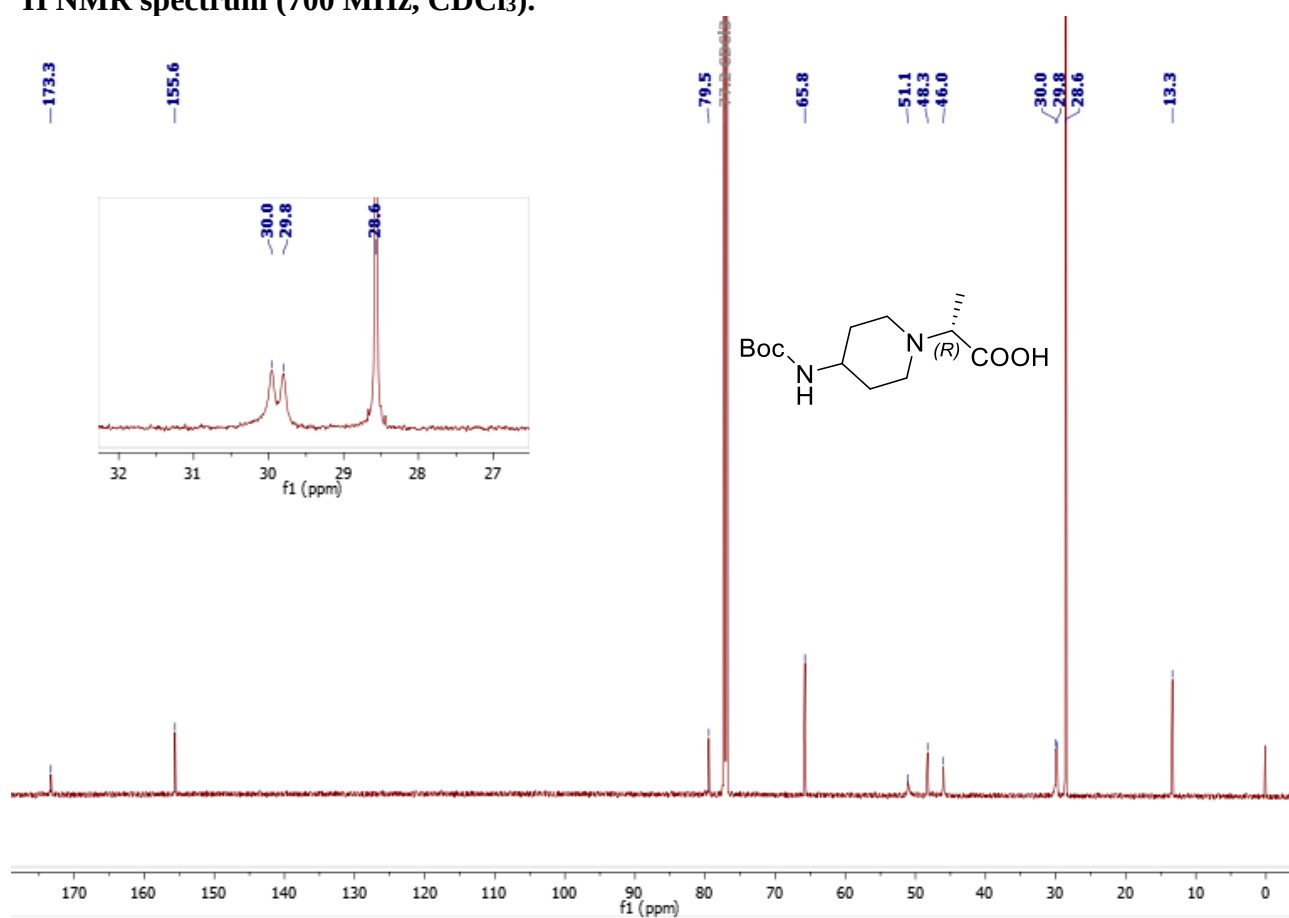


Figure S36. (2R)-2-{4-[(tert-butoxycarbonyl)amino]piperidin-1-yl}propanoic acid ((R)-6). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

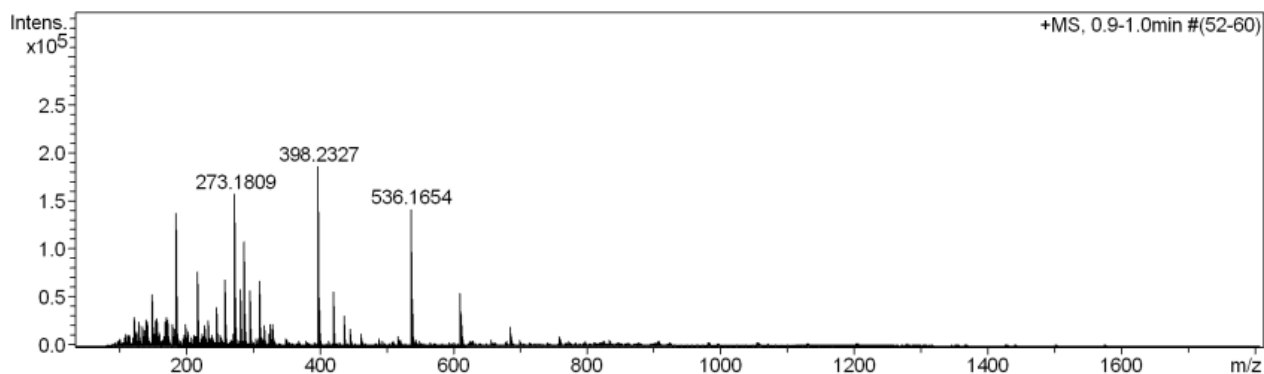
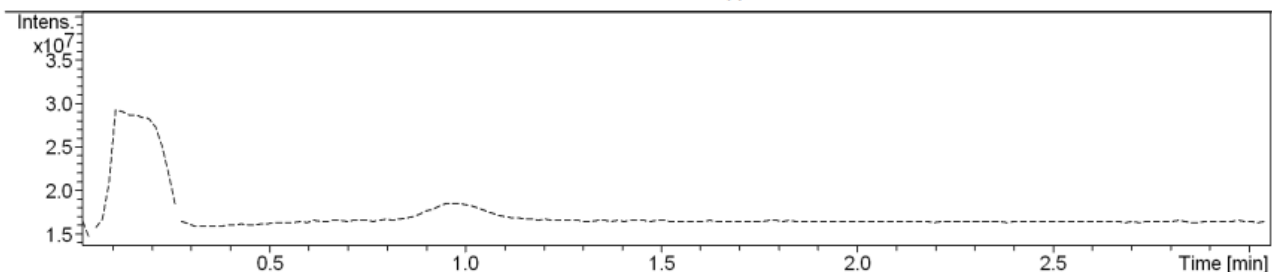
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Method organikai_esi_pos_2013_recover.m
Sample Name GMP_1189
Comment

Acquisition Date 4/21/2023 11:18:48 AM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
273.1809	1	C ₁₃ H ₂₅ N ₂ O ₄	100.00	273.1809	-0.1	0.1	1.0	2.5	even	ok

Figure S37. (2R)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoic acid ((R)-6). HRMS (ESI-TOF).

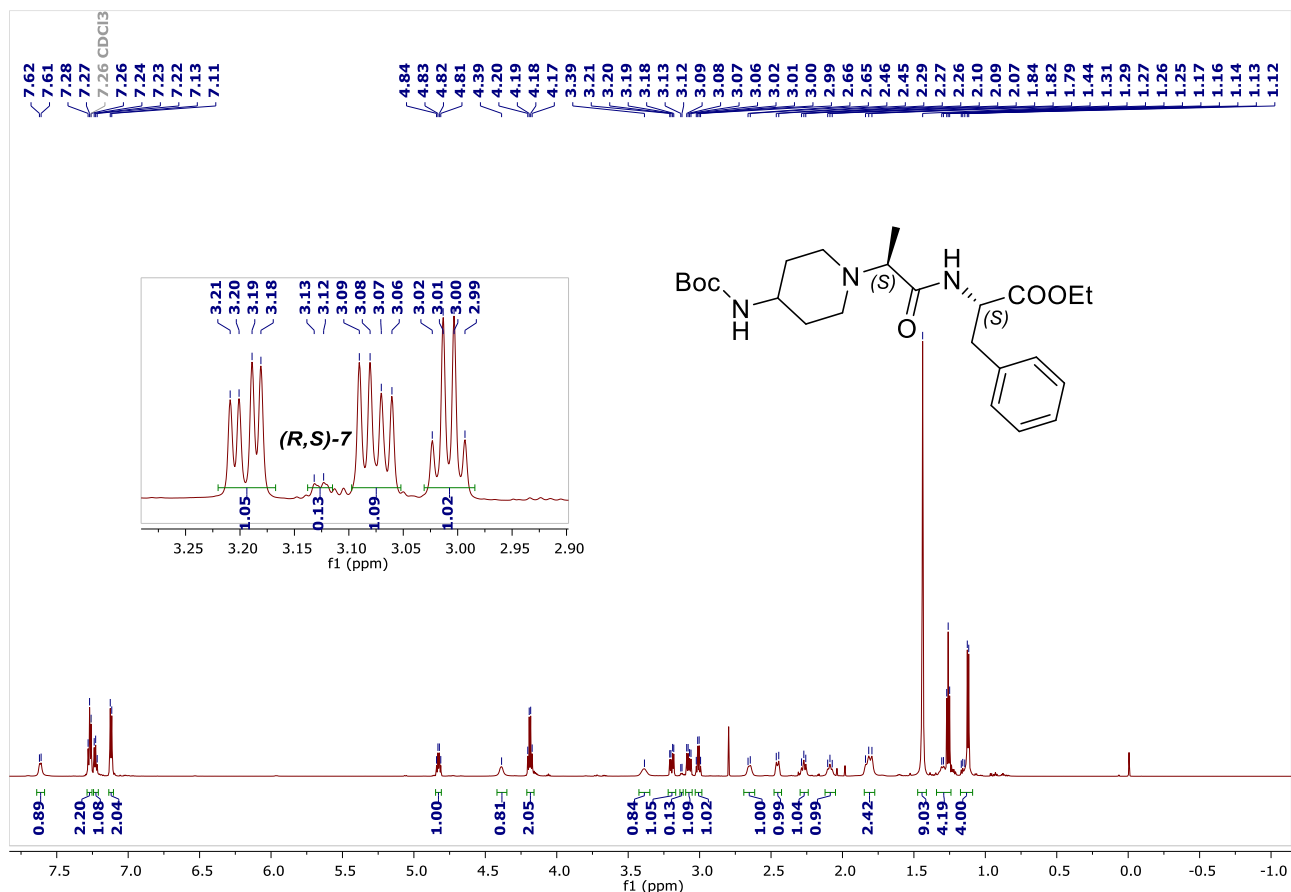


Figure S38. Ethyl *N*-[(2*S*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*S,S*)-7). ¹H NMR spectrum (700 MHz, CDCl₃).

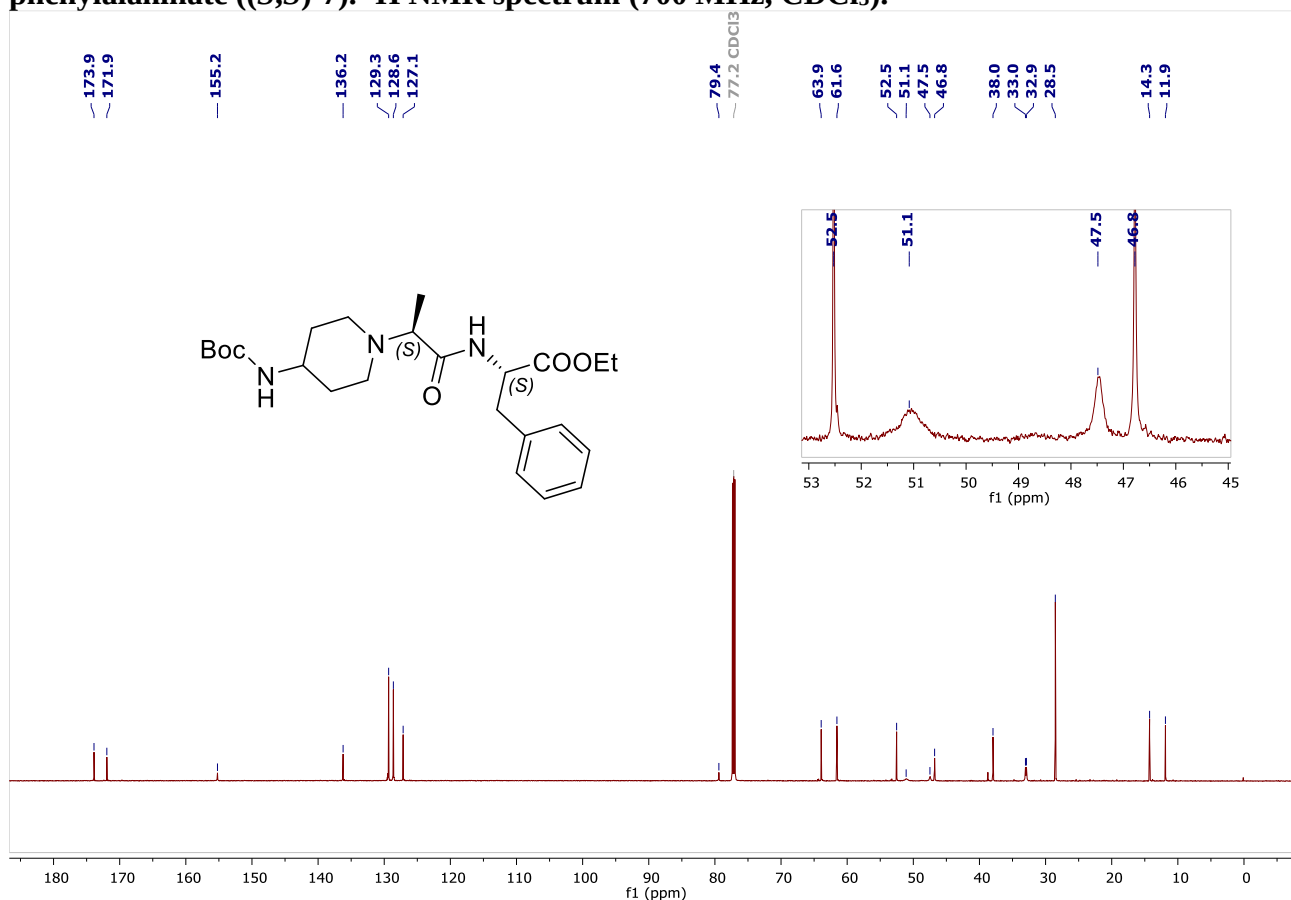


Figure S39. Ethyl *N*-[(2*S*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*S,S*)-7). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

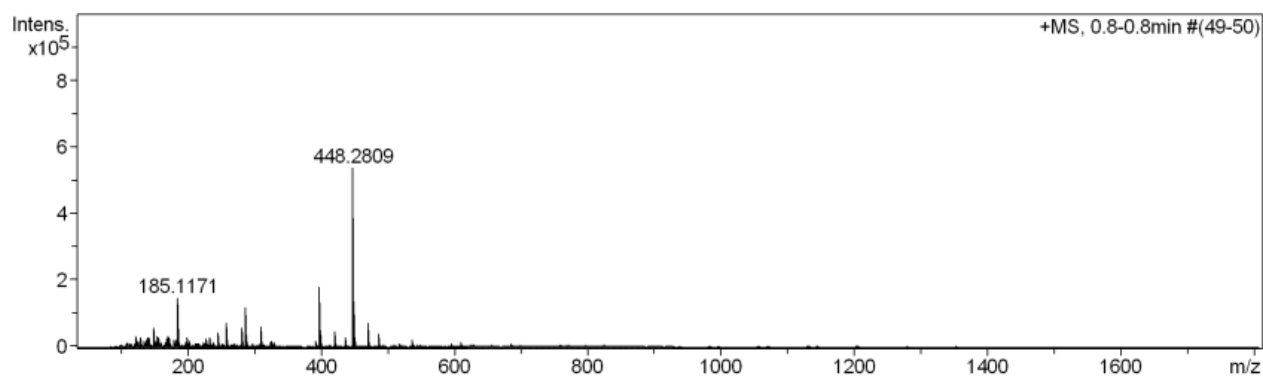
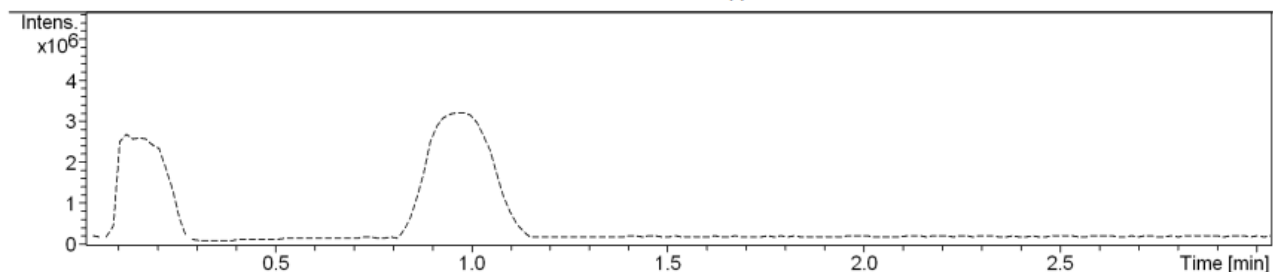
Analysis Info

Analysis Name D:\Data\Organikai\2023_04_11\GMP_1215_1-C,5_01_10227.d
Method organikai_esi_pos_2013_recover.m
Sample Name GMP_1215
Comment

Acquisition Date 4/21/2023 1:47:44 PM
Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
448.2809	1	C ₂₄ H ₃₈ N ₃ O ₅	100.00	448.2806	-0.6	-0.3	14.8	7.5	even	ok

Figure S40. Ethyl *N*-[(2*S*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*S,S*)-7). HRMS (ESI-TOF).

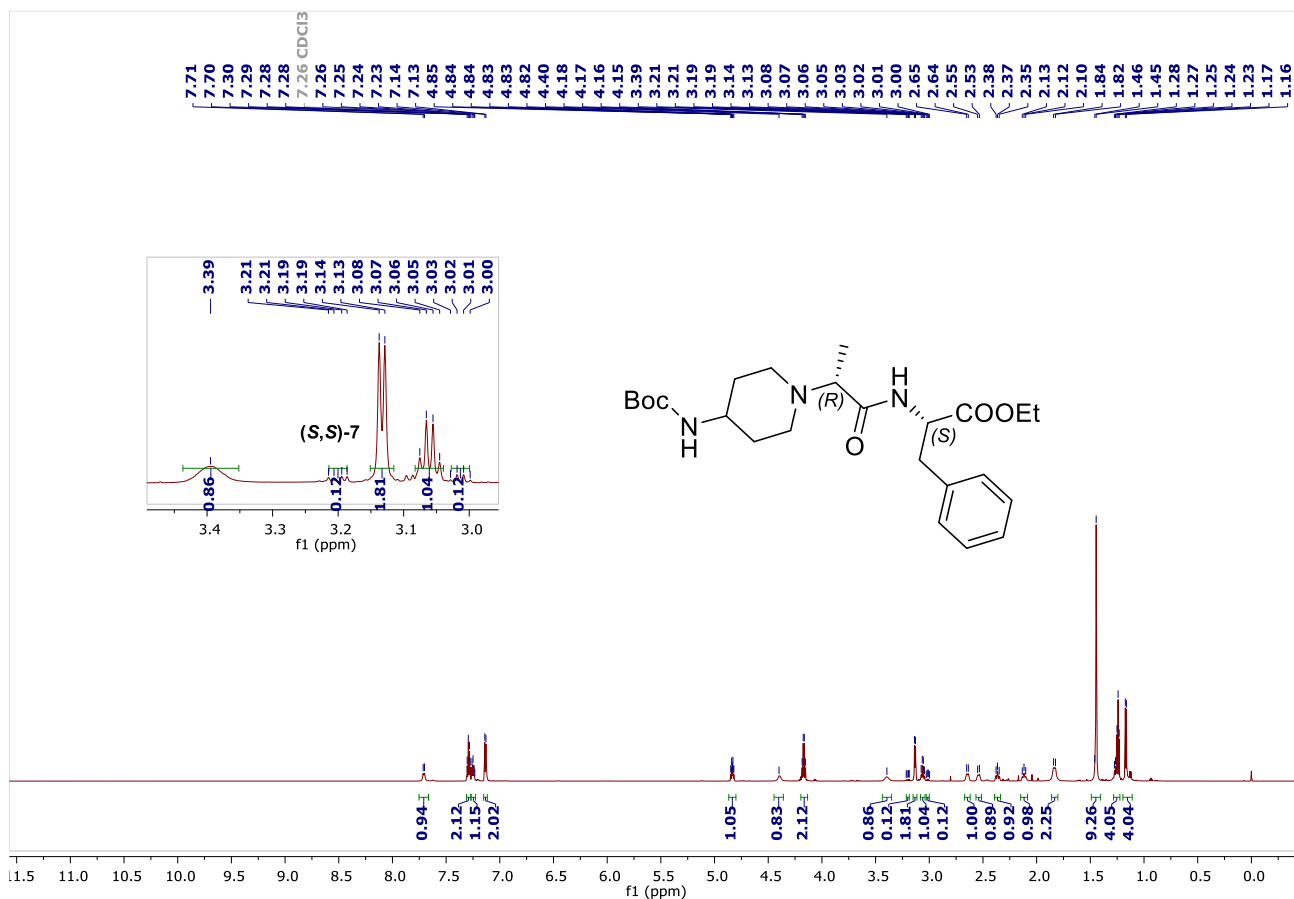


Figure S41. Ethyl *N*-[(2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*R,S*)-7). ¹H NMR spectrum (700 MHz, CDCl₃).

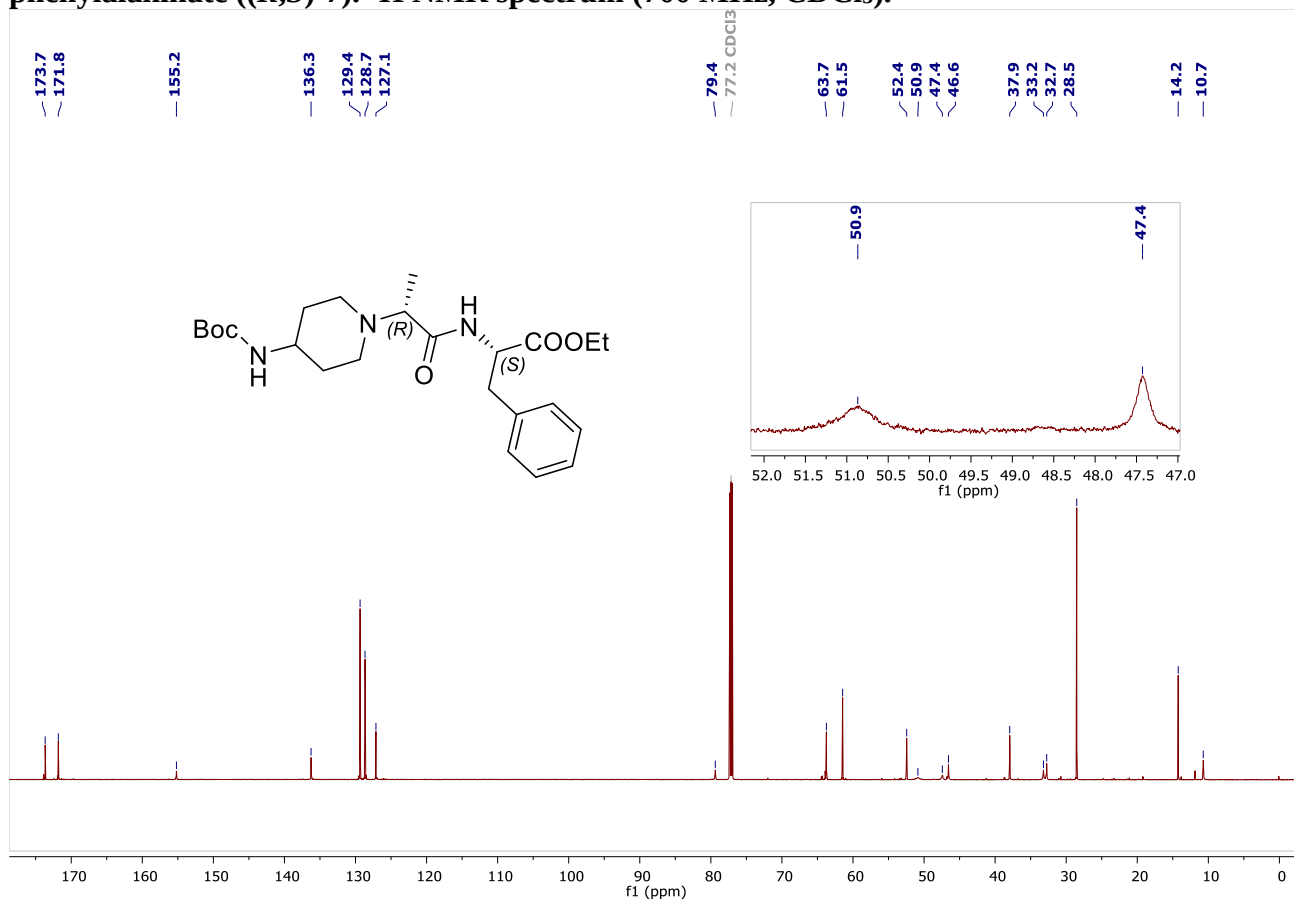


Figure S42. Ethyl *N*-[(2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*R,S*)-7). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

Analysis Name D:\Data\Organikai\2023_04_11\GMP_1216_1-C,4_01_10226.d
Method organikai_esi_pos_2013_recover.m
Sample Name GMP_1216
Comment

Acquisition Date 4/21/2023 1:43:17 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
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Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste

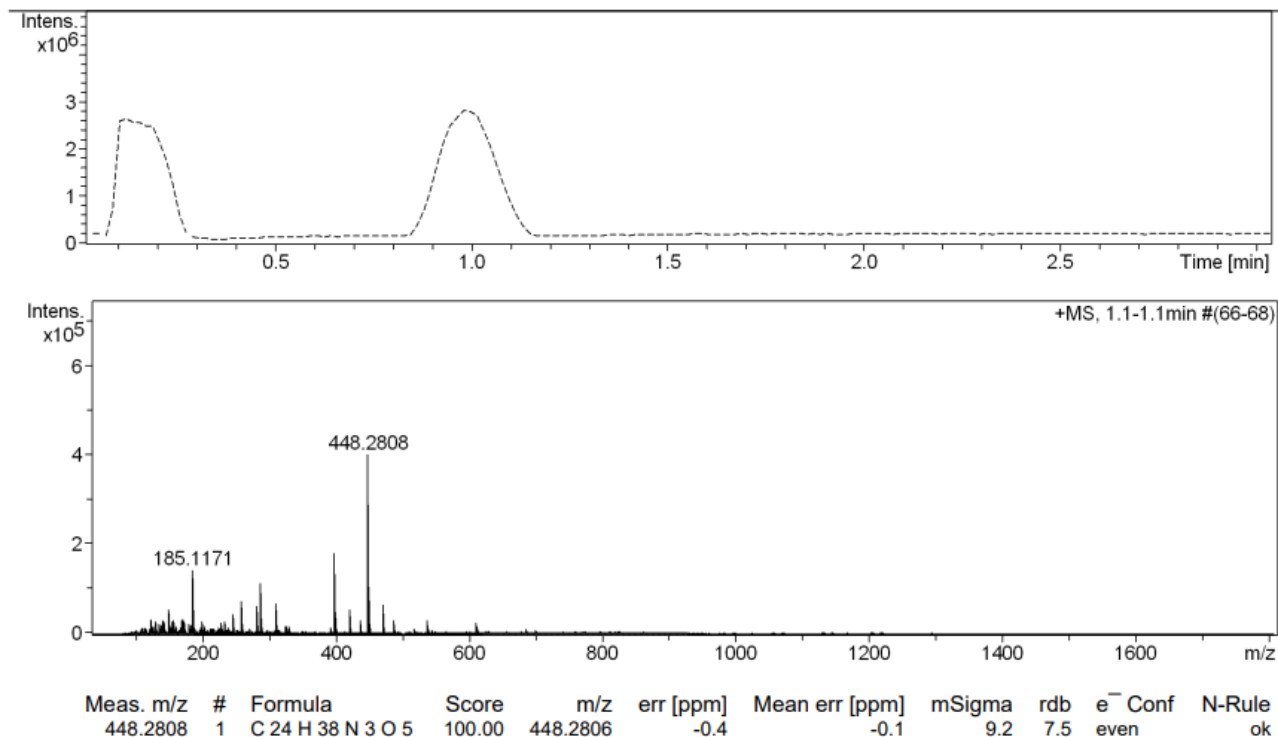


Figure S43. Ethyl *N*-[(2*R*)-2-{4-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*R,S*)-7). HRMS (ESI-TOF).

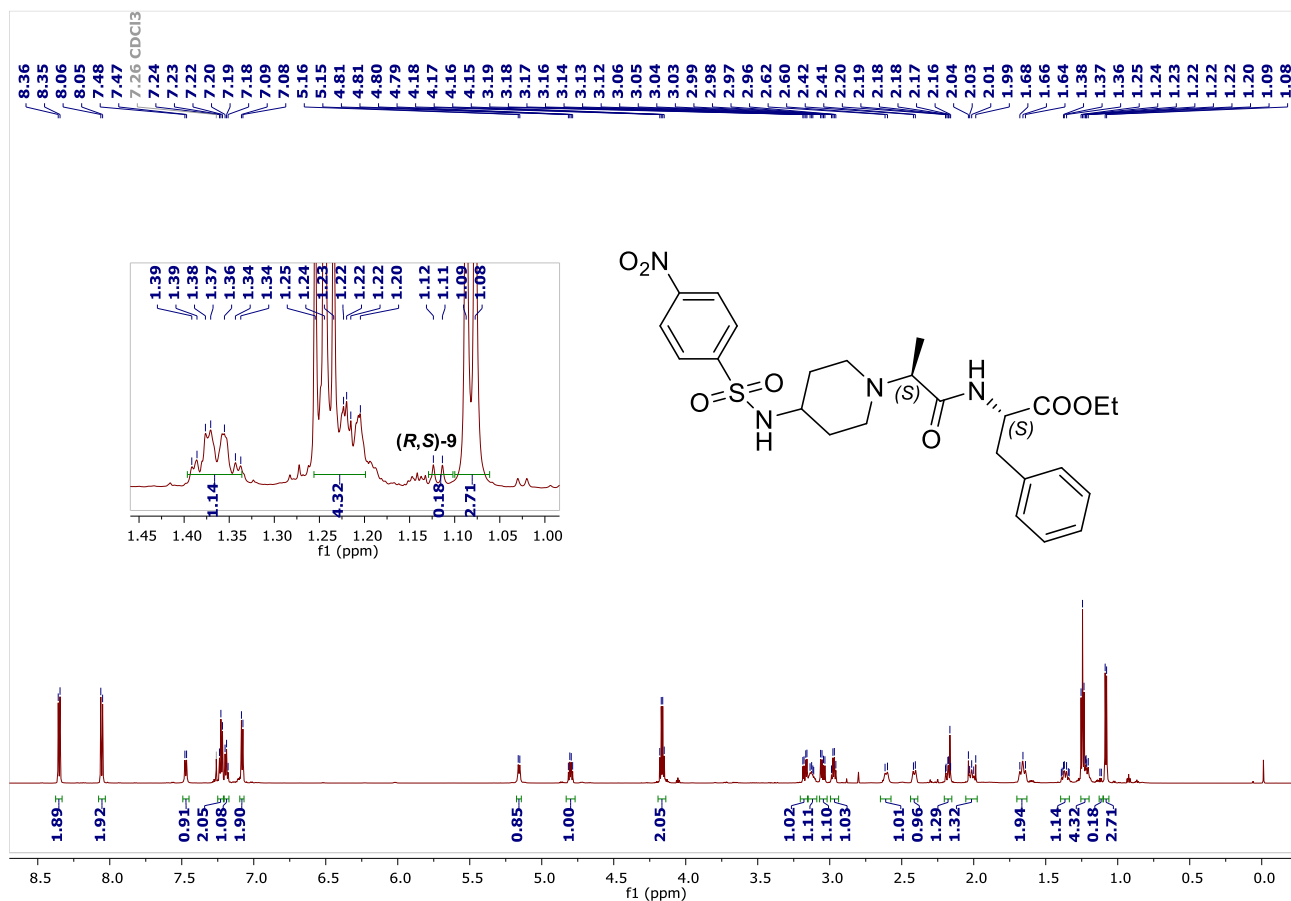


Figure S44. Ethyl *N*-[(2*S*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*S,S*)-9). ¹H NMR spectrum (700 MHz, CDCl₃).

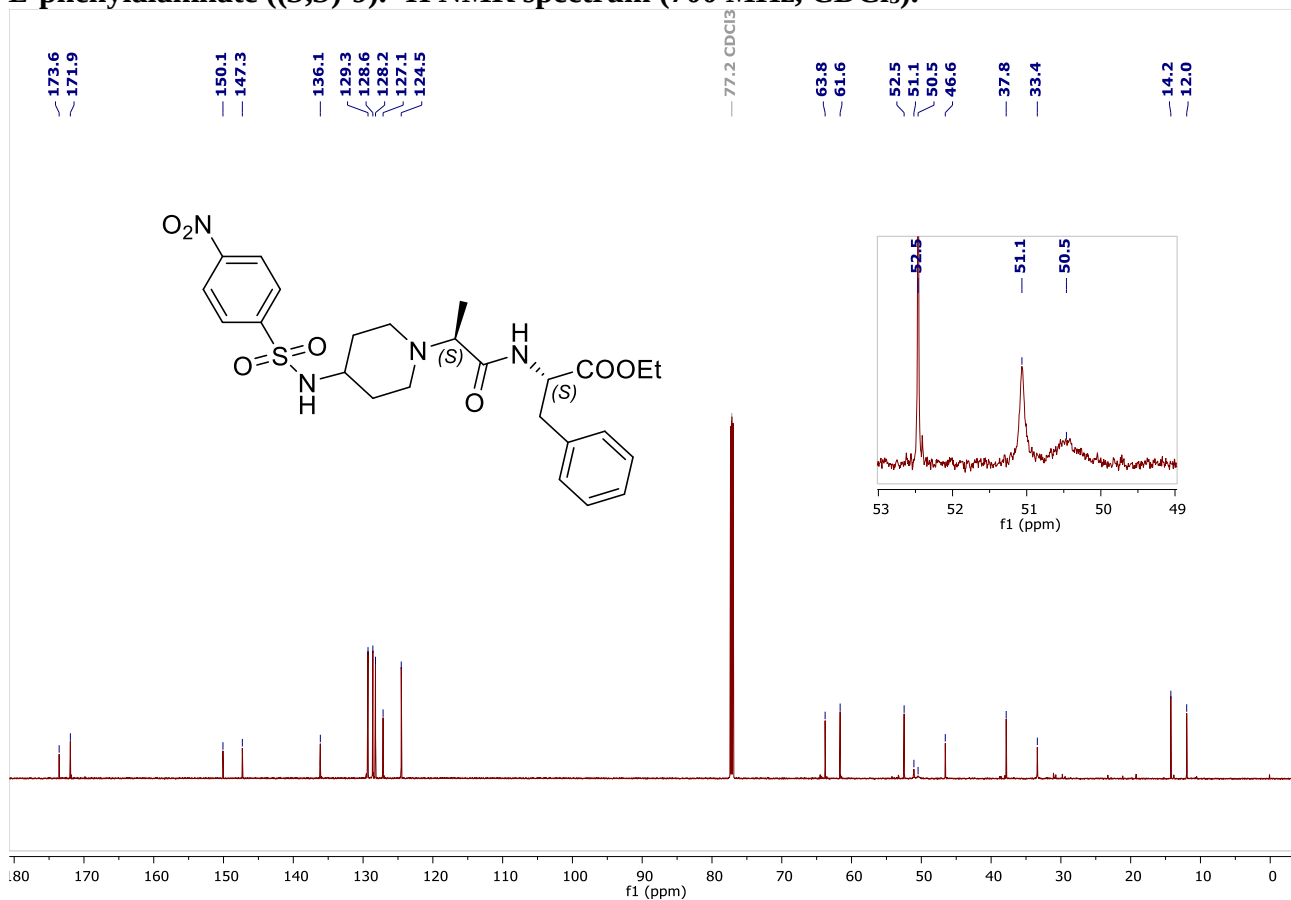


Figure S45. Ethyl *N*-[(2*S*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*S,S*)-9). ¹³C NMR spectrum (176 MHz, CDCl₃).

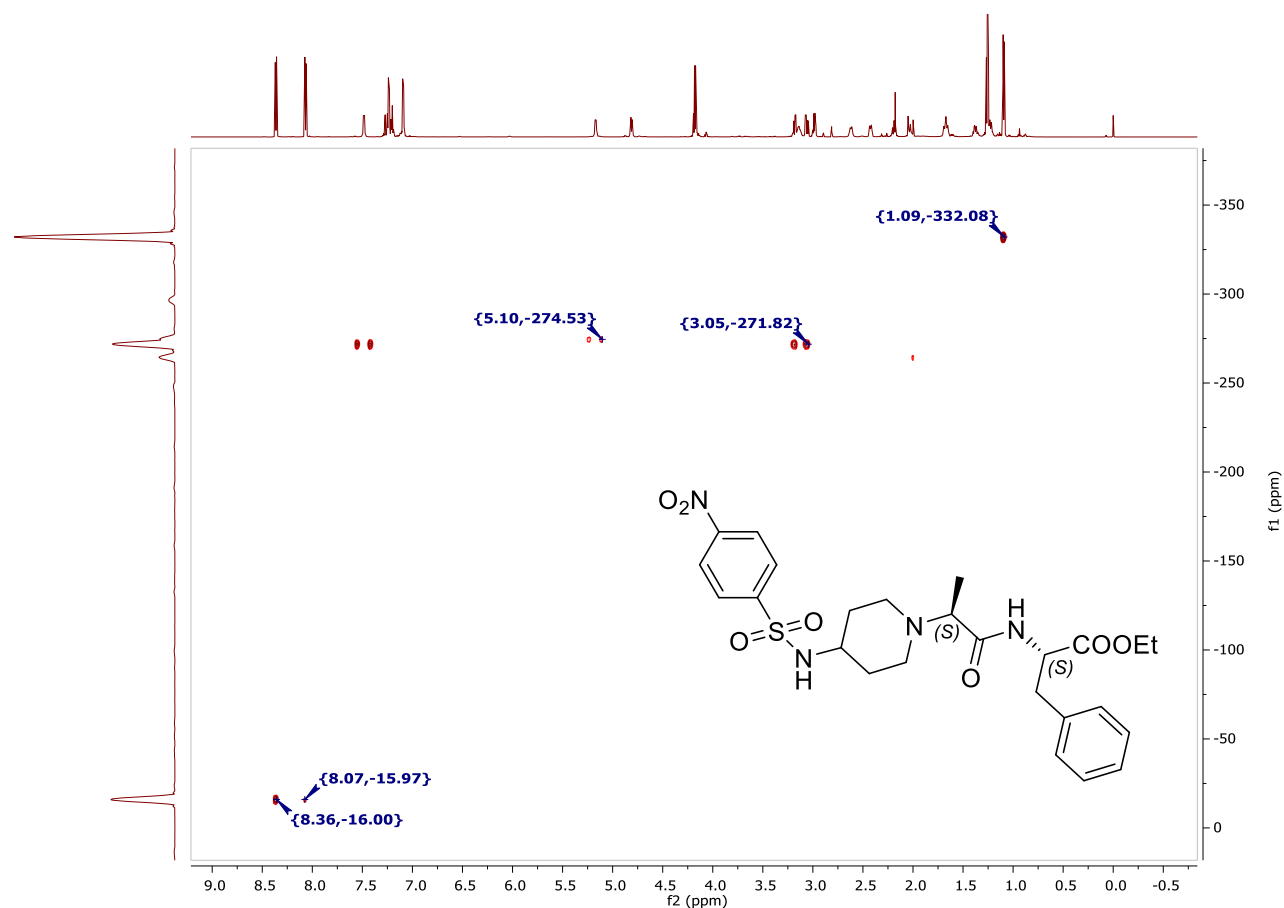


Figure S46. Ethyl *N*-[(2*S*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*S,S*)-9). ¹H-¹⁵N HMBC spectrum (71 MHz, CDCl₃).

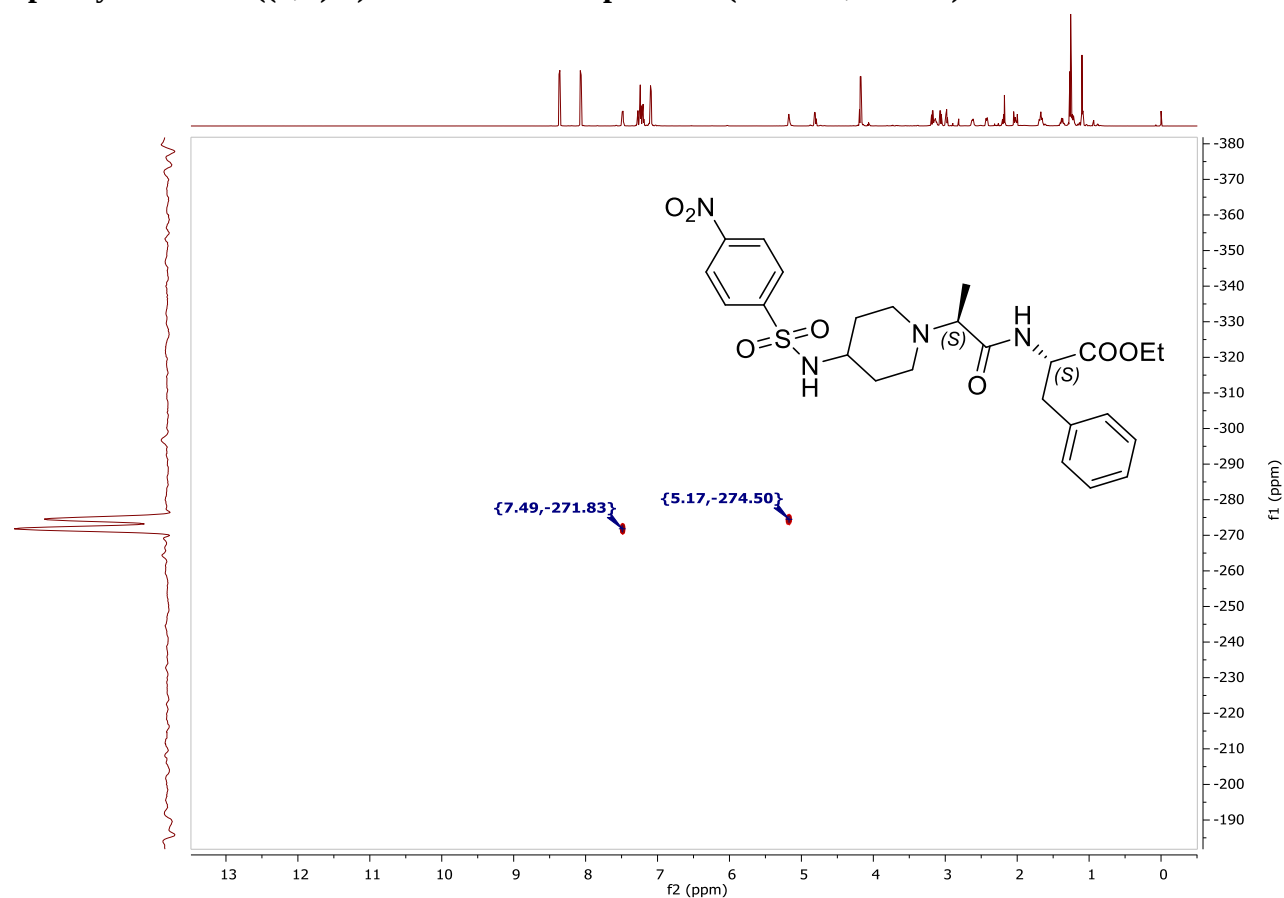


Figure S47. Ethyl *N*-[(2*S*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*S,S*)-9). ¹H-¹⁵N HSQC spectrum (71 MHz, CDCl₃).

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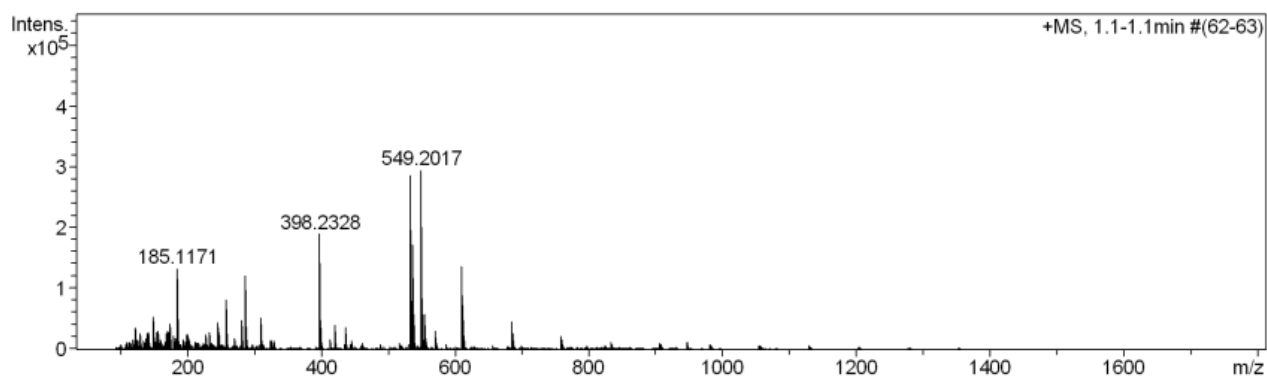
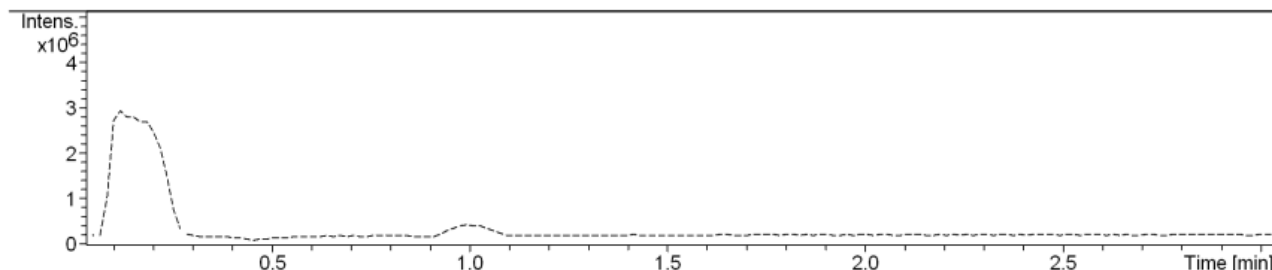
Analysis Info

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 Sample Name GMP_1217
 Comment

Acquisition Date 4/21/2023 1:56:42 PM
 Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSig ma	rdb	e ⁻ Conf	N-R ule
533.2068	1	C 24 H 37 O 11 S	49.10	533.2051	-3.1	-3.4	15.3	6.5	even	ok
	2	C 25 H 33 N 4 O 7 S	100.00	533.2064	-0.6	-0.9	17.9	11.5	even	ok
	3	C 20 H 37 O 16	63.13	533.2076	1.6	2.2	27.1	2.5	even	ok
	4	C 33 H 29 N 2 O 5	51.17	533.2071	0.6	1.5	44.9	20.5	even	ok

Figure S48. Ethyl *N*-[(2*S*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*S,S*)-9). HRMS (ESI-TOF).

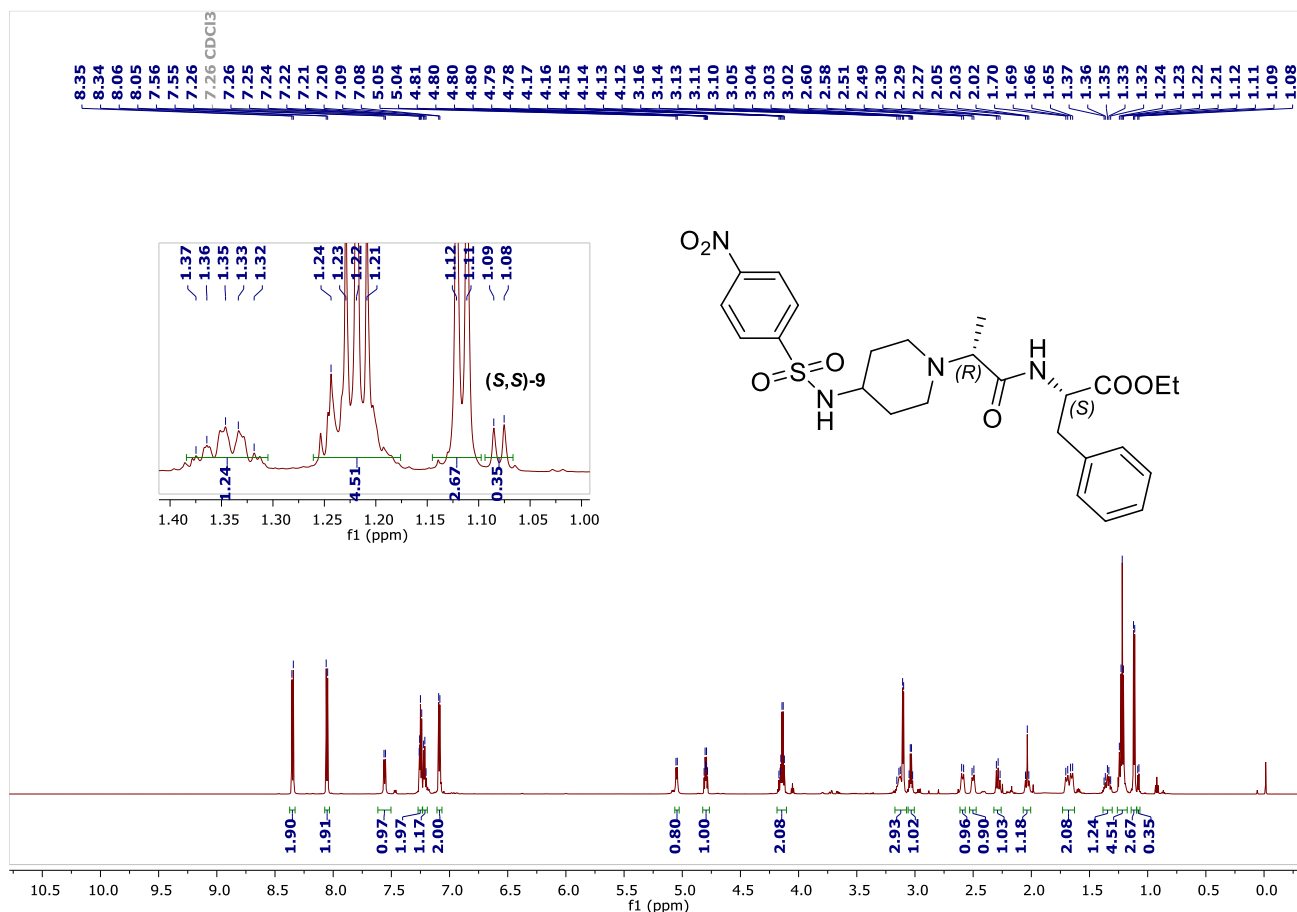


Figure S49. Ethyl *N*-[(2*R*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*R,S*)-9). ¹H NMR spectrum (700 MHz, CDCl₃).

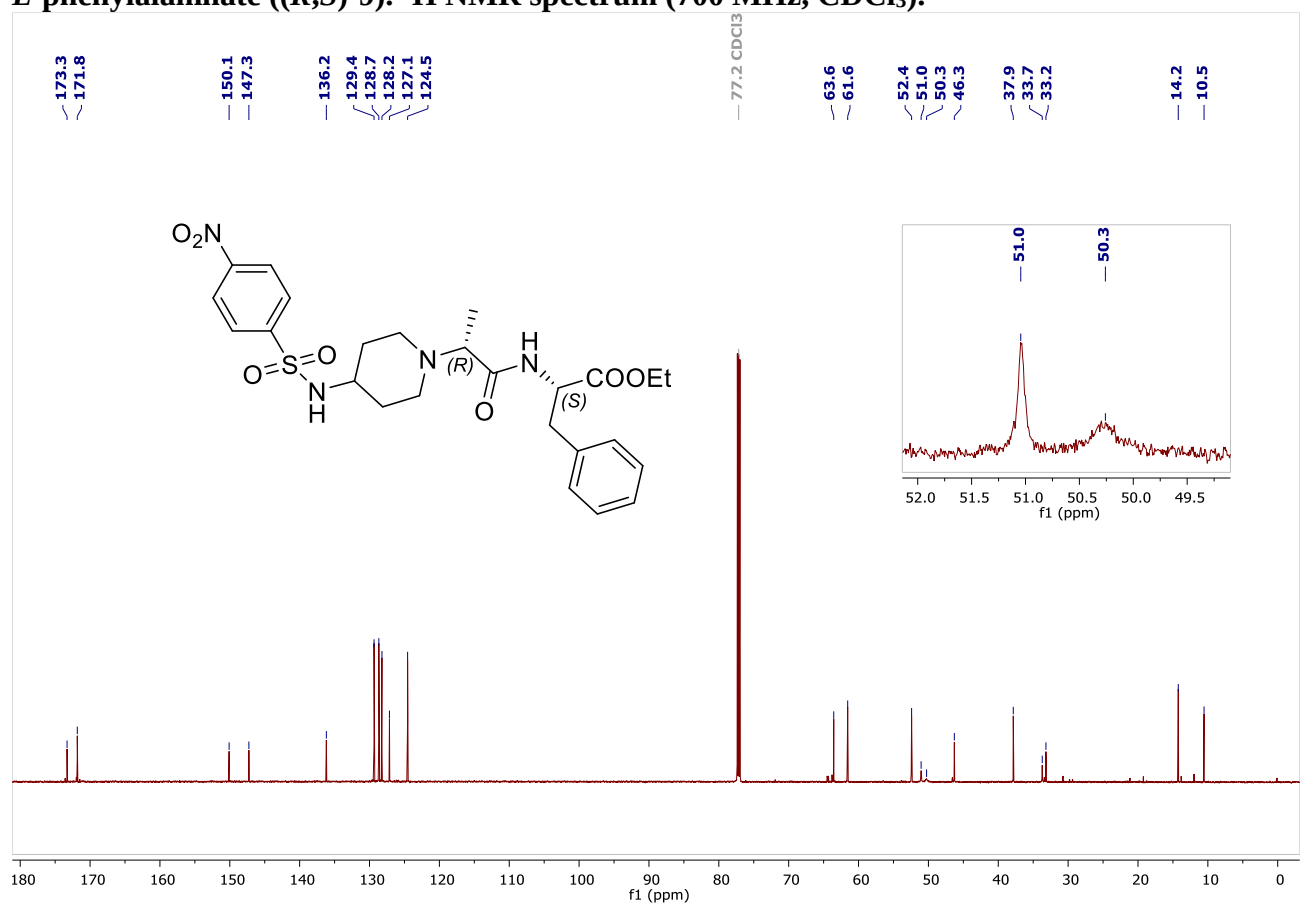


Figure S50. Ethyl *N*-[(2*R*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*R,S*)-9). ¹³C NMR spectrum (176 MHz, CDCl₃).

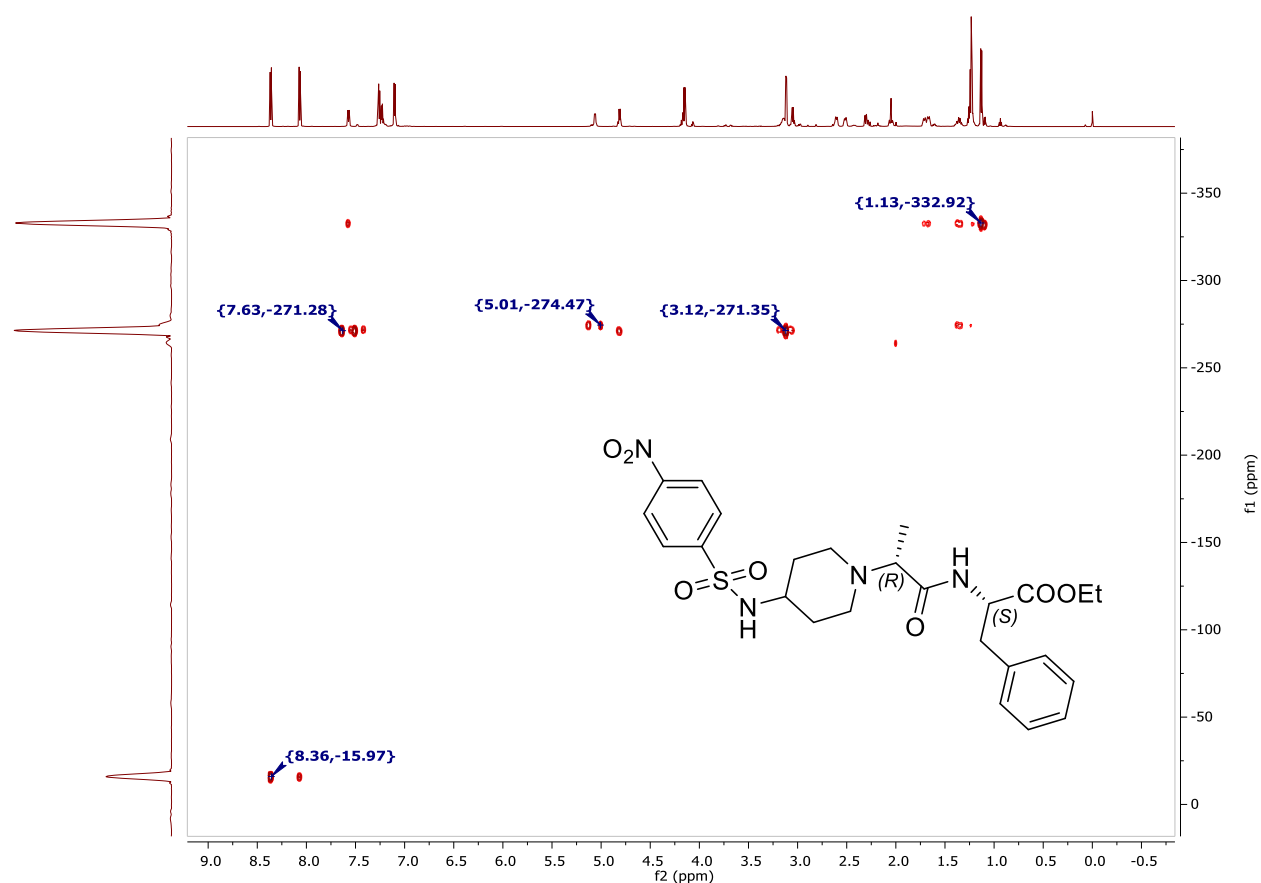


Figure S51. Ethyl *N*-[(2*R*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*R,S*)-9). ^1H - ^{15}N HMBC spectrum (71 MHz, CDCl_3).

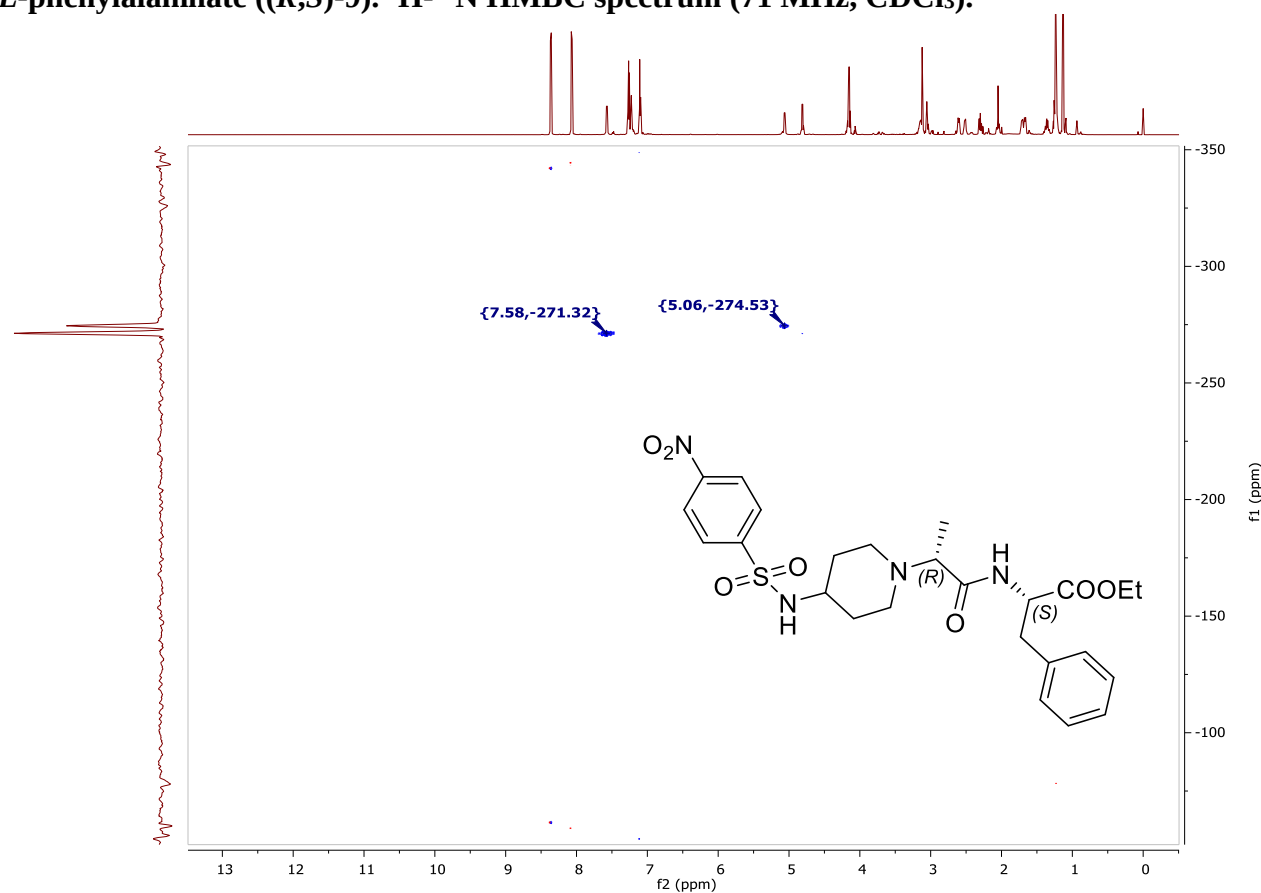


Figure S52. Ethyl *N*-[(2*R*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*R,S*)-9). ^1H - ^{15}N HSQC spectrum (71 MHz, CDCl_3).

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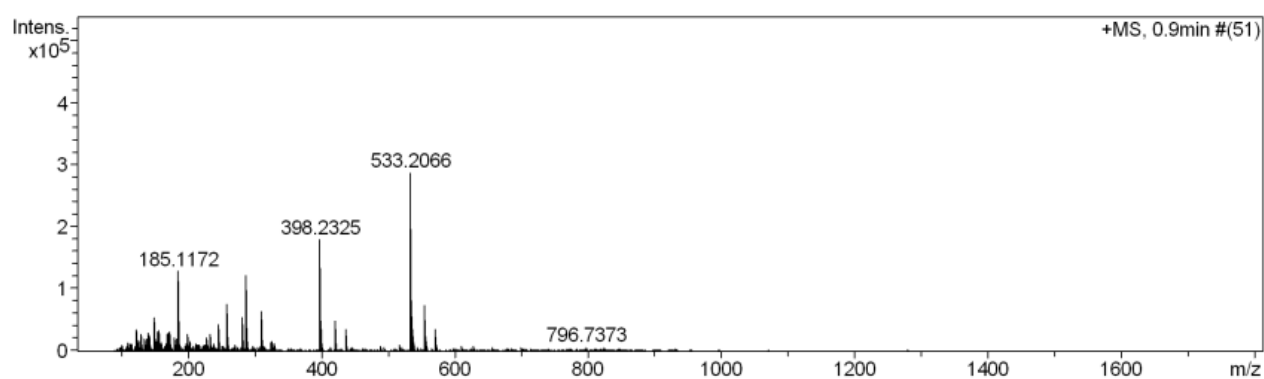
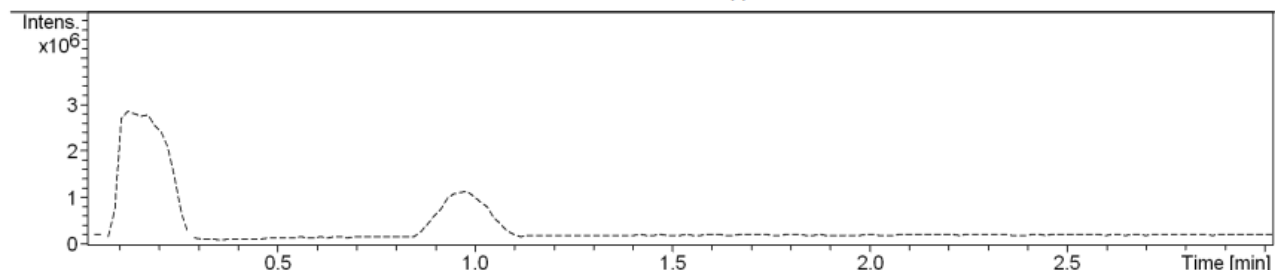
Analysis Info

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 Comment

Acquisition Date 4/21/2023 1:52:14 PM
 Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSig ma	rdb	e ⁻ Conf	N-R ule
533.2066	1	C 24 H 37 O 11 S	48.79	533.2051	-2.8	-3.1	17.8	6.5	even	ok
	2	C 25 H 33 N 4 O 7 S	100.00	533.2064	-0.3	-0.6	18.0	11.5	even	ok
	3	C 20 H 37 O 16	50.43	533.2076	1.9	2.4	29.4	2.5	even	ok
	4	C 33 H 29 N 2 O 5	48.33	533.2071	0.9	1.7	41.4	20.5	even	ok

Figure S53. Ethyl *N*-[(2*R*)-2-{4-[(4-nitrobenzene-1-sulfonyl)amino]piperidin-1-yl}propanoyl]-*L*-phenylalaninate ((*R,S*)-9). HRMS (ESI-TOF).

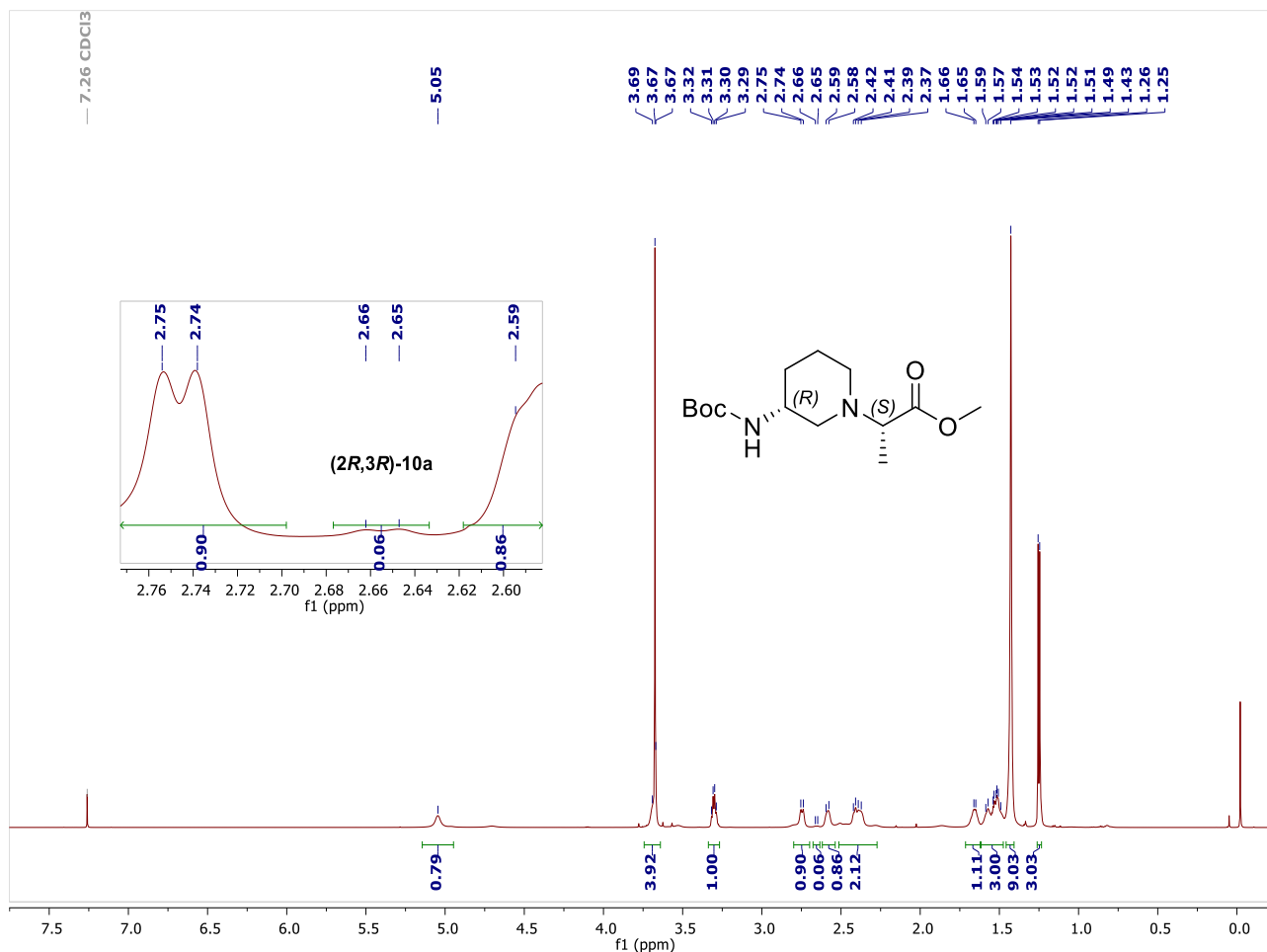


Figure S54. Methyl (2S)-2-((3R)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl)propanoate ((2S,3R)-10a). ¹H NMR spectrum (700 MHz, CDCl₃).

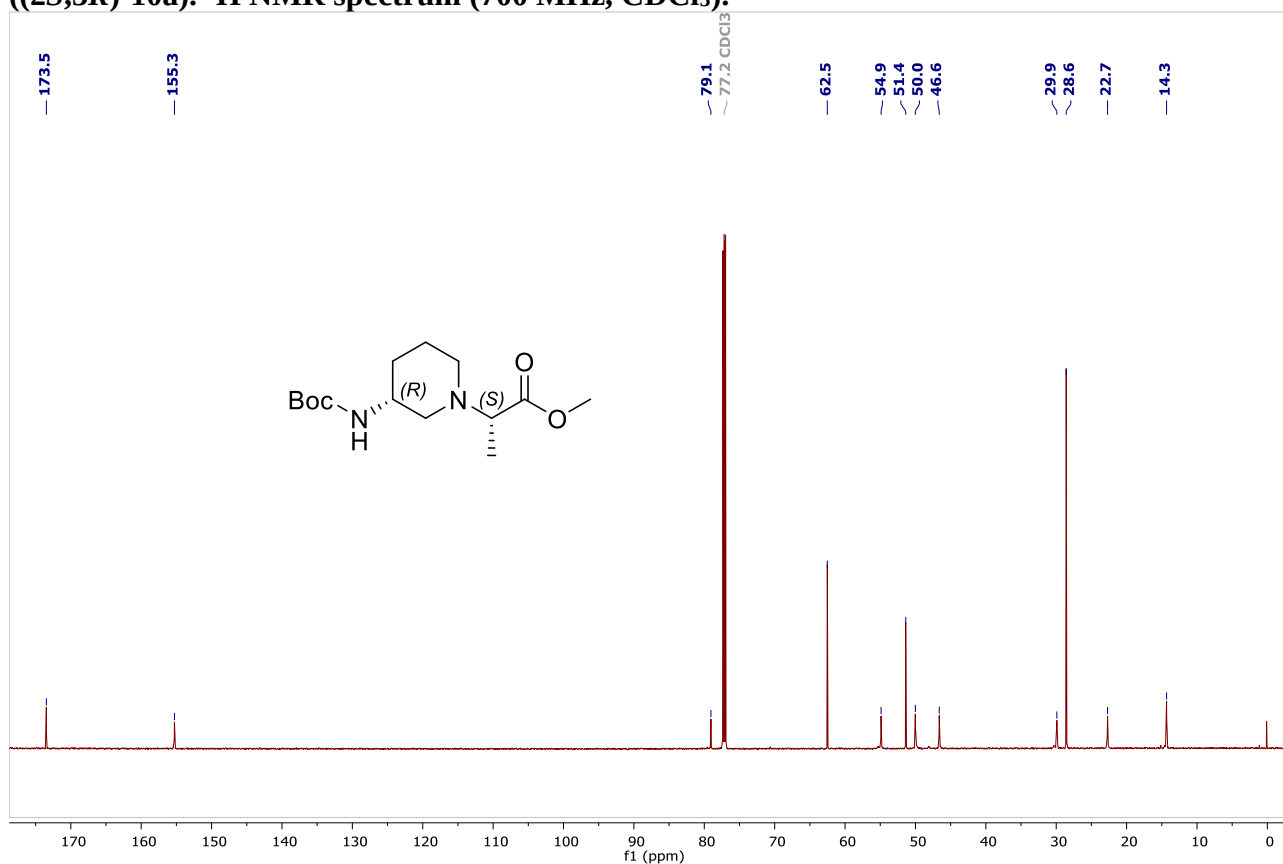
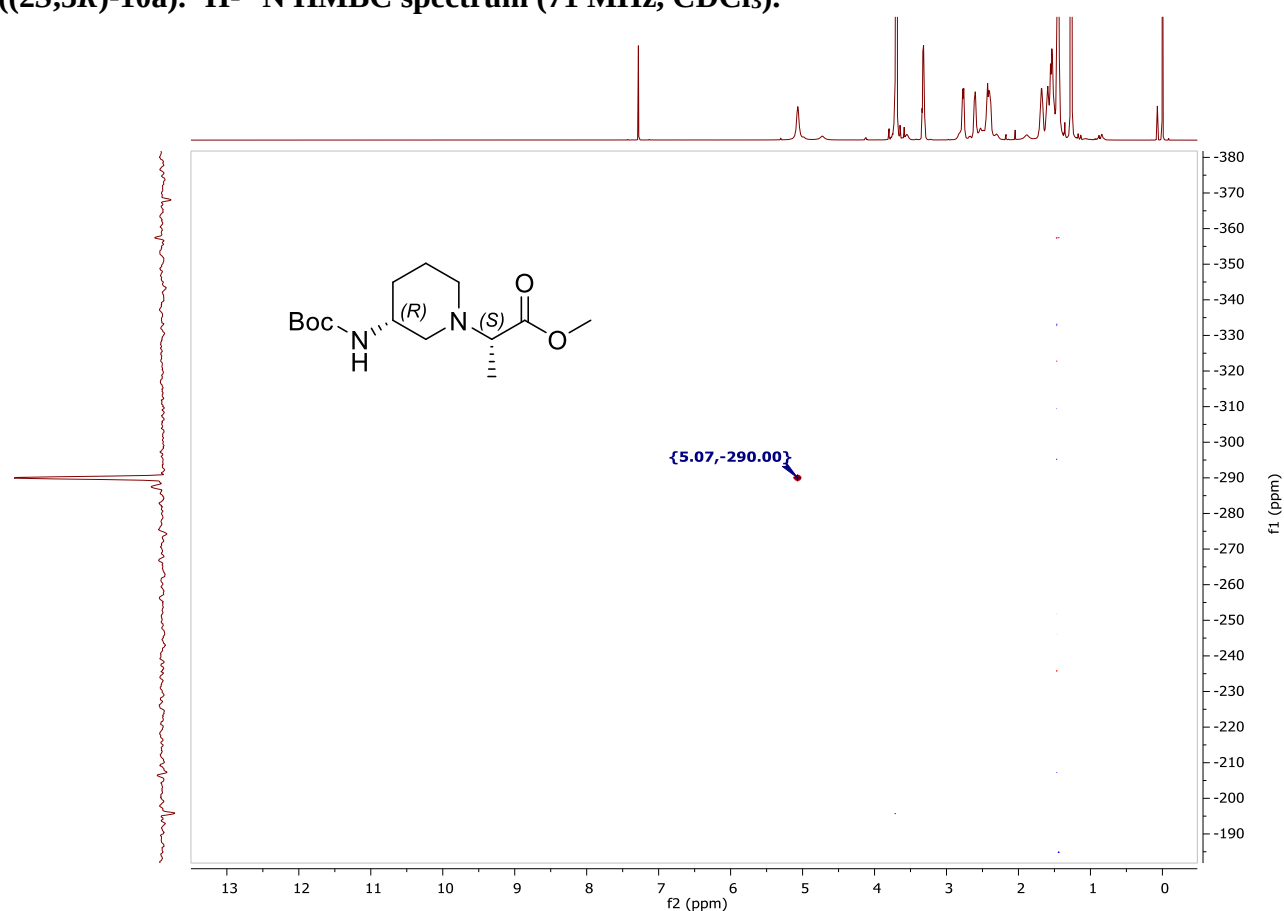
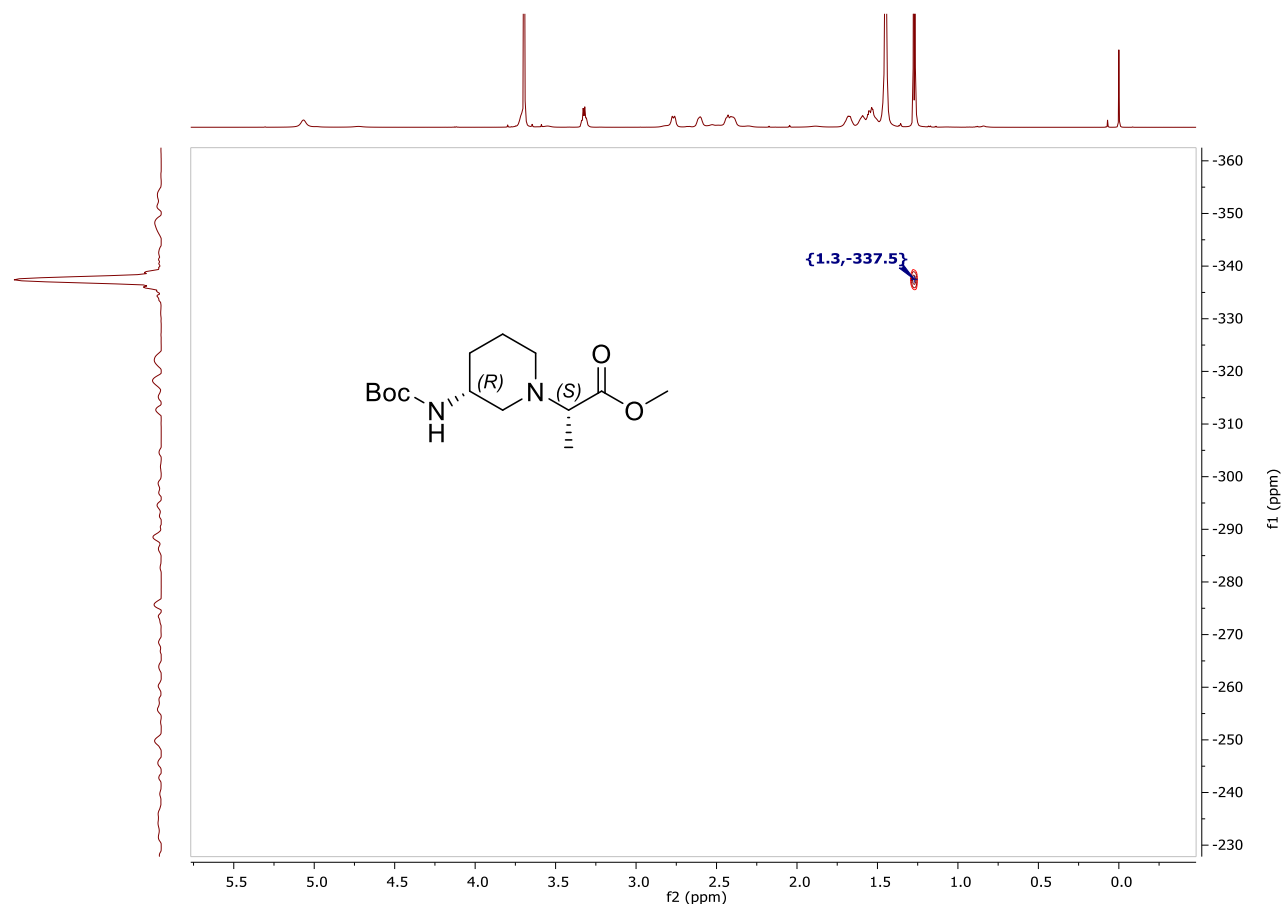


Figure S55. Methyl (2S)-2-((3R)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl)propanoate ((2S,3R)-10a). ¹³C NMR spectrum (176 MHz, CDCl₃).



Mass Spectrum SmartFormula Report

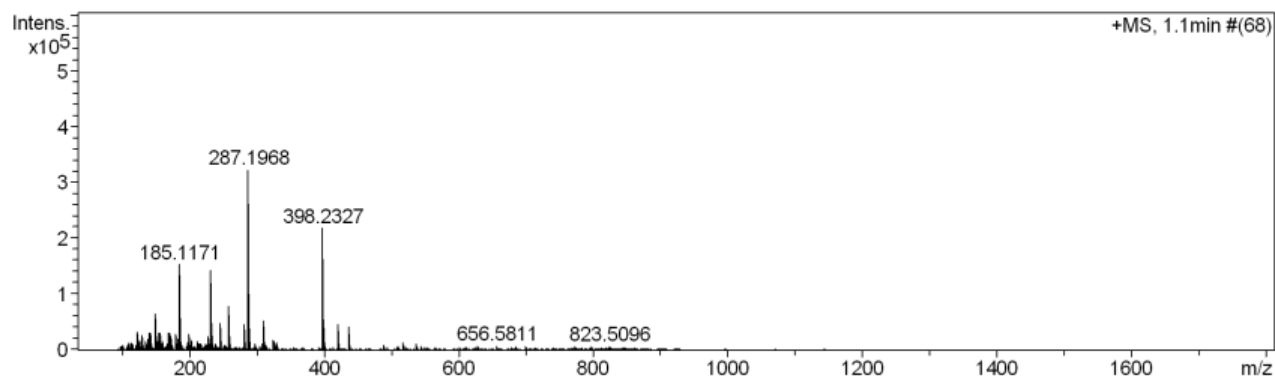
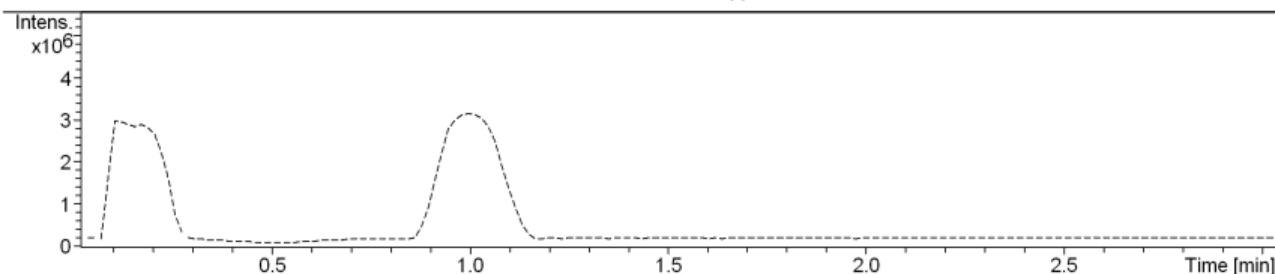
Analysis Info

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 Sample Name GMP_367
 Comment

Acquisition Date 4/21/2023 2:01:08 PM
 Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
287.1968	1	C ₁₄ H ₂₇ N ₂ O ₄	100.00	287.1965	-0.9	-0.7	13.3	2.5	even	ok

Figure S58. Methyl (2*S*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]propanoate ((2*S*,3*R*)-10a). HRMS (ESI-TOF).

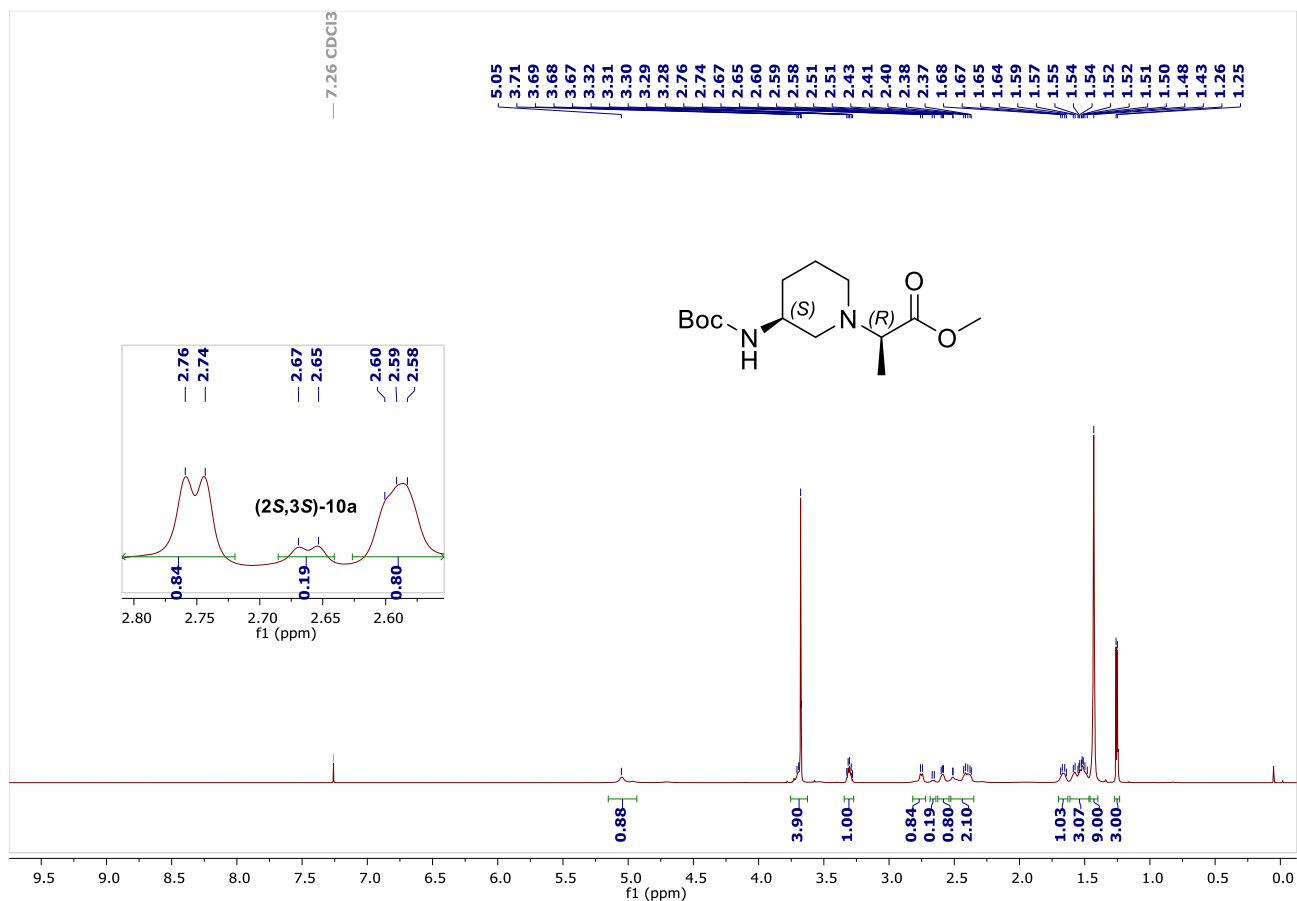


Figure S59. Methyl (2R)-2-((3S)-3-((tert-butoxycarbonyl)amino)piperidin-1-yl)propanoate ((2R,3S)-10a). ¹H NMR spectrum (700 MHz, CDCl₃).

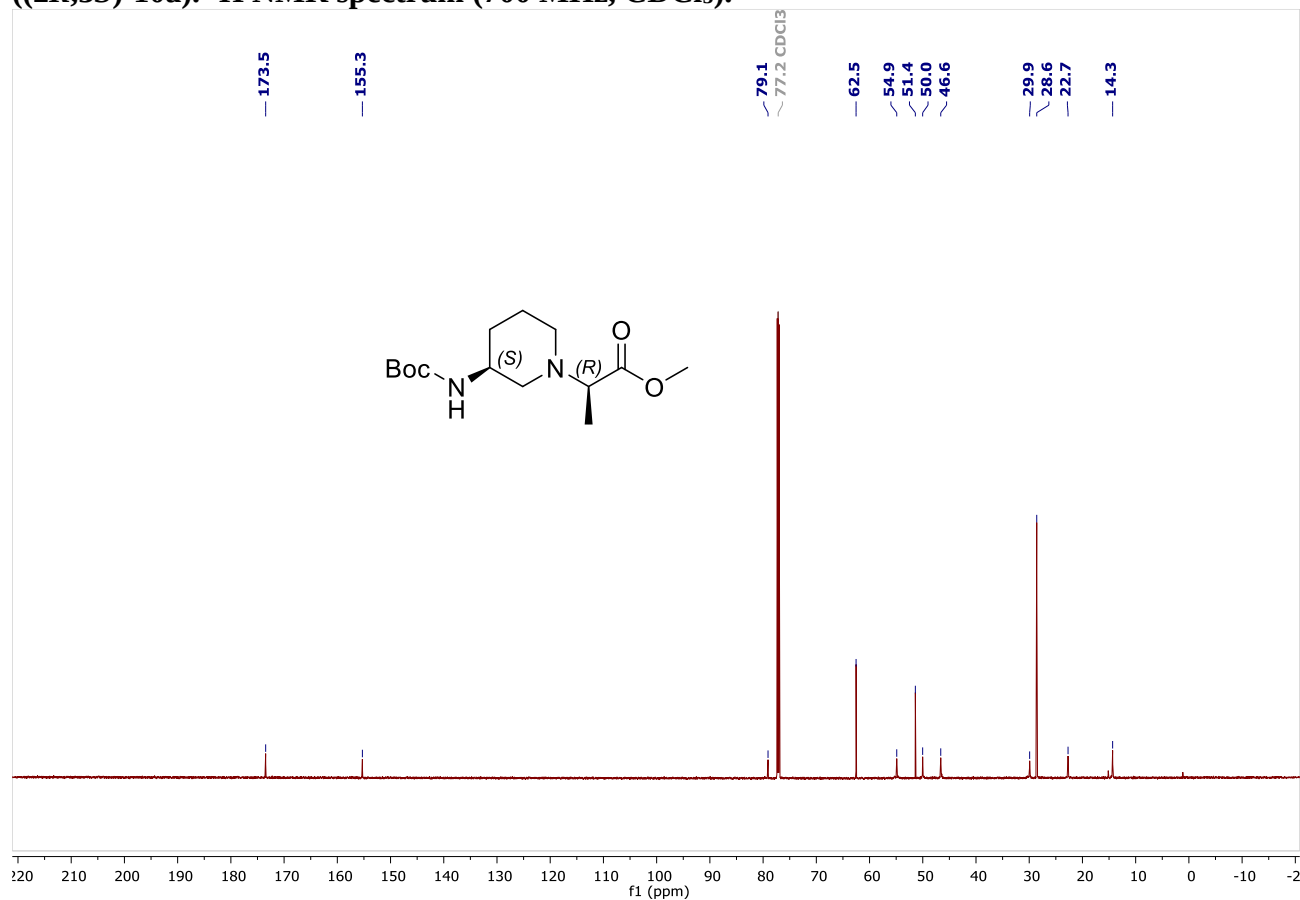


Figure S60. Methyl (2R)-2-((3S)-3-((tert-butoxycarbonyl)amino)piperidin-1-yl)propanoate ((2R,3S)-10a). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

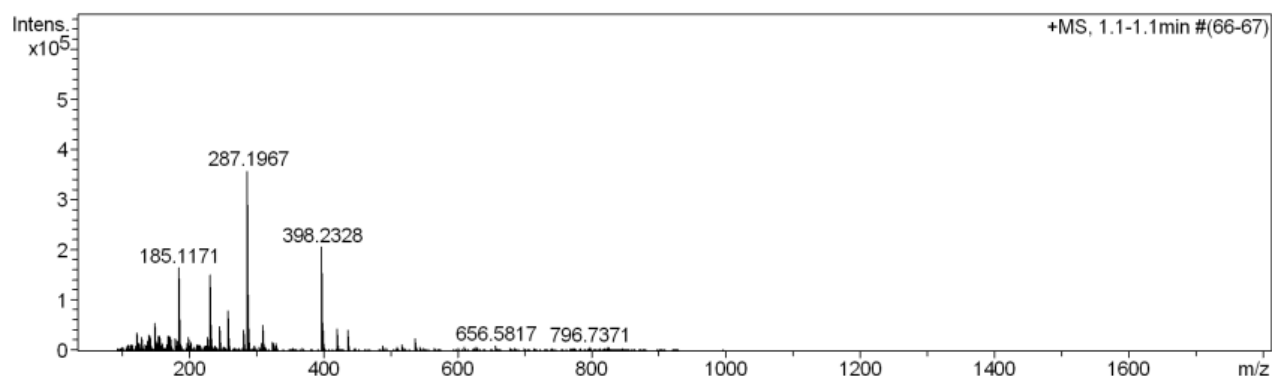
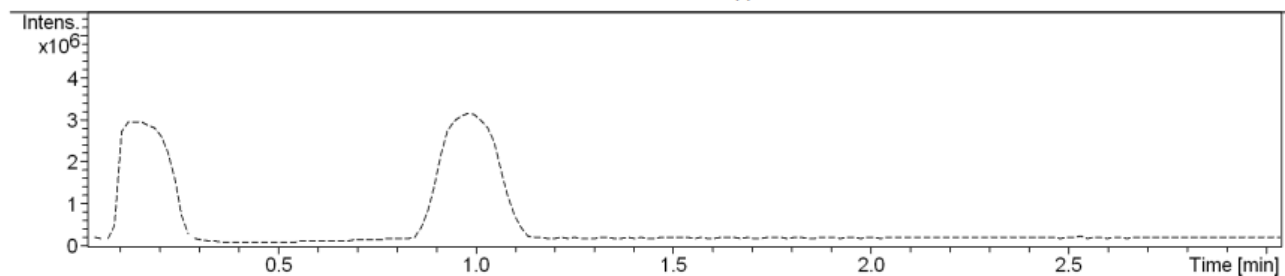
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Method organikai_esi_pos_2013_recover.m
Sample Name GMP_462
Comment

Acquisition Date 4/21/2023 2:14:25 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
287.1967	1	C ₁₄ H ₂₇ N ₂ O ₄	100.00	287.1965	-0.6	-0.2	1.0	2.5	even	ok

Figure S61. Methyl (2R)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl)propanoate ((2R,3S)-10a). HRMS (ESI-TOF).

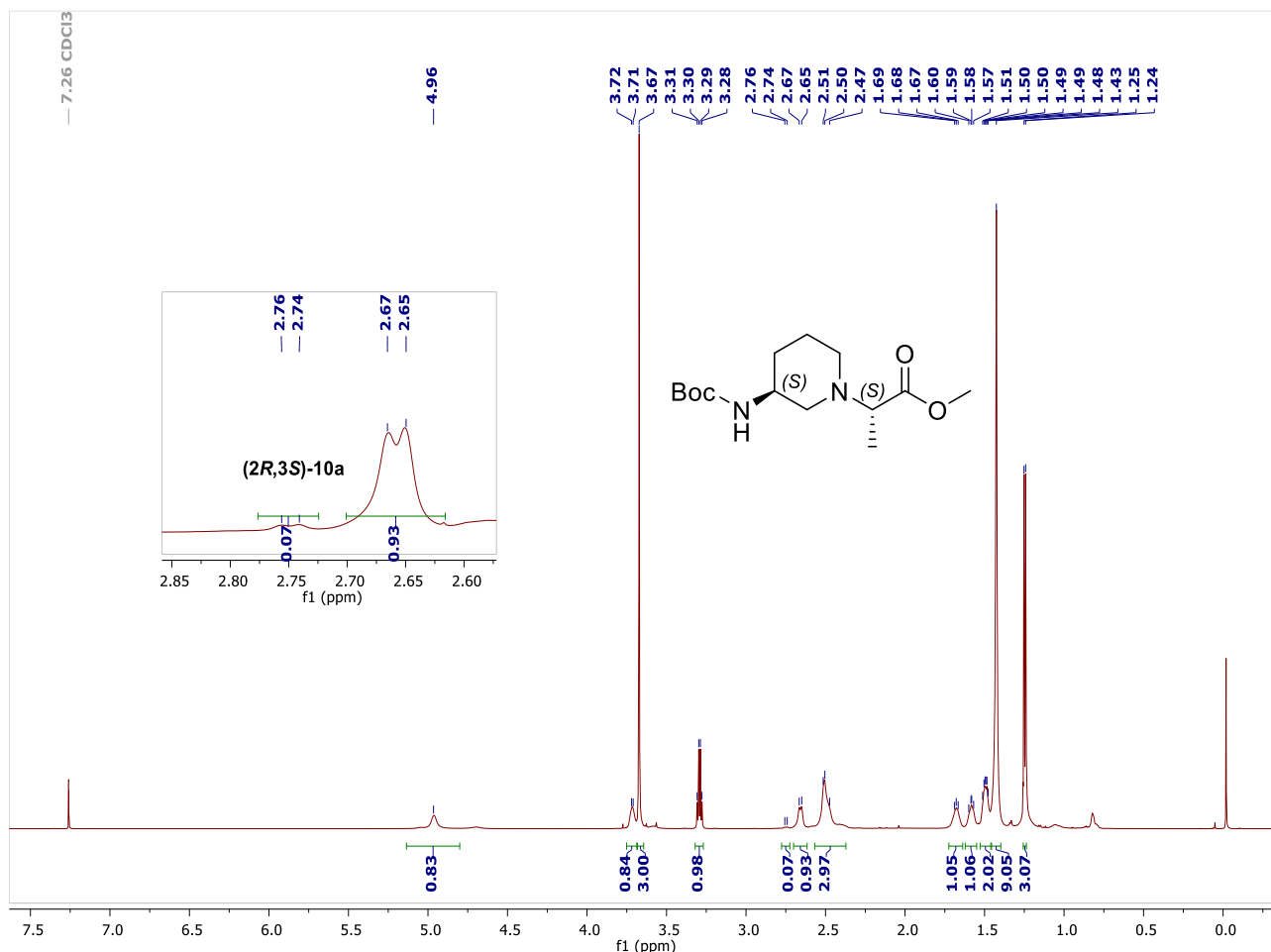


Figure S62. Methyl (2S)-2-((3S)-3-((tert-butoxycarbonyl)amino)piperidin-1-yl)propanoate ((2S,3S)-10a). ¹H NMR spectrum (700 MHz, CDCl₃).

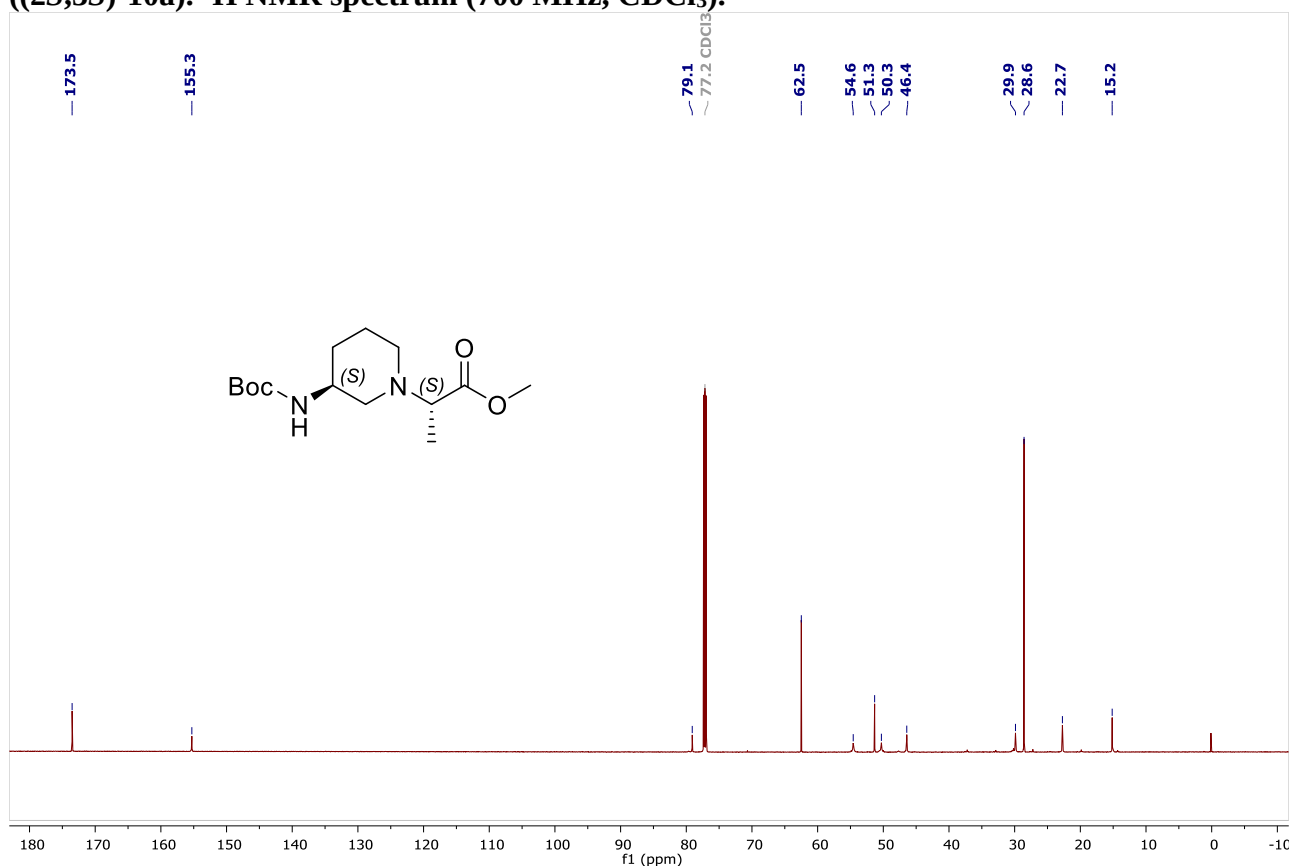


Figure S63. Methyl (2S)-2-((3S)-3-((tert-butoxycarbonyl)amino)piperidin-1-yl)propanoate ((2S,3S)-10a). ¹³C NMR spectrum (176 MHz, CDCl₃).

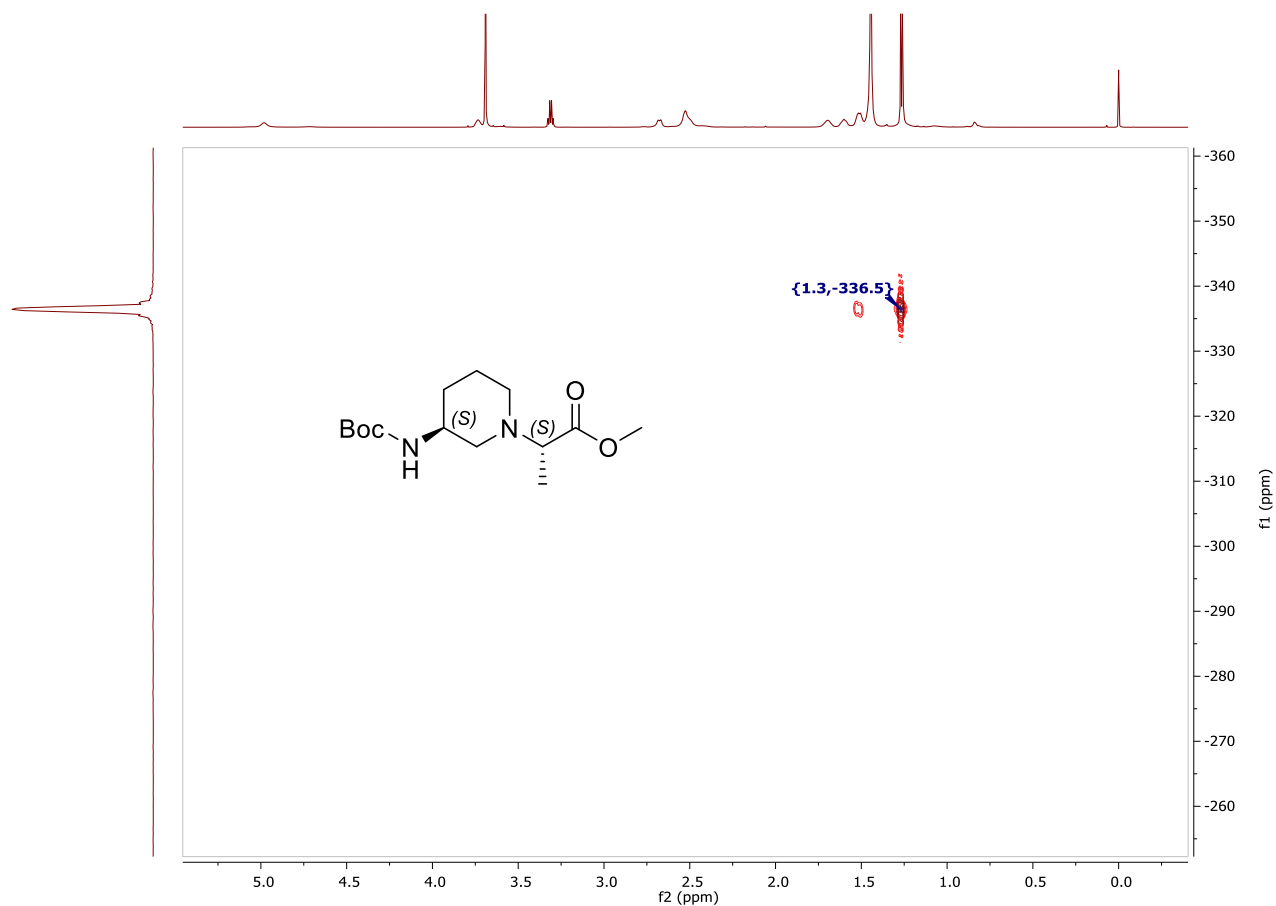


Figure S64. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl)propanoate ((2S,3S)-10a). ^1H - ^{15}N HMBC spectrum (71 MHz, CDCl_3).

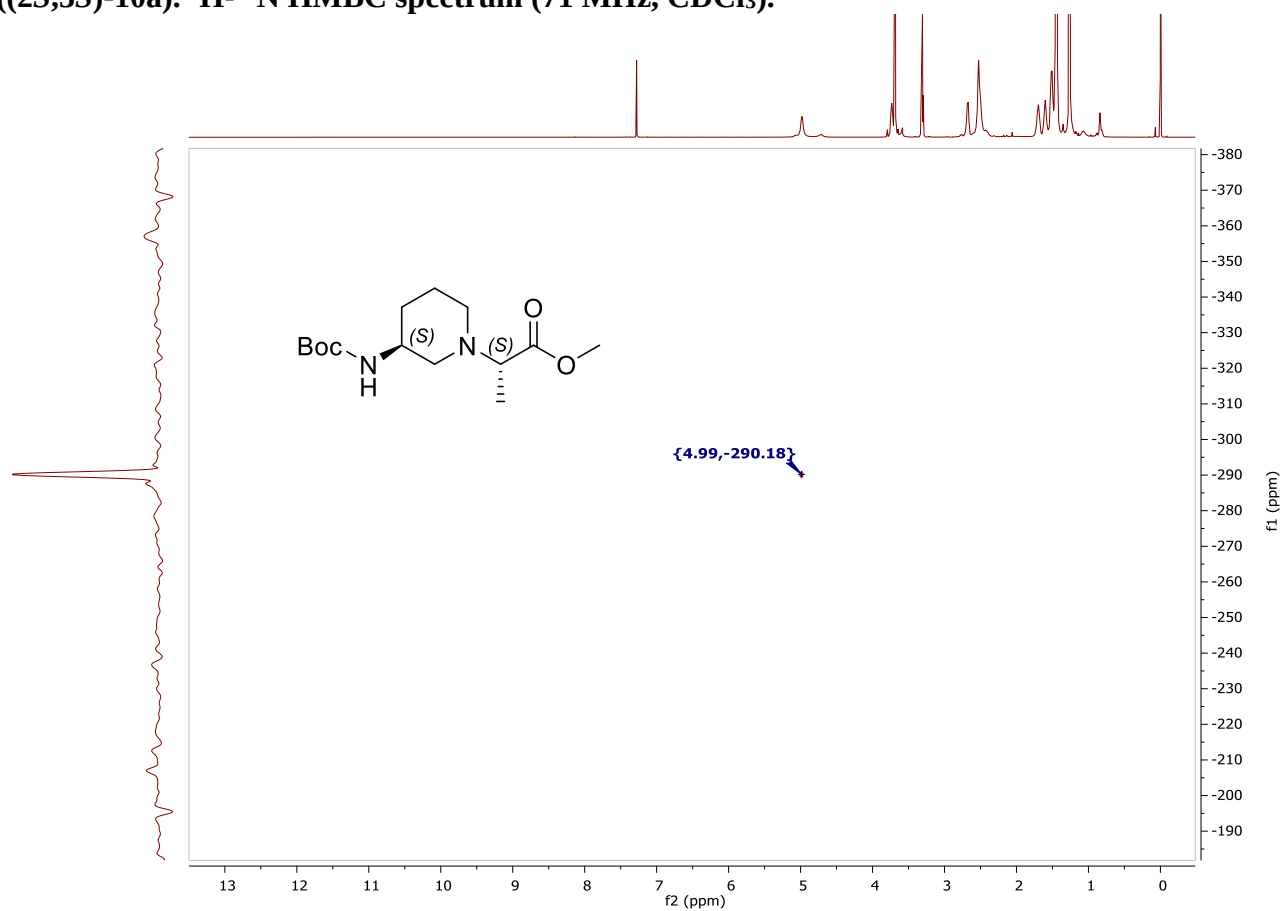


Figure S65. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl)propanoate ((2S,3S)-10a). ^1H - ^{15}N HSQC spectrum (71 MHz, CDCl_3).

Mass Spectrum SmartFormula Report

Analysis Info

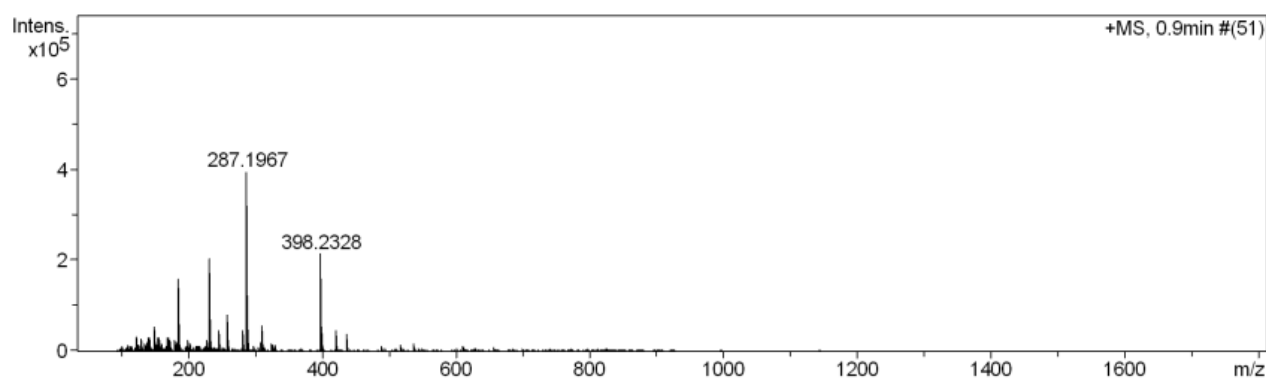
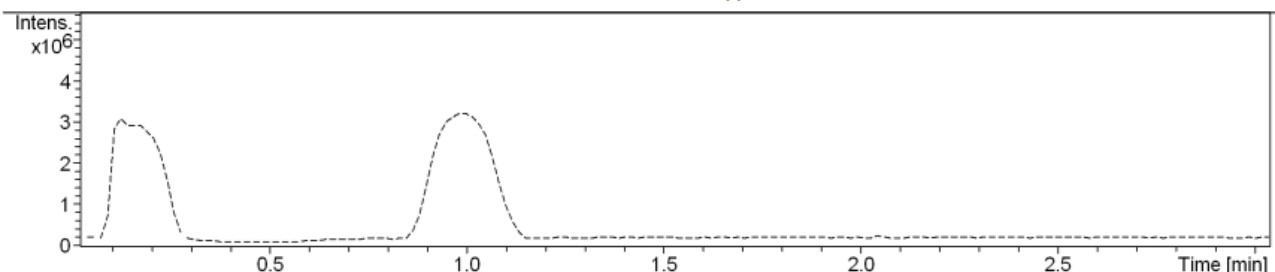
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Sample Name GMP_372
Comment

Acquisition Date 4/21/2023 2:09:59 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

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Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
287.1967	1	C 14 H 27 N 2 O 4	100.00	287.1965	-0.6	-0.5	13.0	2.5	even	ok

Figure S66. Methyl (2S)-2-[(3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]propanoate ((2S,3S)-10a). HRMS (ESI-TOF).

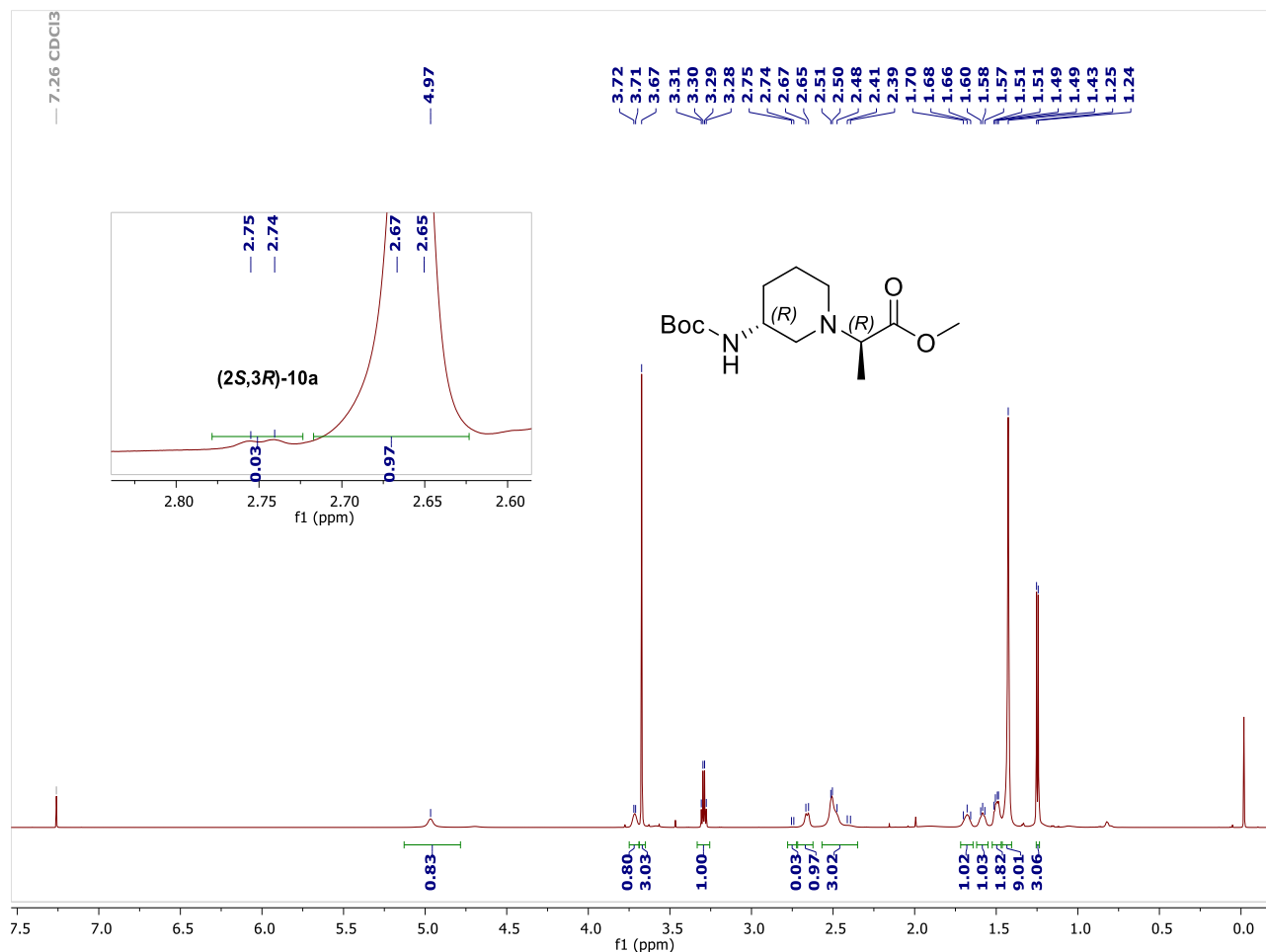


Figure S67. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]propanoate ((2*R*,3*R*)-10a). ¹H NMR spectrum (700 MHz, CDCl₃).

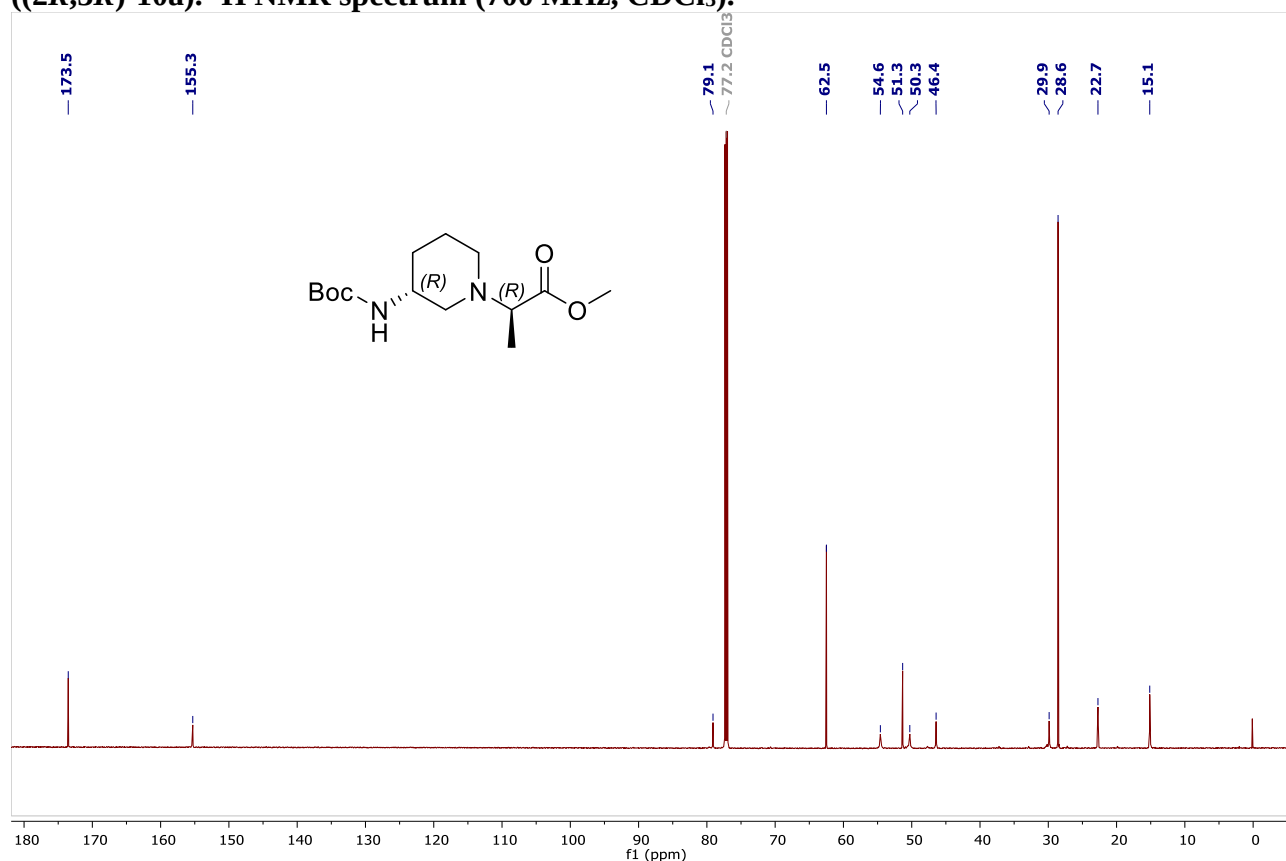


Figure S68. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]propanoate ((2*R*,3*R*)-10a). ¹³C NMR spectrum (176 MHz, CDCl₃).

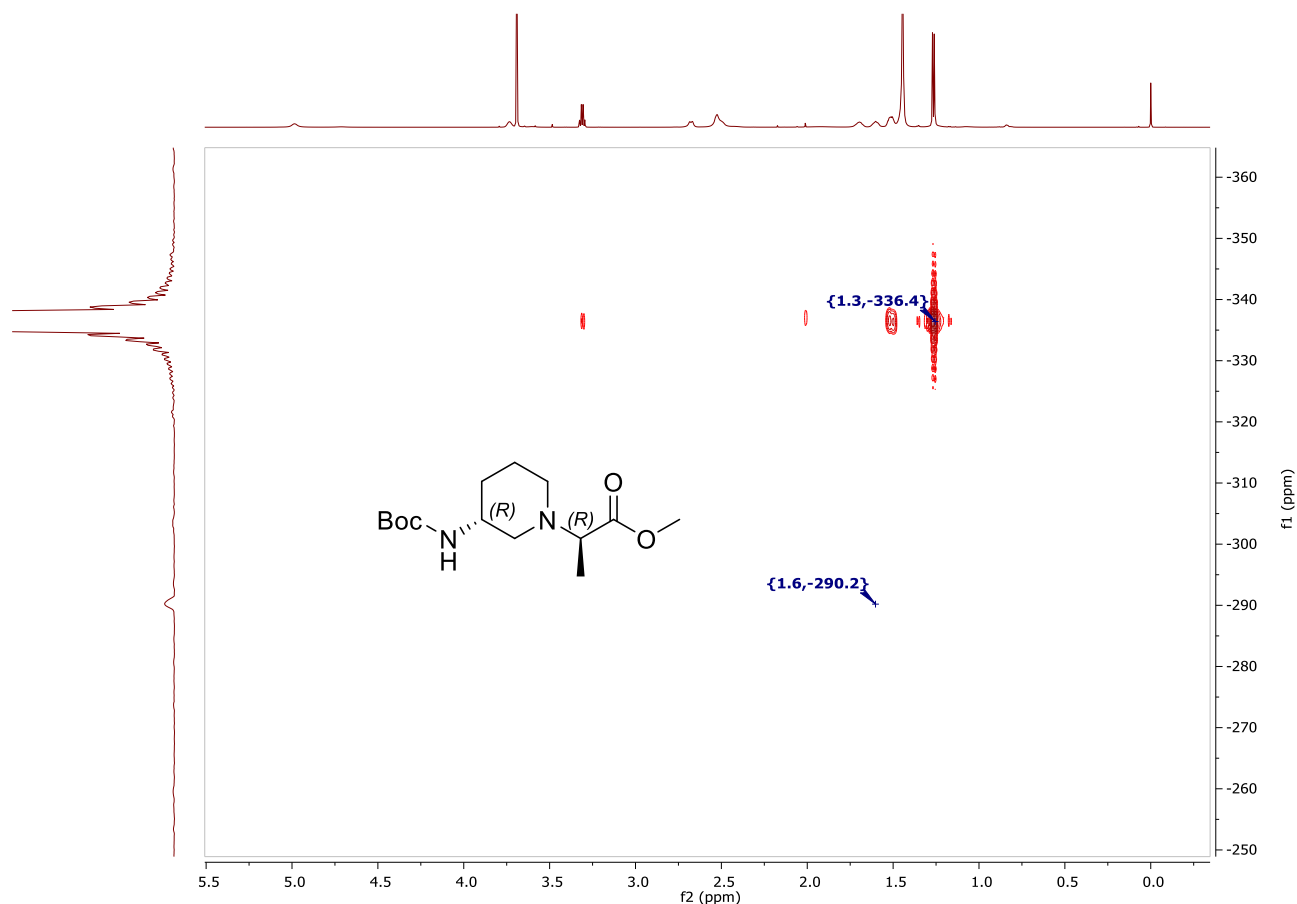


Figure S69. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]propanoate ((2*R*,3*R*)-10a). ^1H - ^{15}N HMBC spectrum (71 MHz, CDCl_3).

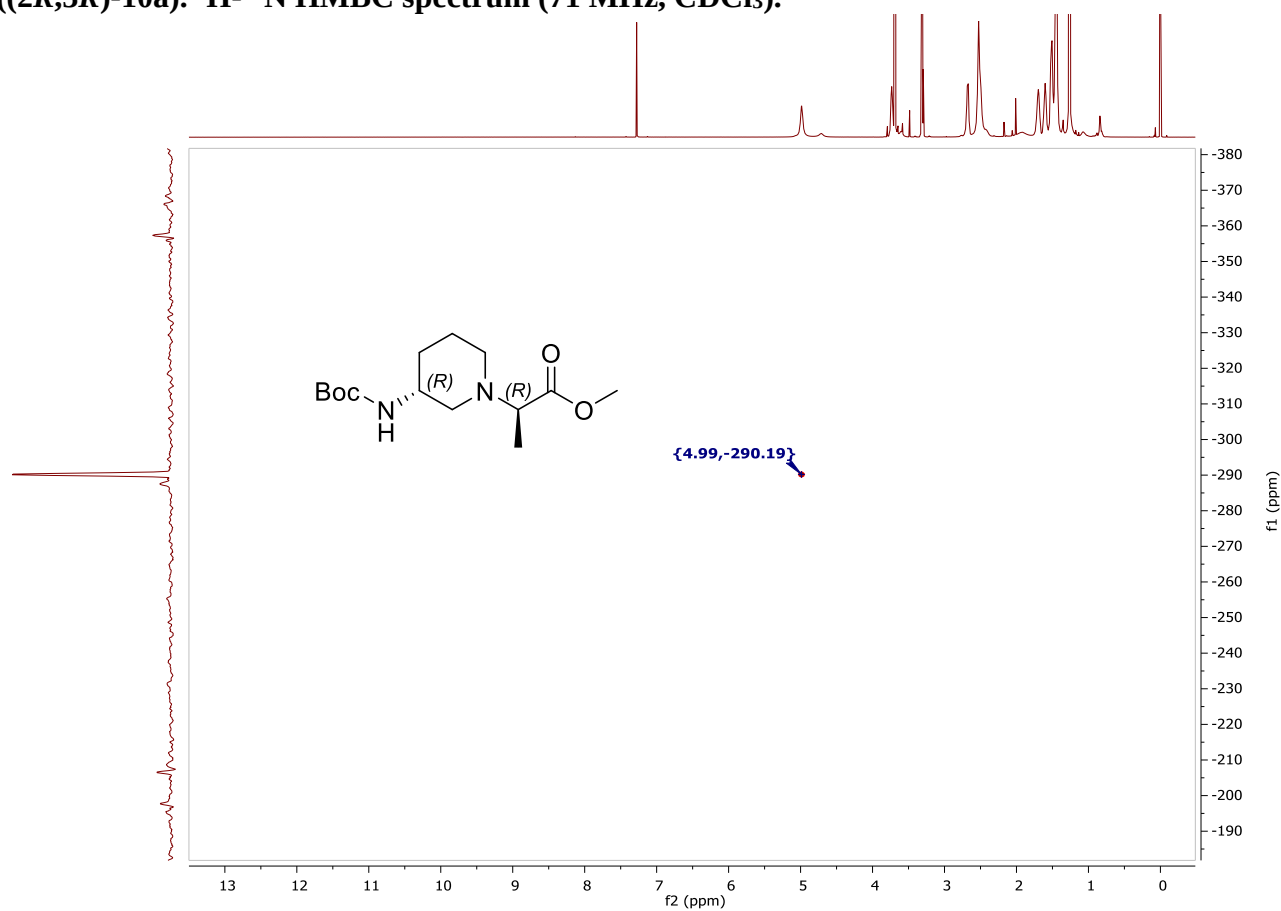


Figure S70. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]propanoate ((2*R*,3*R*)-10a). ^1H - ^{15}N HSQC spectrum (71 MHz, CDCl_3).

Mass Spectrum SmartFormula Report

Analysis Info

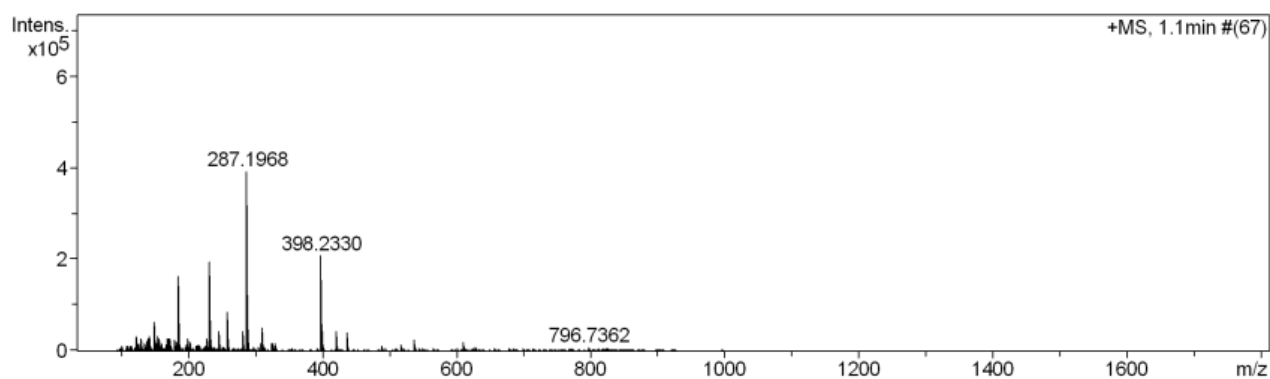
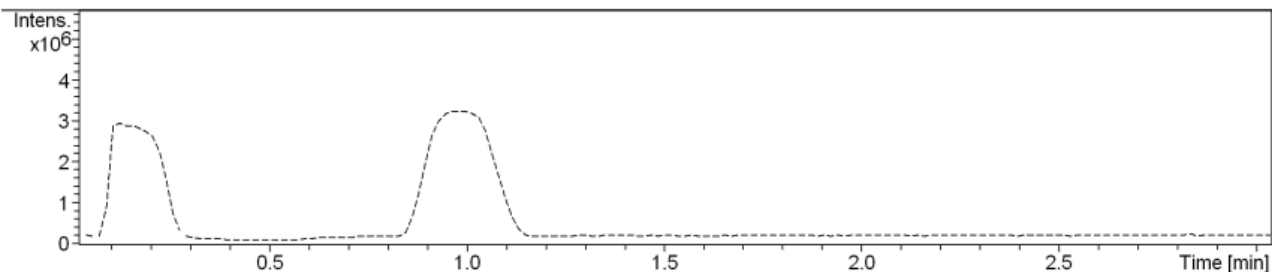
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Acquisition Date 4/21/2023 2:05:33 PM

Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
287.1968	1	C 14 H 27 N 2 O 4	100.00	287.1965	-1.0	-0.9	13.4	2.5	even	ok

Figure S71. Methyl (2R)-2-[(3R)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]propanoate ((2R,3R)-10a). HRMS (ESI-TOF).

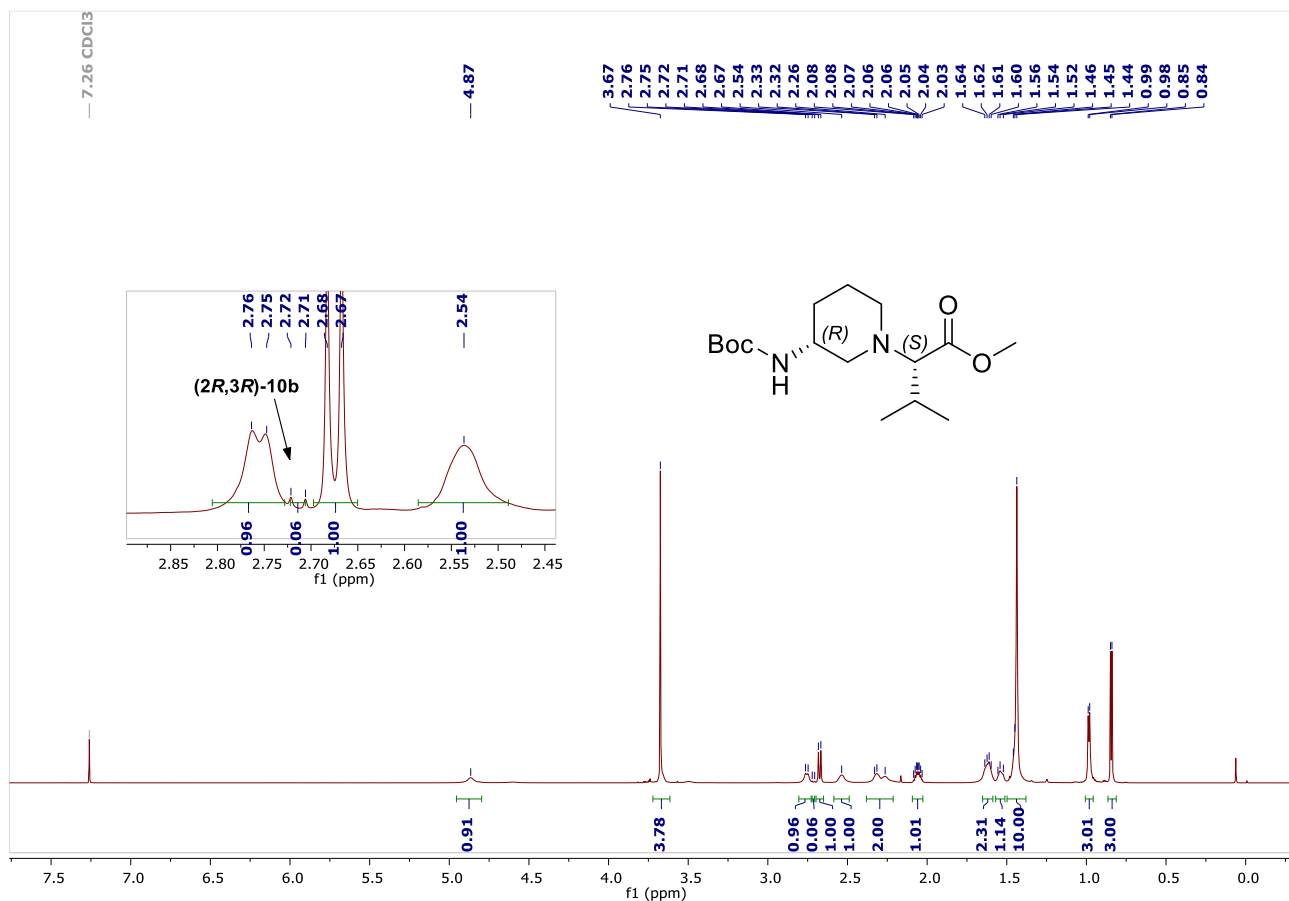


Figure S72. Methyl (2*S*)-2-((3*R*)-3-((*tert*-butoxycarbonyl)amino)piperidin-1-yl)-3-methylbutanoate ((2*S*,3*R*)-10b). ¹H NMR spectrum (700 MHz, CDCl₃).

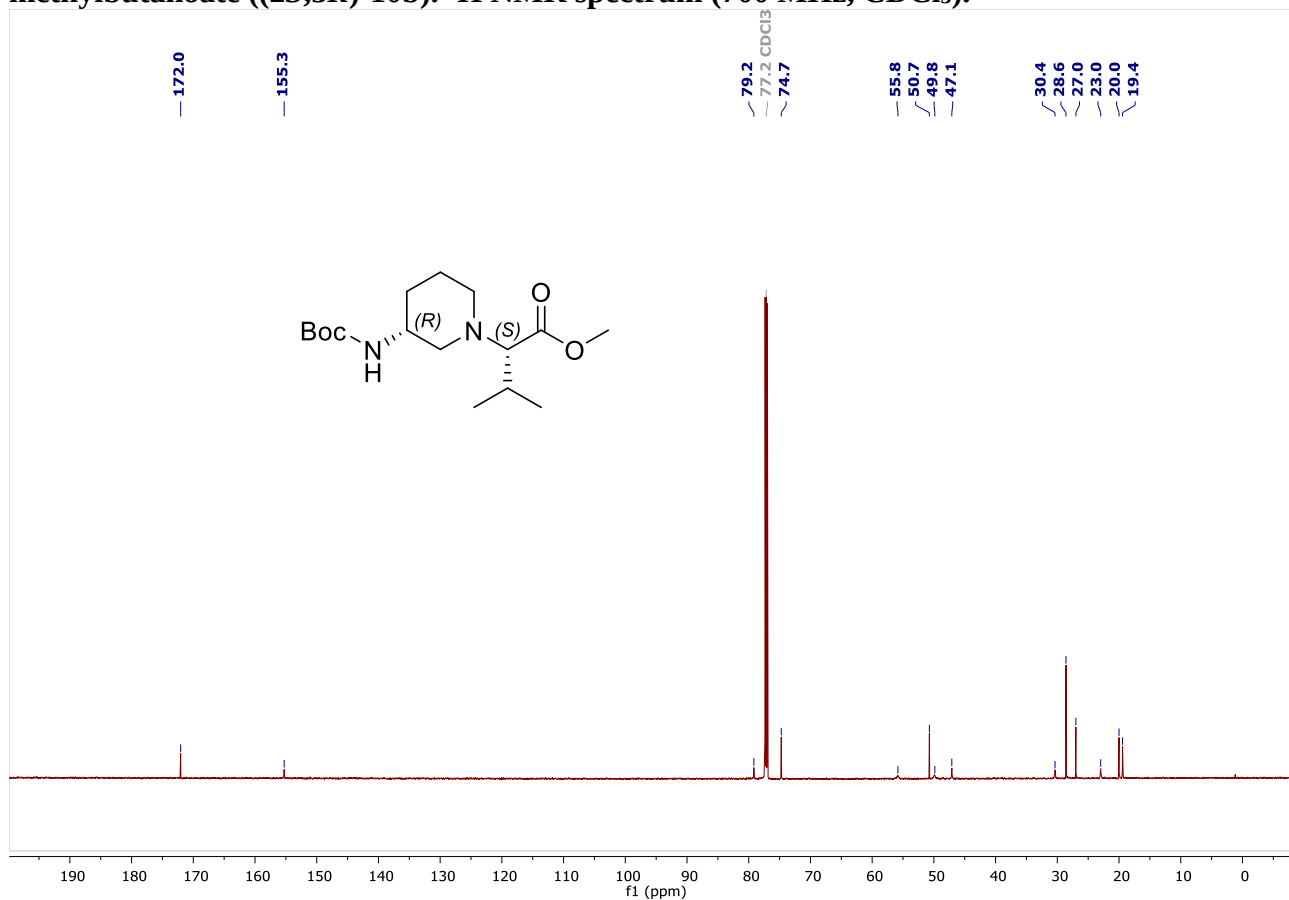


Figure S73. Methyl (2*S*)-2-((3*R*)-3-((*tert*-butoxycarbonyl)amino)piperidin-1-yl)-3-methylbutanoate ((2*S*,3*R*)-10b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

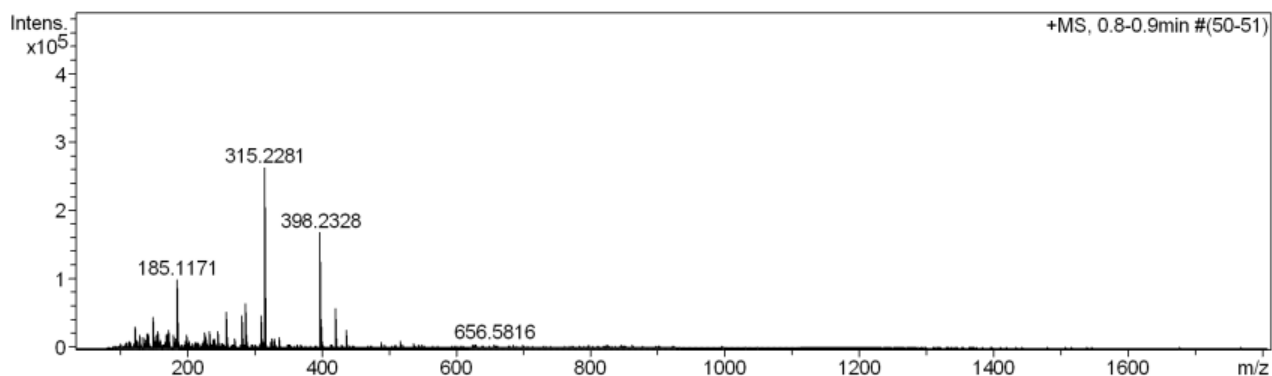
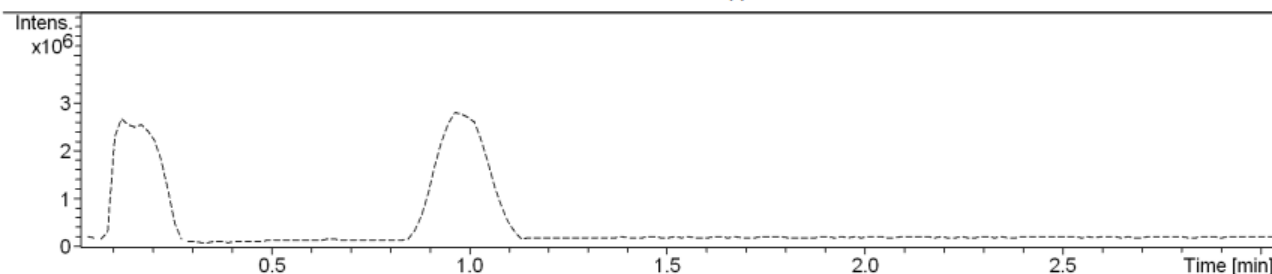
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Acquisition Date 4/14/2023 1:10:31 PM

Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
315.2281	1	C ₁₆ H ₃₁ N ₂ O ₄	100.00	315.2278	-0.9	-0.5	9.9	2.5	even	ok

Figure S74. Methyl (2S)-2-((3R)-3-((*tert*-butoxycarbonyl)amino)piperidin-1-yl)-3-methylbutanoate ((2S,3R)-10b). HRMS (ESI-TOF).

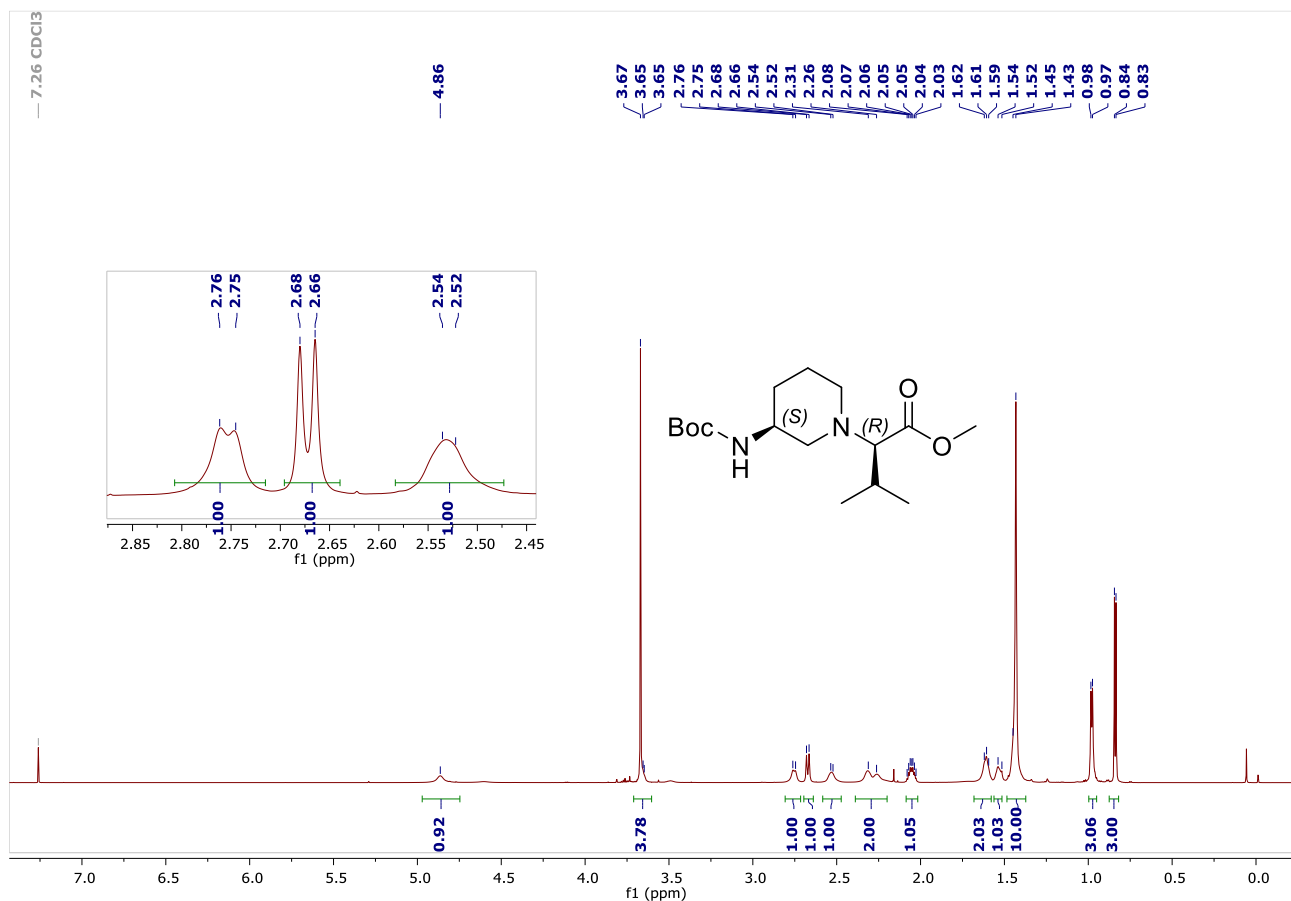


Figure S75. Methyl (2*R*)-2-[(3*S*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-3-methylbutanoate ((2*R*,3*S*)-10b). ¹H NMR spectrum (700 MHz, CDCl₃).

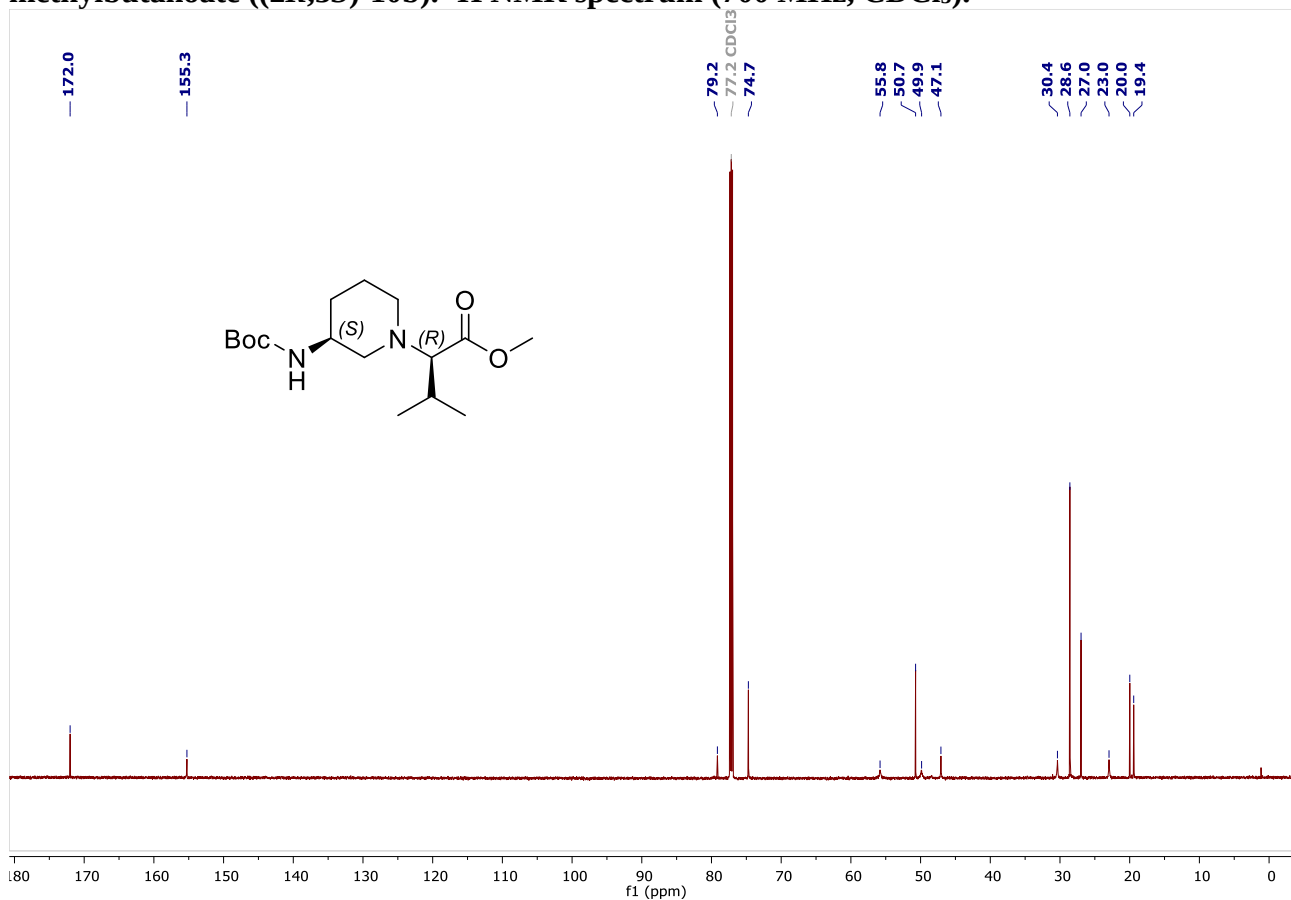


Figure S76. Methyl (2*R*)-2-[(3*S*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-3-methylbutanoate ((2*R*,3*S*)-10b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

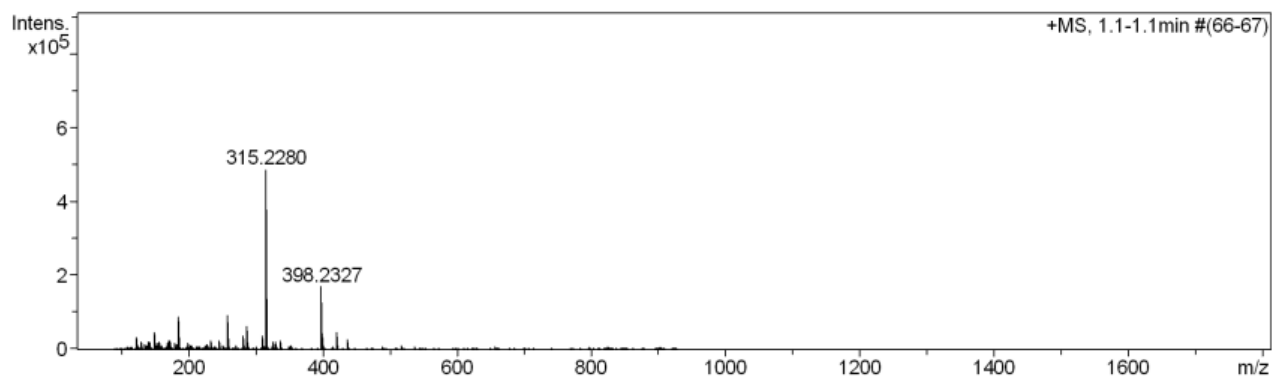
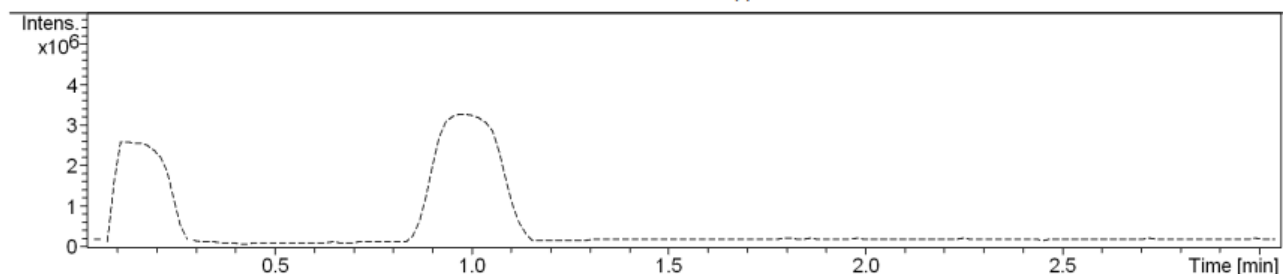
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Comment

Acquisition Date 4/14/2023 12:57:05 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
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Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
315.2280	1	C ₁₆ H ₃₁ N ₂ O ₄	100.00	315.2278	-0.7	-0.4	14.0	2.5	even	ok

Figure S77. Methyl (2R)-2-[(3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-3-methylbutanoate ((2R,3S)-10b). HRMS (ESI-TOF).

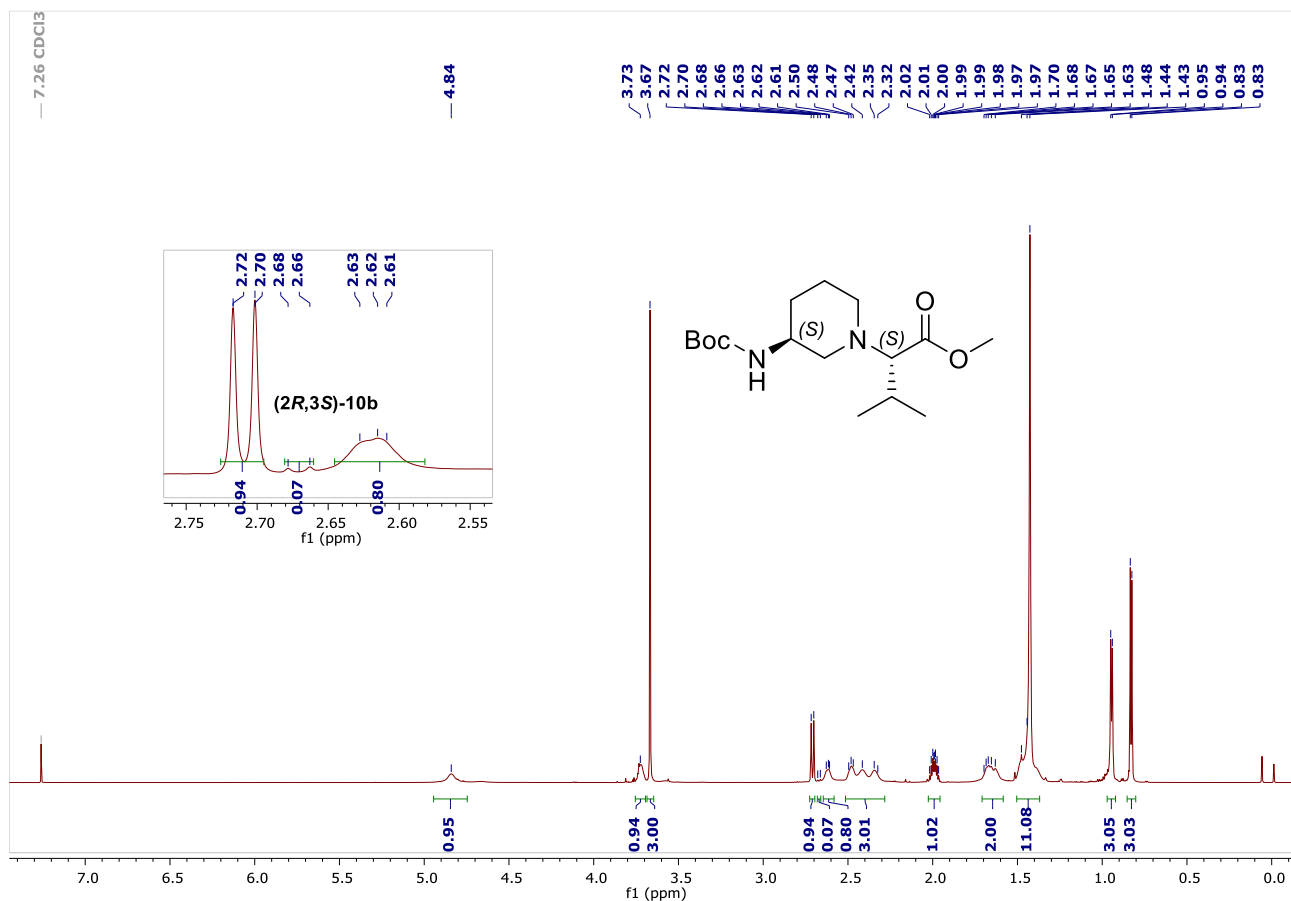


Figure S78. Methyl (2S)-2-[(3S)-3-[(tert-butoxycarbonyl)amino]piperidin-1-yl]-3-methylbutanoate ((2S,3S)-10b). ¹H NMR spectrum (700 MHz, CDCl₃).

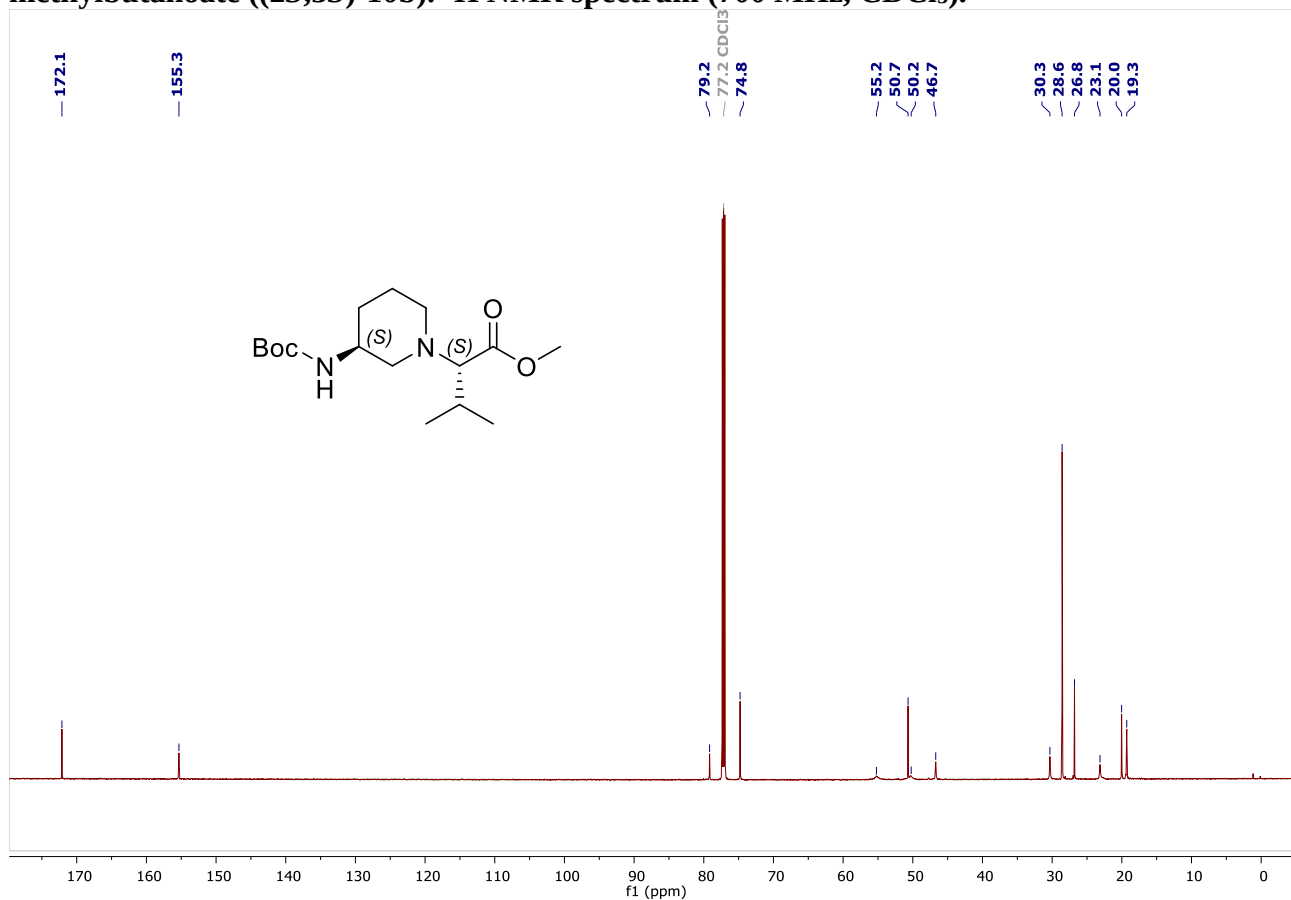


Figure S79. Methyl (2S)-2-[(3S)-3-[(tert-butoxycarbonyl)amino]piperidin-1-yl]-3-methylbutanoate ((2S,3S)-10b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

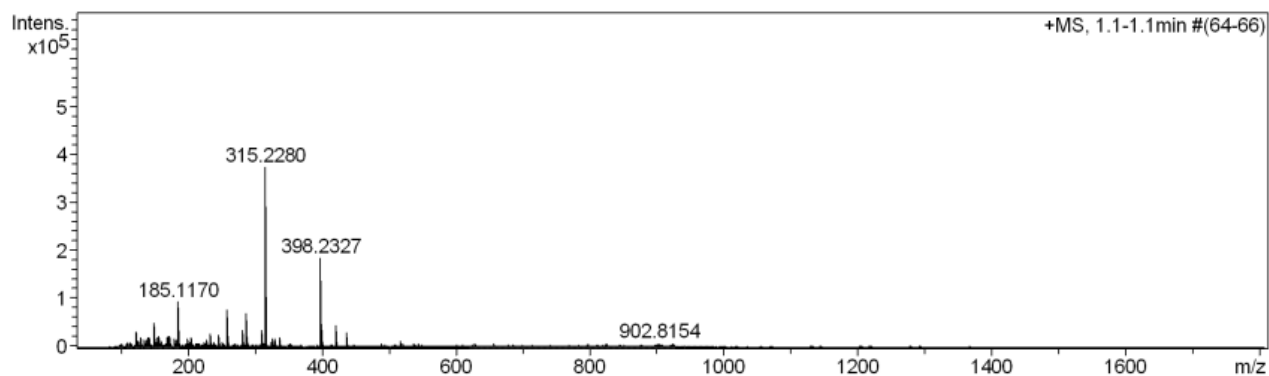
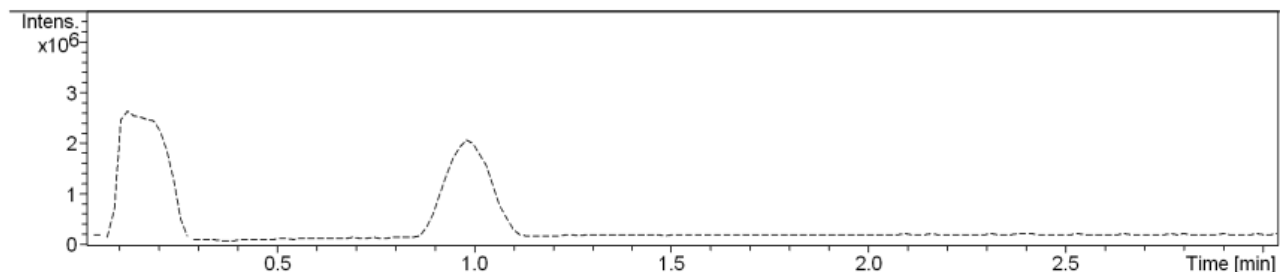
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Sample Name GMP_567
Comment

Acquisition Date 4/14/2023 1:01:35 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
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Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
315.2280	1	C ₁₆ H ₃₁ N ₂ O ₄	100.00	315.2278	-0.6	-0.3	19.2	2.5	even	ok

Figure S80. Methyl (2S)-2-[(3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-3-methylbutanoate ((2S,3S)-10b). HRMS (ESI-TOF).

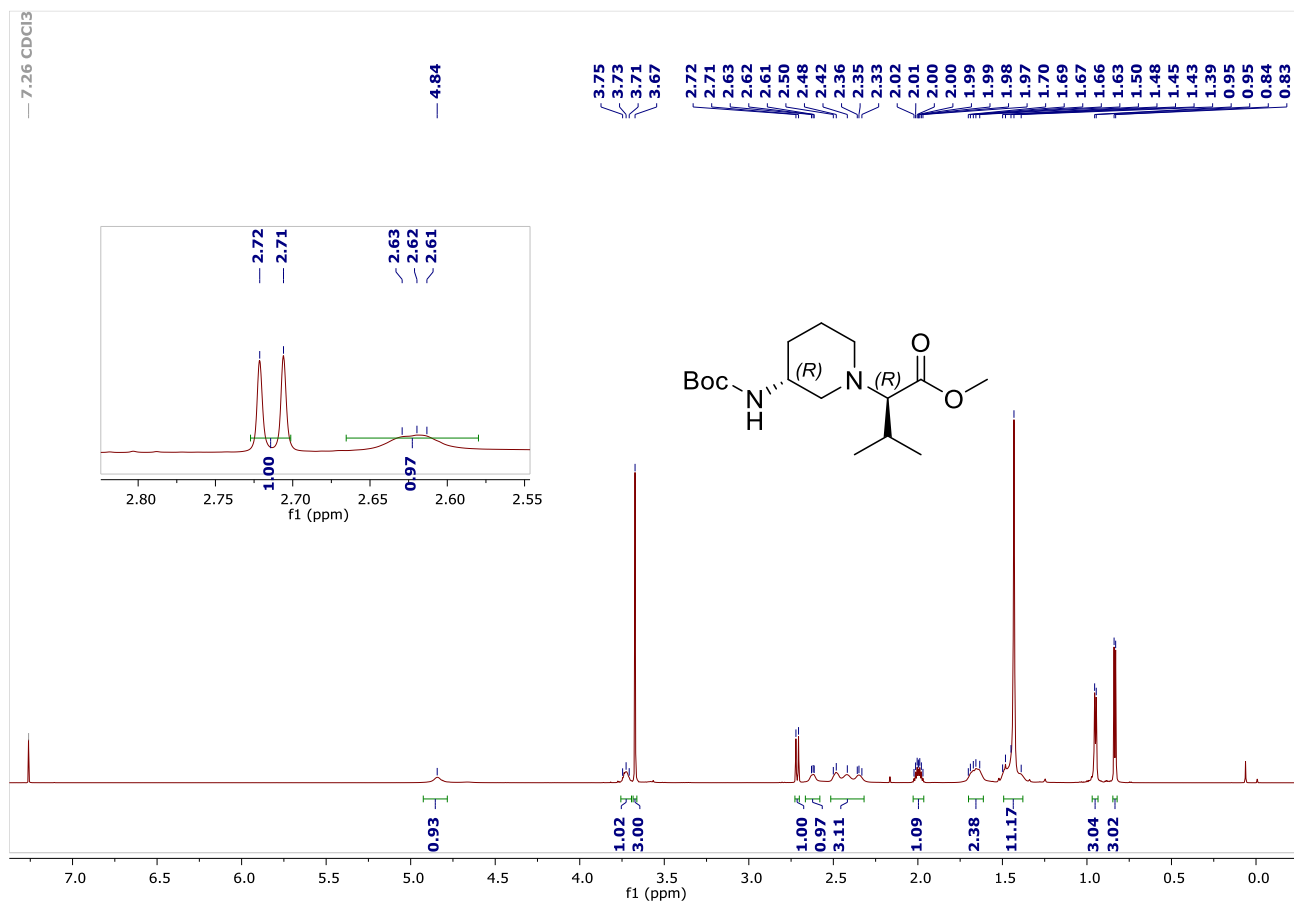


Figure S81. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-3-methylbutanoate ((2*R*,3*R*)-10b). ¹H NMR spectrum (700 MHz, CDCl₃).

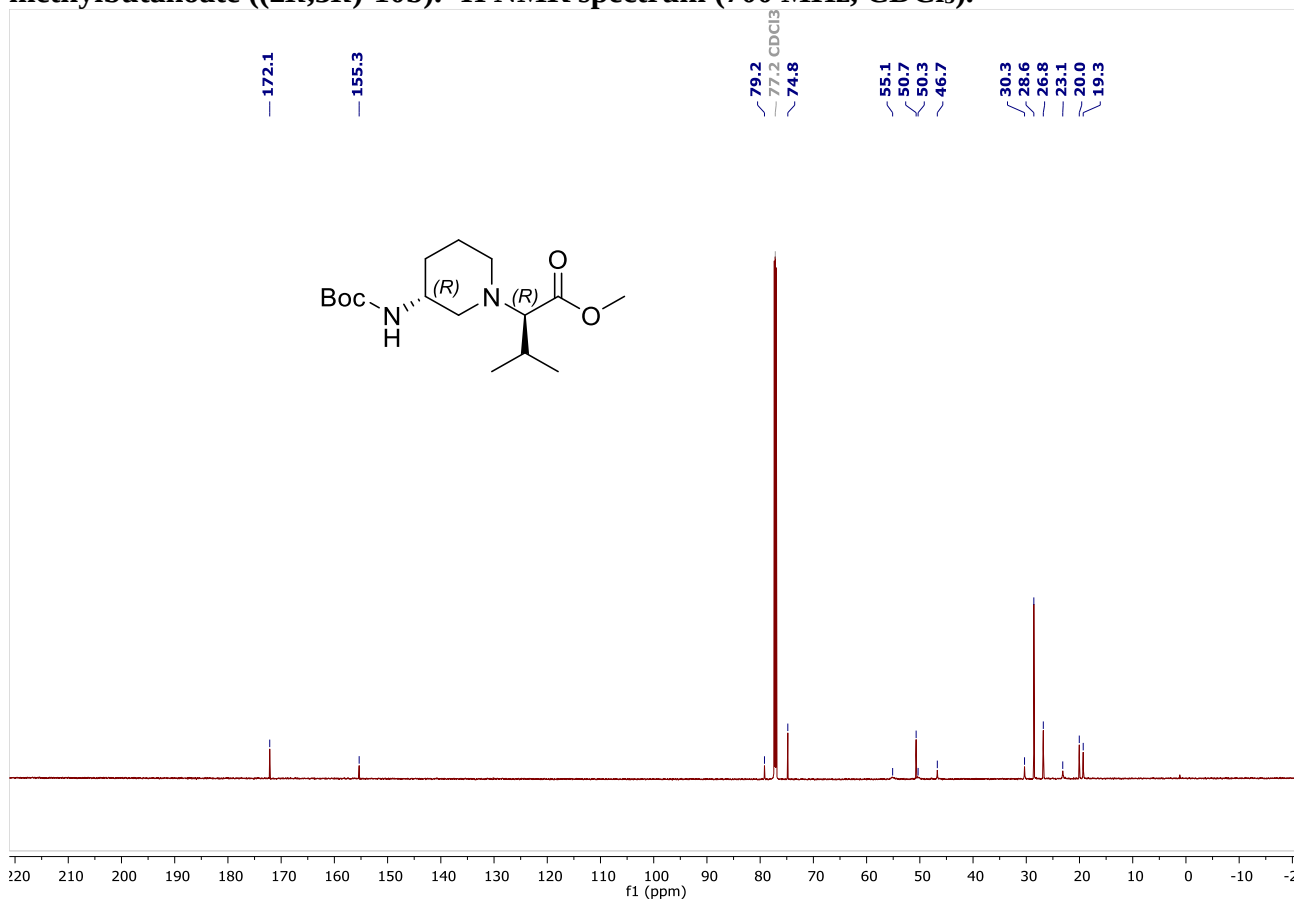


Figure S82. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-3-methylbutanoate ((2*R*,3*R*)-10b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

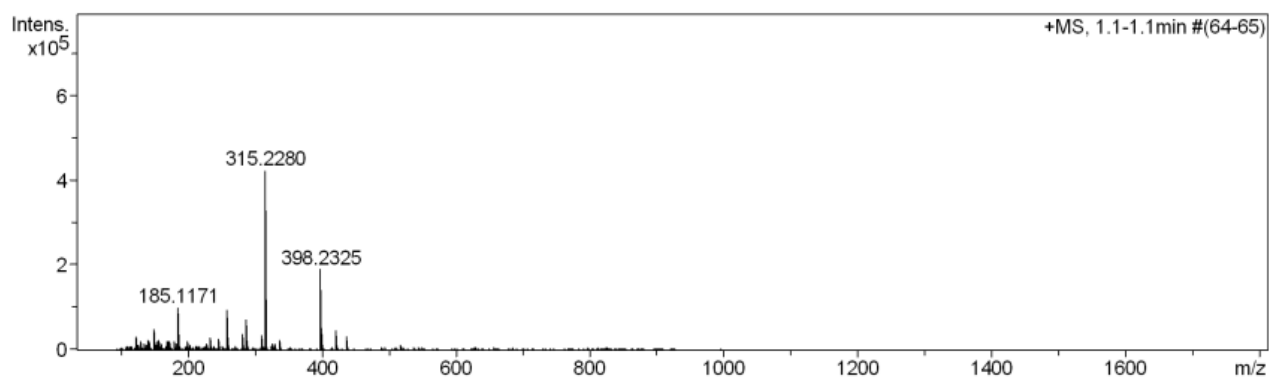
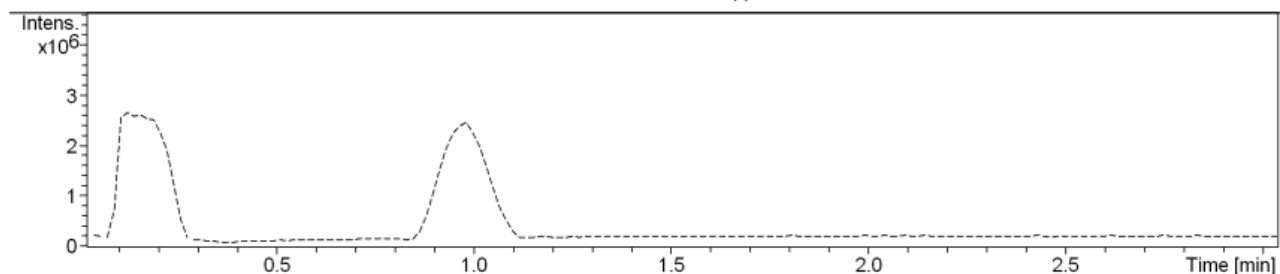
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Sample Name GMP_570
Comment

Acquisition Date 4/14/2023 1:06:04 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
315.2280	1	C ₁₆ H ₃₁ N ₂ O ₄	100.00	315.2278	-0.6	-0.3	11.3	2.5	even	ok

Figure S83. Methyl (2R)-2-[(3R)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-3-methylbutanoate ((2R,3R)-10b). HRMS (ESI-TOF).

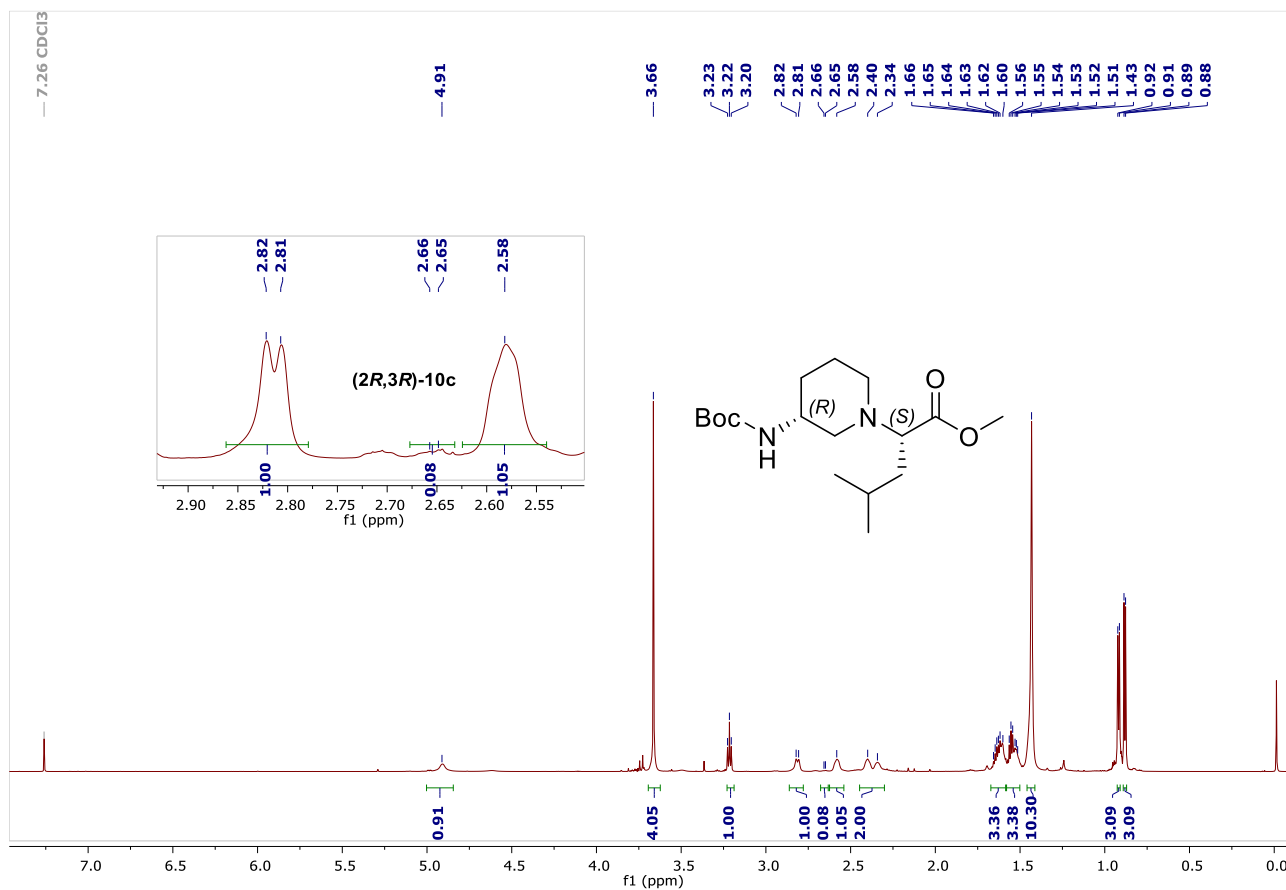


Figure S84. Methyl (2S)-2-((3R)-3-((tert-butoxycarbonyl)amino)piperidin-1-yl)-4-methylpentanoate ((2S,3R)-10c). ¹H NMR spectrum (700 MHz, CDCl₃).

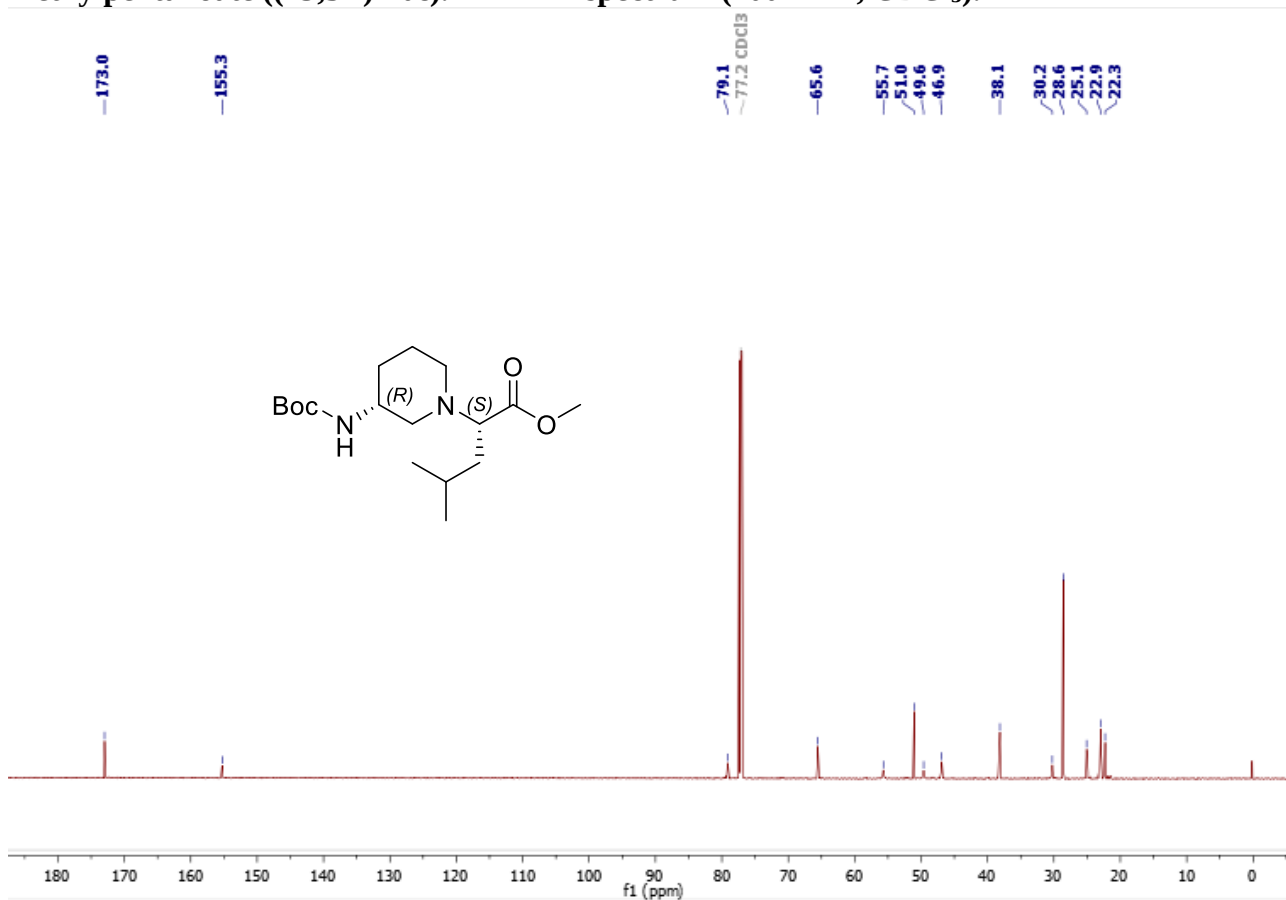


Figure S85. Methyl (2S)-2-((3R)-3-((tert-butoxycarbonyl)amino)piperidin-1-yl)-4-methylpentanoate ((2S,3R)-10c). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

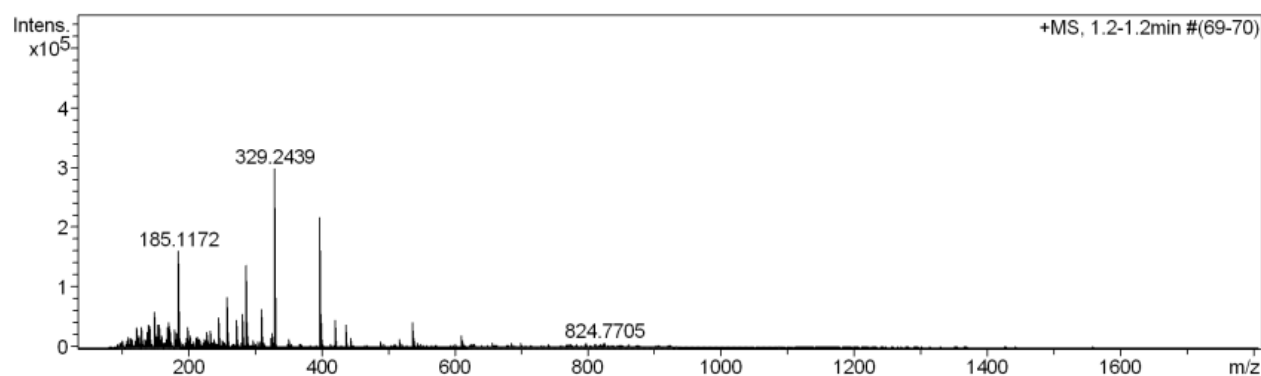
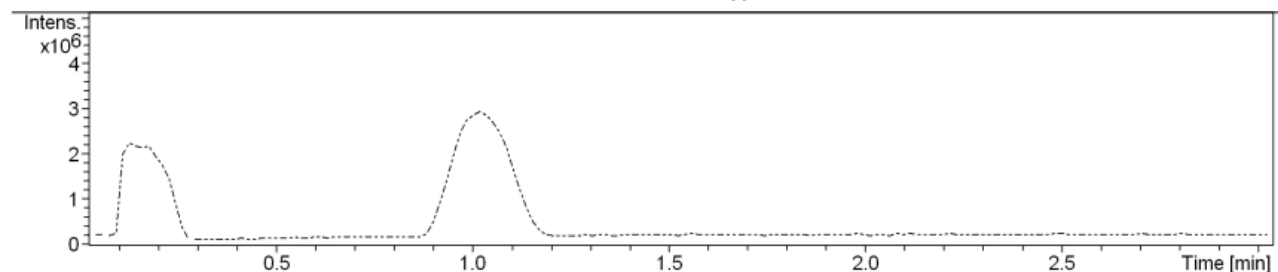
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Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

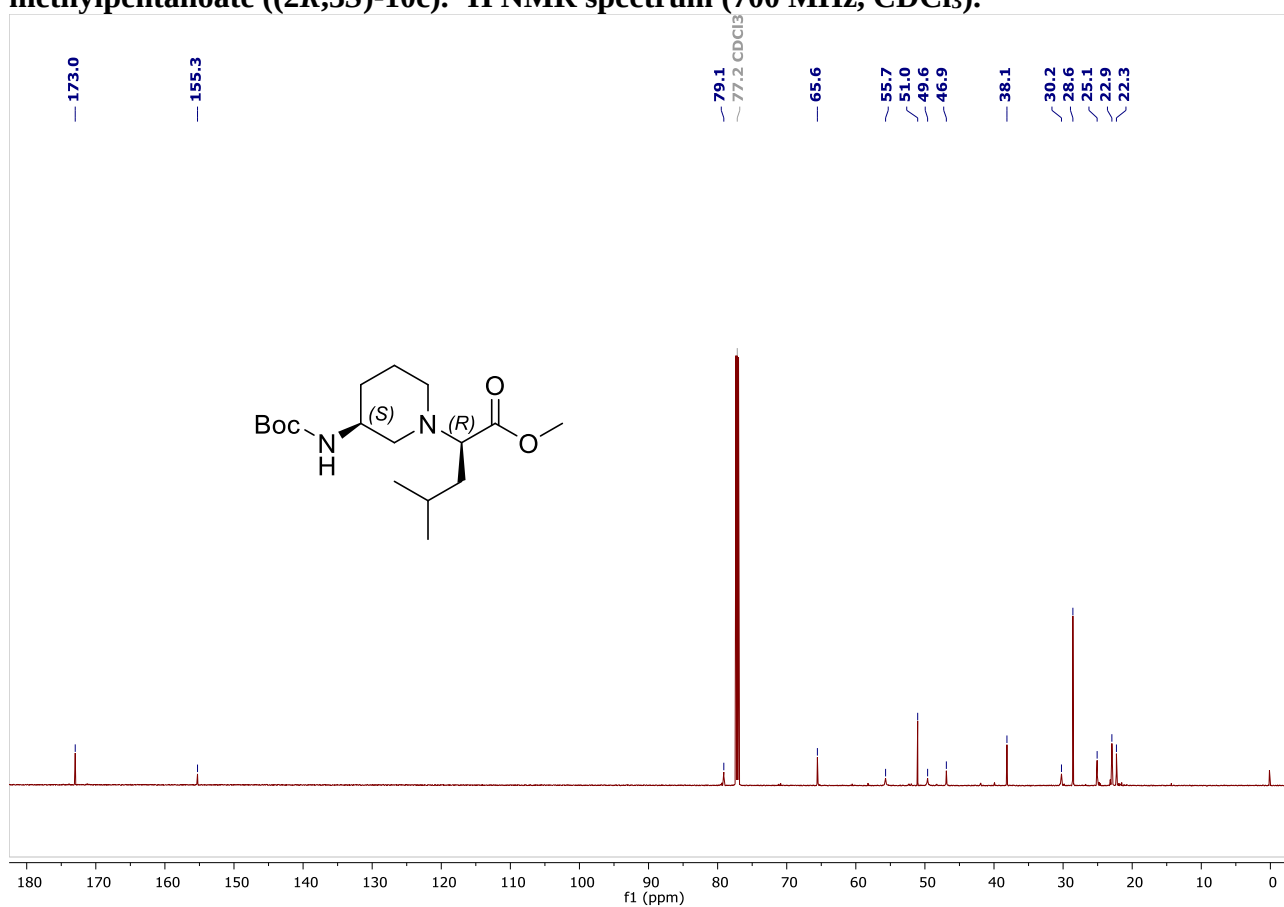
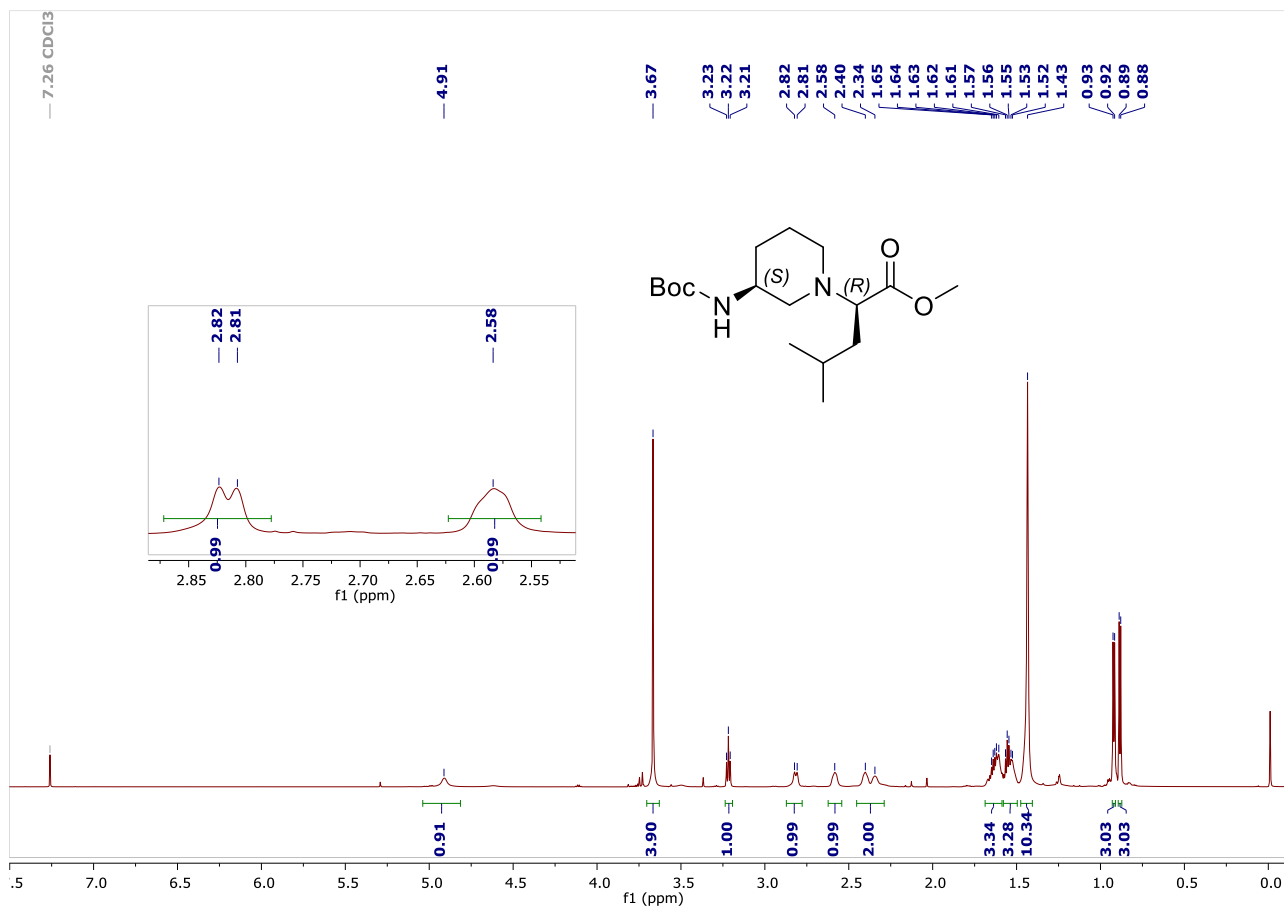
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Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
329.2439	1	C ₁₇ H ₃₃ N ₂ O ₄	100.00	329.2435	-1.2	-1.0	15.2	2.5	even	ok

Figure S86. Methyl (2S)-2-[(3R)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-4-methylpentanoate ((2S,3R)-10c). HRMS (ESI-TOF).



Mass Spectrum SmartFormula Report

Analysis Info

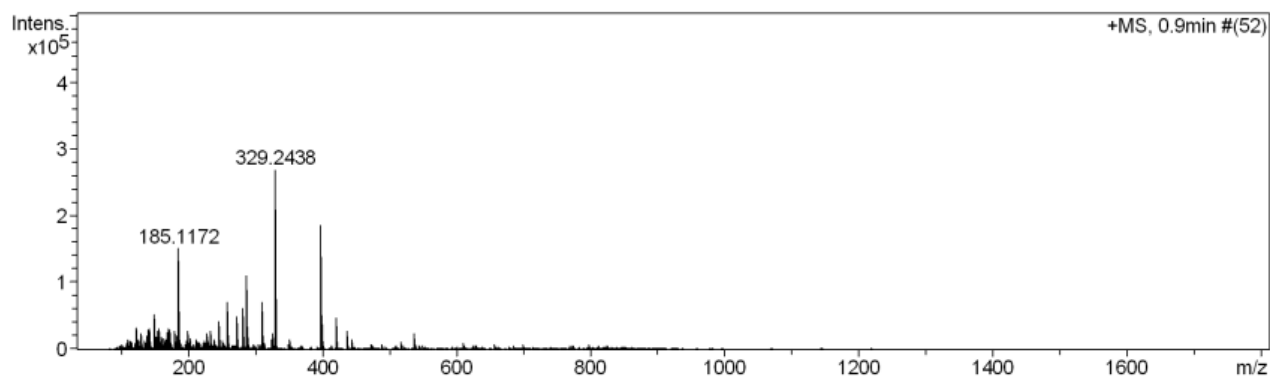
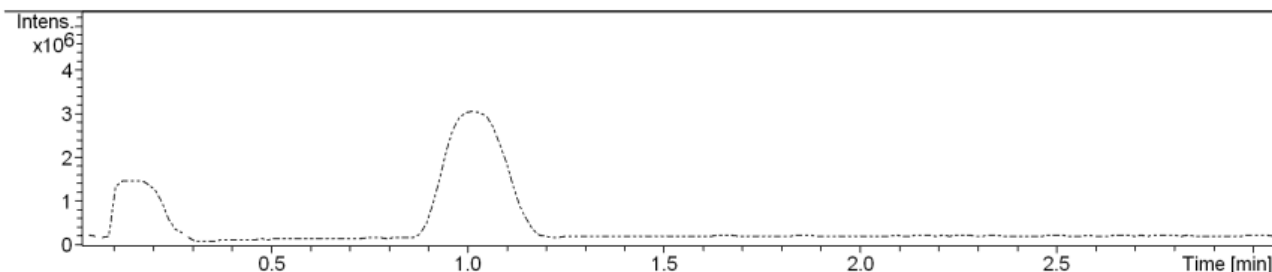
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Comment

Acquisition Date 4/27/2023 3:44:18 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
329.2438	1	C 17 H 33 N 2 O 4	100.00	329.2435	-1.1	-0.9	15.2	2.5	even	ok

Figure S89. Methyl (2R)-2-[(3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-4-methylpentanoate ((2R,3S)-10c). HRMS (ESI-TOF).

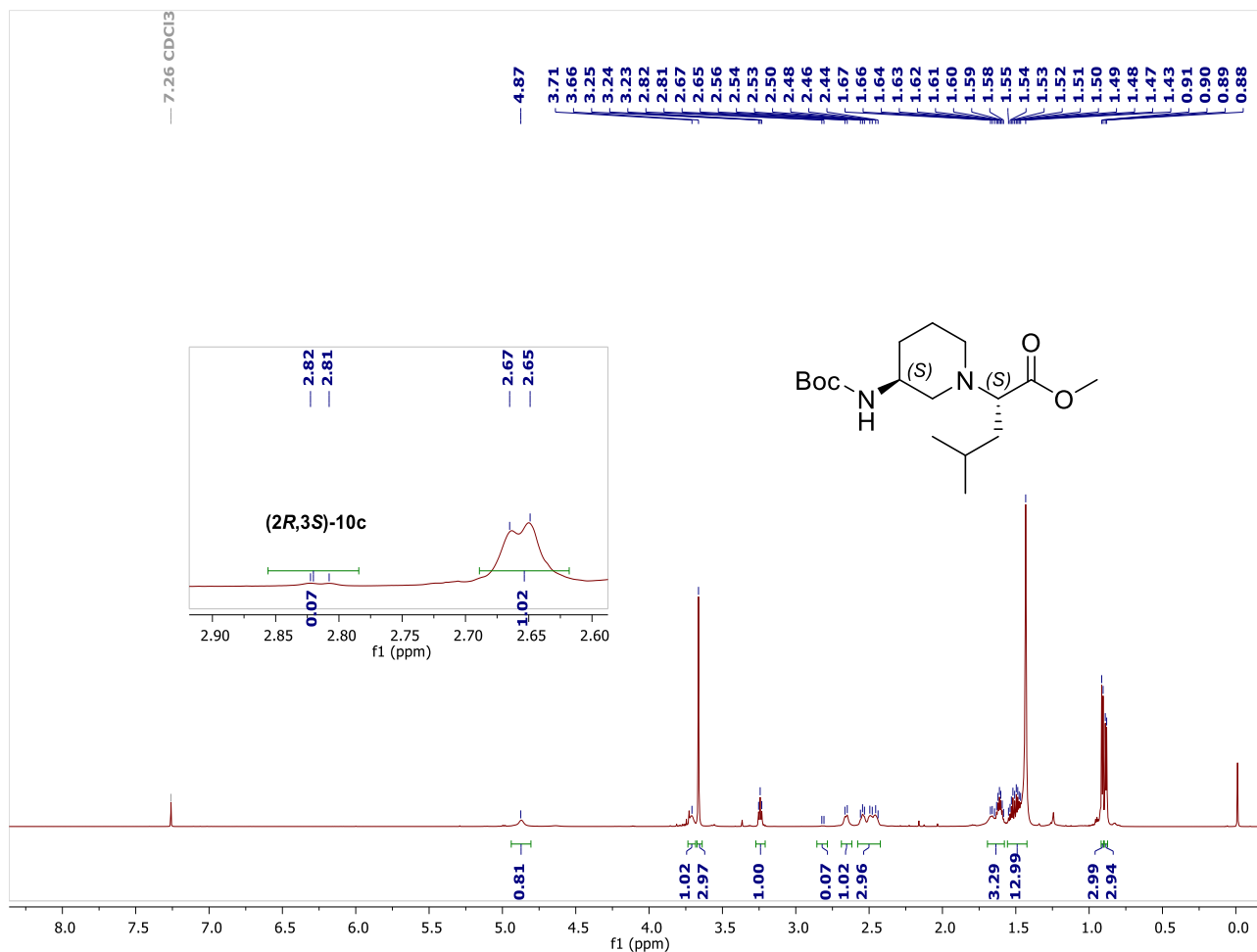


Figure S90. Methyl (2S)-2-[(3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-4-methylpentanoate ((2S,3S)-10c). ¹H NMR spectrum (700 MHz, CDCl₃).

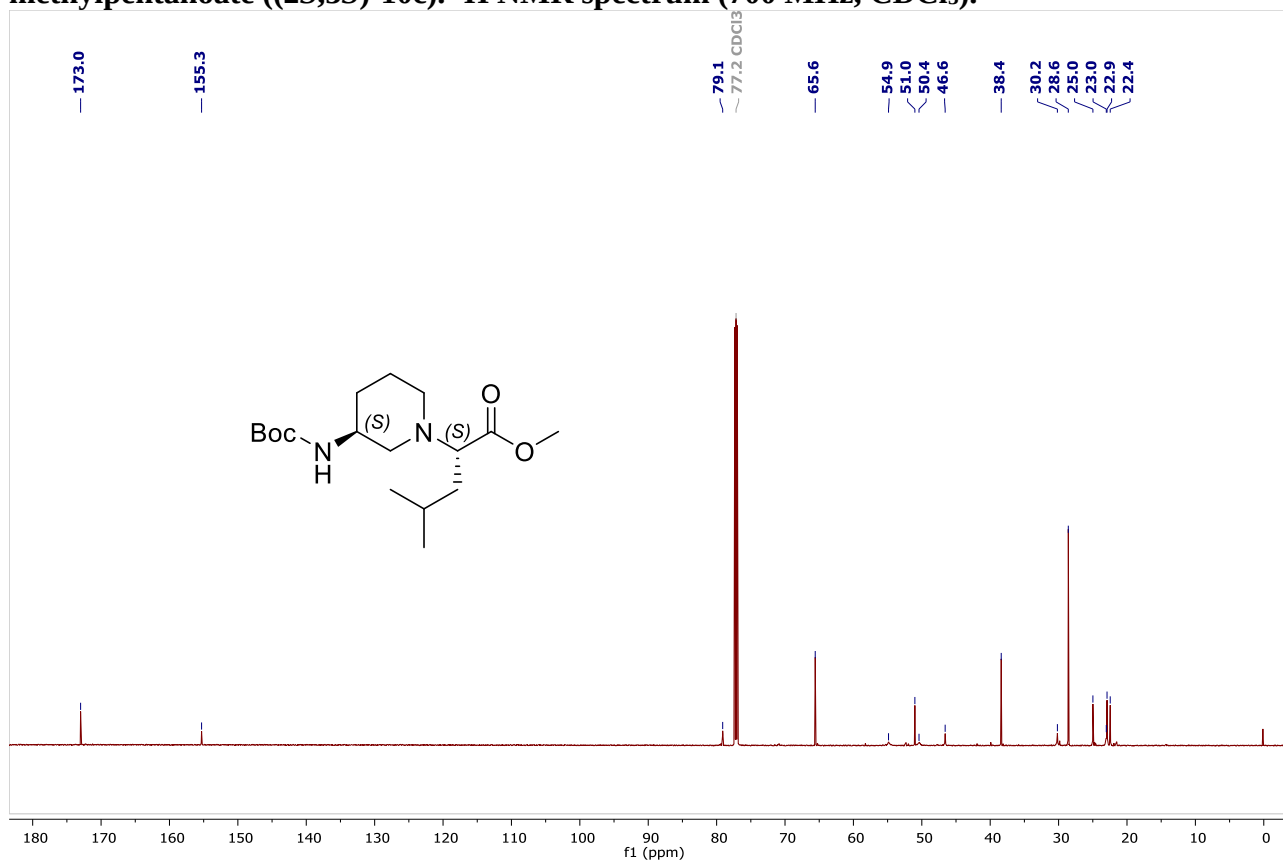


Figure S91. Methyl (2S)-2-[(3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-4-methylpentanoate ((2S,3S)-10c). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

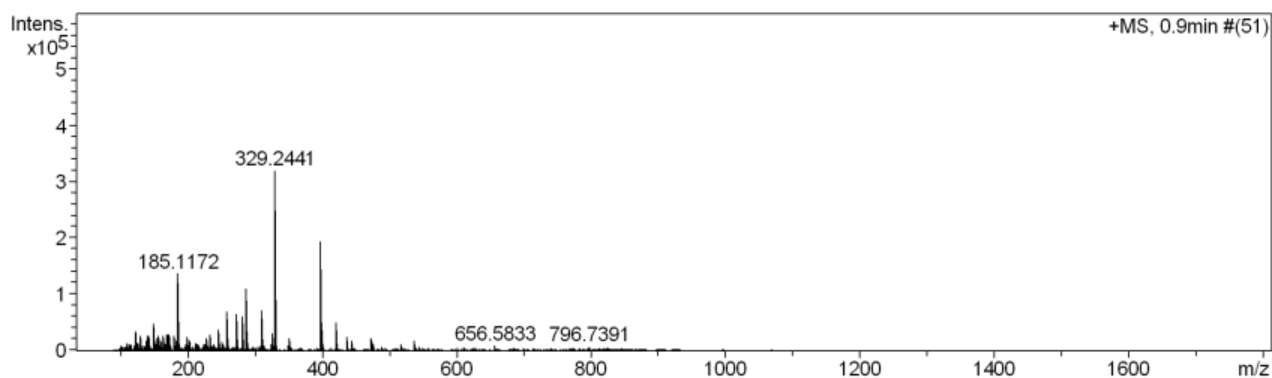
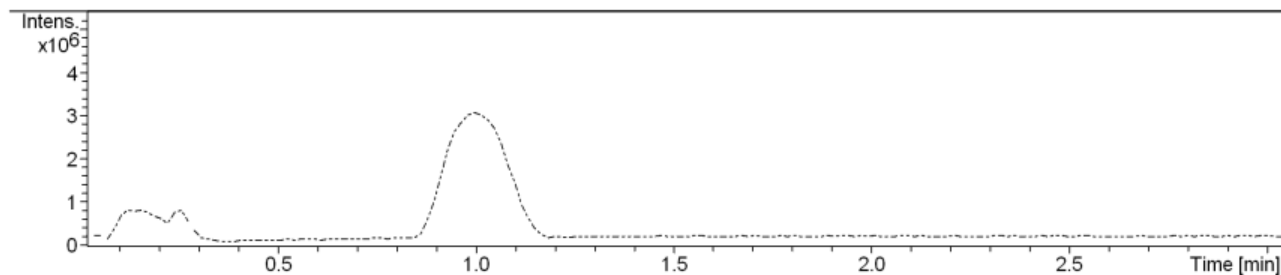
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Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
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Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
329.2441	1	C 17 H 33 N 2 O 4	100.00	329.2435	-1.8	-1.6	16.5	2.5	even	ok

Figure S92. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl)-4-methylpentanoate ((2S,3S)-10c). HRMS (ESI-TOF).

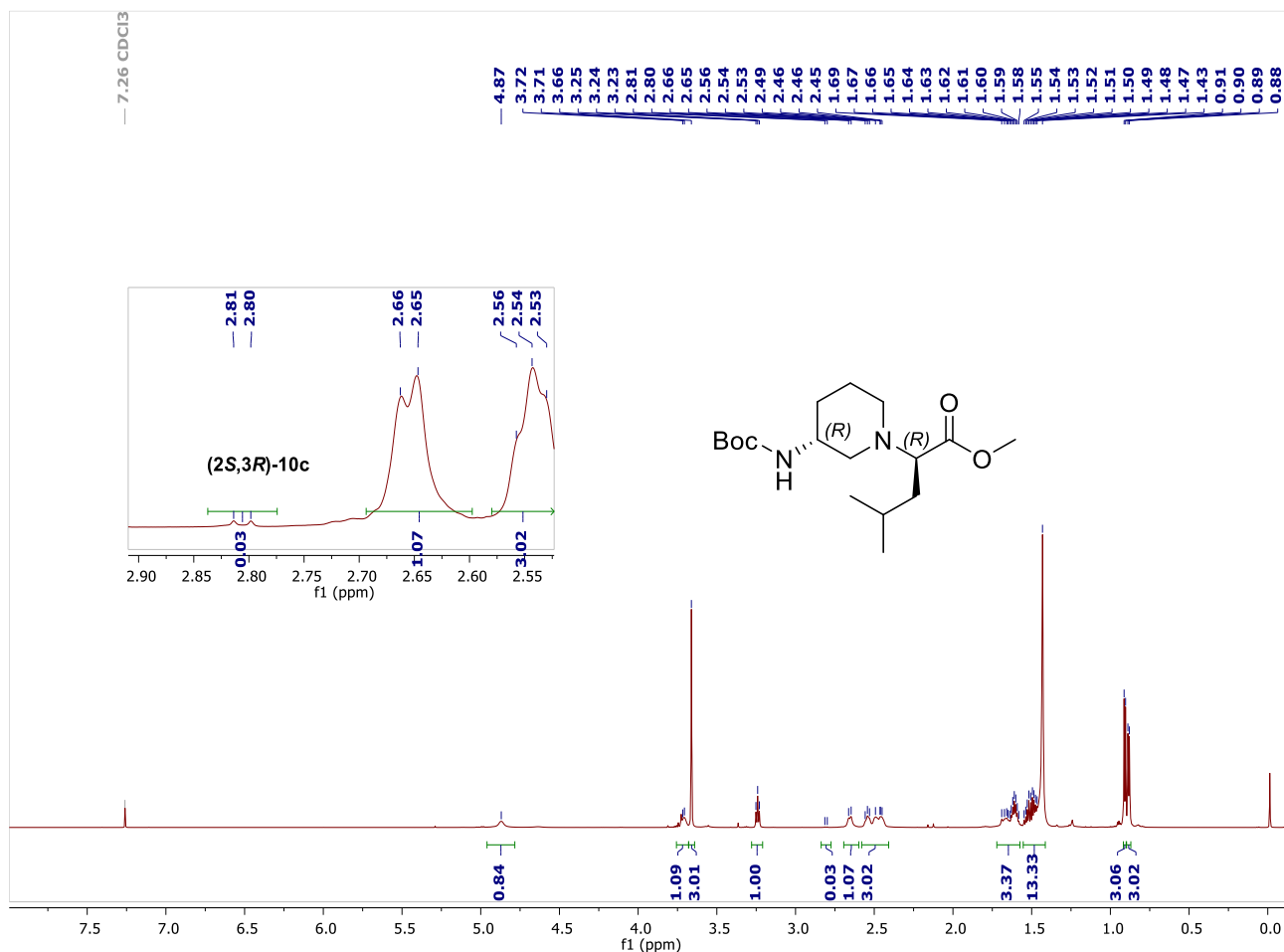


Figure S93. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-4-methylpentanoate ((2*R*,3*R*)-10c). ¹H NMR spectrum (700 MHz, CDCl₃).

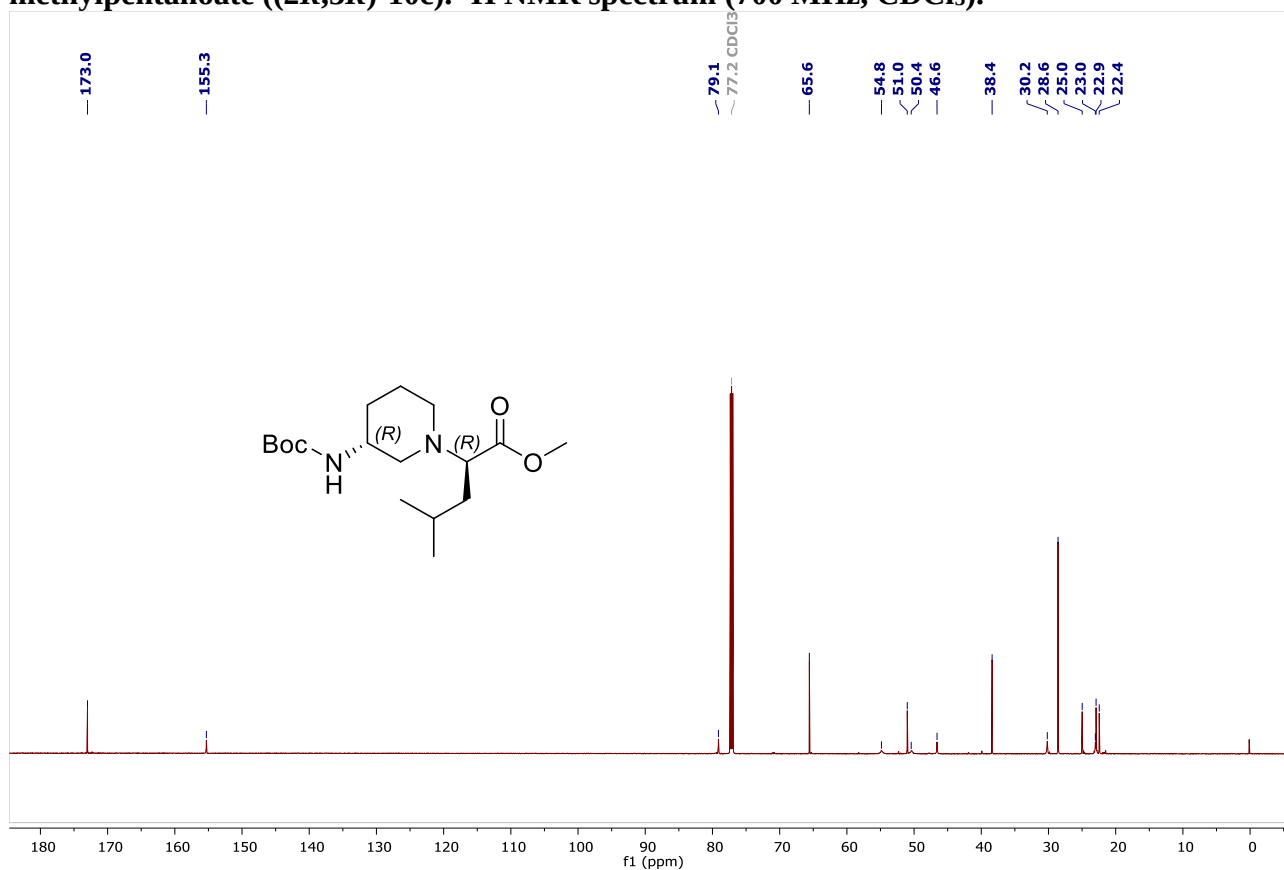


Figure S94. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-4-methylpentanoate ((2*R*,3*R*)-10c). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

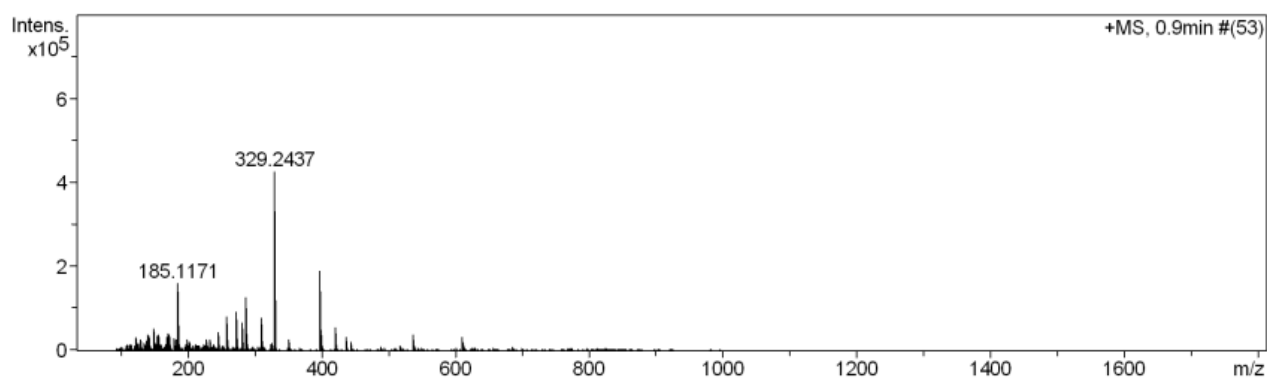
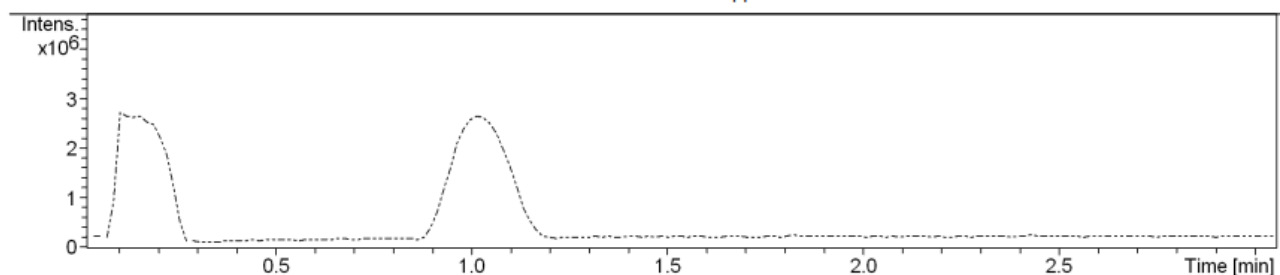
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Method organikai_esi_pos_2013_recover.m
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Comment

Acquisition Date 4/27/2023 3:53:15 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

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Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
329.2437	1	C 17 H 33 N 2 O 4	100.00	329.2435	-0.8	-0.6	15.6	2.5	even	ok

Figure S95. Methyl (2R)-2-[(3R)-3-[(*tert*-butoxycarbonyl)amino]piperidin-1-yl]-4-methylpentanoate ((2R,3R)-10c). HRMS (ESI-TOF).

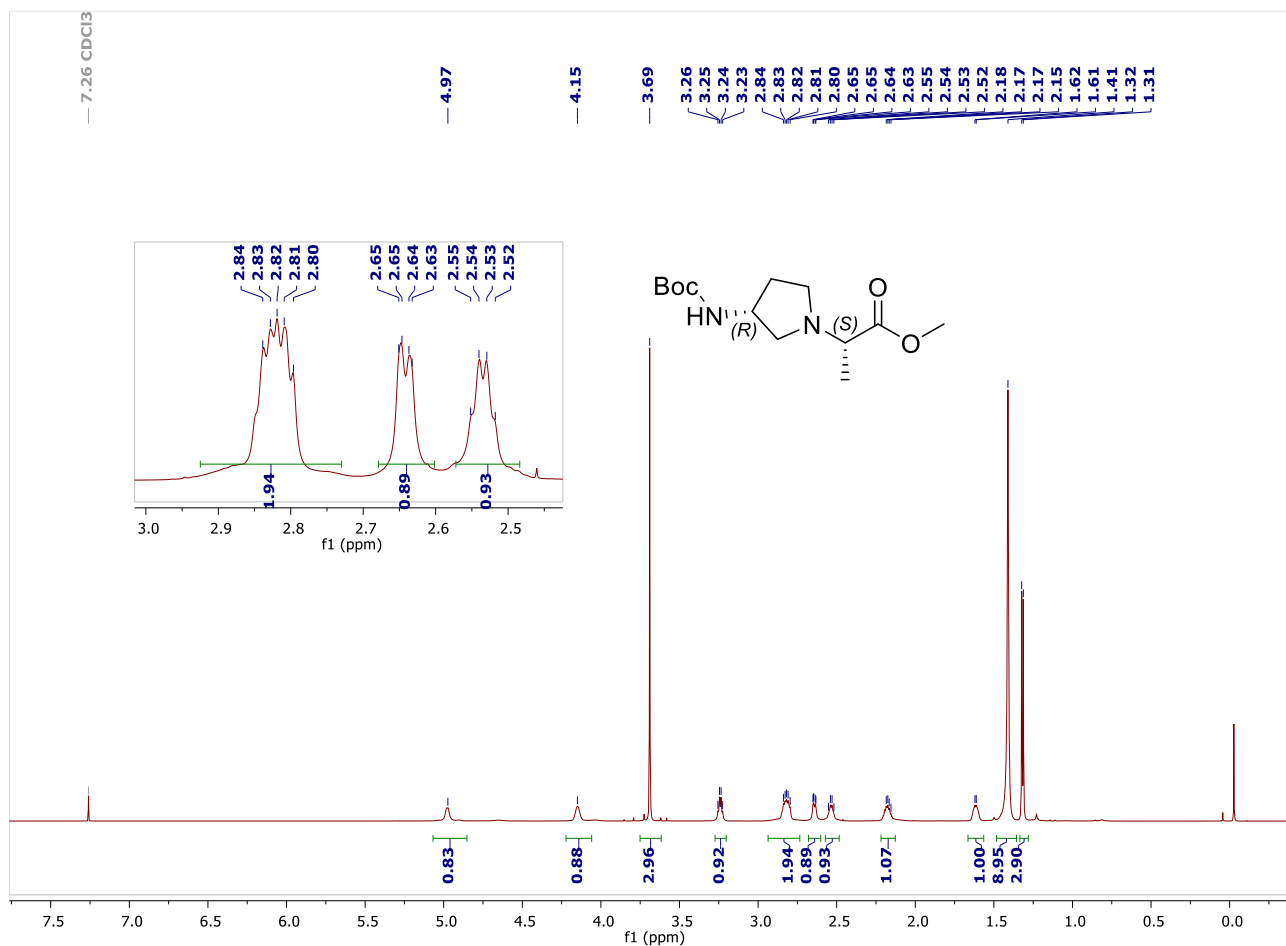


Figure S96. Methyl (2S)-2-((3R)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)propanoate ((2S,3R)-11a). ¹H NMR spectrum (700 MHz, CDCl₃).

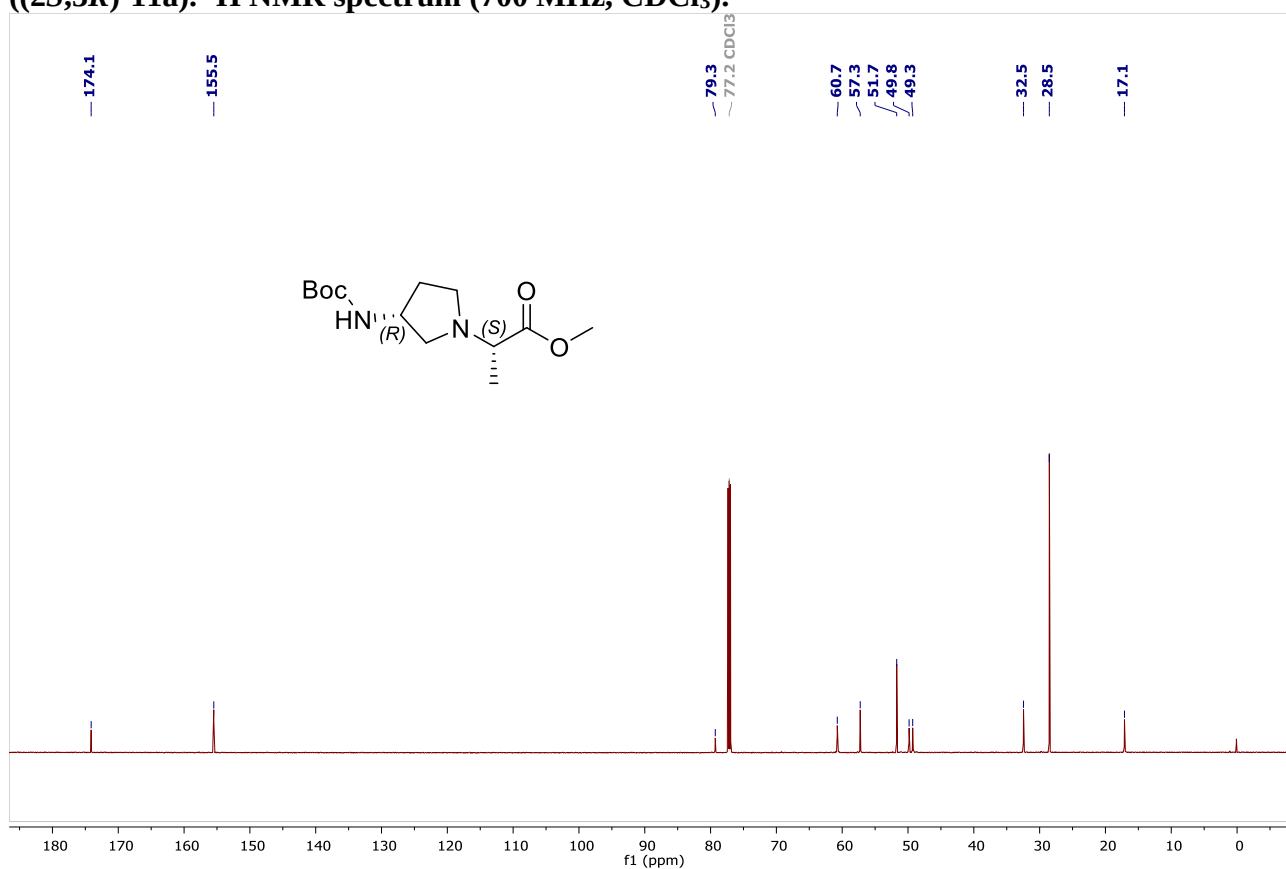
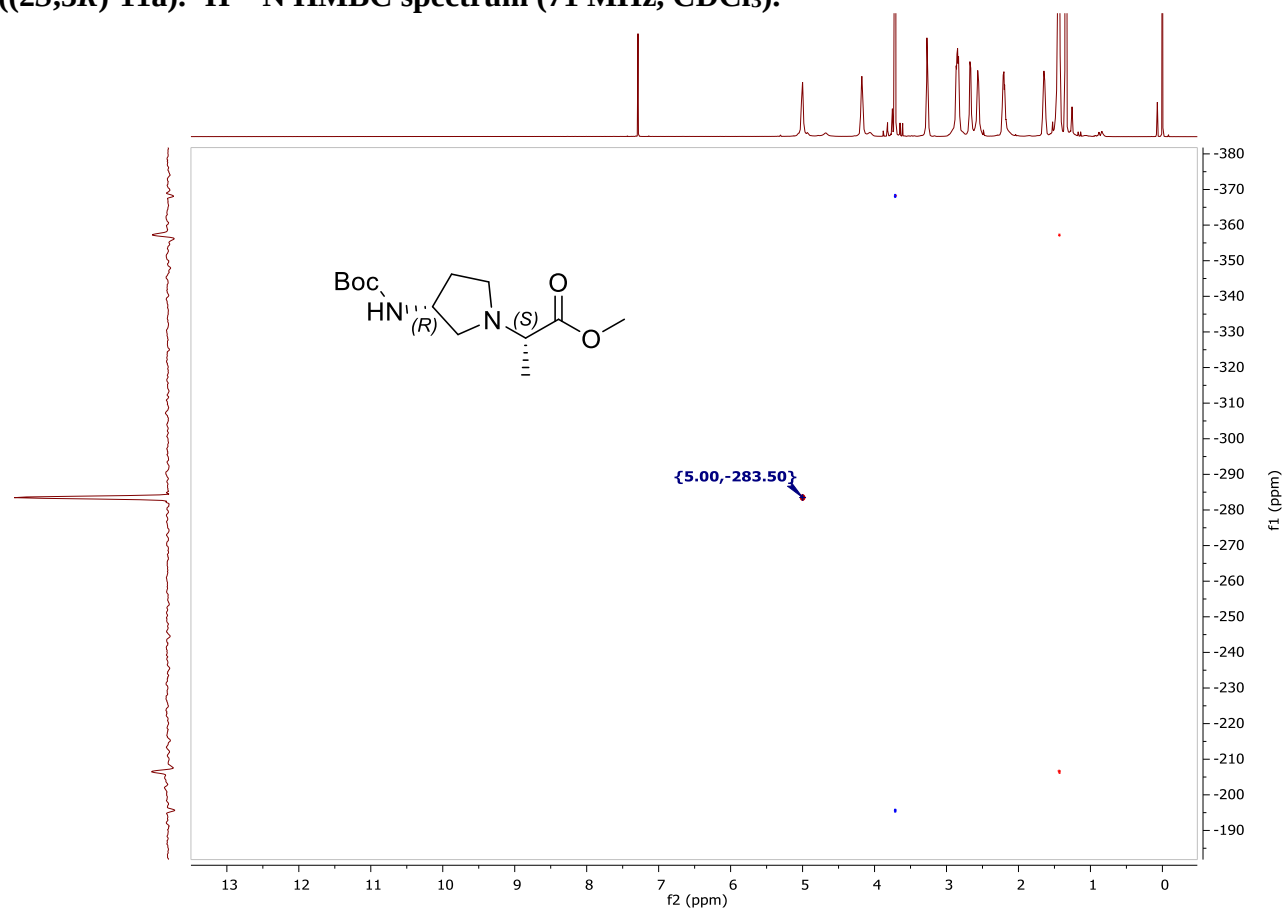
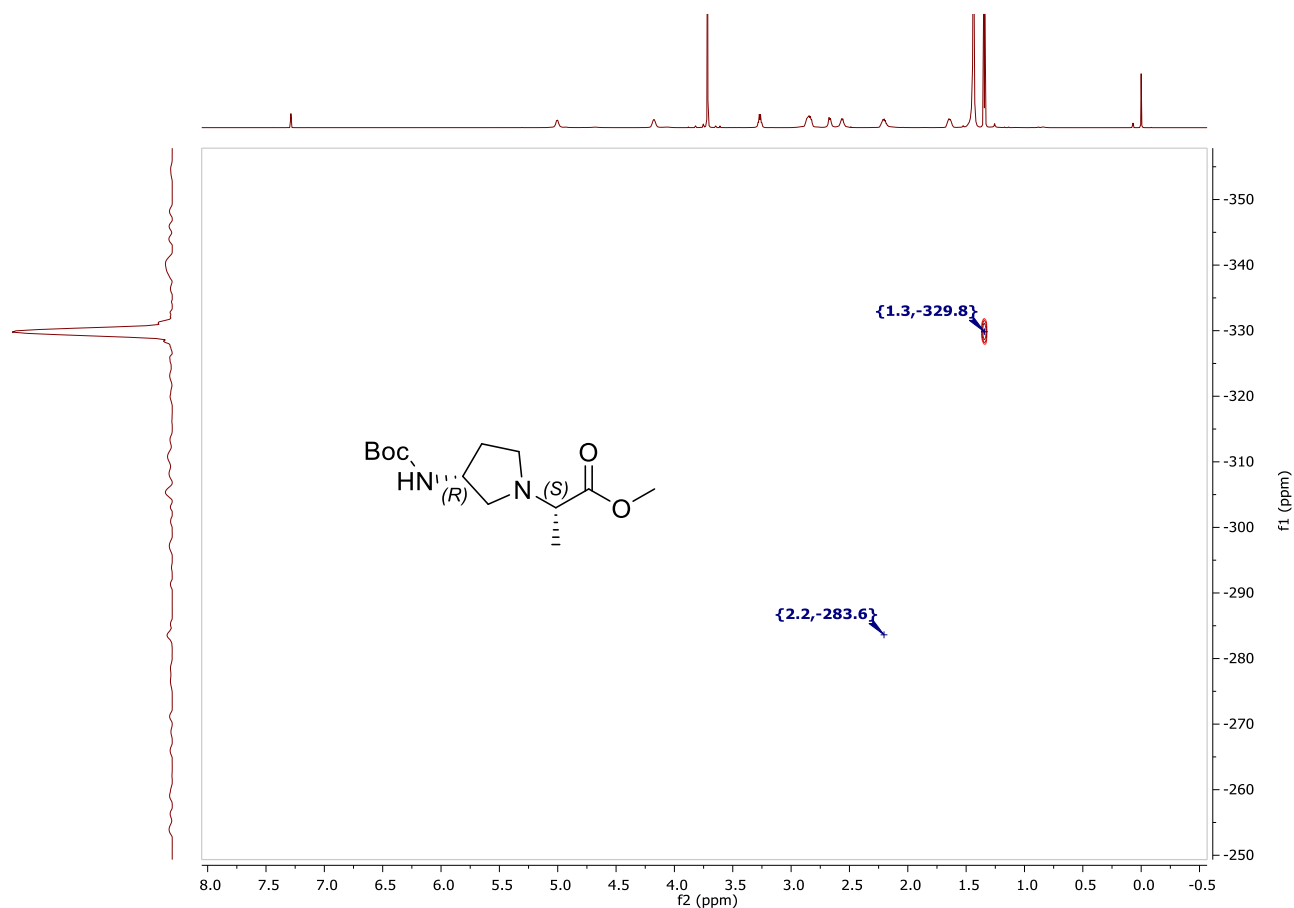


Figure S97. Methyl (2S)-2-((3R)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)propanoate ((2S,3R)-11a). ¹³C NMR spectrum (176 MHz, CDCl₃).



Mass Spectrum SmartFormula Report

Analysis Info

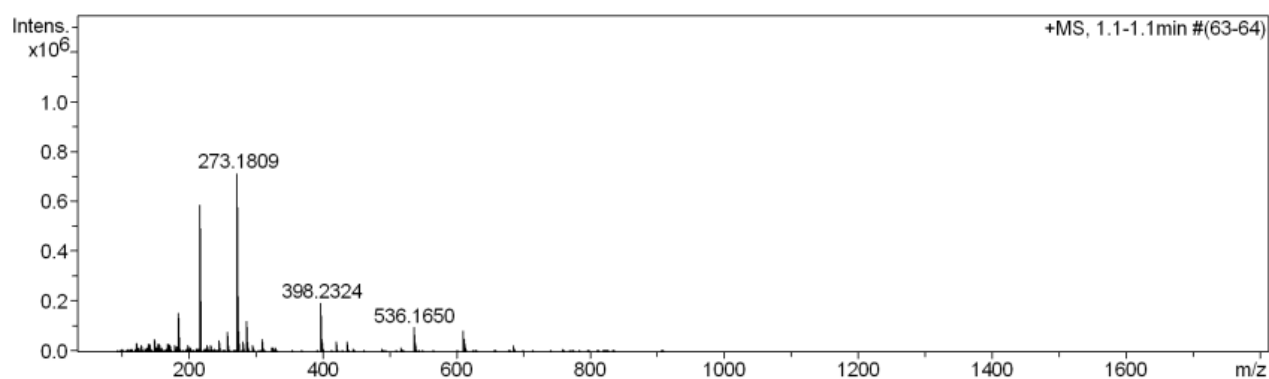
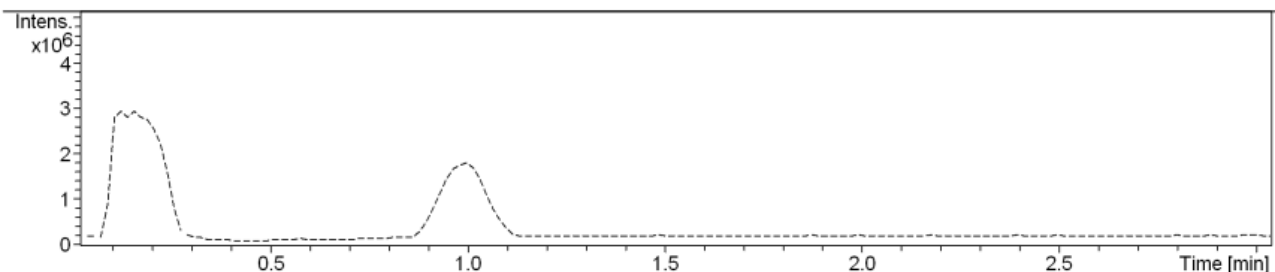
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Method organikai_esi_pos_2013_recover.m
Sample Name GMP_368
Comment

Acquisition Date 4/21/2023 2:18:52 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

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Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
273.1809	1	C ₁₃ H ₂₅ N ₂ O ₄	100.00	273.1809	-0.1	0.2	3.1	2.5	even	ok

Figure S100. Methyl (2*S*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]propanoate ((2*S*,3*R*)-11a). HRMS (ESI-TOF).

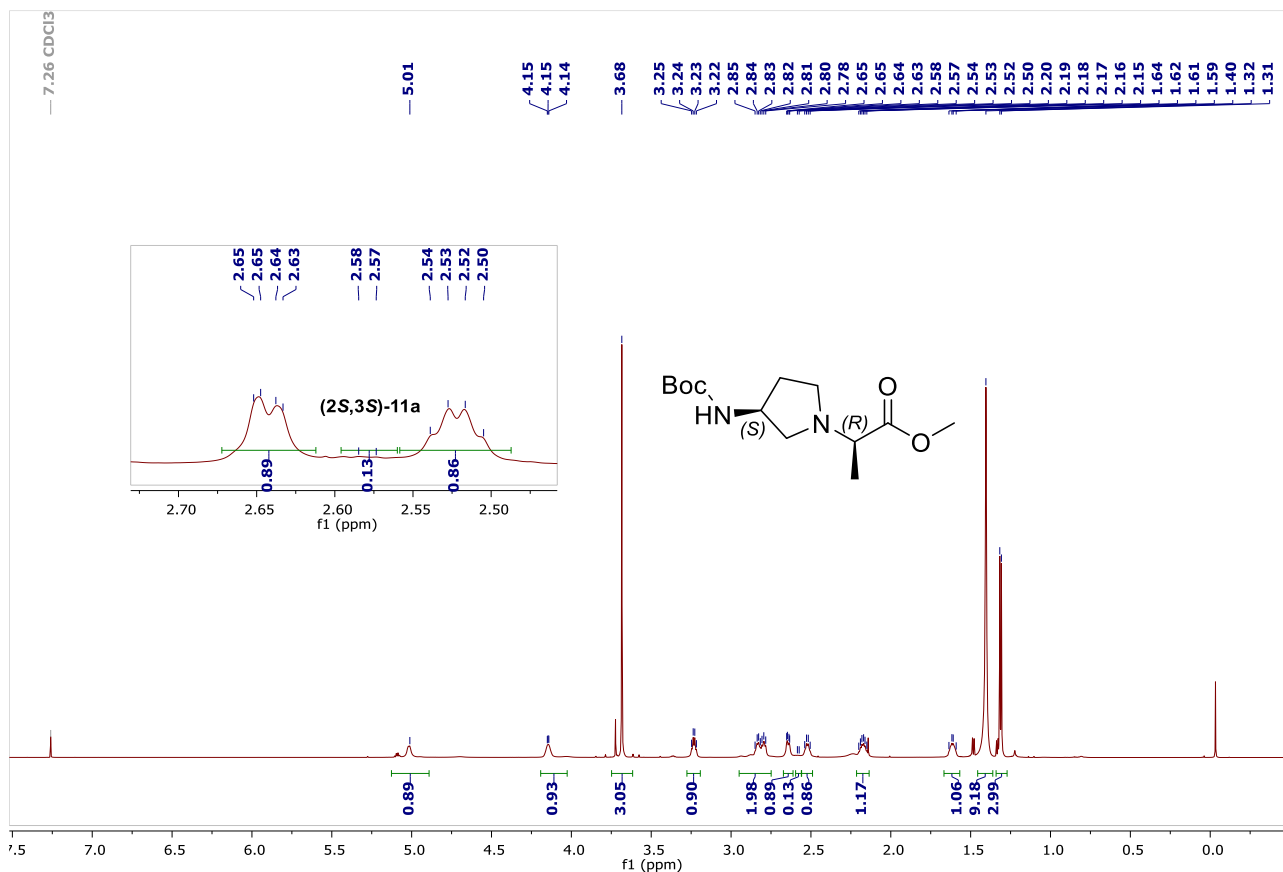


Figure S101. Methyl (2R)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)propanoate ((2R,3S)-11a). ¹H NMR spectrum (700 MHz, CDCl₃).

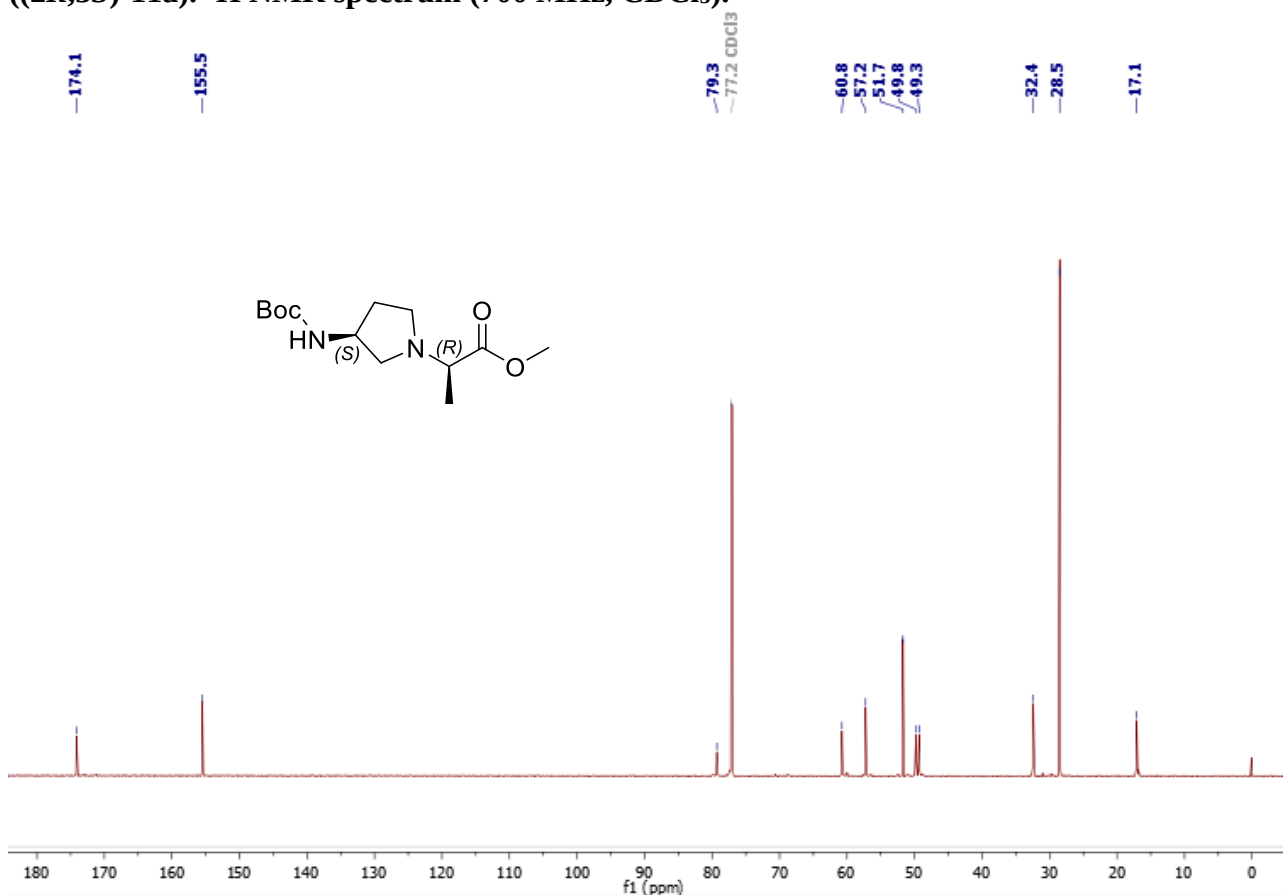


Figure S102. Methyl (2R)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)propanoate ((2R,3S)-11a). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

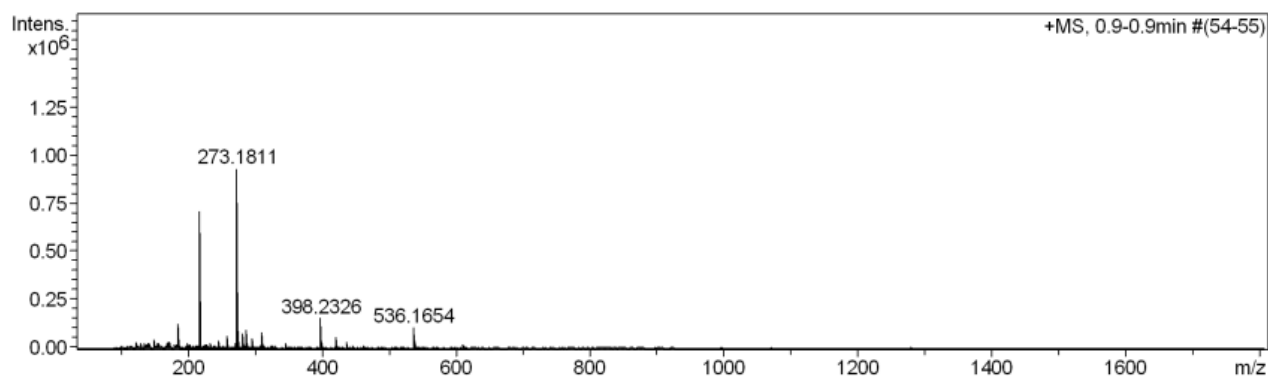
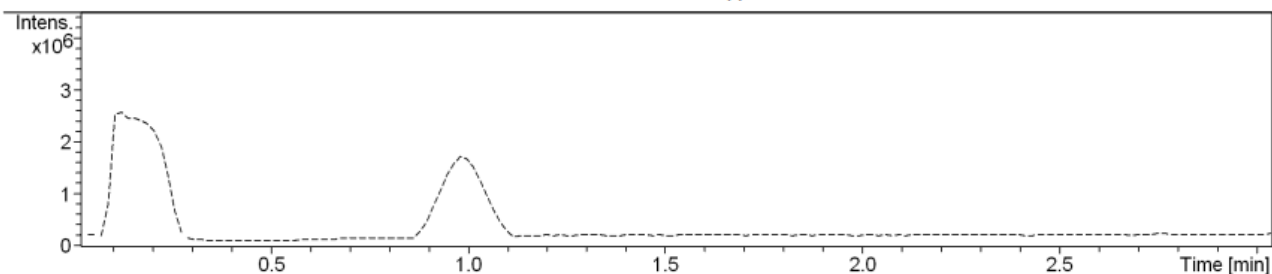
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Method organikai_esi_pos_2013_recover.m
Sample Name GMP_482
Comment

Acquisition Date 4/21/2023 2:32:19 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
273.1811	1	C ₁₃ H ₂₅ N ₂ O ₄	100.00	273.1809	-0.8	-0.5	18.3	2.5	even	ok

Figure S103. Methyl (2R)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)propanoate ((2R,3S)-11a). HRMS (ESI-TOF).

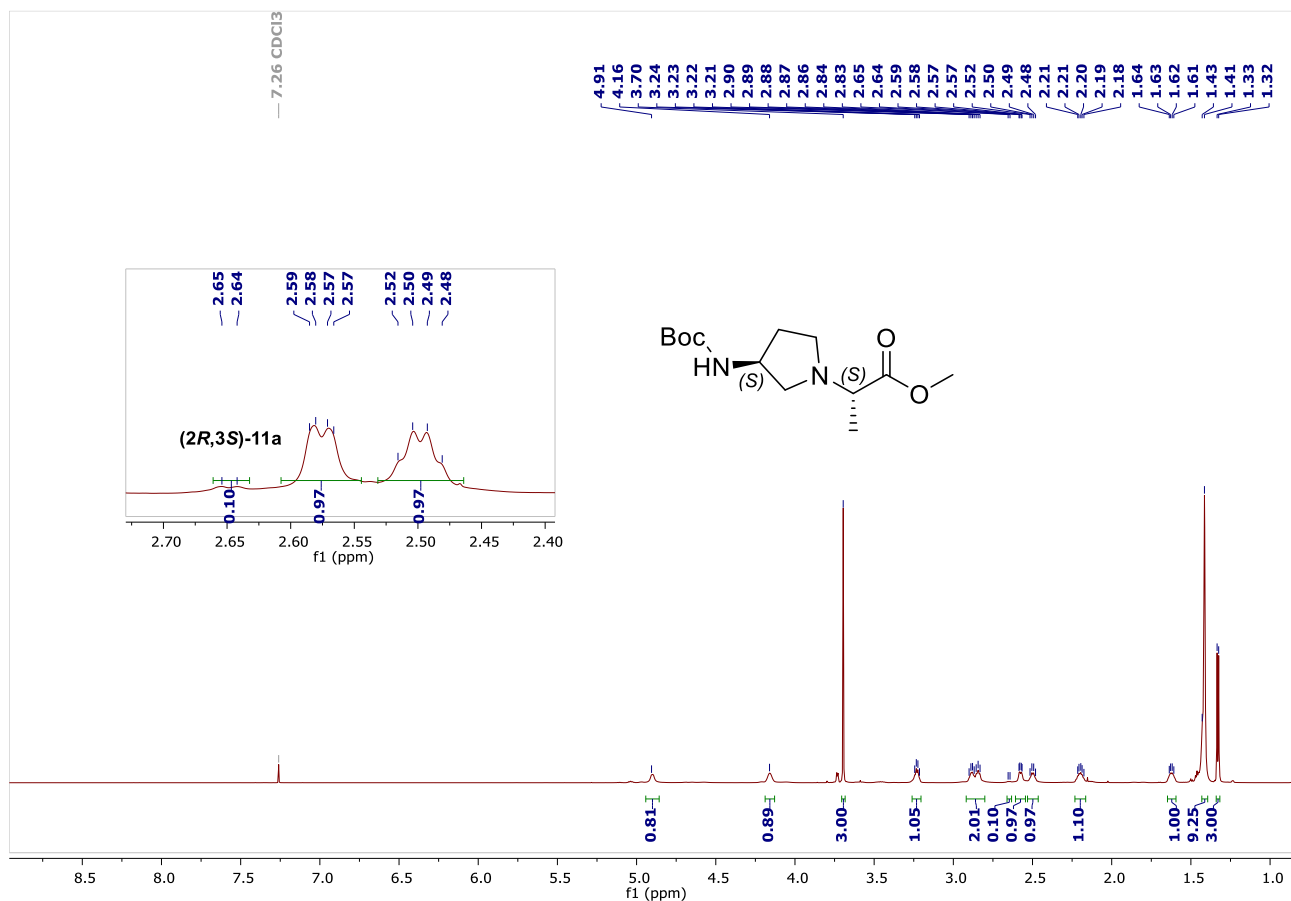


Figure S104. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)propanoate ((2S,3S)-11a). ¹H NMR spectrum (700 MHz, CDCl₃).

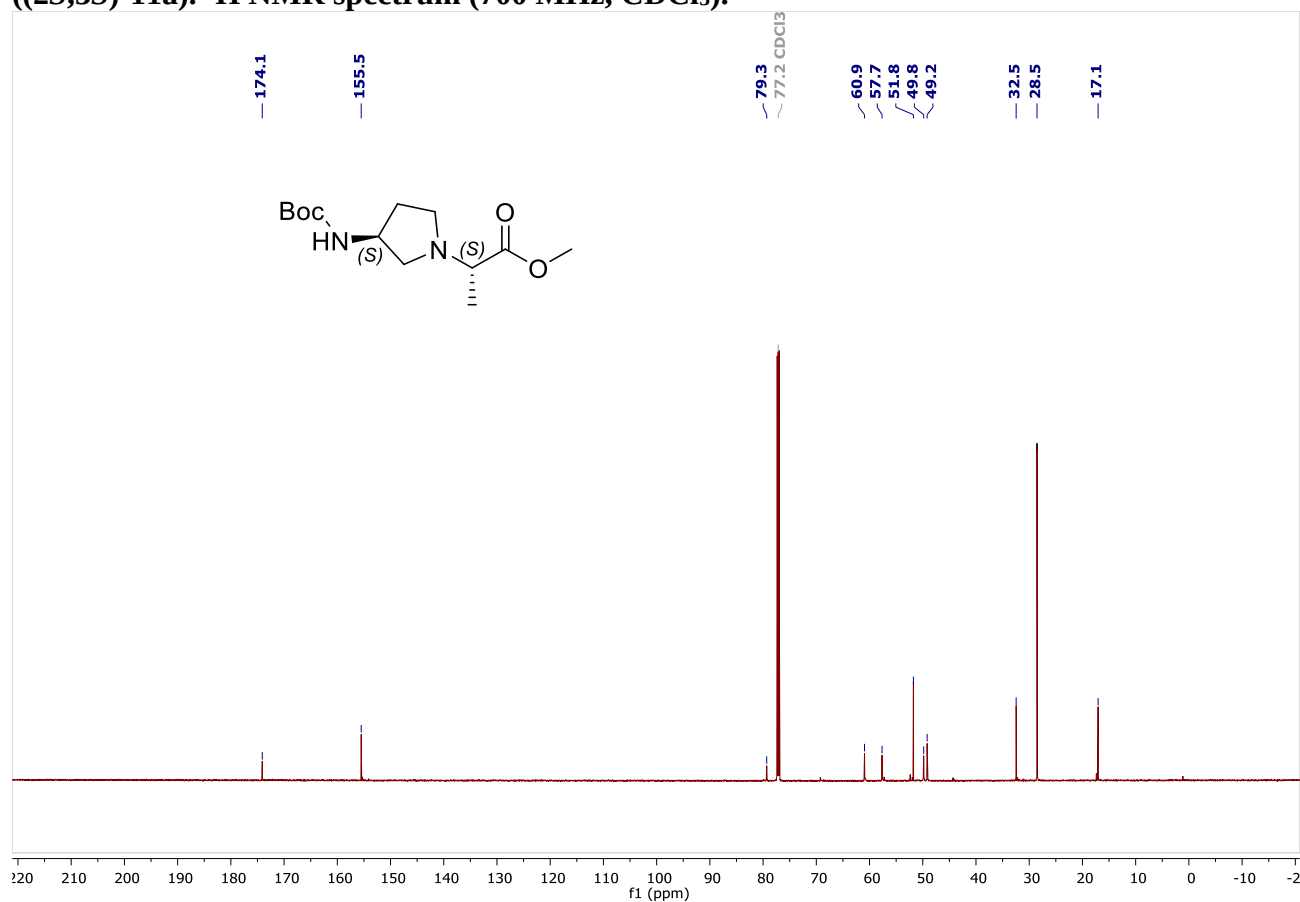


Figure S105. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)propanoate ((2S,3S)-11a). ¹³C NMR spectrum (176 MHz, CDCl₃).

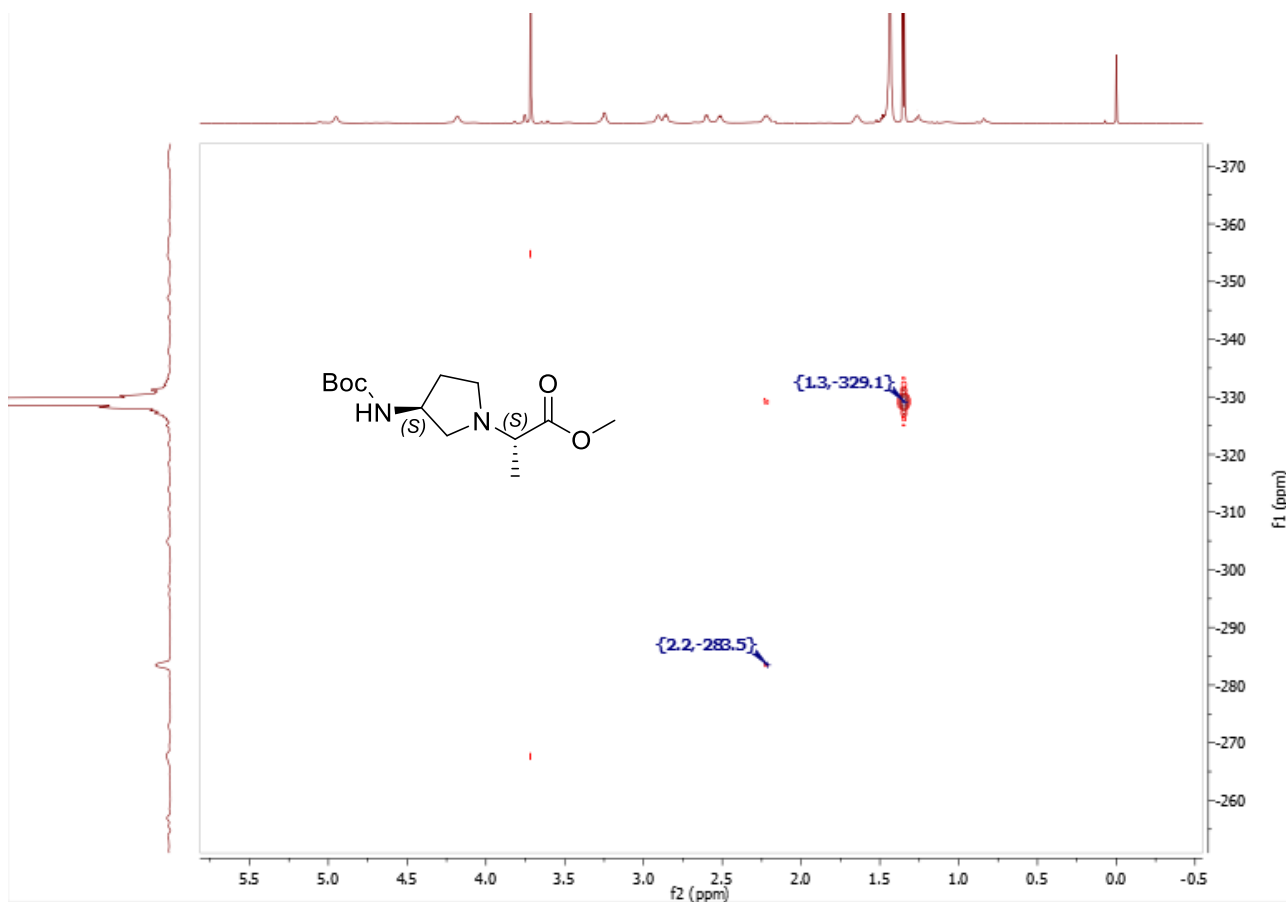


Figure S106. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)propanoate ((2S,3S)-11a). ^1H - ^{15}N HMBC spectrum (71 MHz, CDCl_3).

Mass Spectrum SmartFormula Report

Analysis Info

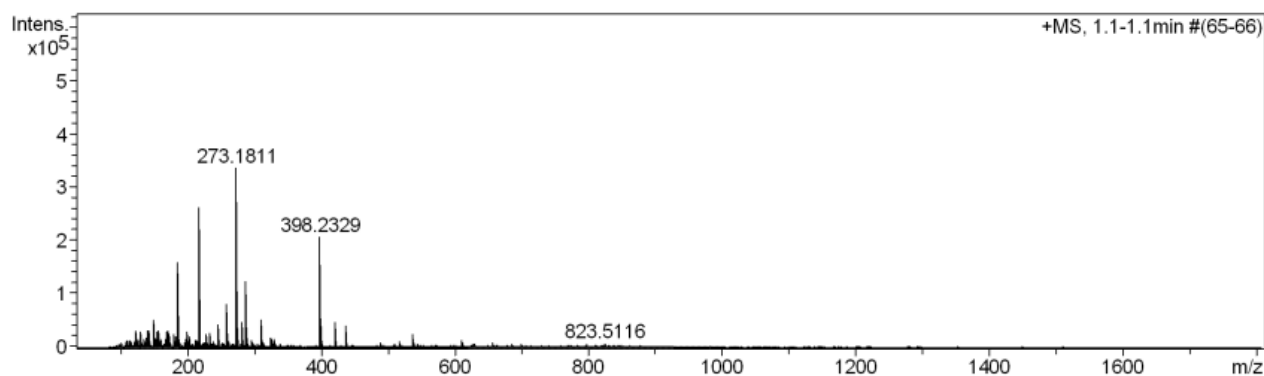
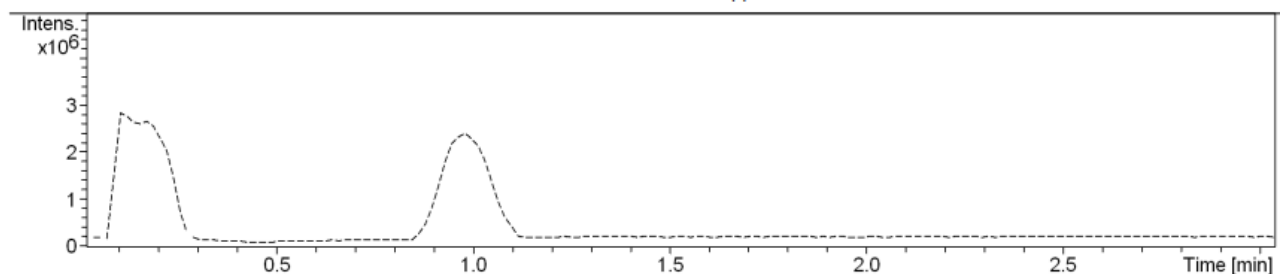
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Method organikai_esi_pos_2013_recover.m
Sample Name GMP_373
Comment

Acquisition Date 4/21/2023 2:27:50 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
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Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
273.1811	1	C ₁₃ H ₂₅ N ₂ O ₄	100.00	273.1809	-0.7	-0.5	8.1	2.5	even	ok

Figure S107. Methyl (2S)-2-[(3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]propanoate ((2S,3S)-11a). HRMS (ESI-TOF).

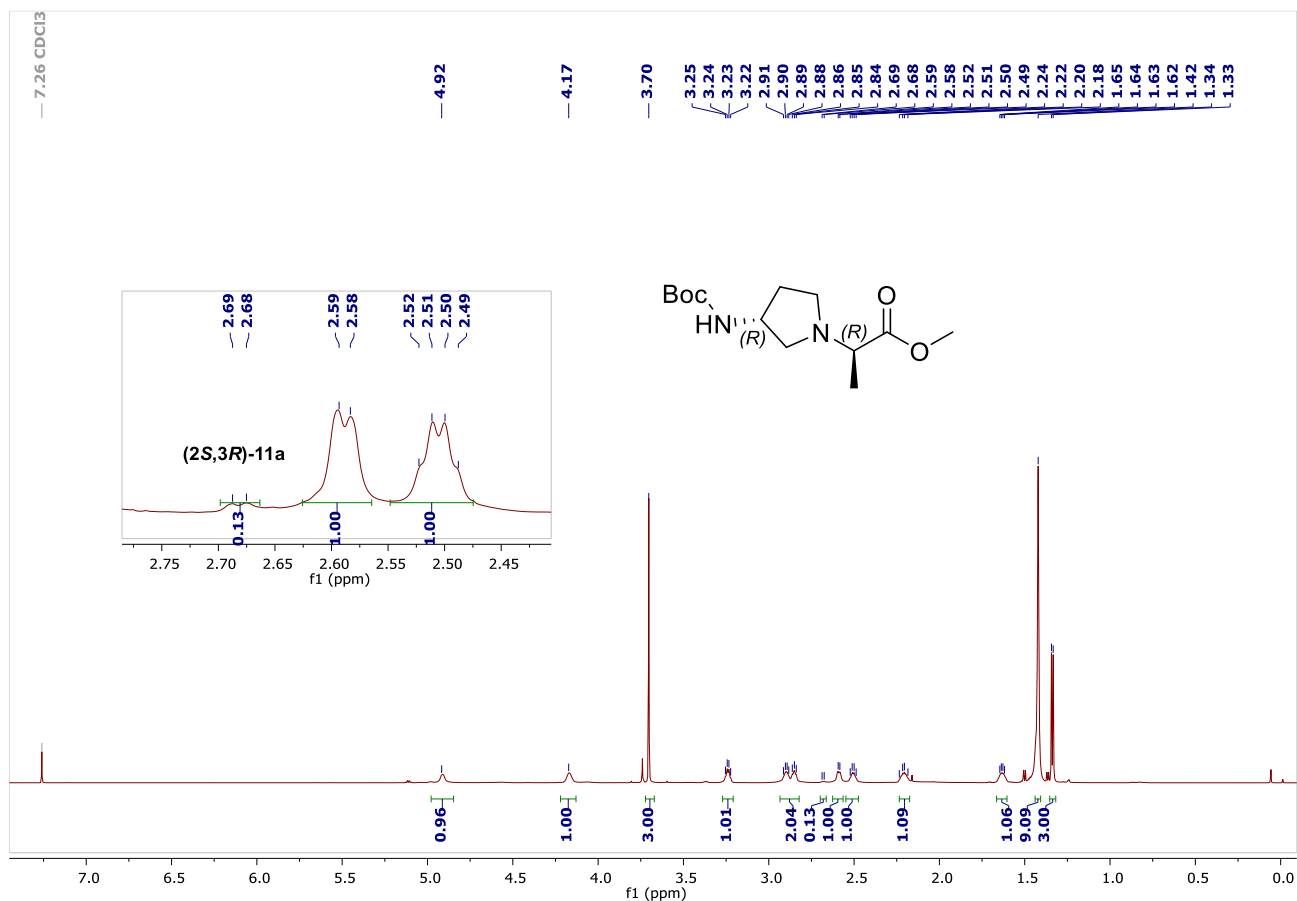


Figure S108. Methyl (2R)-2-[(3R)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]propanoate ((2R,3R)-11a). ¹H NMR spectrum (700 MHz, CDCl₃).

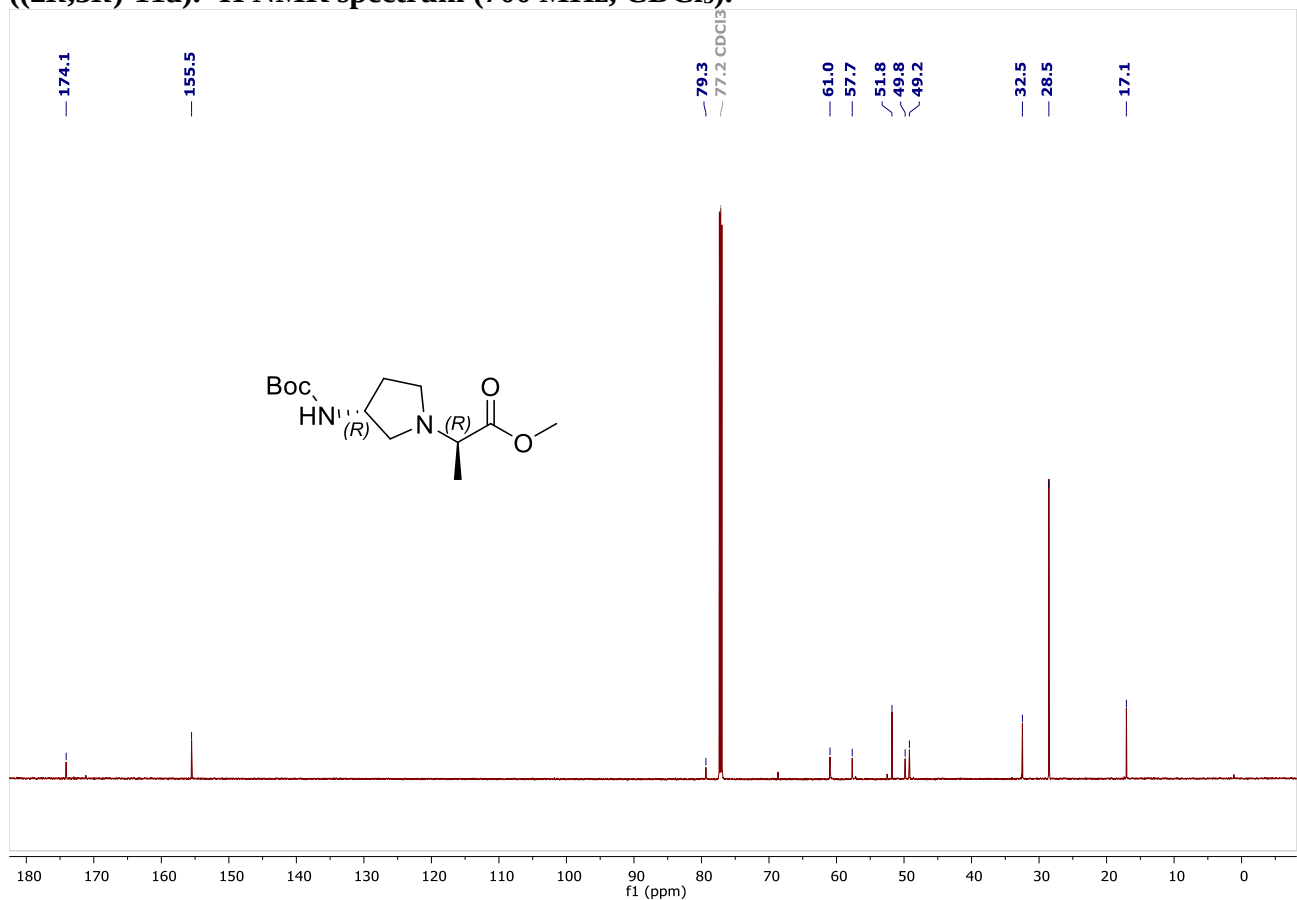


Figure S109. Methyl (2R)-2-[(3R)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]propanoate ((2R,3R)-11a). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

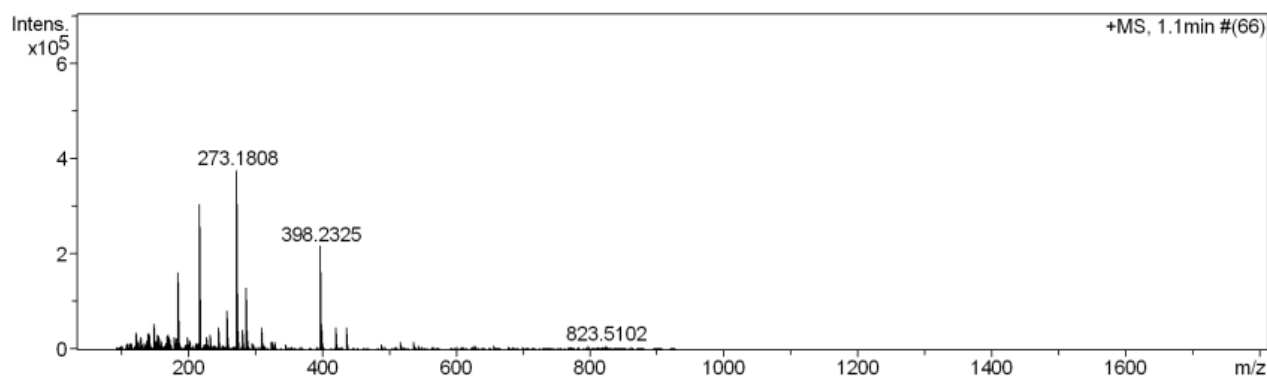
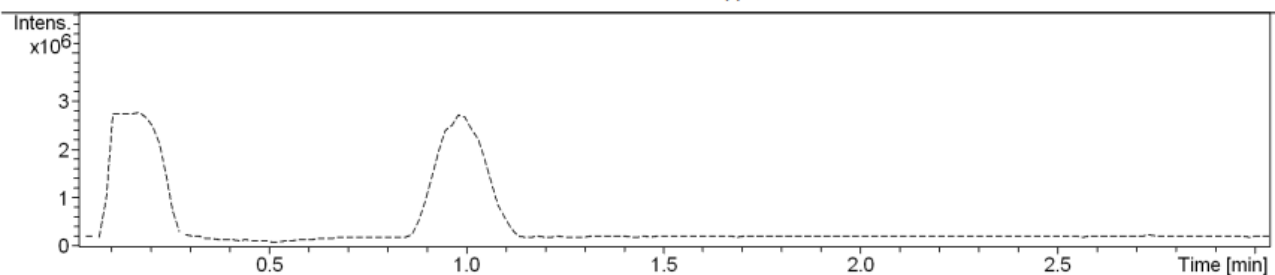
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 Method organikai_esi_pos_2013_recover.m
 Sample Name GMP_476
 Comment

Acquisition Date 4/21/2023 2:23:19 PM

Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
273.1808	1	C ₁₃ H ₂₅ N ₂ O ₄	100.00	273.1809	0.2	0.4	14.7	2.5	even	ok

Figure S110. Methyl (2R)-2-[(3R)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]propanoate ((2R,3R)-11a). HRMS (ESI-TOF).

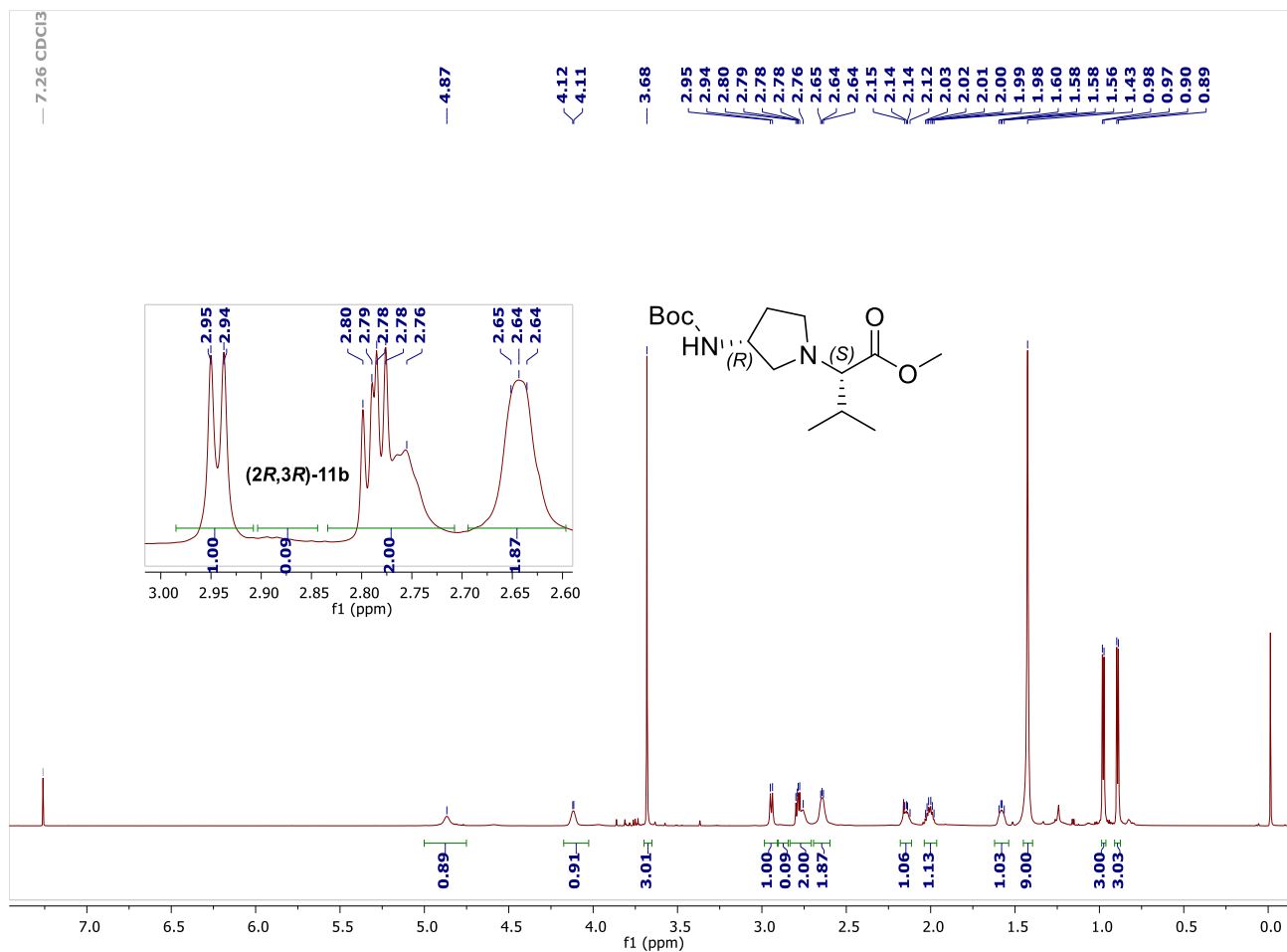


Figure S111. Methyl (2S)-2-((3R)-3-((tert-butoxycarbonyl)amino)pyrrolidin-1-yl)-3-methylbutanoate ((2S,3R)-11b). ¹H NMR spectrum (700 MHz, CDCl₃).

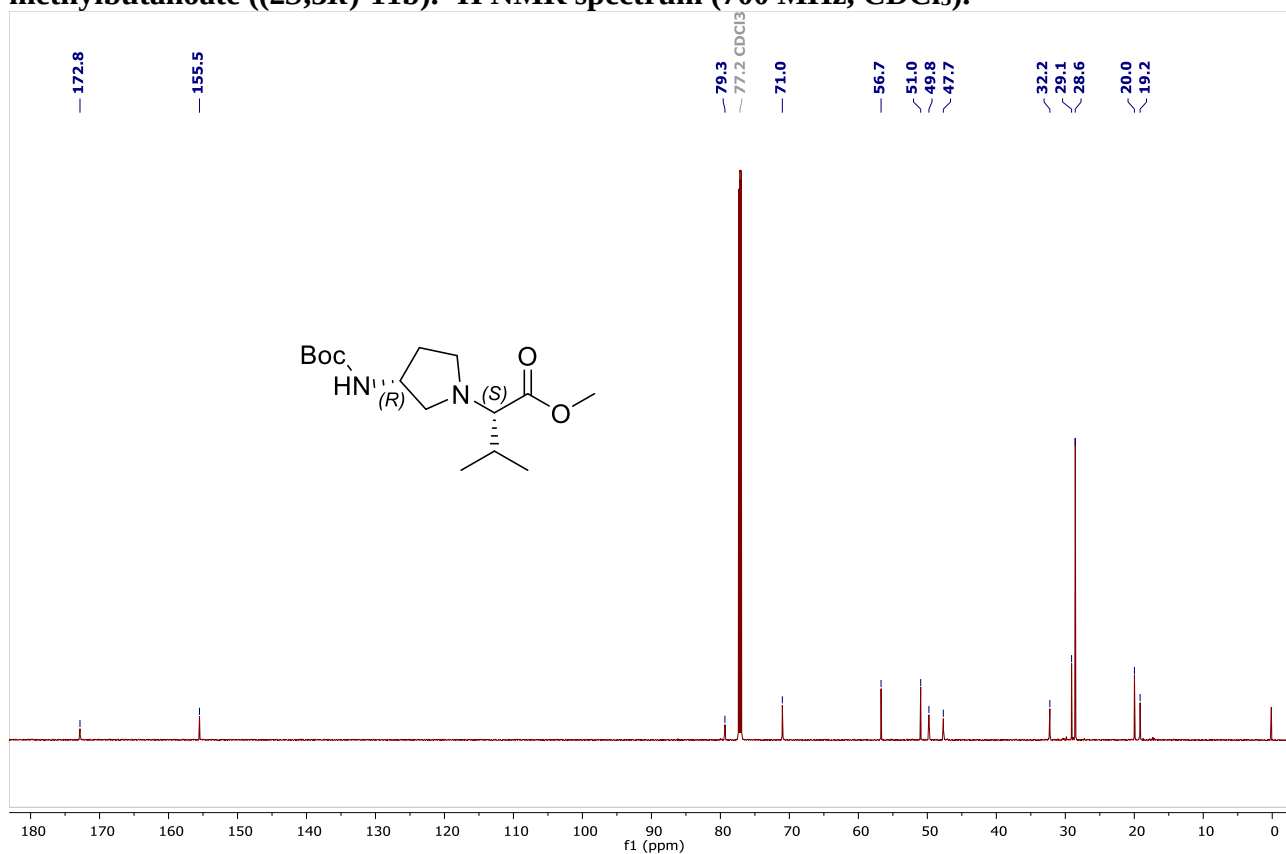


Figure S112. Methyl (2S)-2-((3R)-3-((tert-butoxycarbonyl)amino)pyrrolidin-1-yl)-3-methylbutanoate ((2S,3R)-11b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

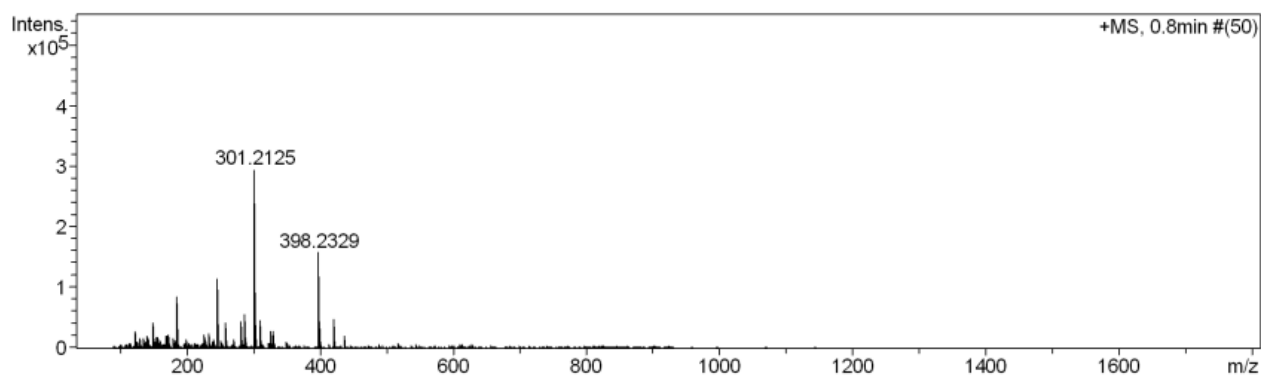
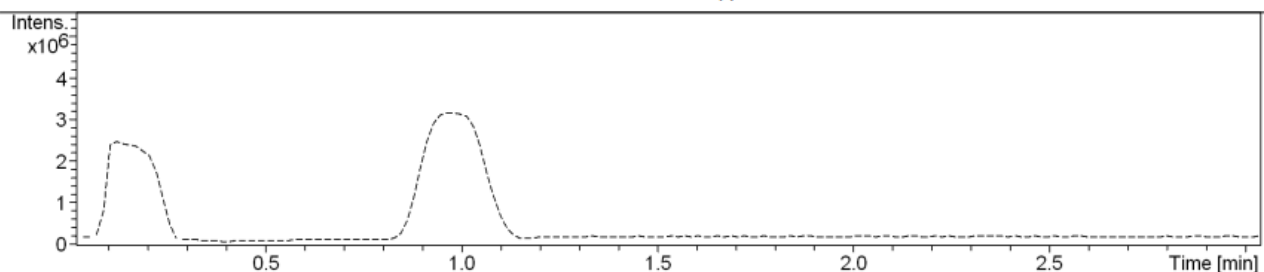
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Method organikai_esi_pos_2013_recover.m
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Comment

Acquisition Date 4/14/2023 12:52:37 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

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Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
301.2125	1	C ₁₅ H ₂₉ N ₂ O ₄	100.00	301.2122	-0.9	-0.7	13.2	2.5	even	ok

Figure S113. Methyl (2S)-2-[(3R)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]-3-methylbutanoate ((2S,3R)-11b). HRMS (ESI-TOF).

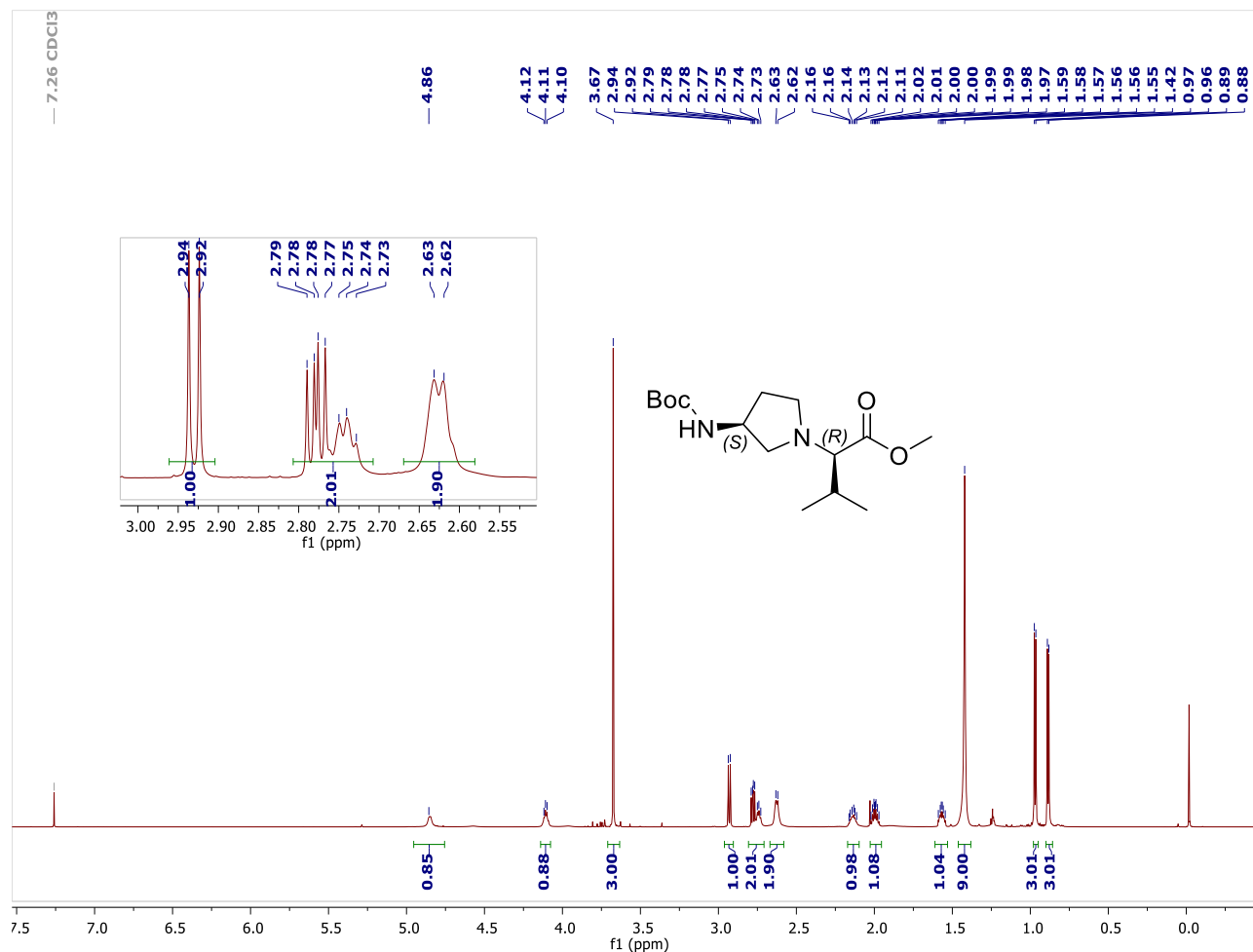


Figure S114. Methyl (2R)-2-[(3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]-3-methylbutanoate ((2R,3S)-11b). ¹H NMR spectrum (700 MHz, CDCl₃).

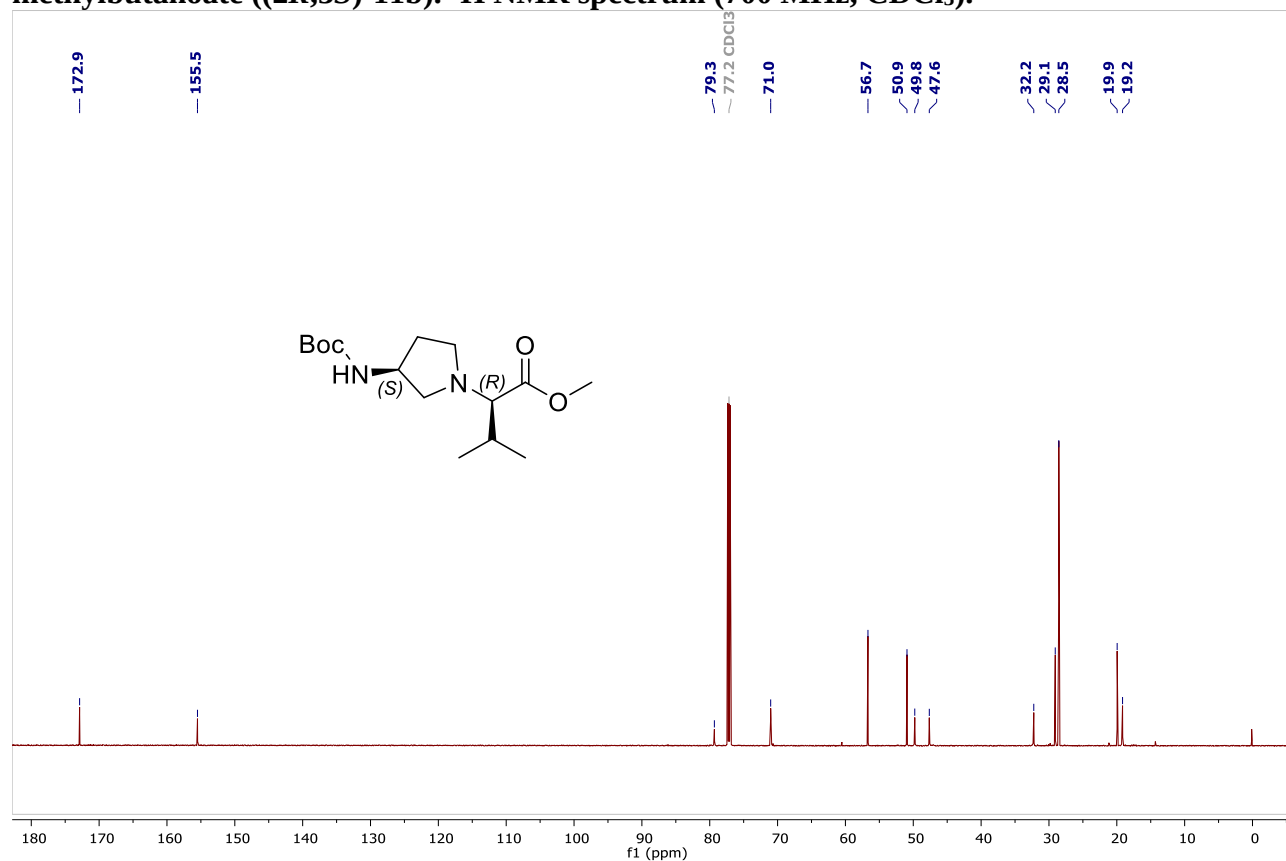


Figure S115. Methyl (2R)-2-[(3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]-3-methylbutanoate ((2R,3S)-11b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

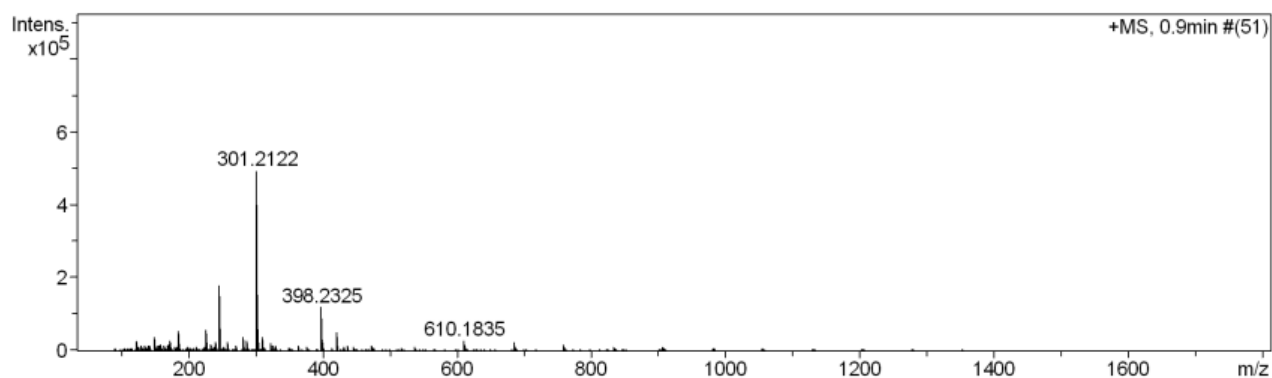
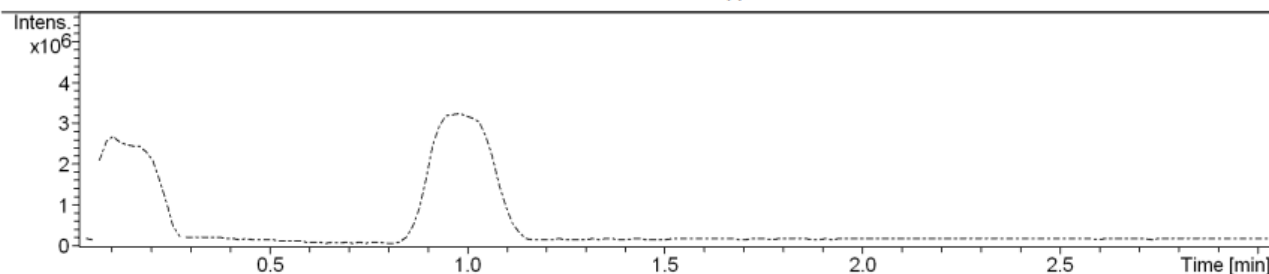
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Method organikai_esi_pos_2013_recover.m
Sample Name GMP_566
Comment

Acquisition Date 4/13/2023 11:07:30 AM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
301.2122	1	C ₁₅ H ₂₉ N ₂ O ₄	100.00	301.2122	0.0	0.2	16.8	2.5	even	ok

Figure S116. Methyl (2R)-2-((3S)-3-((tert-butoxycarbonyl)amino)pyrrolidin-1-yl)-3-methylbutanoate ((2R,3S)-11b). HRMS (ESI-TOF).

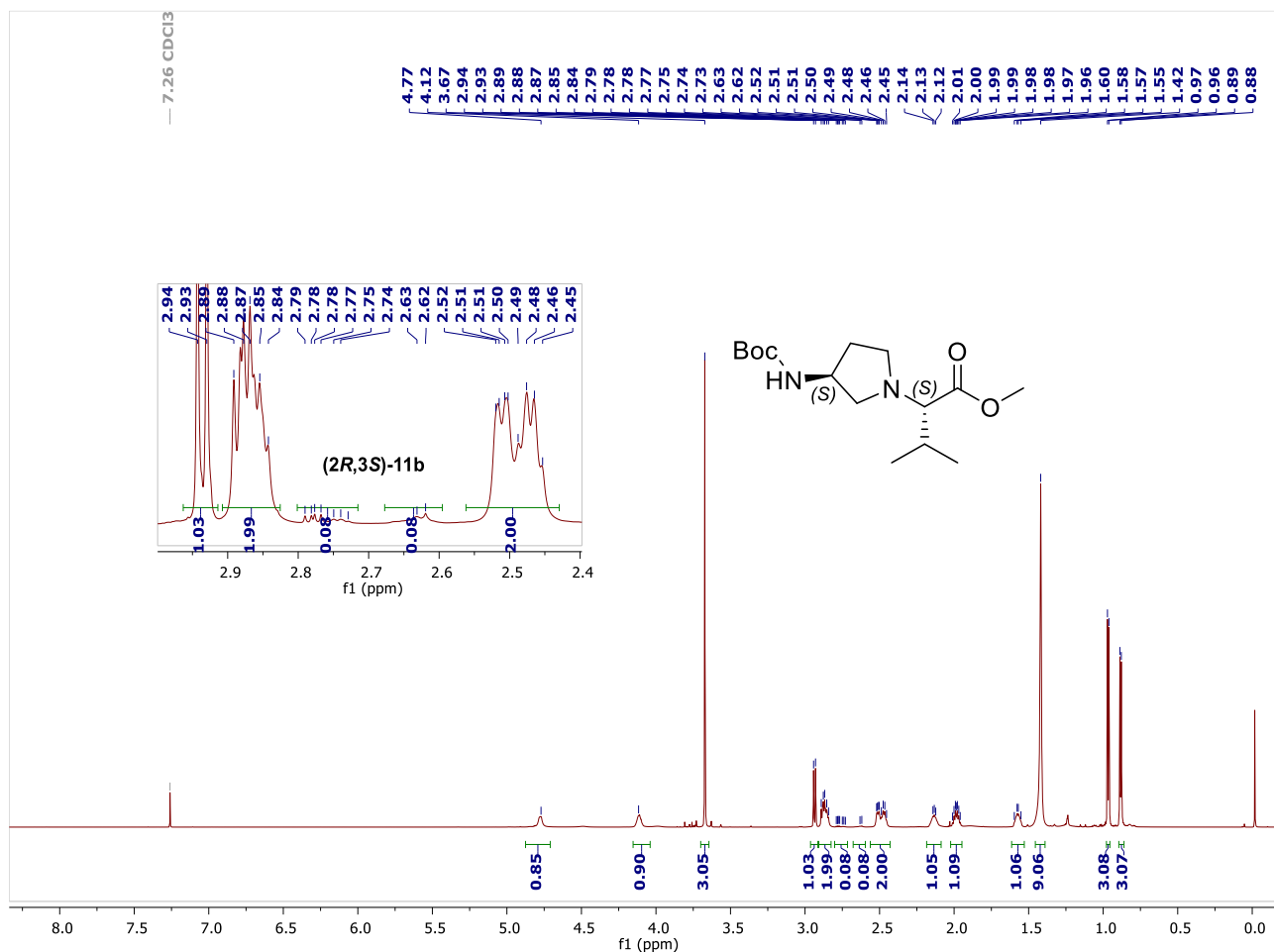


Figure S117. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)-3-methylbutanoate ((2S,3S)-11b). ¹H NMR spectrum (700 MHz, CDCl₃).

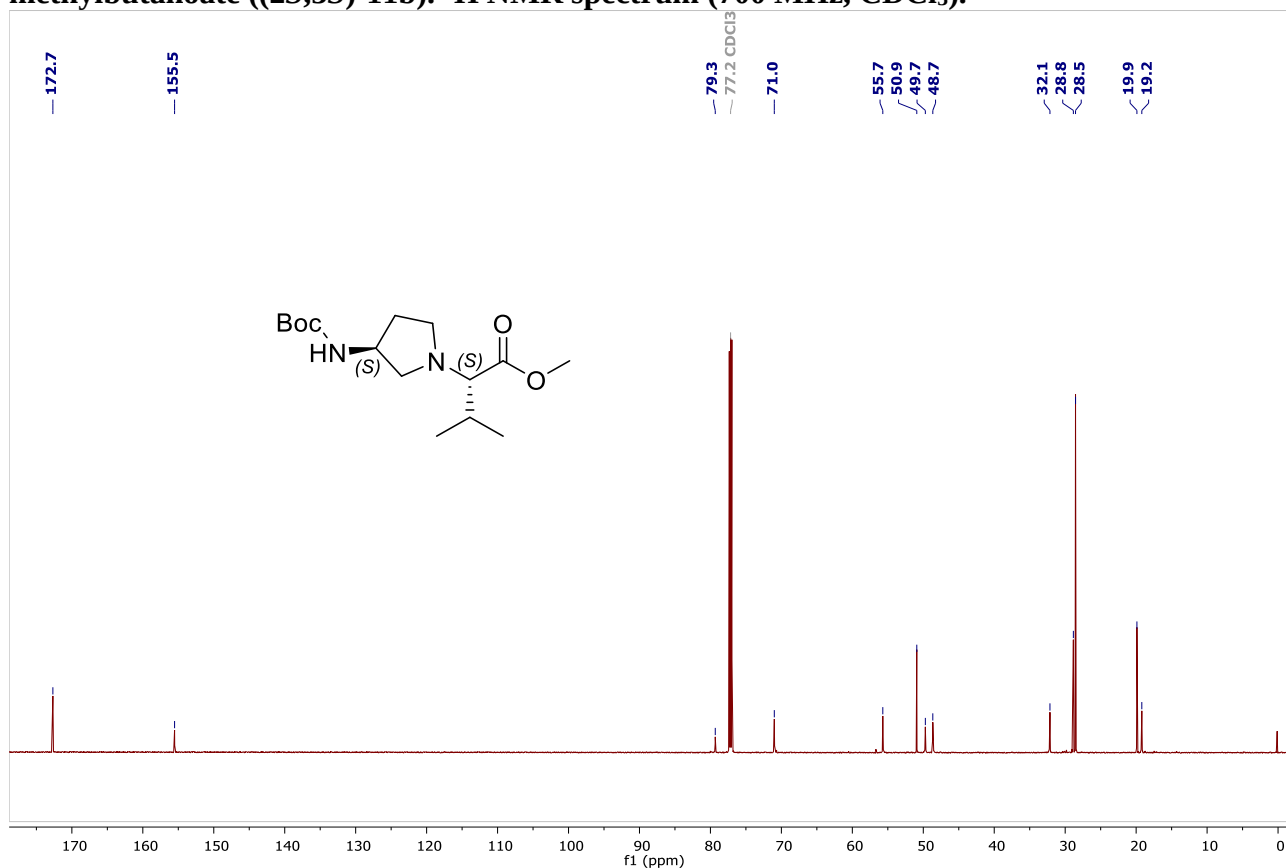


Figure S118. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)-3-methylbutanoate ((2S,3S)-11b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Compound Spectrum SmartFormula Report

Analysis Info

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Sample Name GMP-569
Comment AB

Acquisition Date 5/2/2023 7:57:07 PM

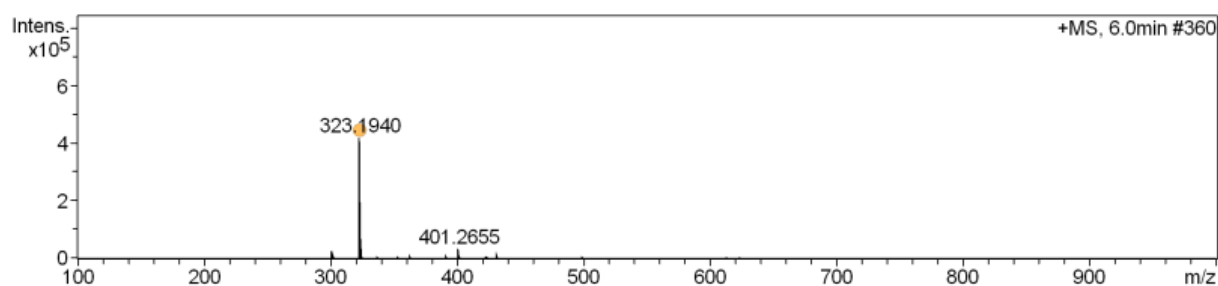
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Instrument micrOTOF-Q III 8228888.20448

Acquisition Parameter

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Scan Begin	50 m/z	Set End Plate Offset	-500 V	Set Dry Gas	4.0 l/min
Scan End	1000 m/z	Set Collision Cell RF	140.0 Vpp	Set Divert Valve	Waste

#	RT [min]	Area	Int. Type	I	S/N	Chromatogram	Max. m/z	FWHM [min]
n.a.	6.0	n.a.	Single spectrum	n.a.	n.a.	n.a.	323.1940	n.a.

+MS, 6.0min #360



Meas. m/z	#	Ion Formula	m/z	err [ppm]	mSigma	# Sigma	Score	rdb	e ⁻ Conf	N-Rule
323.1940	1	C15H28N2NaO4	323.1941	-0.5	2.4	1	100.00	2.5	even	ok

Figure S119. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)-3-methylbutanoate ((2S,3S)-11b). HRMS (ESI-TOF).

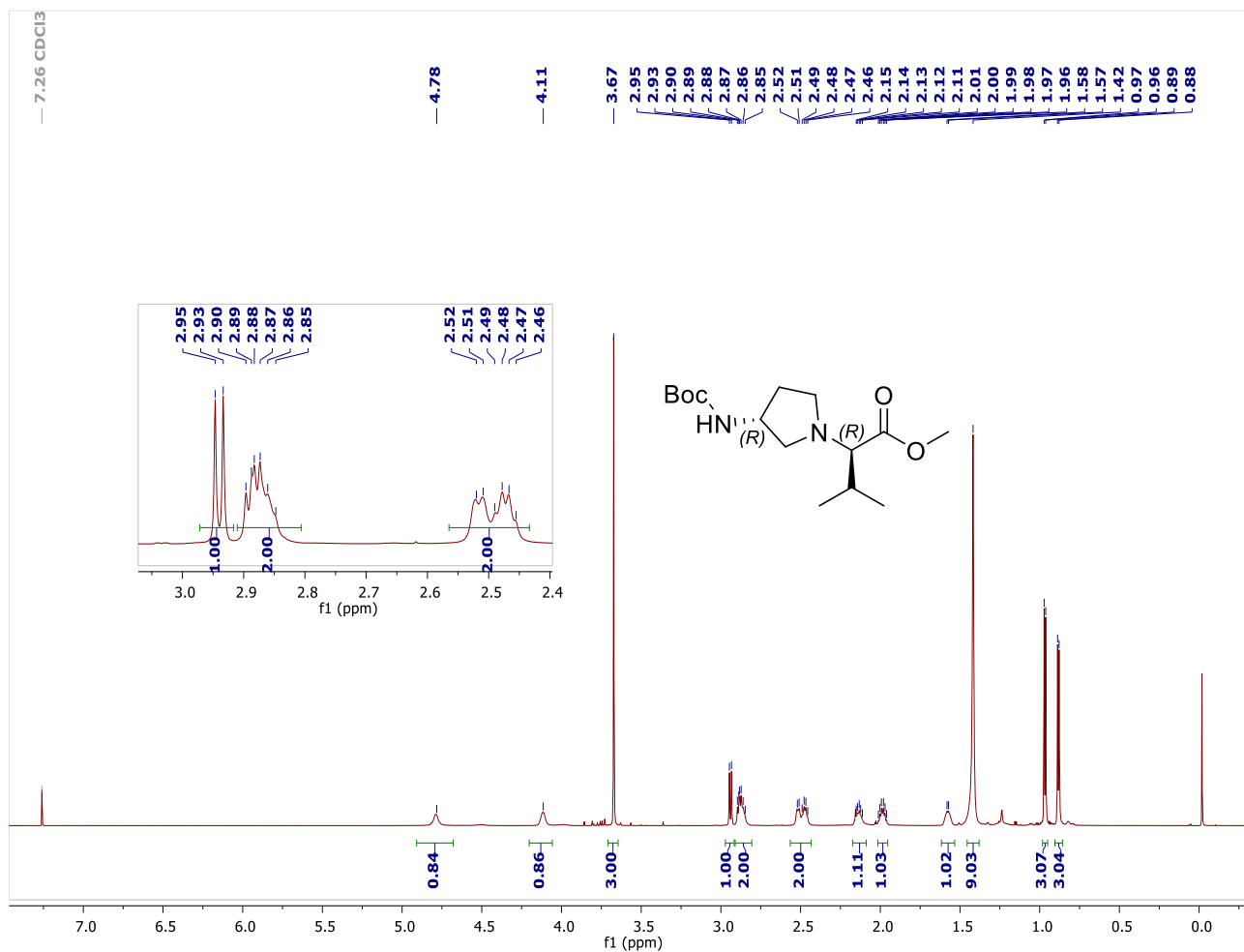


Figure S120. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]-3-methylbutanoate ((2*R*,3*R*)-11b). ¹H NMR spectrum (700 MHz, CDCl₃).

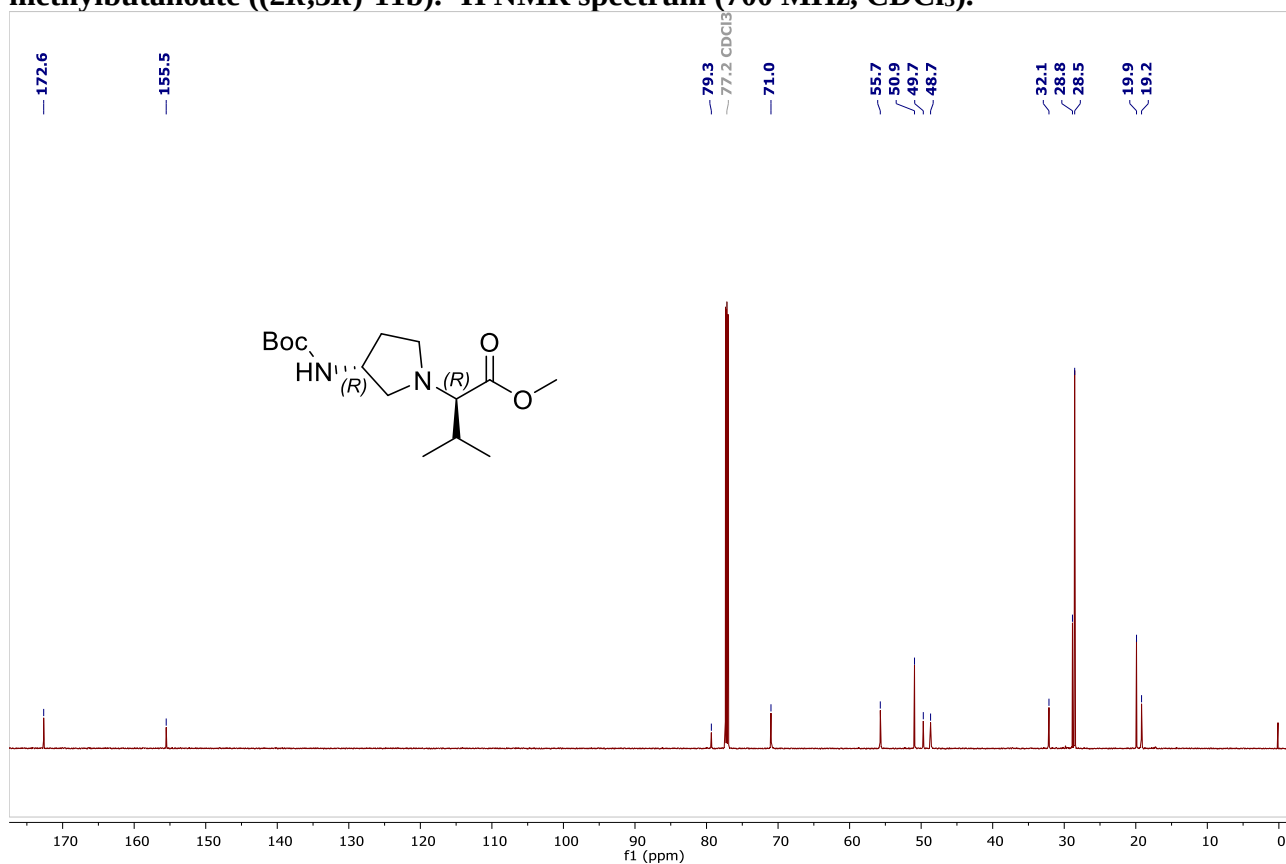


Figure S121. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]-3-methylbutanoate ((2*R*,3*R*)-11b). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

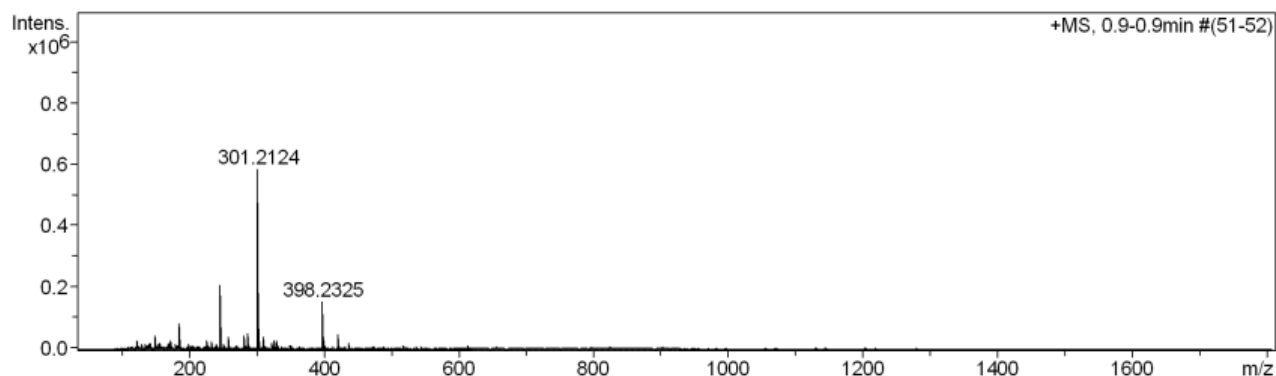
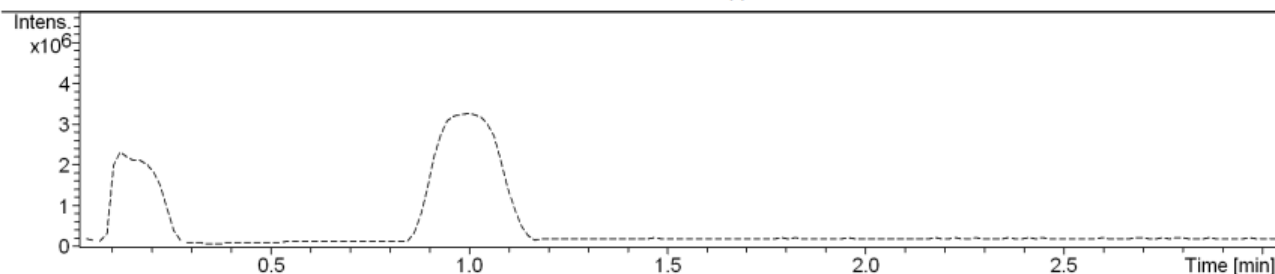
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Sample Name GMP_572
Comment

Acquisition Date 4/14/2023 12:48:09 PM

Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	1.5 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
301.2124	1	C ₁₅ H ₂₉ N ₂ O ₄	100.00	301.2122	-0.6	-0.3	15.7	2.5	even	ok

Figure S122. Methyl (2R)-2-((3R)-3-((tert-butoxycarbonyl)amino)pyrrolidin-1-yl)-3-methylbutanoate ((2R,3R)-11b). HRMS (ESI-TOF).

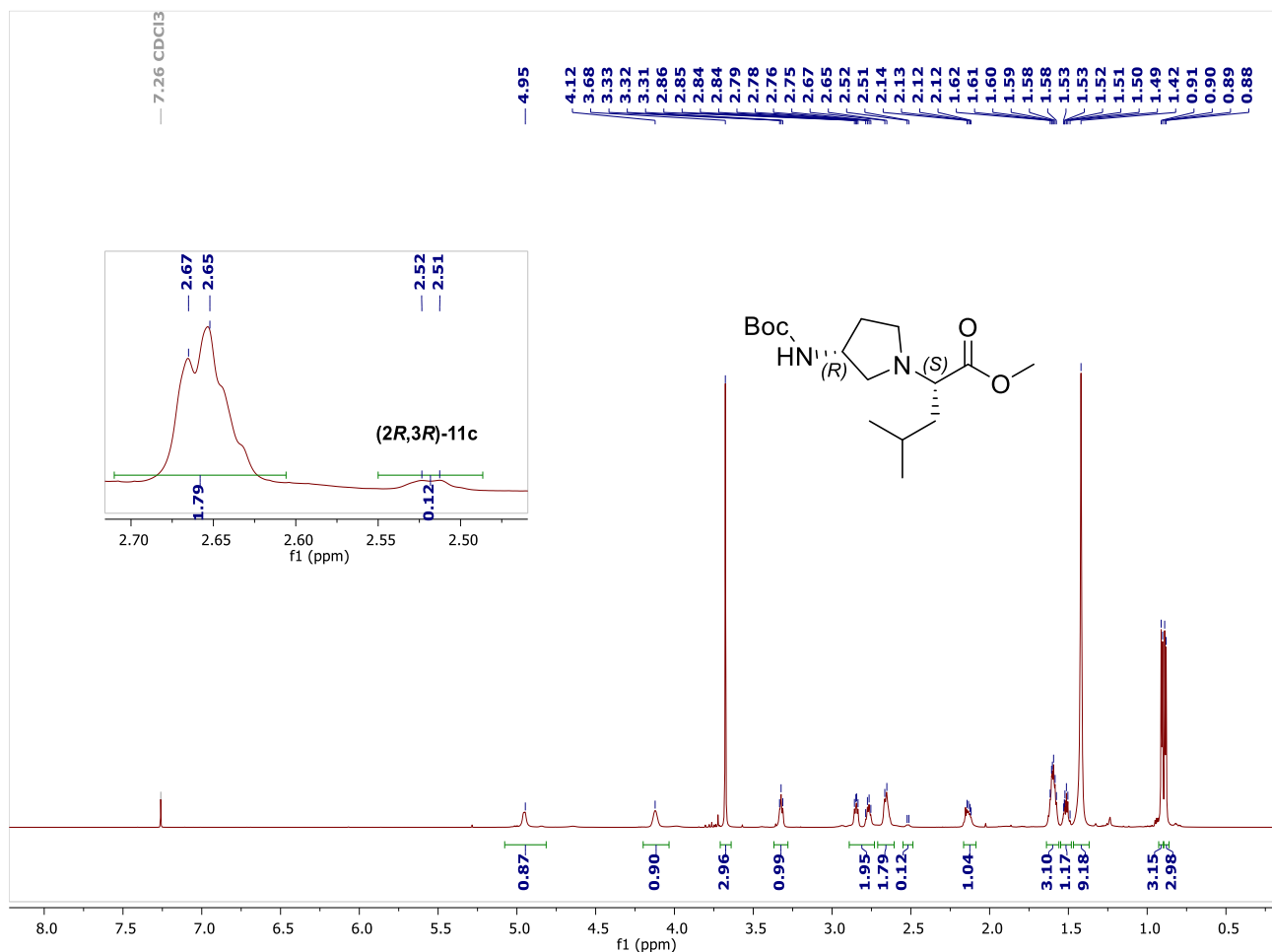


Figure S123. Methyl (2*S*)-2-((3*R*)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)-4-methylpentanoate ((2*S*,3*R*)-11c). ¹H NMR spectrum (700 MHz, CDCl₃).

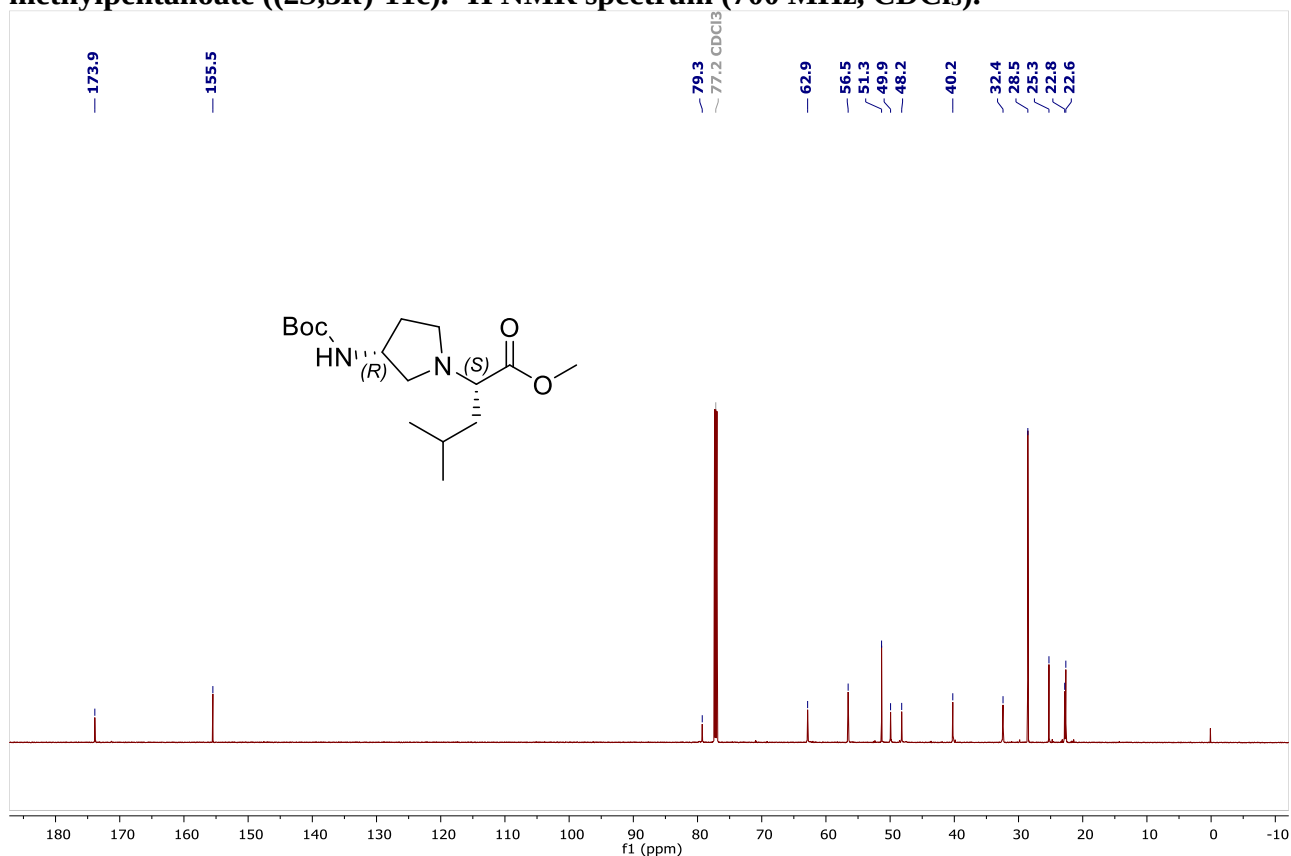


Figure S124. Methyl (2*S*)-2-((3*R*)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)-4-methylpentanoate ((2*S*,3*R*)-11c). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

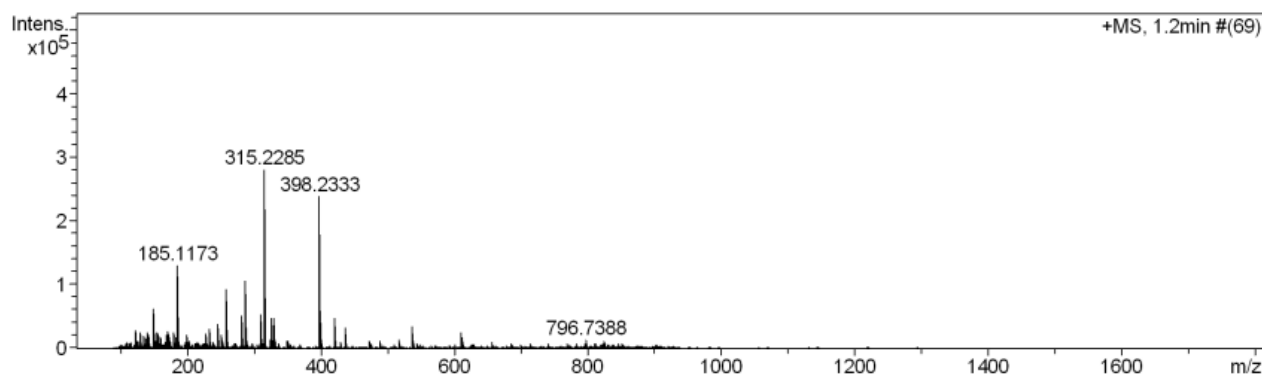
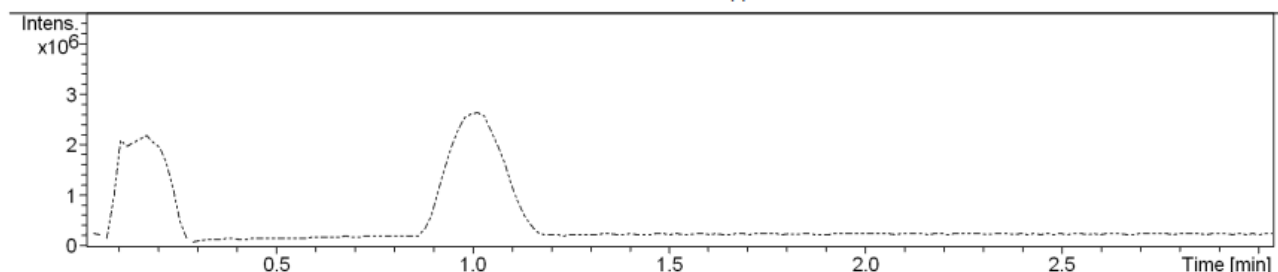
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Acquisition Date 4/27/2023 3:30:54 PM

Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

Acquisition Parameter

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Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
315.2285	1	C ₁₆ H ₃₁ N ₂ O ₄	100.00	315.2278	-2.0	-1.8	15.0	2.5	even	ok

Figure S125. Methyl (2S)-2-((3R)-3-((tert-butoxycarbonyl)amino)pyrrolidin-1-yl)-4-methylpentanoate ((2S,3R)-11c). HRMS (ESI-TOF).

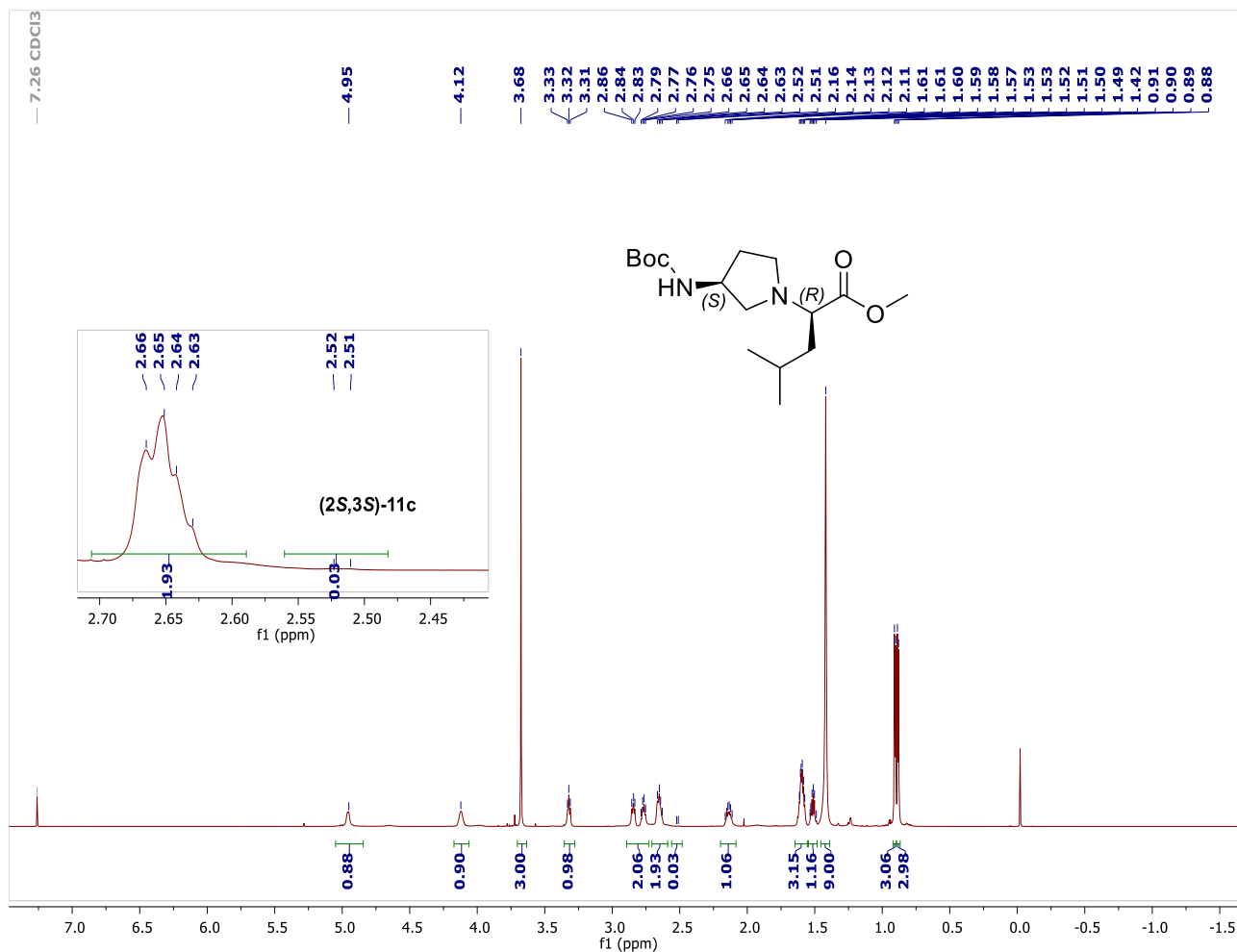


Figure S126. Methyl (2R)-2-[(3S)-3-[(tert-butoxycarbonyl)amino]pyrrolidin-1-yl]-4-methylpentanoate ((2R,3S)-11c). ¹H NMR spectrum (700 MHz, CDCl₃).

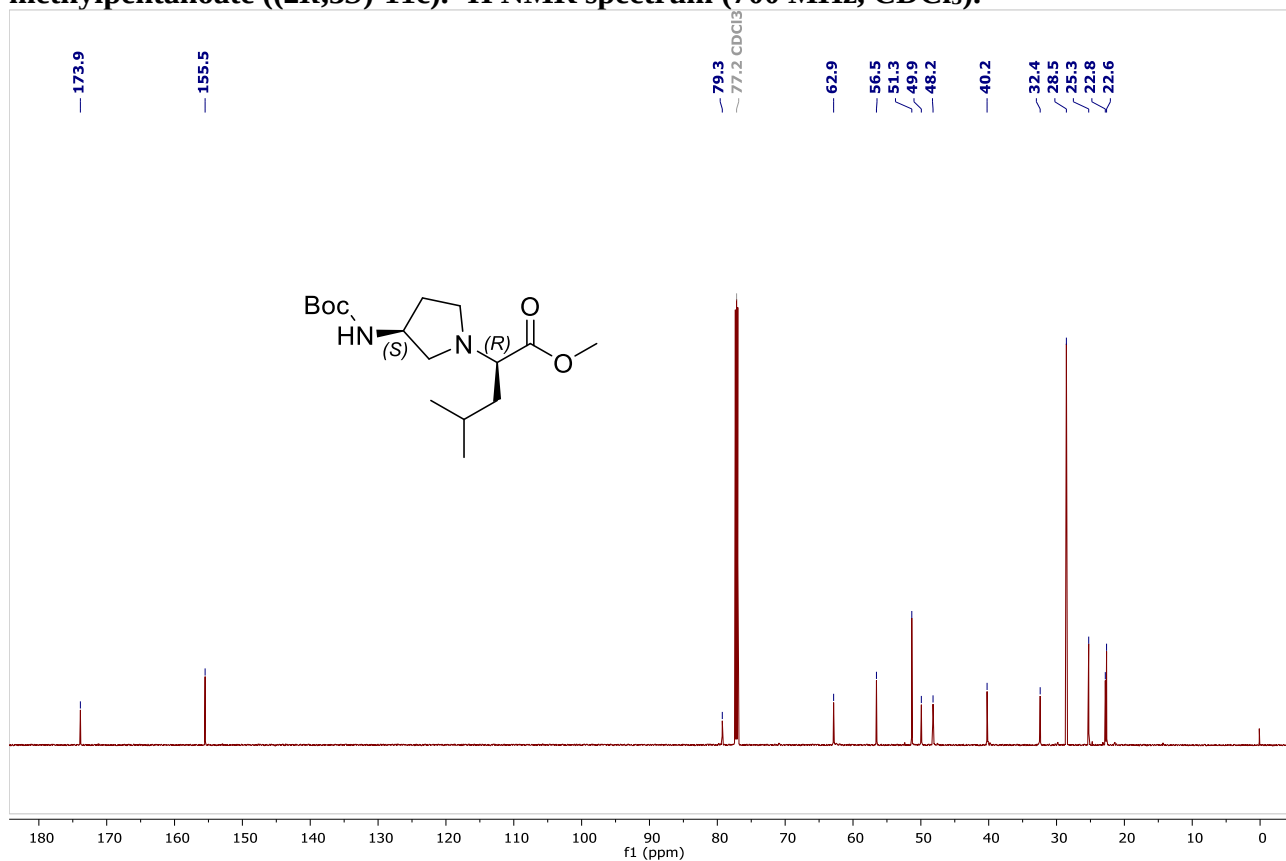


Figure S127. Methyl (2R)-2-[(3S)-3-[(tert-butoxycarbonyl)amino]pyrrolidin-1-yl]-4-methylpentanoate ((2R,3S)-11c). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

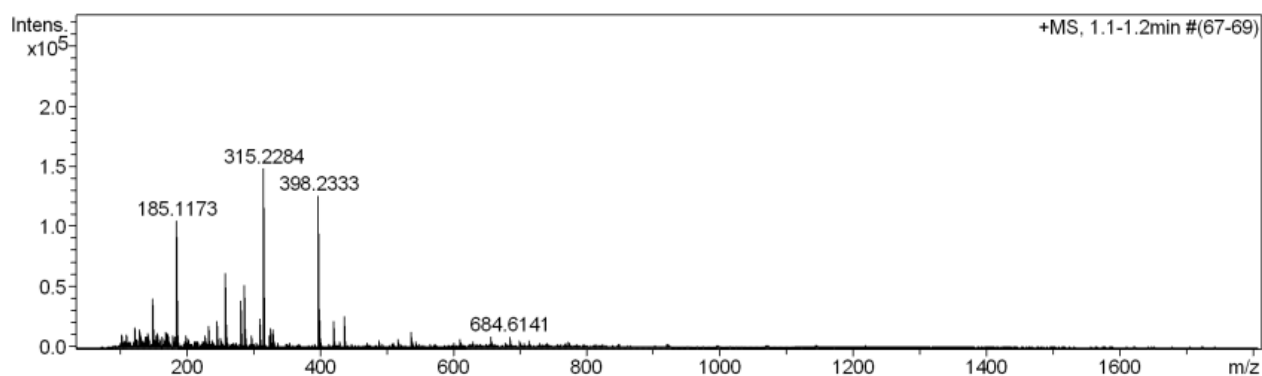
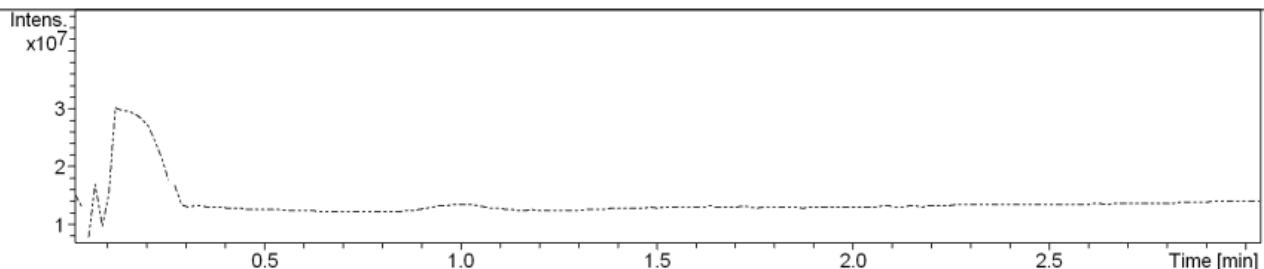
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Acquisition Date 4/27/2023 3:26:28 PM

Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

Acquisition Parameter

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Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
315.2284	1	C ₁₆ H ₃₁ N ₂ O ₄	100.00	315.2278	-1.8	-1.6	7.3	2.5	even	ok

Figure S128. Methyl (2*R*)-2-[(3*S*)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]-4-methylpentanoate ((2*R*,3*S*)-11c). HRMS (ESI-TOF).

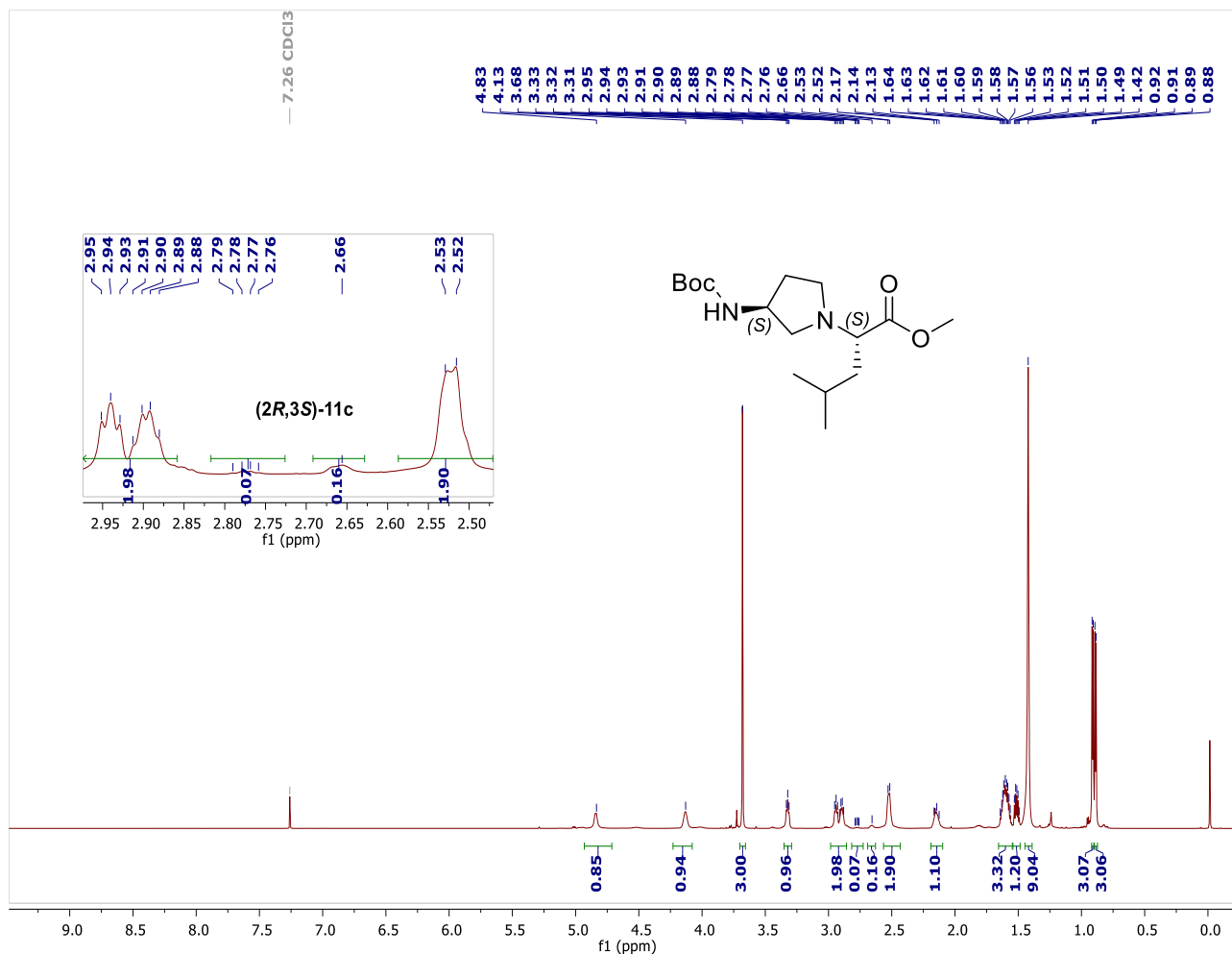


Figure S129. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)-4-methylpentanoate ((2S,3S)-11c). ¹H NMR spectrum (700 MHz, CDCl₃).

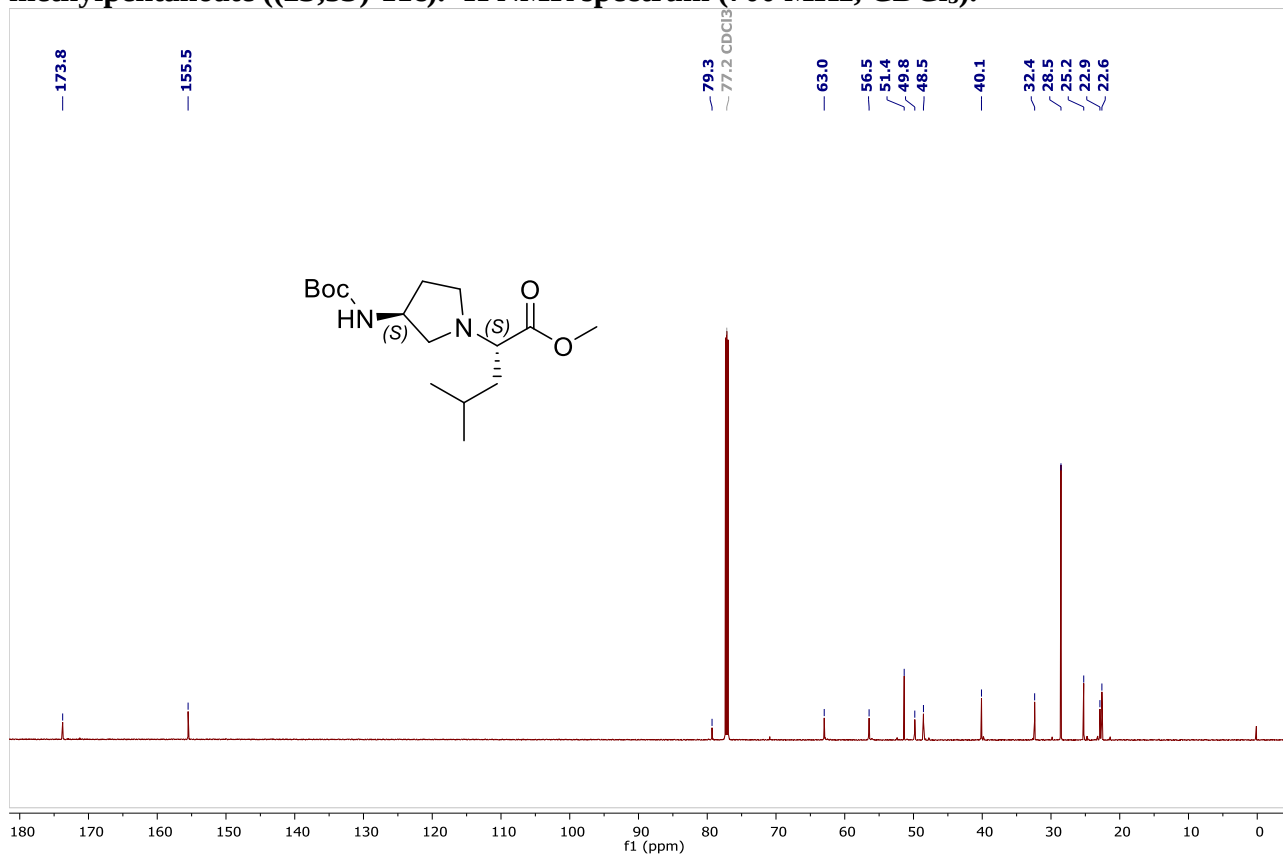


Figure S130. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)-4-methylpentanoate ((2S,3S)-11c). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

Analysis Info

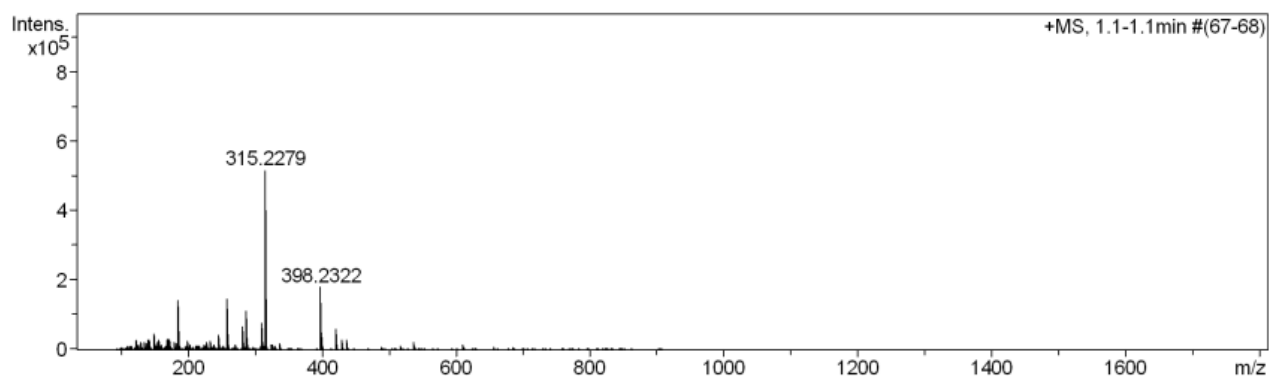
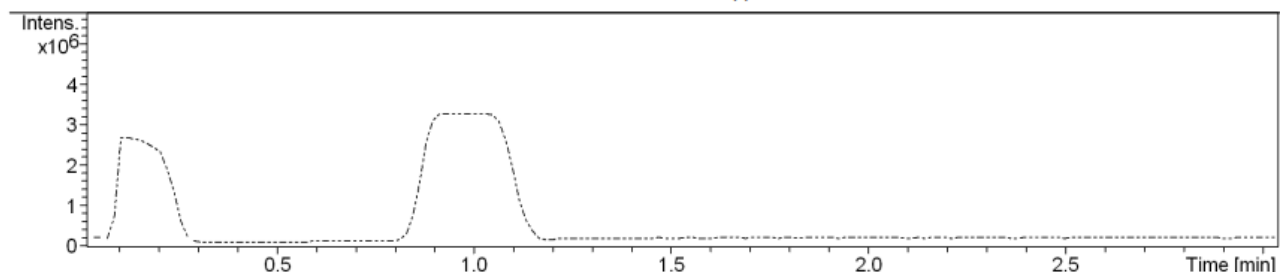
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Acquisition Date 4/21/2023 2:36:49 PM

Operator Milda Pukalskiene
 Instrument / Ser# maXis 4G 20218

Acquisition Parameter

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Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
315.2279	1	C ₁₆ H ₃₁ N ₂ O ₄	100.00	315.2278	-0.3	0.1	16.0	2.5	even	ok
	2	C ₁₇ H ₂₇ N ₆	41.89	315.2292	4.0	4.3	28.5	7.5	even	ok

Figure S131. Methyl (2S)-2-((3S)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl)-4-methylpentanoate ((2S,3S)-11c). HRMS (ESI-TOF).

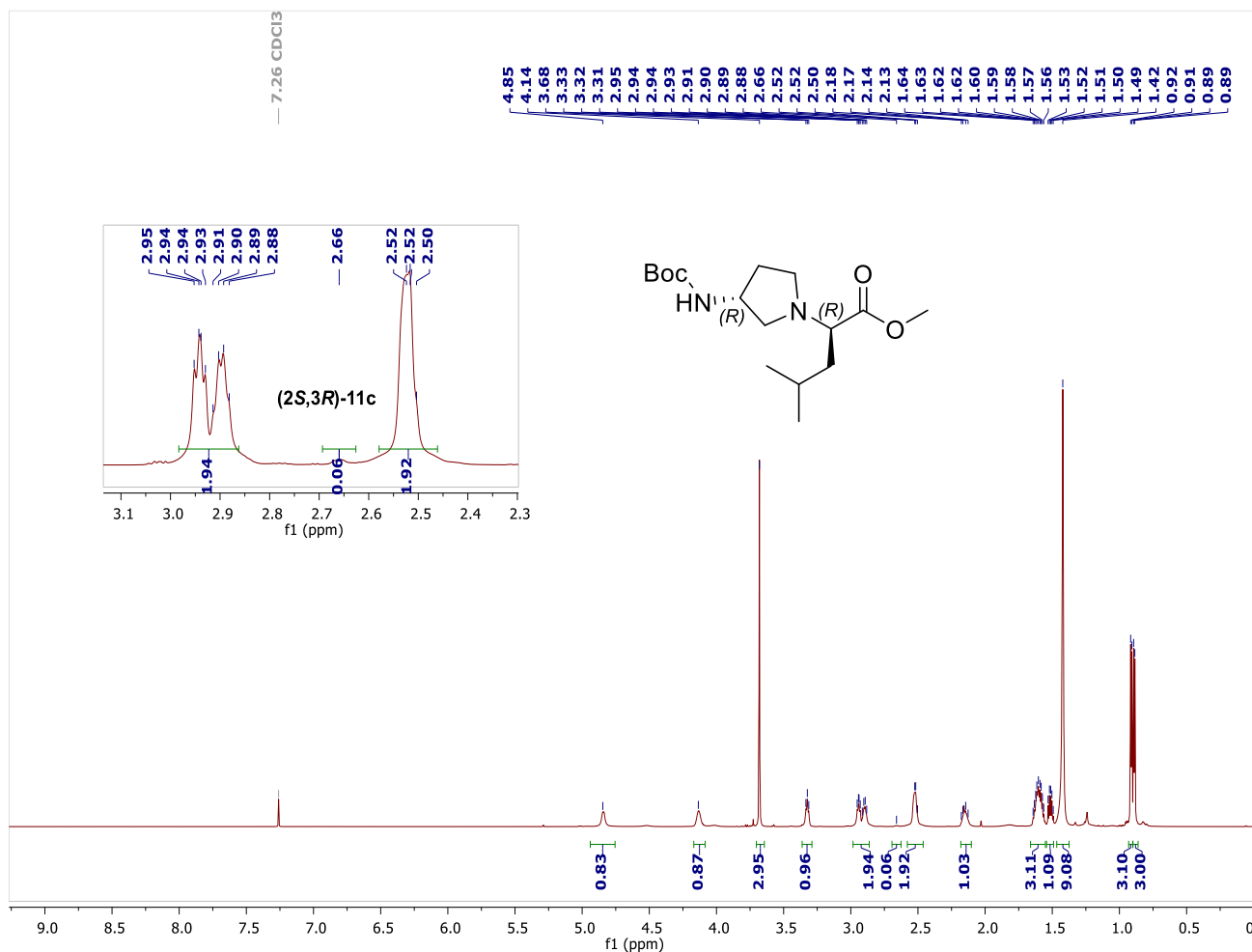


Figure S132. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]-4-methylpentanoate ((2*R*,3*R*)-11c). ¹H NMR spectrum (700 MHz, CDCl₃).

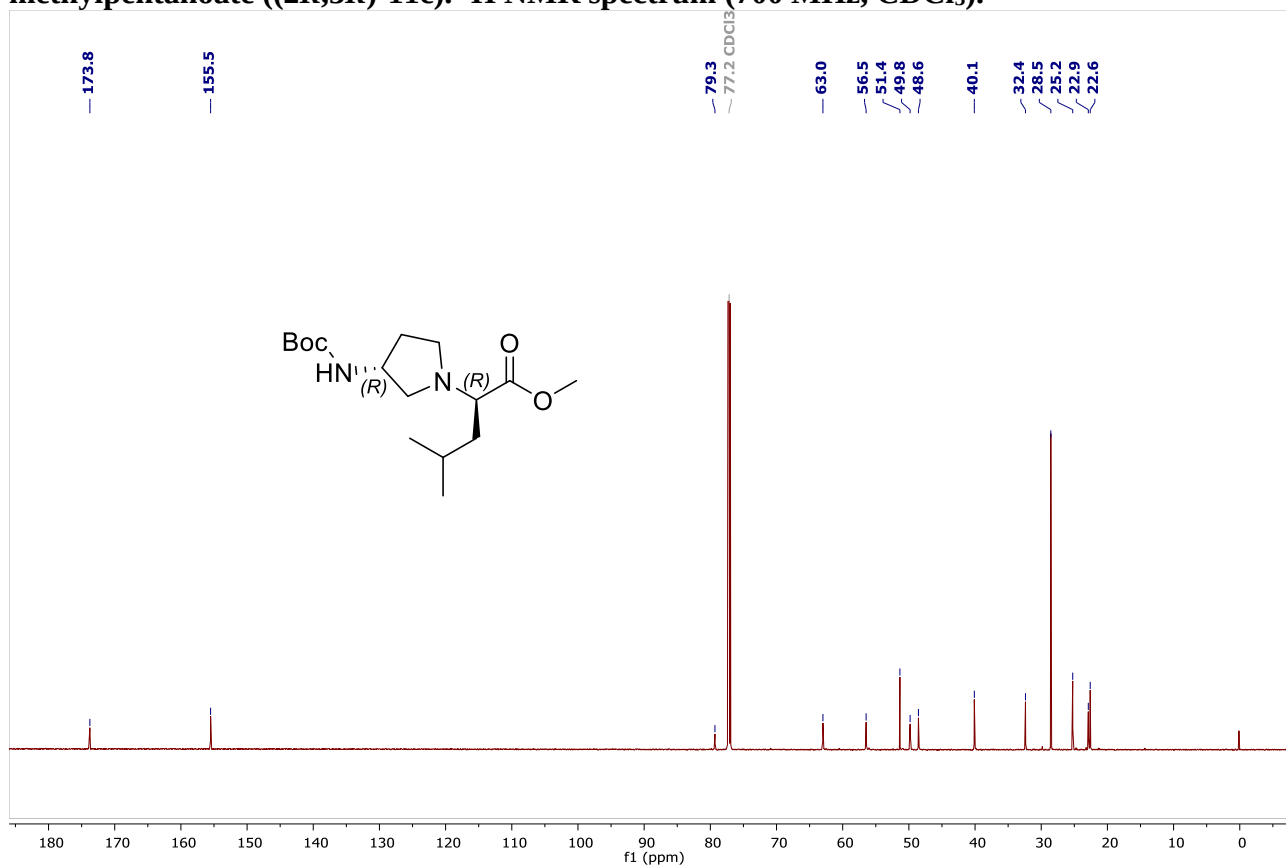


Figure S133. Methyl (2*R*)-2-[(3*R*)-3-[(*tert*-butoxycarbonyl)amino]pyrrolidin-1-yl]-4-methylpentanoate ((2*R*,3*R*)-11c). ¹³C NMR spectrum (176 MHz, CDCl₃).

Mass Spectrum SmartFormula Report

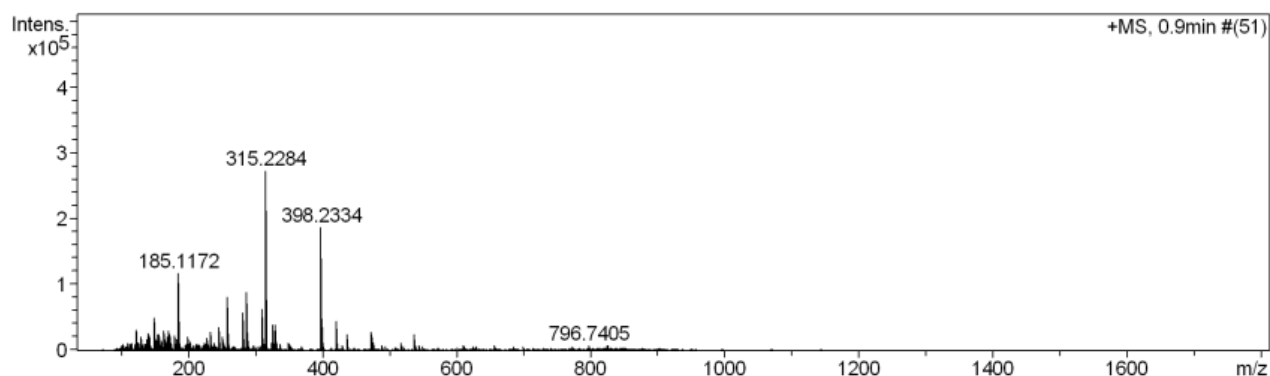
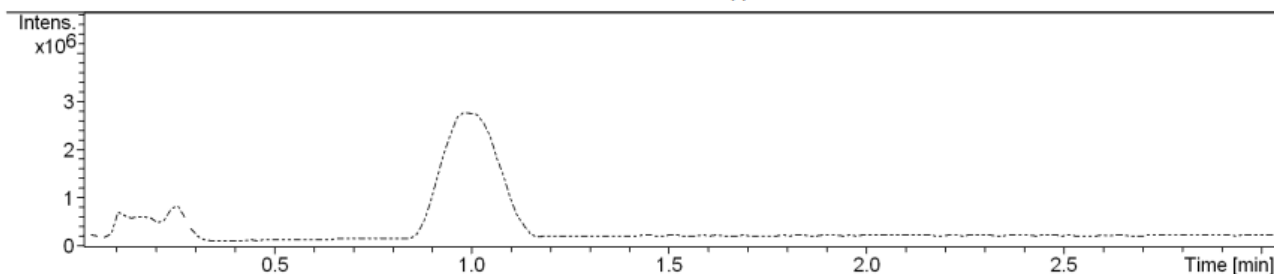
Analysis Info

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Comment

Acquisition Date 4/27/2023 3:35:22 PM
Operator Milda Pukalskiene
Instrument / Ser# maXis 4G 20218

Acquisition Parameter

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Scan Begin	40 m/z	Set End Plate Offset	-500 V	Set Dry Gas	8.0 l/min
Scan End	1800 m/z	Set Collision Cell RF	350.0 Vpp	Set Divert Valve	Waste



Meas. m/z	#	Formula	Score	m/z	err [ppm]	Mean err [ppm]	mSigma	rdb	e ⁻ Conf	N-Rule
315.2284	1	C ₁₆ H ₃₁ N ₂ O ₄	100.00	315.2278	-1.8	-1.7	19.8	2.5	even	ok

Figure S134. Methyl (2R)-2-((3R)-3-((tert-butoxycarbonyl)amino)pyrrolidin-1-yl)-4-methylpentanoate ((2R,3R)-11c). HRMS (ESI-TOF).

- **X-ray analysis of compound (S)-3b:**

Single crystals of $C_{32}H_{60}N_4O_8$ [(S)-3b] were investigated on a Rigaku, XtaLAB Synergy, Dualflex, HyPix diffractometer. The crystal was kept at 150.0(1) K during data collection. Using Olex2 [1], the structure was solved with the ShelXT [2] structure solution program using Intrinsic Phasing and refined with the olex2.refine [3] refinement package using Gauss-Newton minimisation.

Table S1: Experimental parameter and CCDC-2168602.

Sample	Machine	Source	Temp.	Detector distance	Time/Frame	#Frames	Frame width	CCDC
			[K]	[mm]	[s]		[°]	
(S)-3b	Rigaku, XtaLAB Synergy, Dualflex, HyPix	$\mu(\text{Cu K}\alpha) = 0.669\text{mm}^{-1}$	150	34	1.25	5748	0.50	2168602

Table S2: Sample and crystal data of compound (S)-3b.

Chemical formula	$2(C_{16}H_{30}N_2O_4)$	Crystal system	Monoclinic	
Formula weight [g/mol]	628.86	Space group	$P2_1$	
Temperature [K]	150	Z	2	
Measurement method	ϕ and ω scans	Volume [$\text{\AA}^3$]	1805.08(5)	
Radiation wavelength [$\text{\AA}$]	1.54184	Unit cell dimensions [$\text{\AA}^3$ and $^\circ$]	10.0924(2)	90
			12.8242(2)	97.425(2)
			14.0647(2)	90
Crystal size/ [mm^3]	$0.17 \times 0.11 \times 0.08$			
Crystal habit	Block, colourless			
Density (calculated)/ [g/cm^3]	1.1569	Absorption coefficient [mm^{-1}]	0.669	
Abs. Correction $T_{\min}$	0.859	Abs. Correction $T_{\max}$	0.961	
Abs. Correction type	multi-scan <i>CrysAlis PRO</i> 1.171.40.35a (Rigaku Oxford Diffraction, 2018) Empirical absorption correction using spherical harmonics, implemented in SCALE3 ABSPACK scaling algorithm.	F(000) [e$^-$]	688	

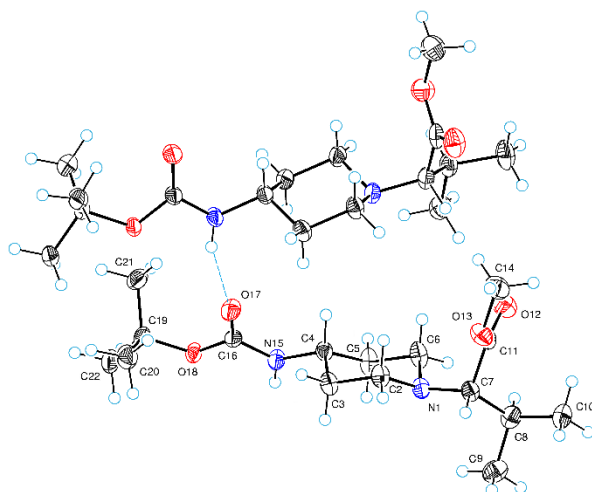


Figure S135. ORTEP view of assymetric unit of crystal **(S)-3b**.

Table S3: Data collection and structure refinement of compound **(S)-3b**.

Index ranges	-12 ≤ h ≤ 12 -16 ≤ k ≤ 13 -17 ≤ l ≤ 17	2θ range for data collection [°]	155.0	
Reflections numbers	37496	Data / restraints / parameters	6342/1/417	
Refinement method	Least squares matrix: full	Final R indices	All data	R1 = 0.0316; wR2 = 0.0840
Function minimized	$\Sigma w[F_o ^2 - (1/k) F_c ^2]$		I > 2σ(I)	R1 = 0.0309; wR2 = 0.0835
Goodness-of-fit on F²	1.052	Weighting scheme	$w = 1/[\sigma^2(F_o^2) + (0.052P)^2 + 0.1876P]$ where $P = (F_o^2 + 2F_c^2)/3$	
Largest diff. peak and hole [e Å⁻³]	+0.20 and -0.12	Flack's x parameter	0.04(9)	

- **X-ray analysis of compound (R)-3b:**

Single crystals of $C_{32}H_{60}N_4O_8$ [(R)-3b] were investigated on a Rigaku, XtaLAB Synergy, Dualflex, HyPix diffractometer. The crystal was kept at 150.0(1) K during data collection. Using Olex2 [1], the structure was solved with the ShelXT [2] structure solution program using Intrinsic Phasing and refined with the olex2.refine [3] refinement package using Gauss-Newton minimisation.

Table S4: Experimental parameter and CCDC-2168596

Sample	Machine	Source	Temp.	Detector distance	Time/Frame	#Frames	Frame width	CCDC
			[K]	[mm]	[s]		[°]	
(R)-3b	Rigaku, XtaLAB Synergy, Dualflex, HyPix	$\mu(\text{Cu K}\alpha) = 0.669 \text{ mm}^{-1}$	150	34	1.25	5748	0.50	2168596

Table S5: Sample and crystal data of compound (R)-3b.

Chemical formula	$2(C_{16}H_{30}N_2O_4)$	Crystal system	Monoclinic	
Formula weight [g/mol]	628.86	Space group	$P2_1$	
Temperature [K]	150	Z	2	
Measurement method	ϕ and ω scans	Volume [$\text{\AA}^3$]	1802.25(4)	
Radiation wavelength [$\text{\AA}$]	1.54184	Unit cell dimensions [$\text{\AA}^3$ and $^\circ$]	10.0869(1) 12.8088(2) 14.0693(2)	90 97.4889(12) 90
Crystal size/ [mm^3]	$0.18 \times 0.11 \times 0.08$			
Crystal habit	Colourless block			
Density (calculated)/ [g/cm^3]	1.1587	Absorption coefficient [mm^{-1}]	0.670	
Abs. Correction $T_{\min}$	0.872	Abs. Correction $T_{\max}$	0.960	
Abs. Correction type	multi-scan <i>CrysAlis PRO</i> 1.171.40.35a (Rigaku Oxford Diffraction, 2018) Empirical absorption correction using spherical harmonics, implemented in SCALE3 ABSPACK scaling algorithm.	F(000) [e^-]	688	

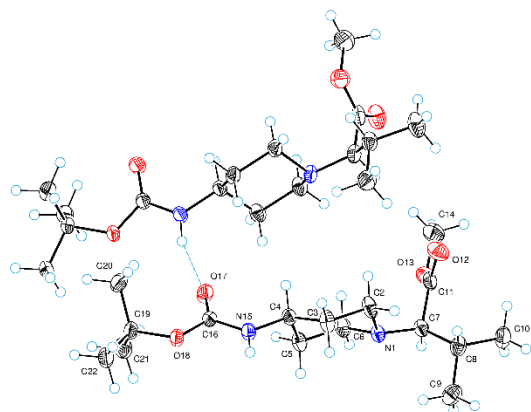


Figure S136. ORTEP view of assymetric unit of crystal **(R)-3b**.

Table S6: Data collection and structure refinement of compound **(R)-3b**.

Index ranges	-12 ≤ h ≤ 9 -16 ≤ k ≤ 16 -17 ≤ l ≤ 17	2θ range for data collection [°]	155.0	
Reflections numbers	22287	Data / restraints / parameters	7121/1/417	
Refinement method	Least squares matrix: full	Final R indices	All data	R1 = 0.0314; wR2 = 0.0818
Function minimized	$\Sigma w[F_o ^2 - (1/k) F_c ^2]$		I > 2σ(I)	R1 = 0.0306; wR2 = 0.0804
Goodness-of-fit on F2	1.051	Weighting scheme	$w = 1/[\sigma^2(F_o^2) + (0.0P)^2 + 0.2072P]$ where $P = (F_o^2 + 2F_c^2)/3$	
Largest diff. peak and hole [e Å⁻³]	+0.14 and -0.19	Flack's x parameter	0.06(8)	

- **Examples of unsuccessful HPLC analysis**

Analysis were performed using Shimadzu LC2030C chromatograph with chiral columns and water + 0.1% formic acid / acetonitrile (various gradient conditions).

Table S7. Experimental parameter.

Mobile phase	A – Acetonitrile B – Water + 0.1% formic acid
Temperature	36 °C
Flow rate	1 mL/min
Detection	UV (254, 210 nm); ELSD (Evaporative light scattering detector)
Injection	10 µL
Sample	Dissolved in methanol

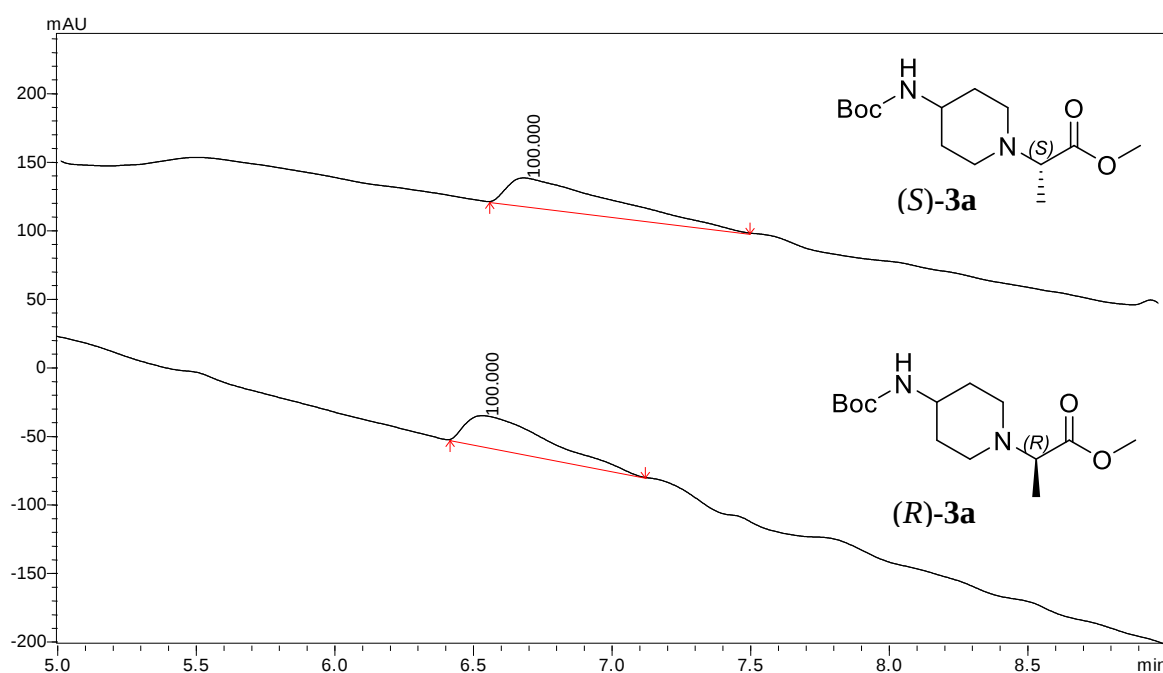


Figure S137. Unsuccessful chiral HPLC analysis of compounds (S)-3a and (R)-3a. Separation was carried out on YMC chiral column CHIRAL NEA (R), gradient conditions water + 0.1% formic acid / acetonitrile 90:10 to 20:80 in 15 minutes. UV detector, 210 nm.

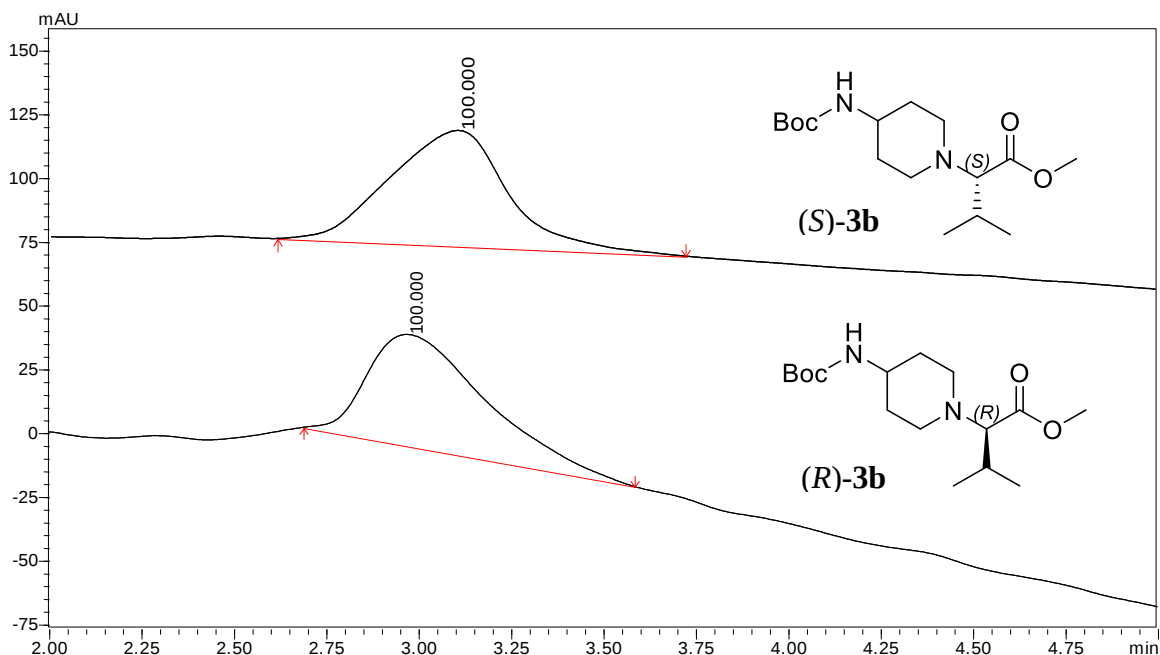


Figure S138. Unsuccessful chiral HPLC analysis of compounds (S)-3b and (R)-3b. Separation was carried out on chiral column CHIRAL ART Cellulose-SB, gradient conditions: water + 0.1% formic acid / acetonitrile 90:10 to 60:40 in 12 minutes. UV detector, 210 nm.

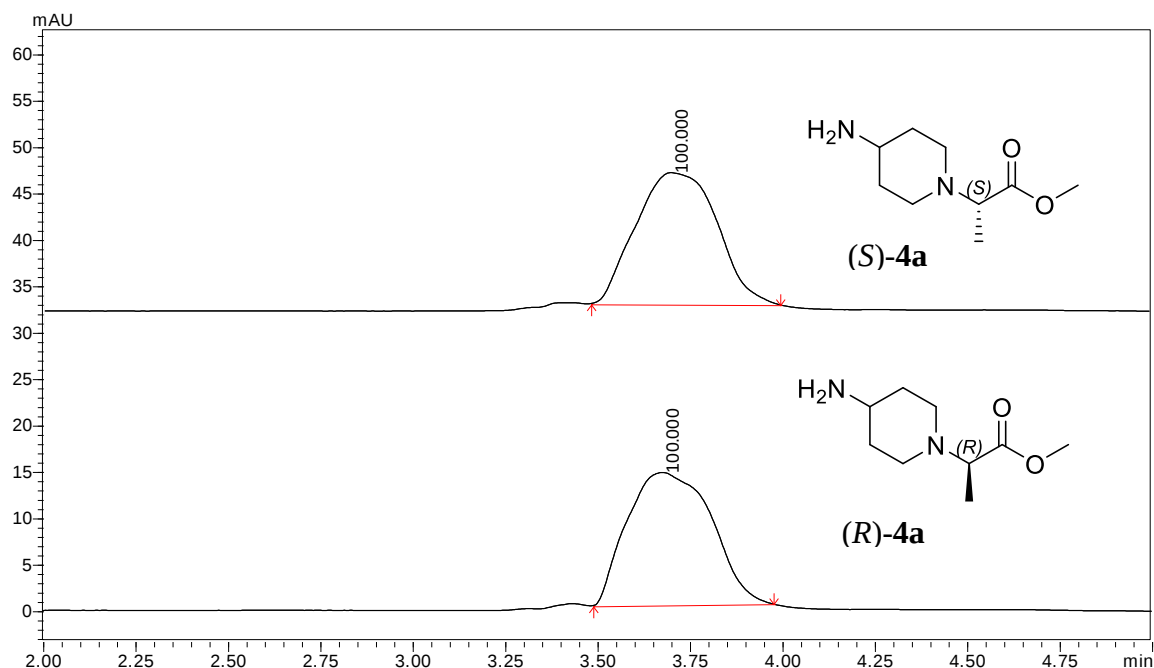


Figure S139. Unsuccessful chiral HPLC analysis of compounds (S)-4a and (R)-4a. Separation was carried out on YMC chiral column CHIRAL NEA (R), gradient conditions: water + 0.1% formic acid / acetonitrile 90:10 to 20:80 in 15 minutes. UV detector, 254 nm.

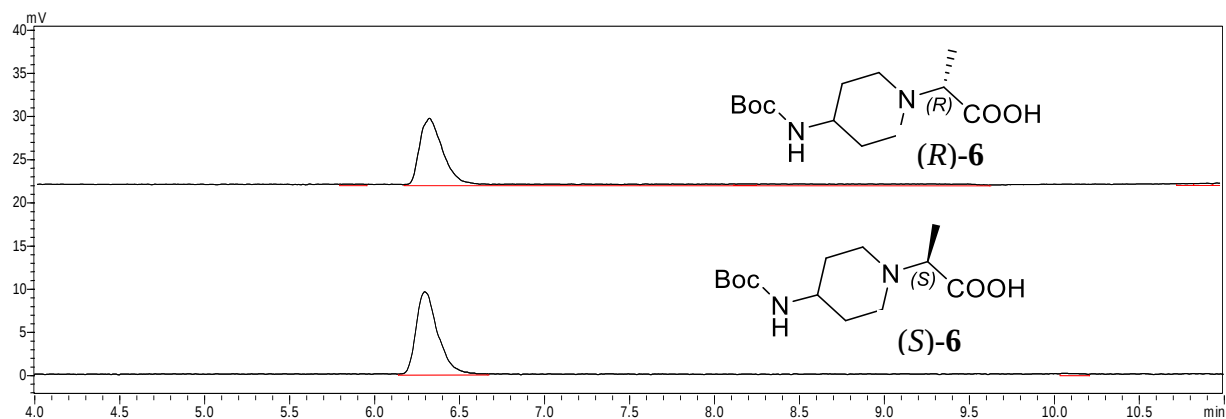


Figure S140. Unsuccessful chiral HPLC analysis of compounds (S)-6 and (R)-6. Separation was carried out on YMC chiral column CHIRAL NEA (R), gradient conditions water + 0.1% formic acid / acetonitrile 90:10 to 20:80 in 15 minutes. Evaporative light scattering detector.

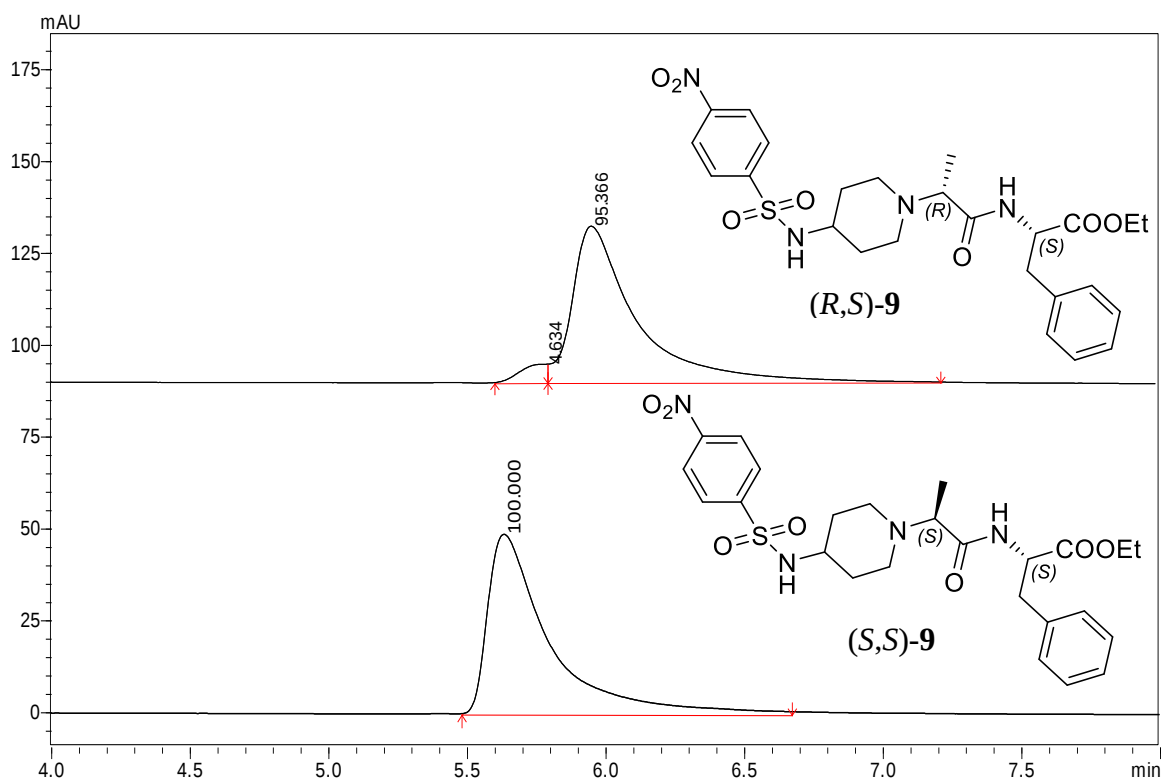


Figure S141. Unsuccessful chiral HPLC analysis of compounds (S,S)-9 and (R,S)-9. Separation was carried out on chiral column CHIRAL ART Cellulose-SB, gradient conditions: water + 0.1% formic acid / acetonitrile 85:15 to 60:40 in 10 minutes. UV detector, 254 nm wavelength.

- **References:**

1. Dolomanov, O.V., Bourhis, L.J., Gildea, R.J, Howard, J.A.K. & Puschmann, H. (2009), J. Appl. Cryst. 42, 339-341.
2. Sheldrick, G.M. (2015). Acta Cryst. A71, 3-8.
3. Bourhis, L.J., Dolomanov, O.V., Gildea, R.J., Howard, J.A.K., Puschmann, H. (2015). Acta Cryst. A71, 59-75.