

---- *Supporting Information* ----

Betti Base Derived P-Stereogenic Phosphine-Diamidophosphite Ligands with a Single Atom Spacer and their Application in Asymmetric Catalysis

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▪ General procedure for the synthesis of acetamidoacrylates

The acetamidoacrylate derivatives were synthesized following the literature procedure from the corresponding aromatic aldehyde and N-acetyl glycine.¹ The analytical data match with the previously reported data.

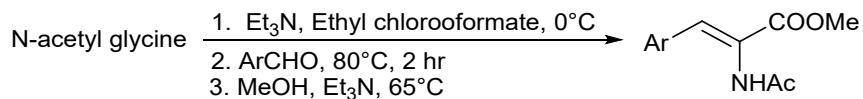


Figure S 1 General procedure for the synthesis of dehydroamino acid derivatives

▪ General procedure for the upscaling

The asymmetric hydrogenation upscaling experiments were performed in a stainless-steel autoclave charged with an insert suitable for up to 5 reaction vessels (10 mL each) with teflon mini stirring bars. In a typical experiment, a reaction vessel is charged with [Rh(cod)₂]BF₄ (1 mol%) and ligand (M/L=1/1.1) and stirred for 10-15 mins in the appropriate solvent. The desired substrates (3.0 mmol) were added to the reaction vessel maintaining the inert atmosphere and the vessels were placed in high pressure autoclave. The autoclave was purged two times with nitrogen and three times with hydrogen. Finally, it was pressurized at the 10 bar of H₂ pressure at 25 ° C for 14 hrs. After the desired reaction time, the autoclave was depressurized, and the reaction mixture was diluted with EtOAc and filtered through a short pad of silica. The conversion was determined by GC and GC-MS measurement and the enantiomeric excess was measured by chiral HPLC.

■ NMR Spectra of the ligands

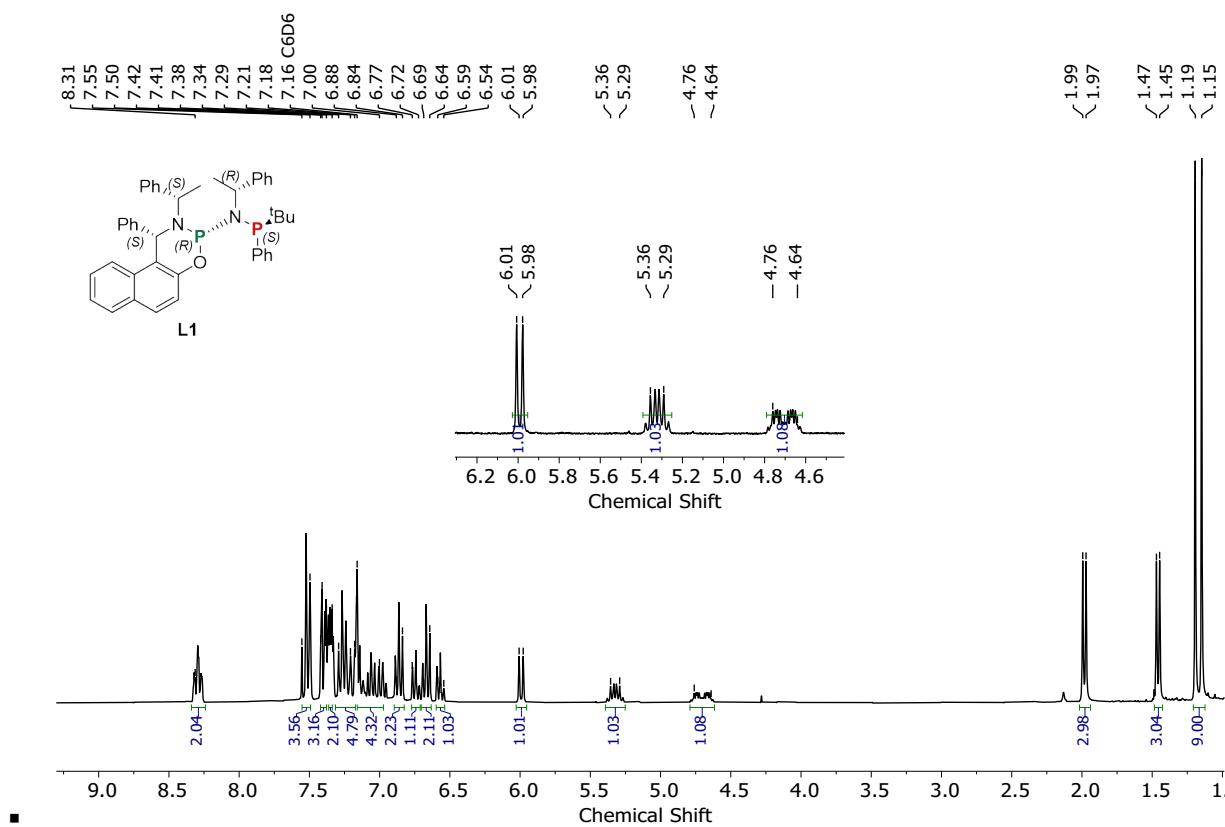


Figure S 2 1H NMR of (*S,S,R_P,R,S_P*)-L1

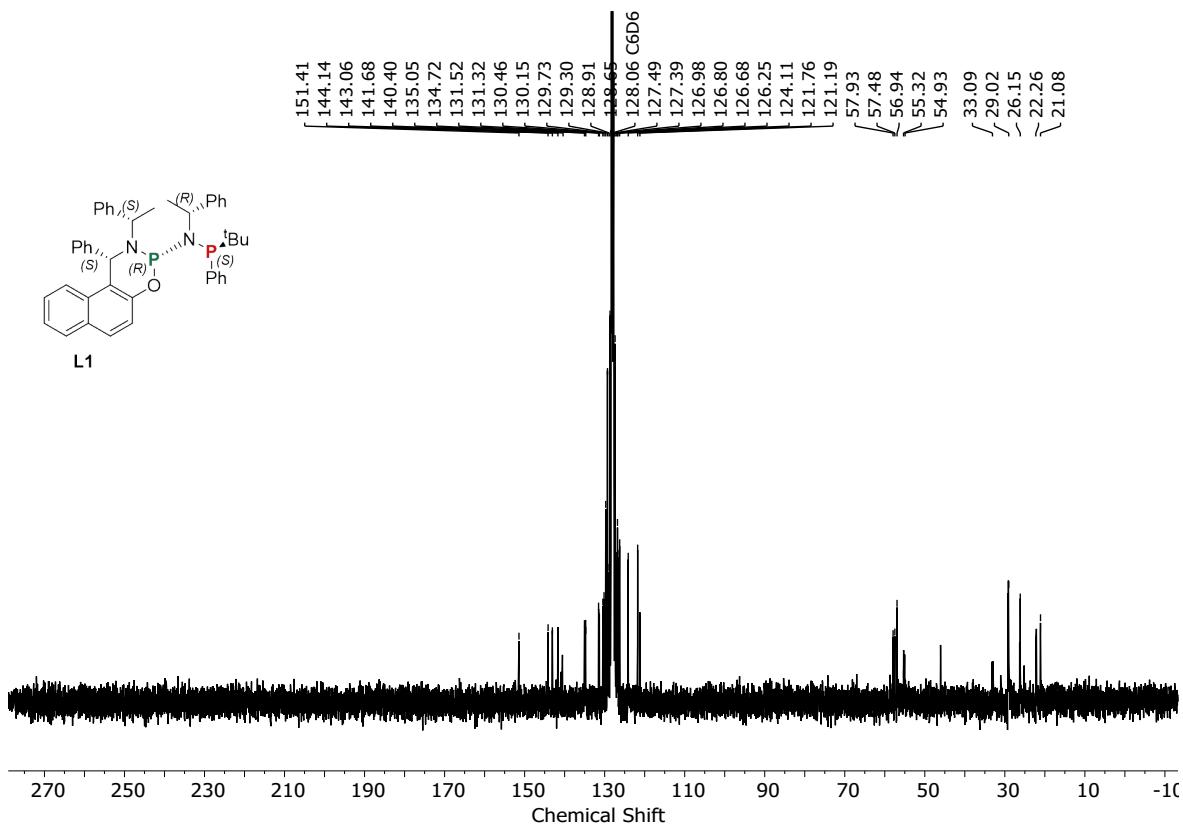


Figure S 3 L1 ^{13}C NMR of (*S,S,R_P,R,S_P*)-L1

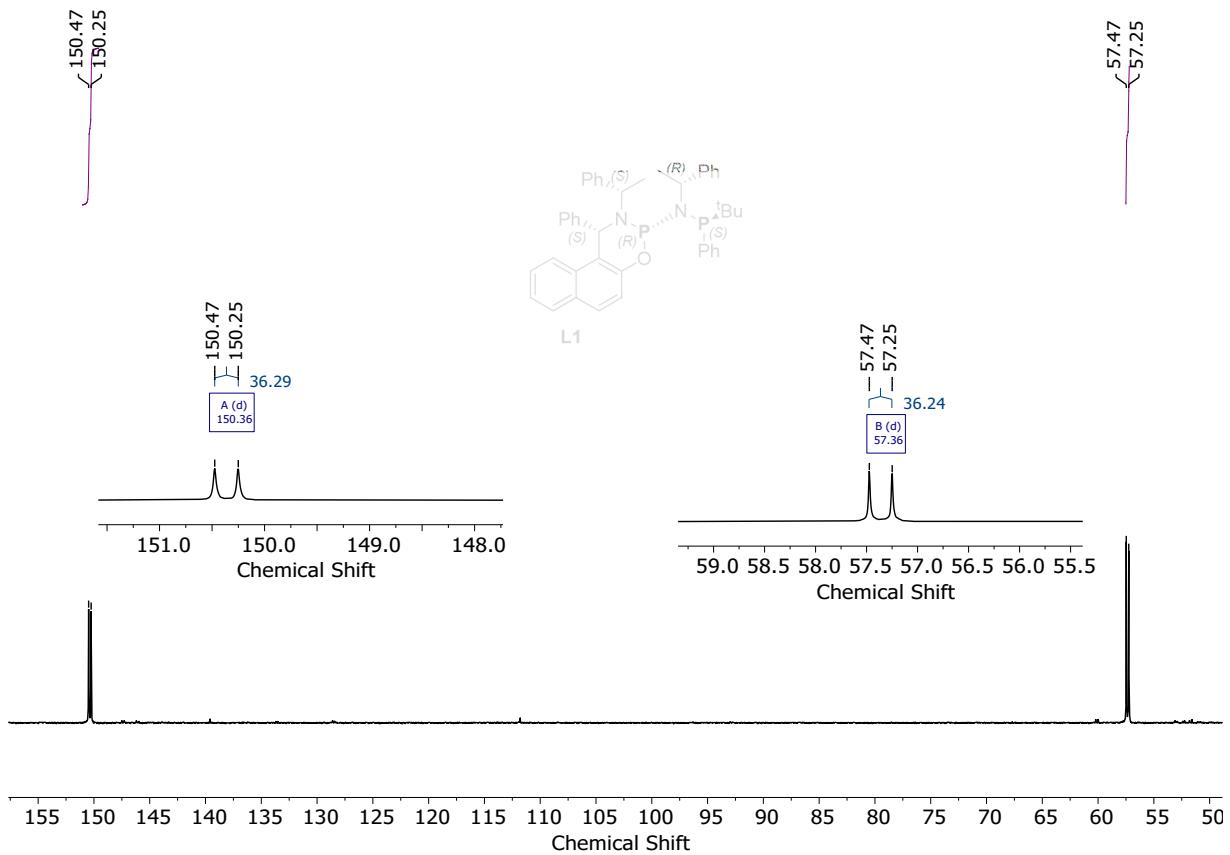


Figure S 4 ^{31}P { H } NMR of (S,S,R_P,R_S)- L1

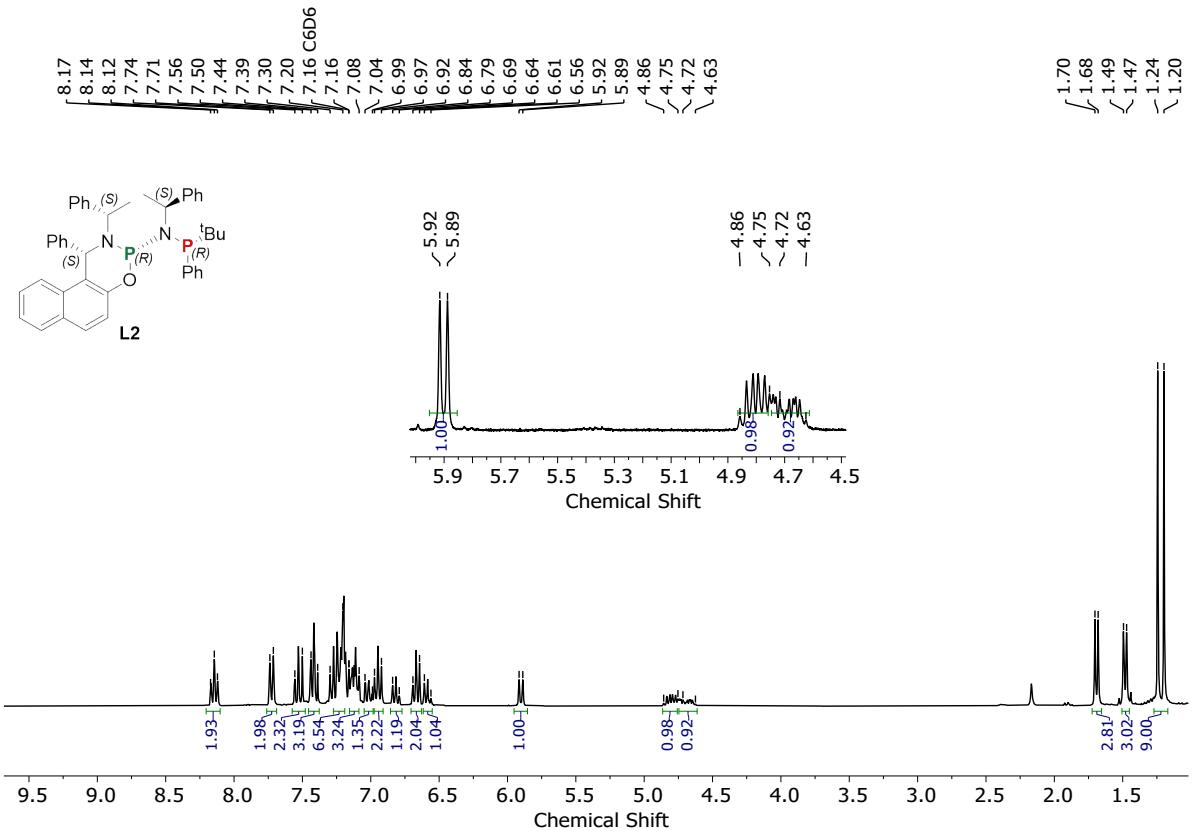


Figure S 5 ^1H NMR of (S,S,R_P,S_R)- L2

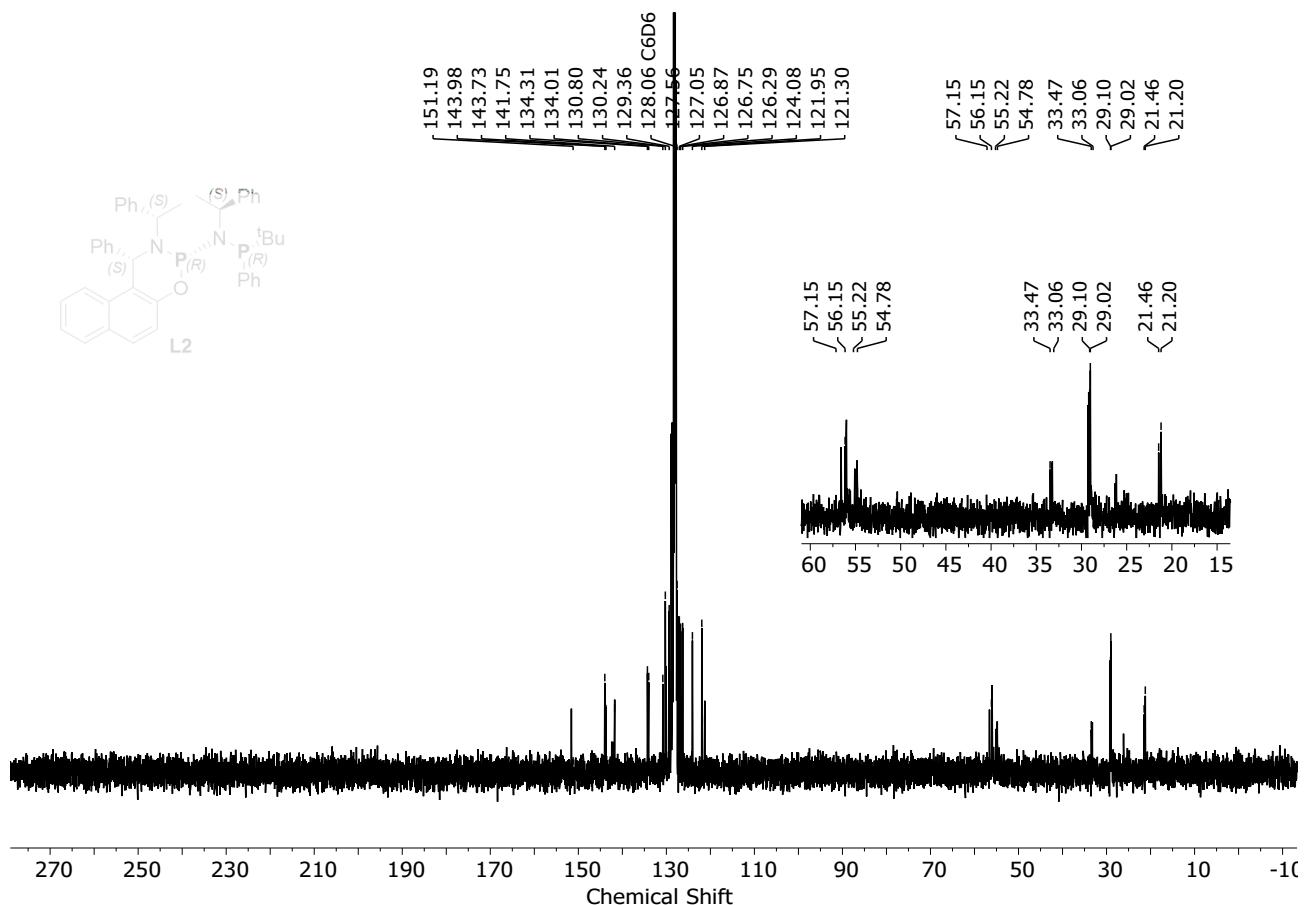


Figure S 6 ^{13}C NMR of $(\text{S},\text{S},\text{R}_P,\text{S},\text{R}_P)$ - L2

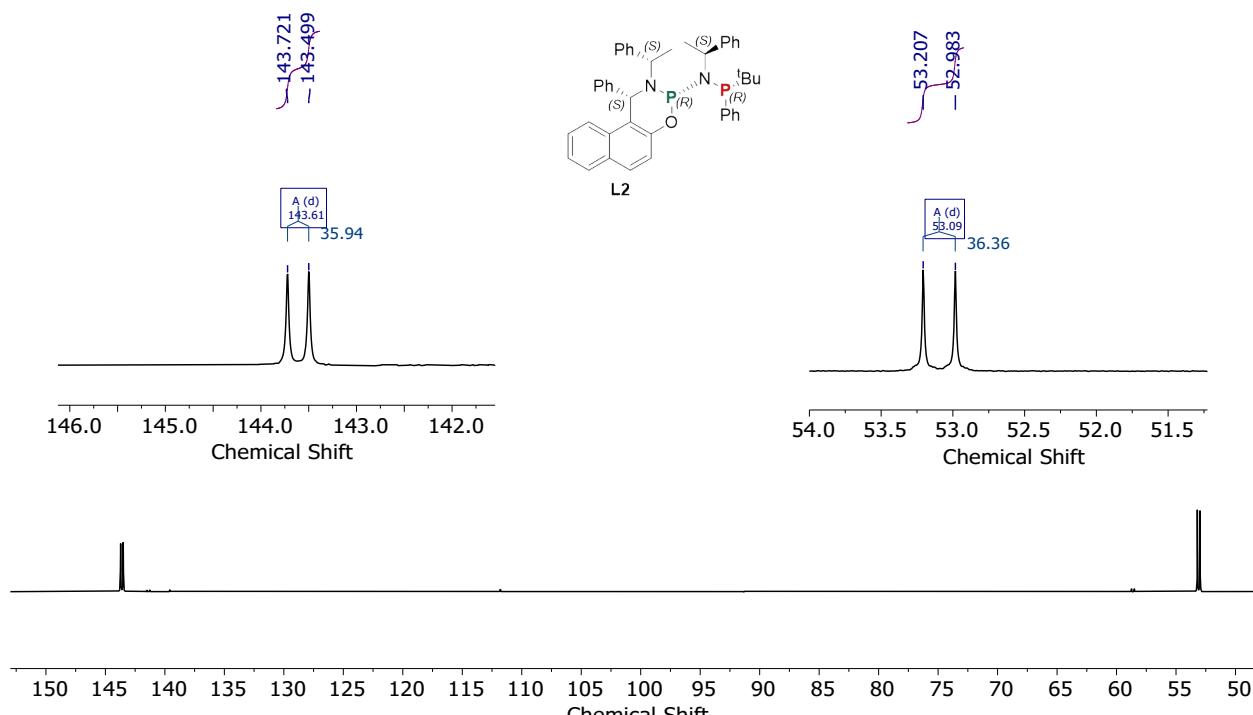


Figure S 7 ^{31}P { H } NMR of $(\text{S},\text{S},\text{R}_P,\text{S},\text{R}_P)$ - L2

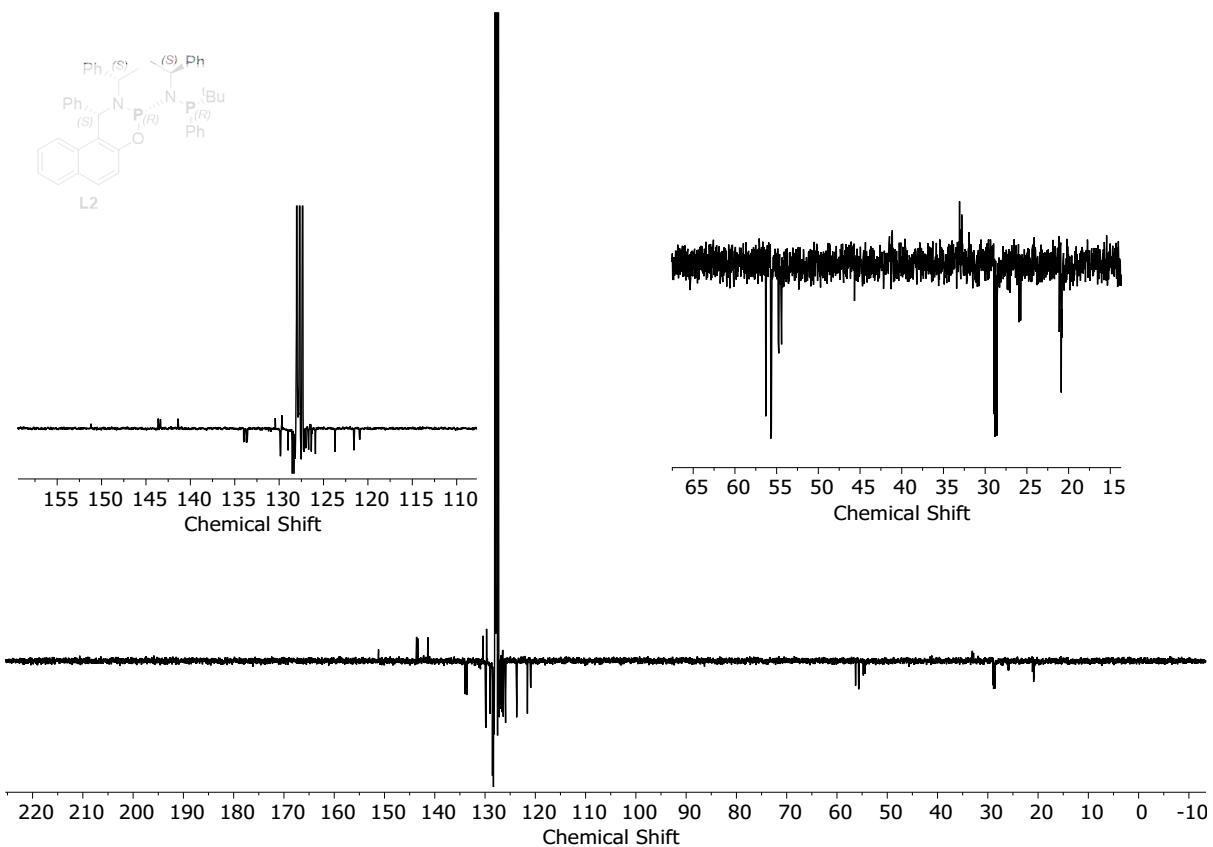


Figure S 8 APT of (*S,S,R_P,S,R_P*)-L2

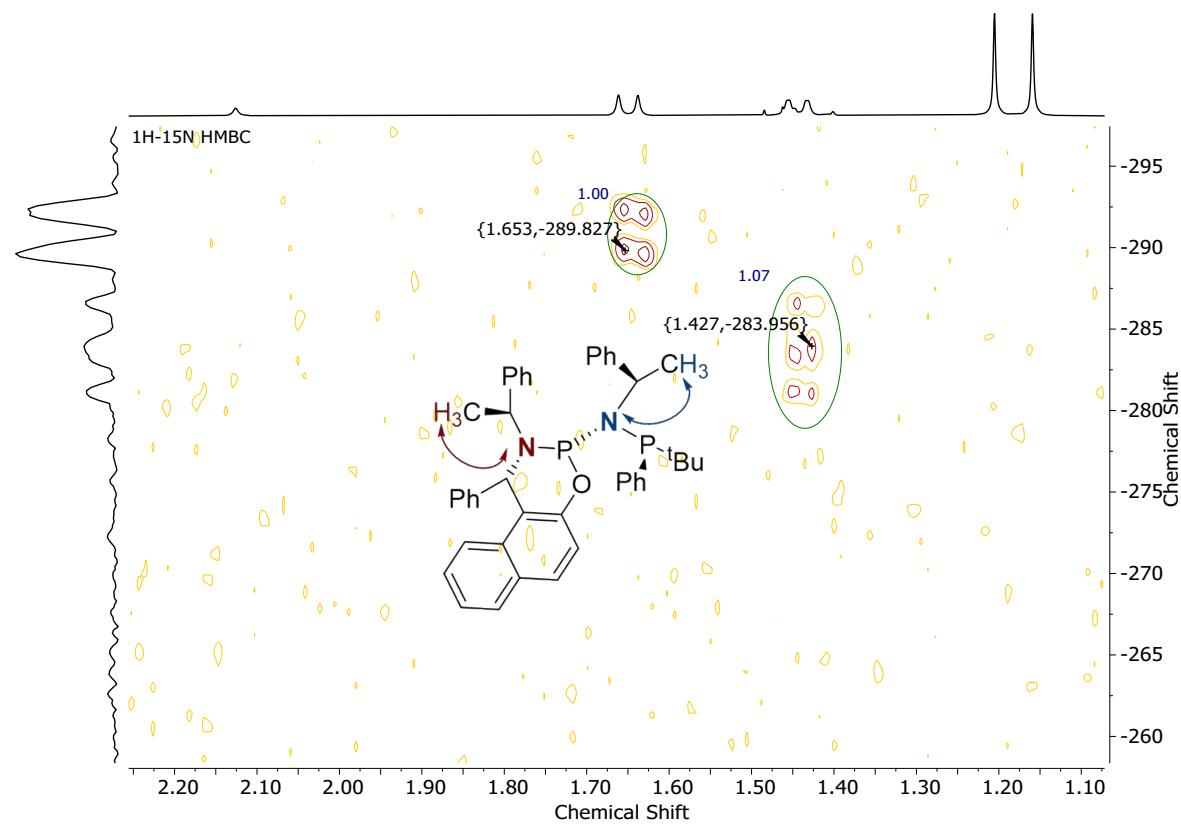


Figure S 9 ¹H-¹⁵N-HMBC of (*S,S,R_P,S,R_P*)-L2

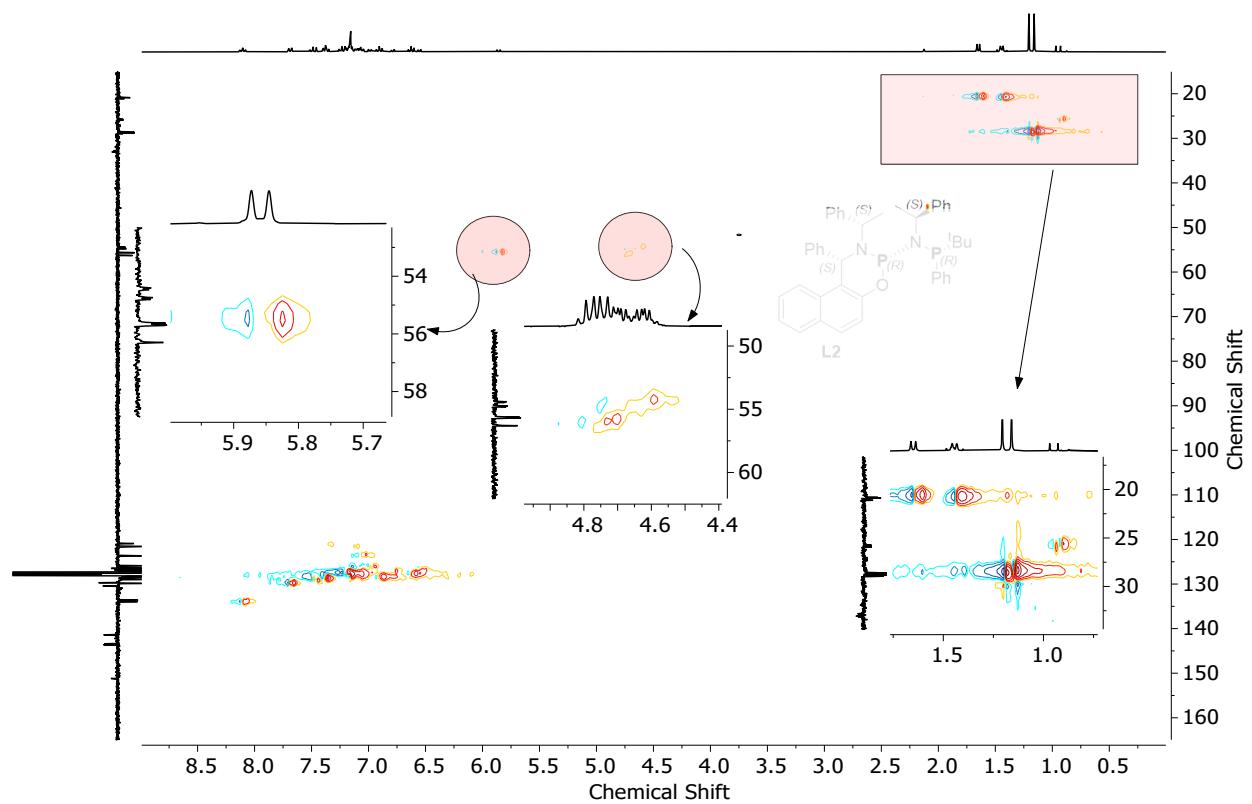


Figure S 10 ^1H - ^{13}C -HSQC of $(S,S,R_p,S,R_p)\text{-L2}$

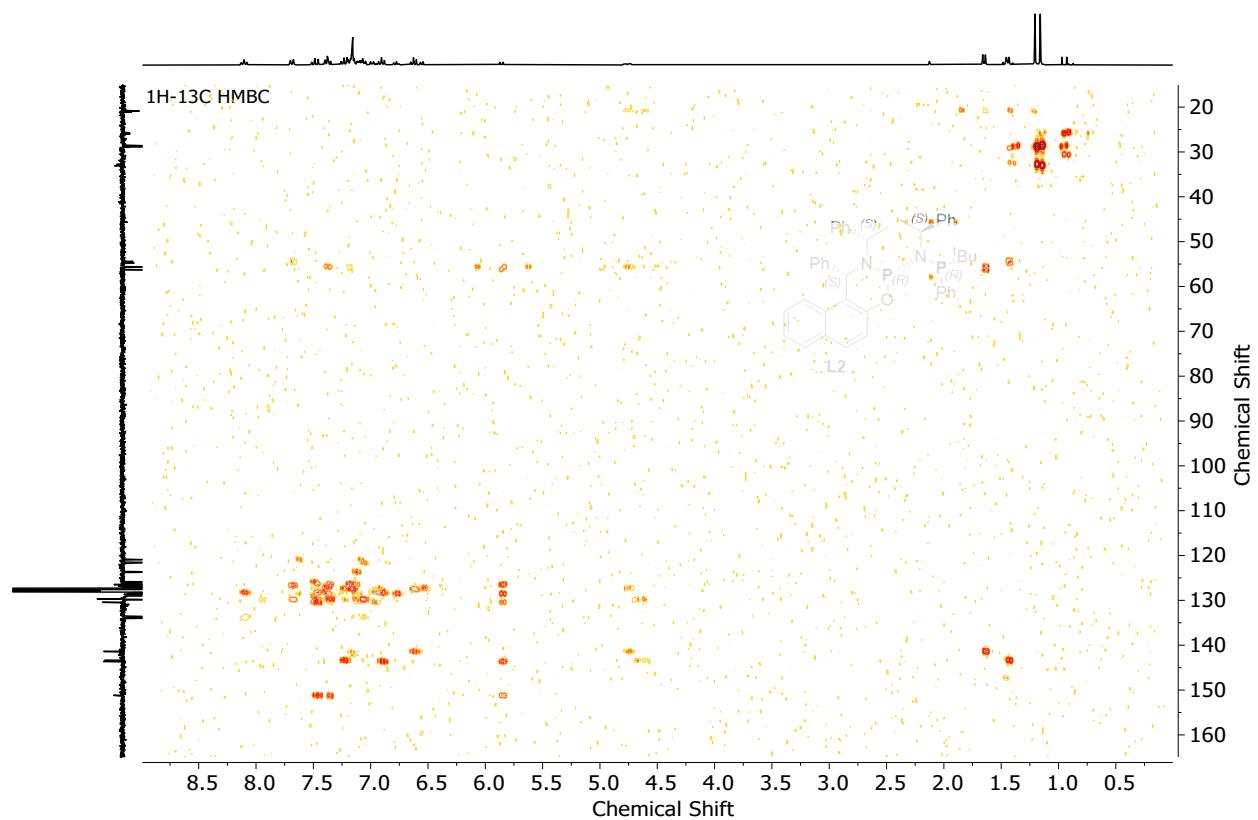


Figure S 11 ^1H - ^{13}C -HMBC of $(S,S,R_p,S,R_p)\text{-L2}$

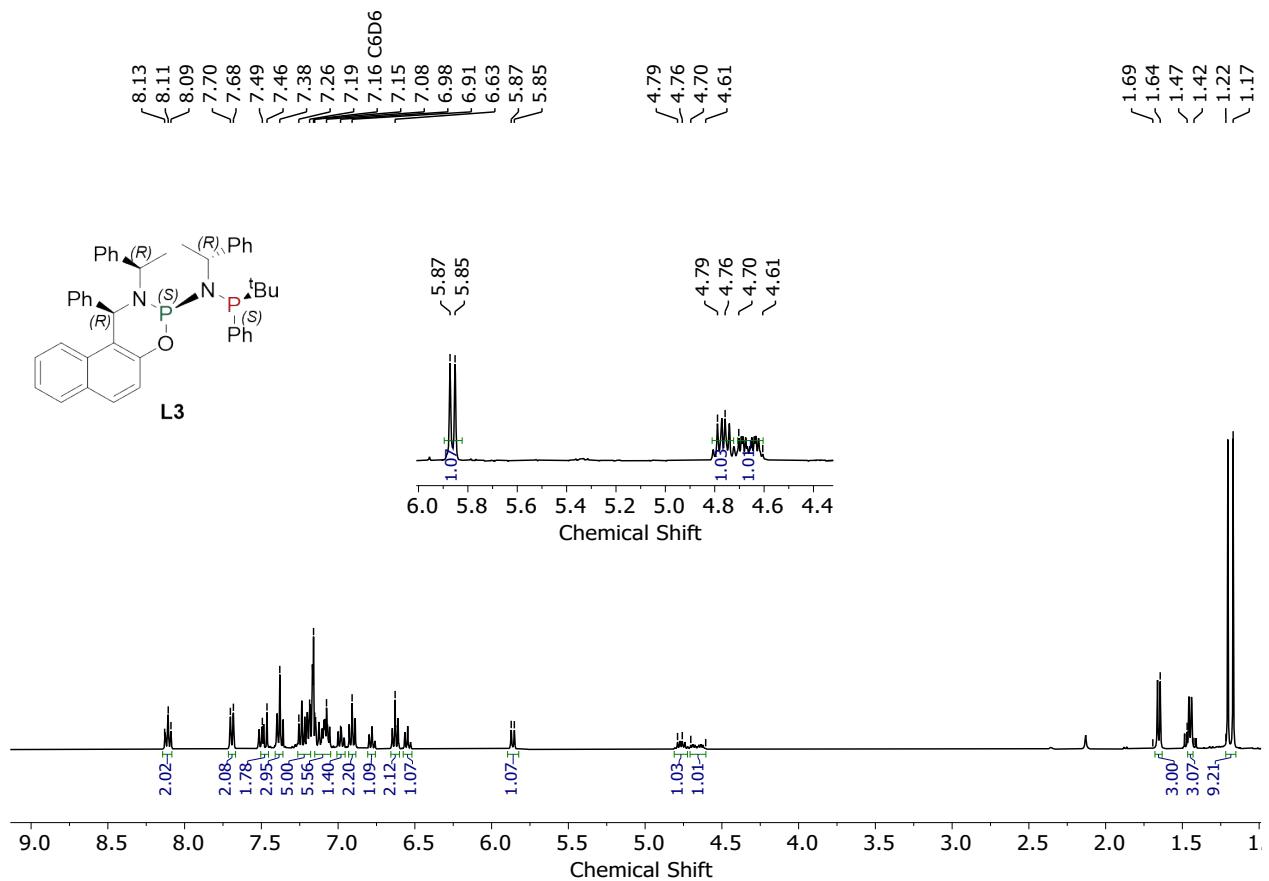


Figure S 12 ^1H NMR of ($\text{R},\text{R},\text{S}_P,\text{R},\text{S}_P$)- L3

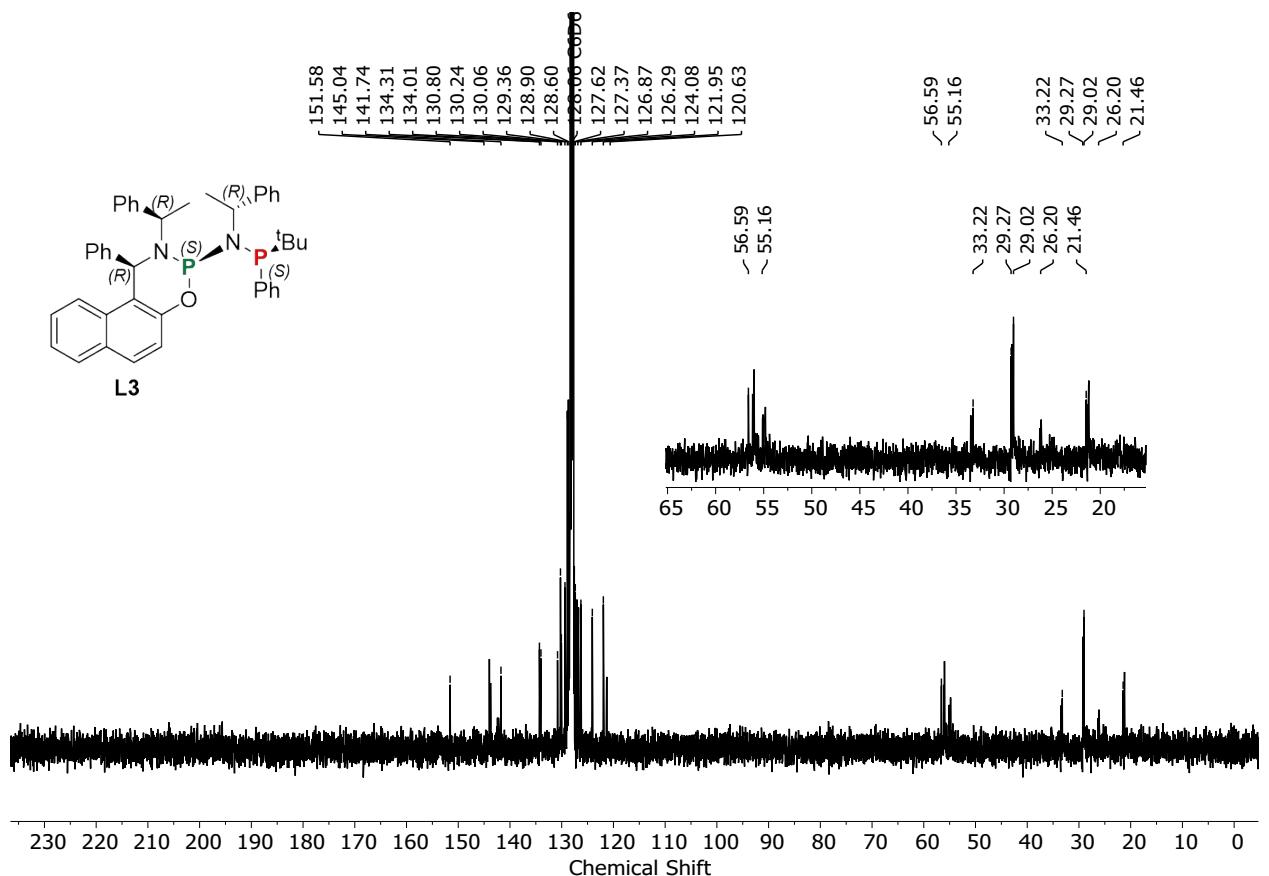


Figure S 13 ^{13}C NMR of ($\text{R},\text{R},\text{S}_P,\text{R},\text{S}_P$)- L3

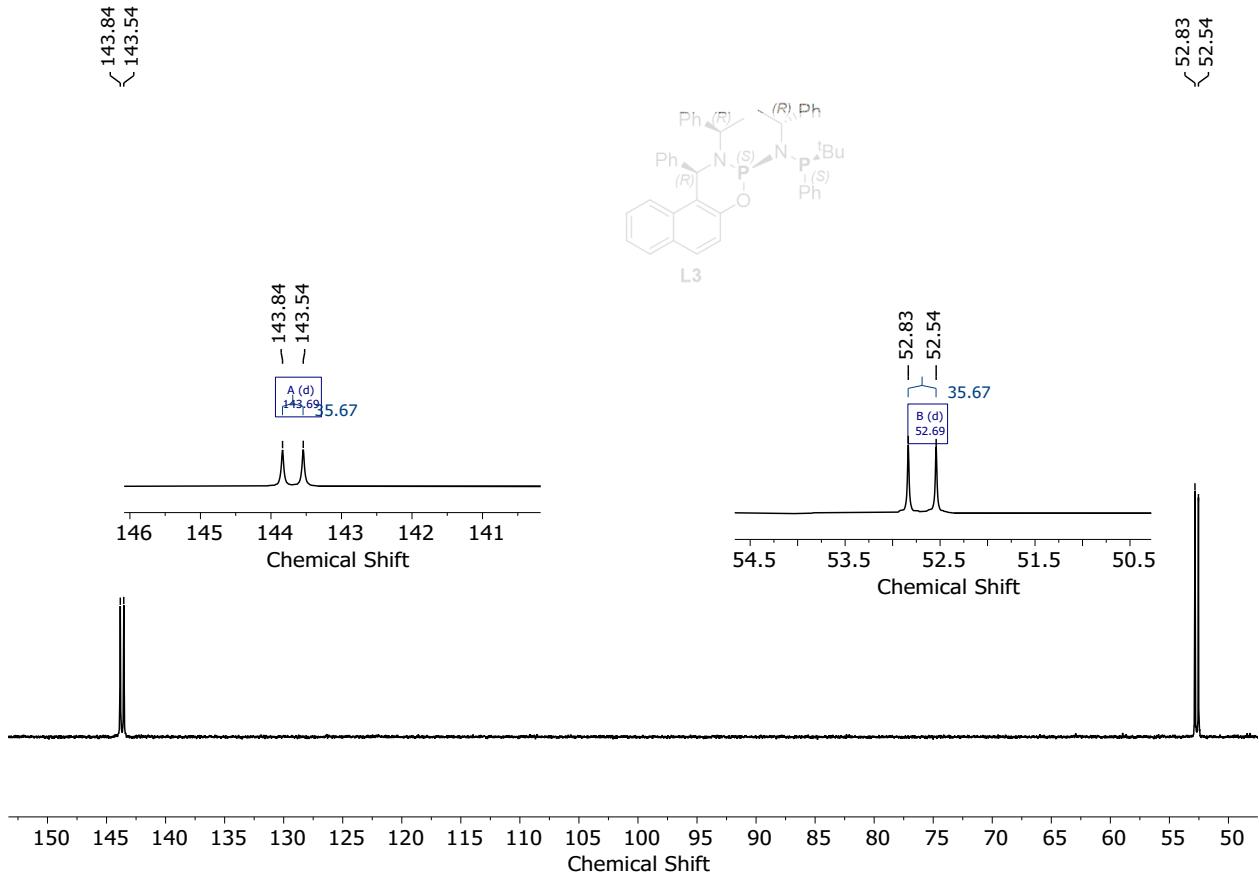


Figure S 14 ^{31}P {H} NMR of (*R,R,S_P,R,S_P*)-L3

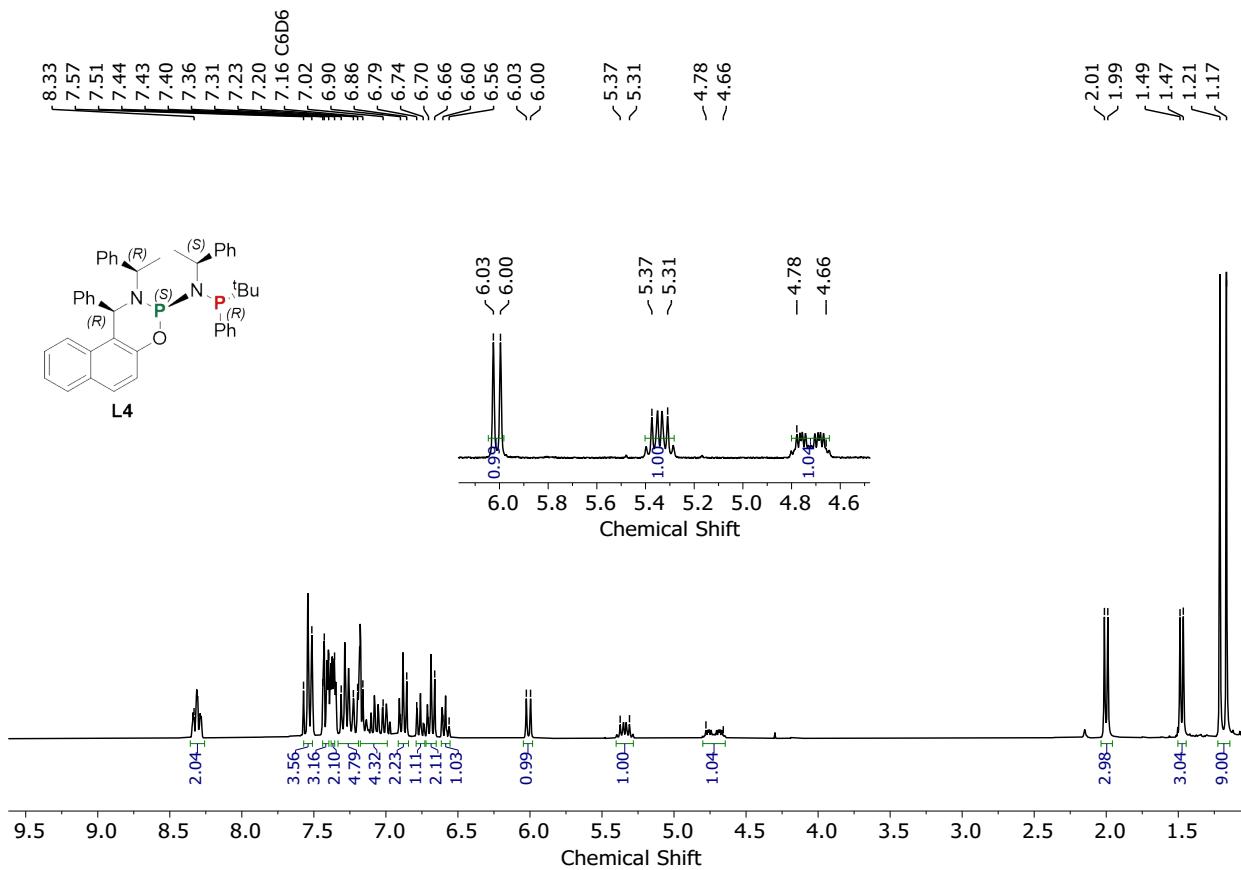


Figure S 15 1H NMR of (*R,R,S_P,S,R_P*)-L4

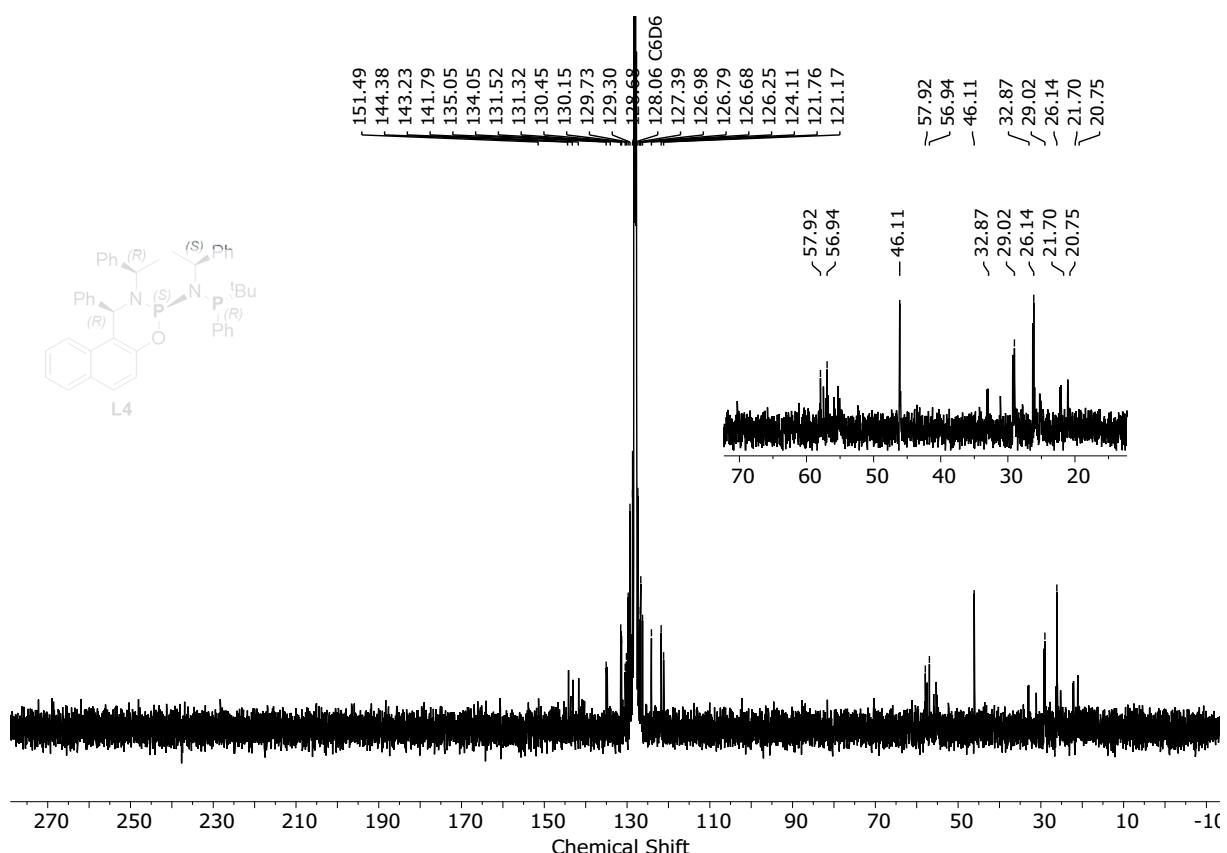


Figure S 16 ^{13}C NMR of ($\text{R},\text{R},\text{S}_P,\text{S},\text{R}_P$)-L4

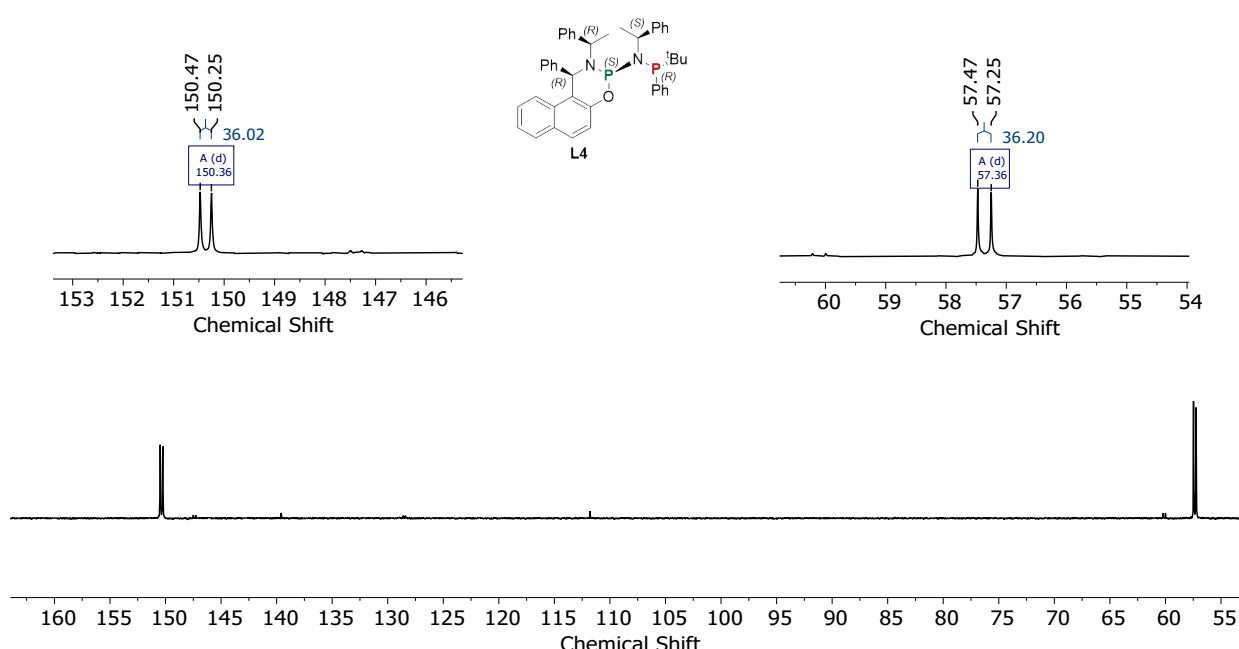


Figure S 17 $^{31}\text{P}\{\text{H}\}$ NMR of ($\text{R},\text{R},\text{S}_P,\text{S},\text{R}_P$)-L4

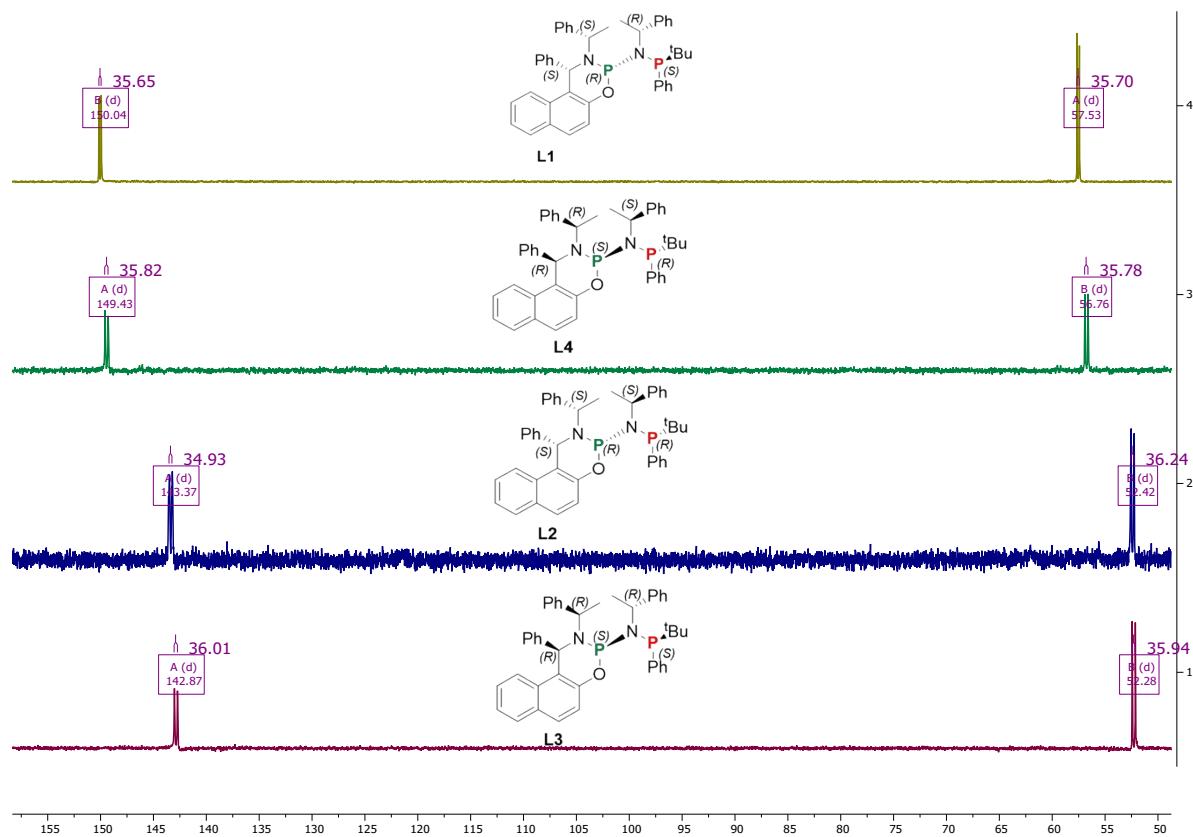


Figure S 18 ^{31}P {H} NMR of the possible ligand combination

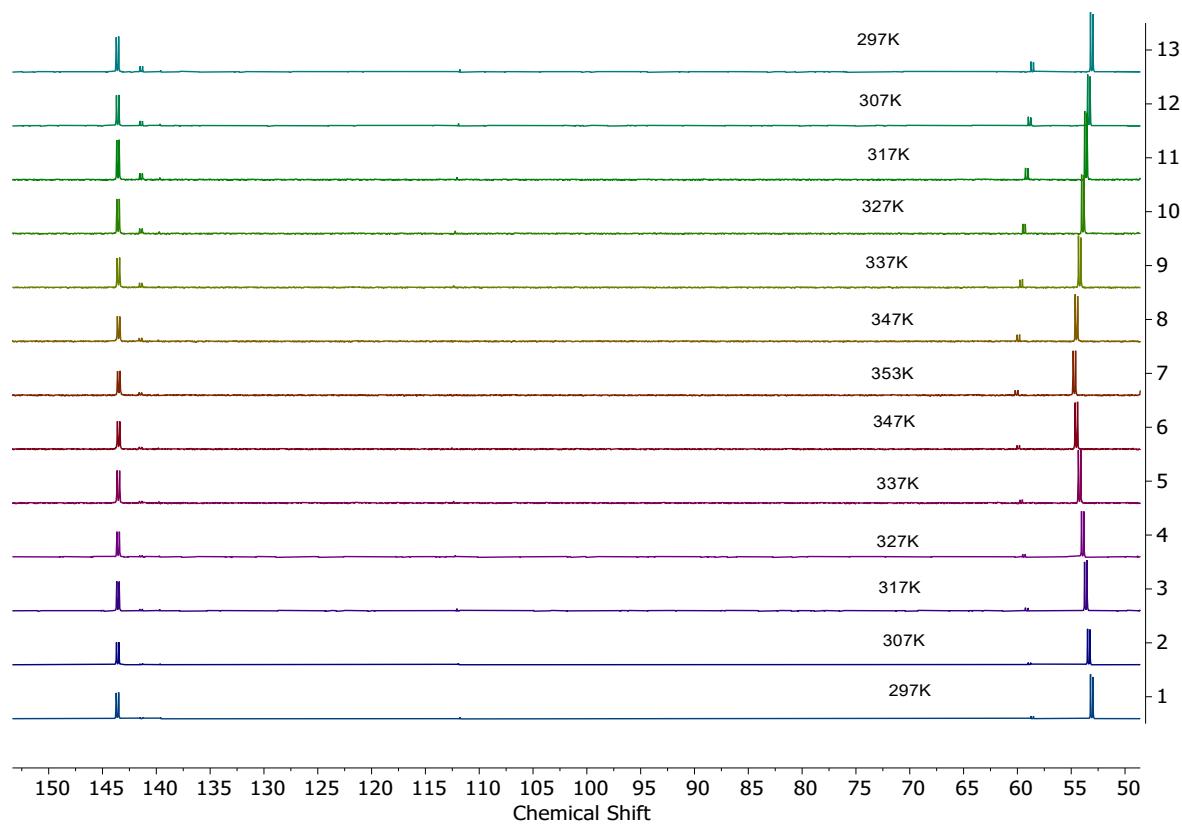


Figure S 19 Temperature dependent epimerization of *S,S,R_P,S,R_P-L2*.

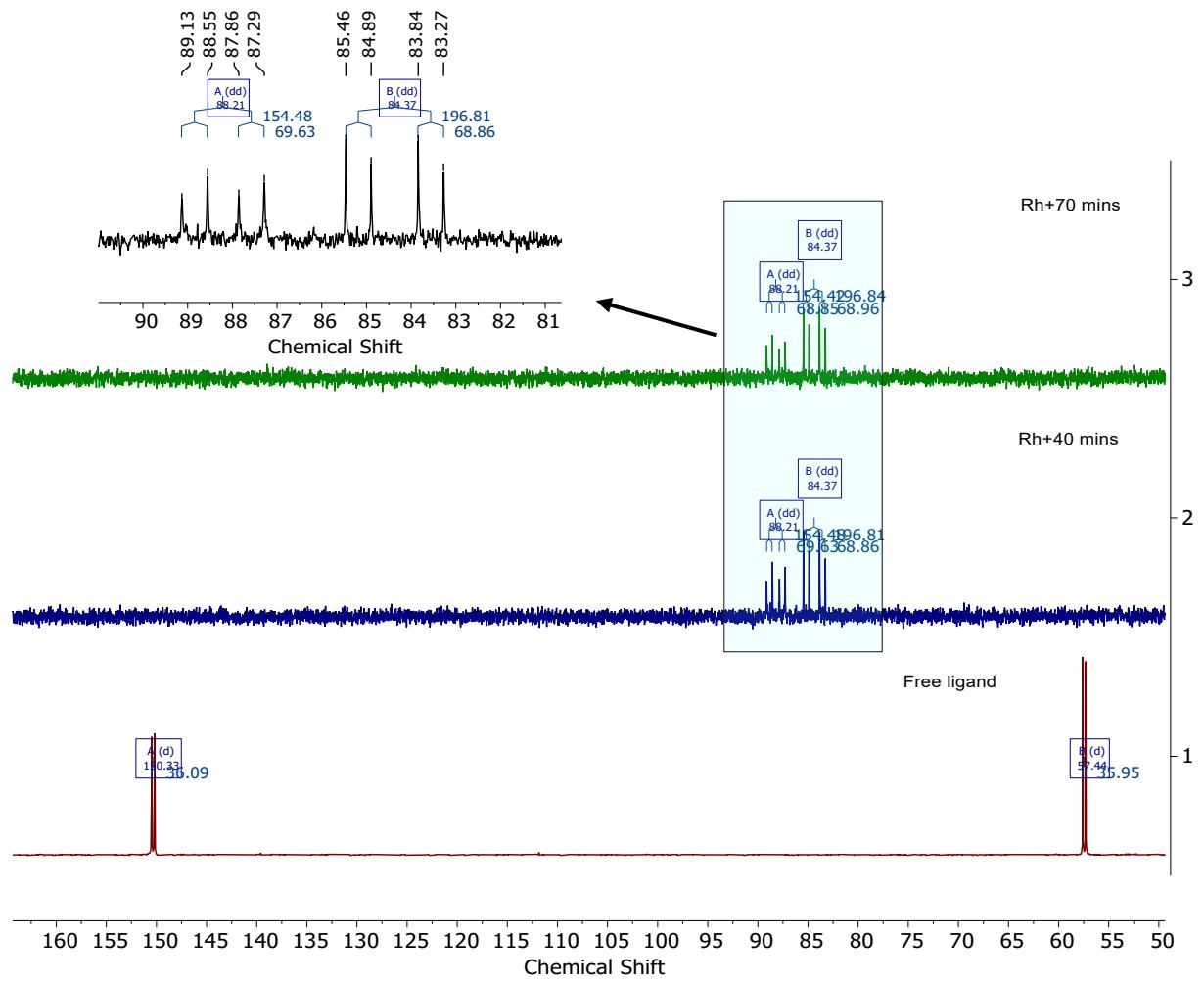


Figure S 20 ^{31}P NMR of In-situ Complexation studies using L2/Rh system (in Methanol- d_4).

- **HRMS of the ligands**

ESI-TOF Accurate Mass Report							Page 1
Results file: E:\Projects\2106.PRO\SampleDB\2106.rpt							
Last modified: Friday, June 04, 2021 08:24:26							
Sample Summary:							
Sample	File	Sample Name	User	Target	Formula	Expected Mass	Observed Mass
58	21060404	SDC-L-1 :hakrabortty	666.2929	C43H44N2O2P2	667.3007	667.3005	-0.3
							-0.2

Figure S 21 HRMS of (S,S,R_P,R,S_P)- L1

ESI-TOF Accurate Mass Report

Results file: E:\Projects\2106.PRO\SampleDB\2106.rpt
Last modified: Friday, June 04, 2021 08:27:01

Page 1

Sample Summary:

Sample	File	Sample Name	User	Target	Expected Mass	Observed Mass	Error PPM	Error mDa
59	21060405	SDC-L-2 \hakrabortty	666.2929	C43H44N2O2P2	667.3007	667.3011	0.6	0.4

Figure S 22 HRMS of (S,S,R_P,S,R_P)- L2

ESI-TOF Accurate Mass Report

Results file: E:\Projects\2006.PRO\SampleDB\2006.rpt
Last modified: Monday, June 22, 2020 16:34:04

Page 1

Sample Summary:

Sample	File	Sample Name	User	Target	Formula	Expected Mass	Observed Mass	Error PPM	Error mDa
172	20062212	SDC-152	Chakrabortty	666.2929	C43H44N2OP2	667.3007	667.3020	1.9	1.3

Figure S 23 HRMS of (R,R,S_P,R,S_P)- L3

ESI-TOF Accurate Mass Report

Results file: E:\Projects\2106\PRO\SampleDB\2106.rpt
Last modified: Friday, June 04, 2021 08:32:18

Page 1

Sample Summary:

Sample	File	Sample Name	User	Target	Formula	Expected Mass	Observed Mass	Error PPM	Error mDa
61	21060407	SDC-L-4 :hakraborttty		666.2929	C43H44N2O2	667.3007	667.3016	1.3	0.9

Figure S 24 HRMS of (R, R, S_p, S, R_p)- L4

■ NMR of the hydrogenated products

All the hydrogenated products have been characterized and matched with the literature reports.^{2,3}

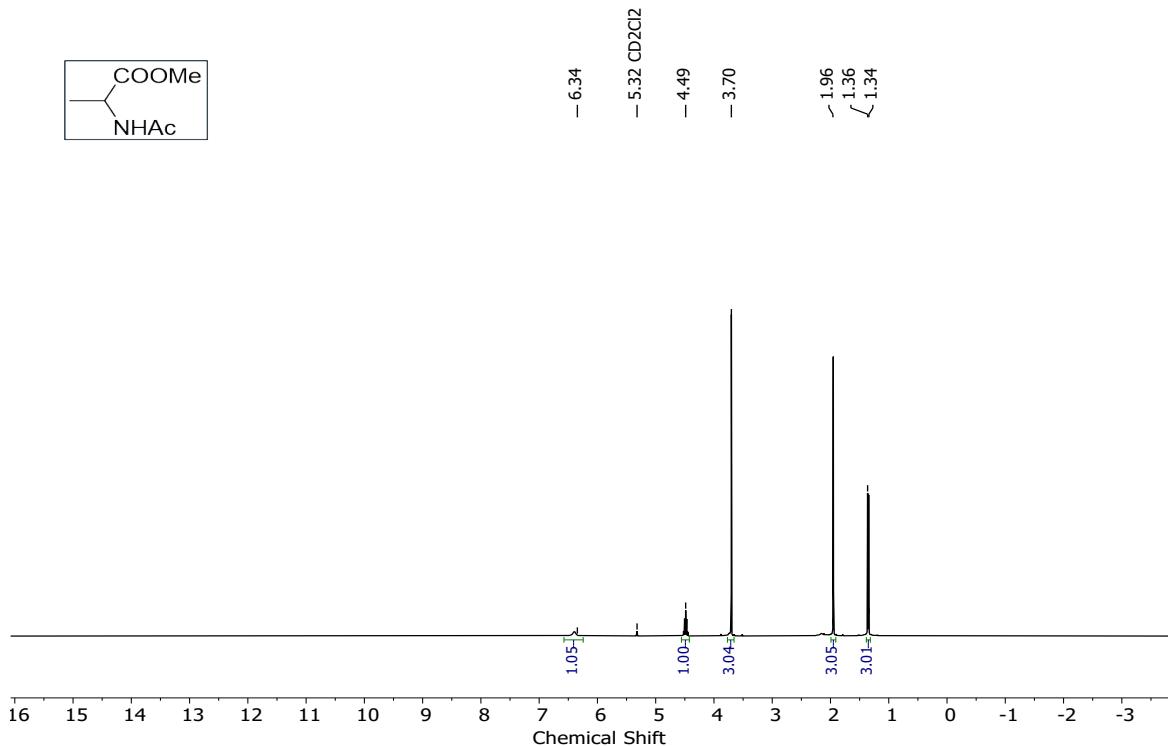


Figure S 25 ¹H NMR of 1b

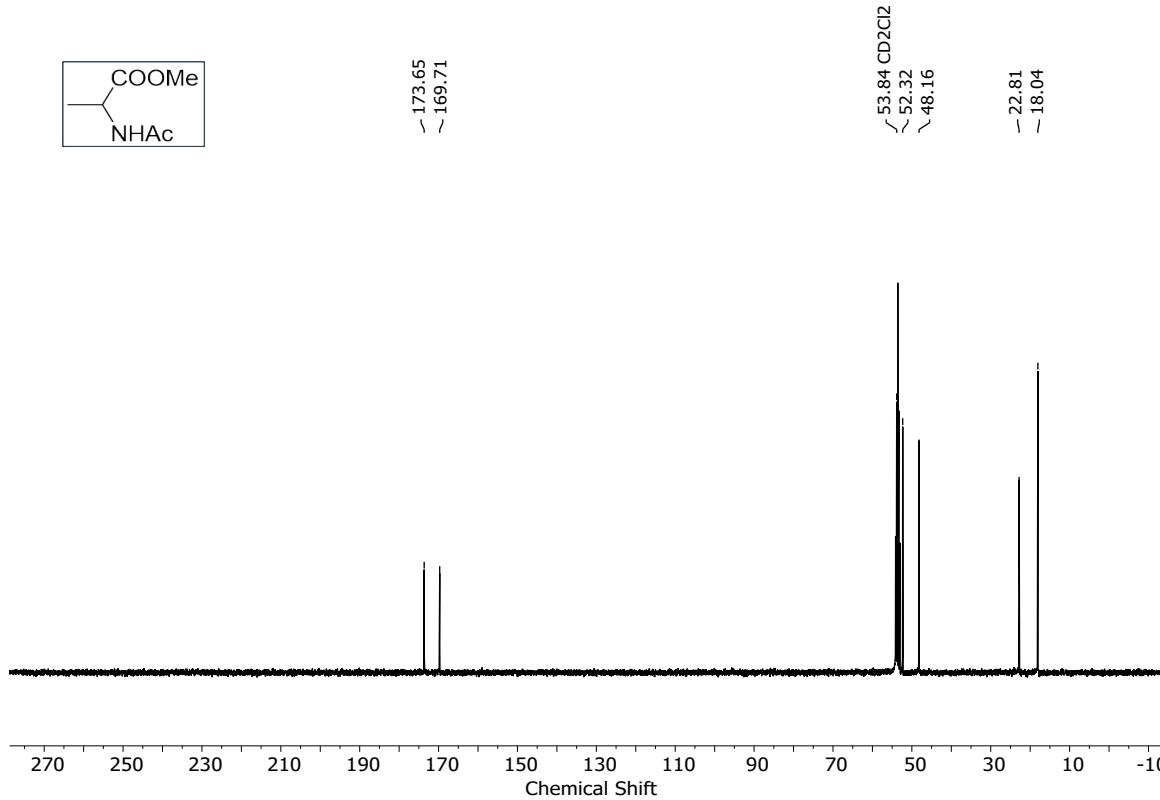


Figure S 26 ¹³C NMR of 1b

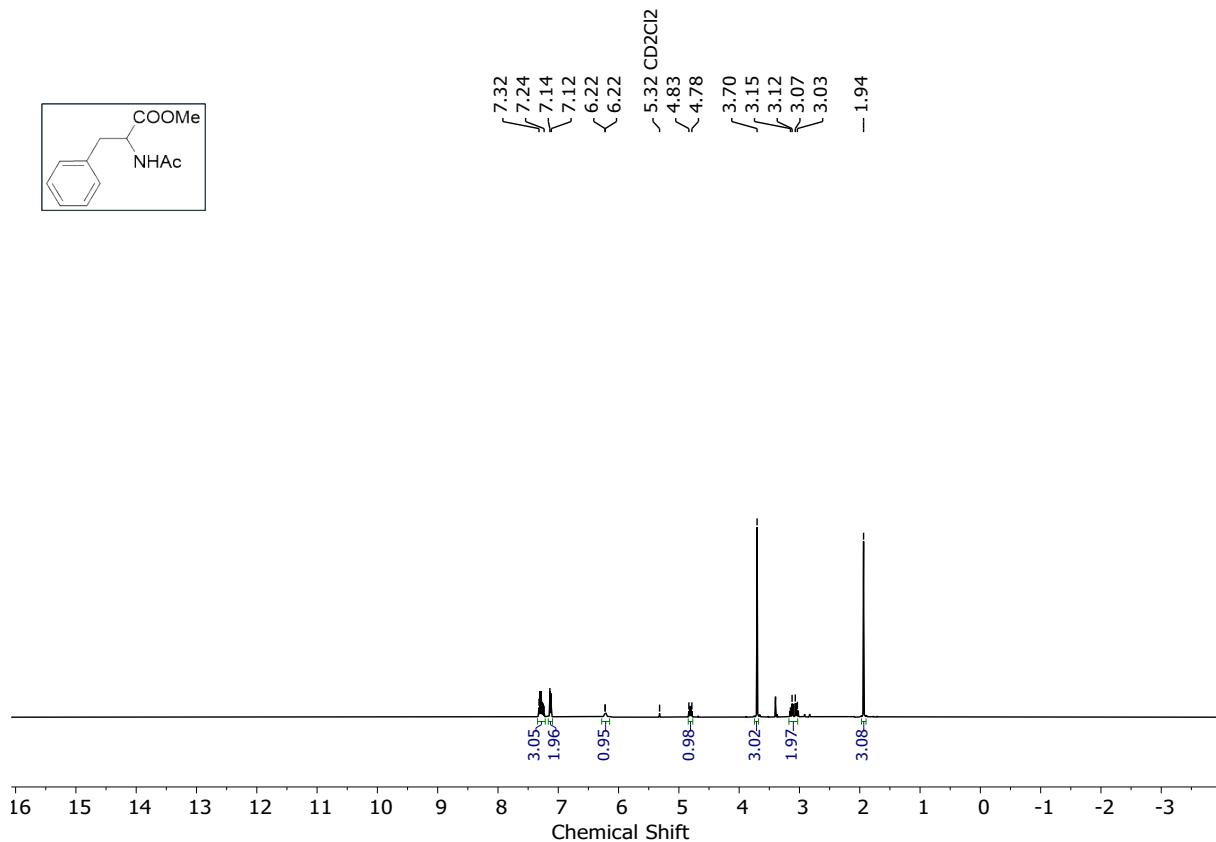


Figure S 27 ¹H NMR of **Ic**

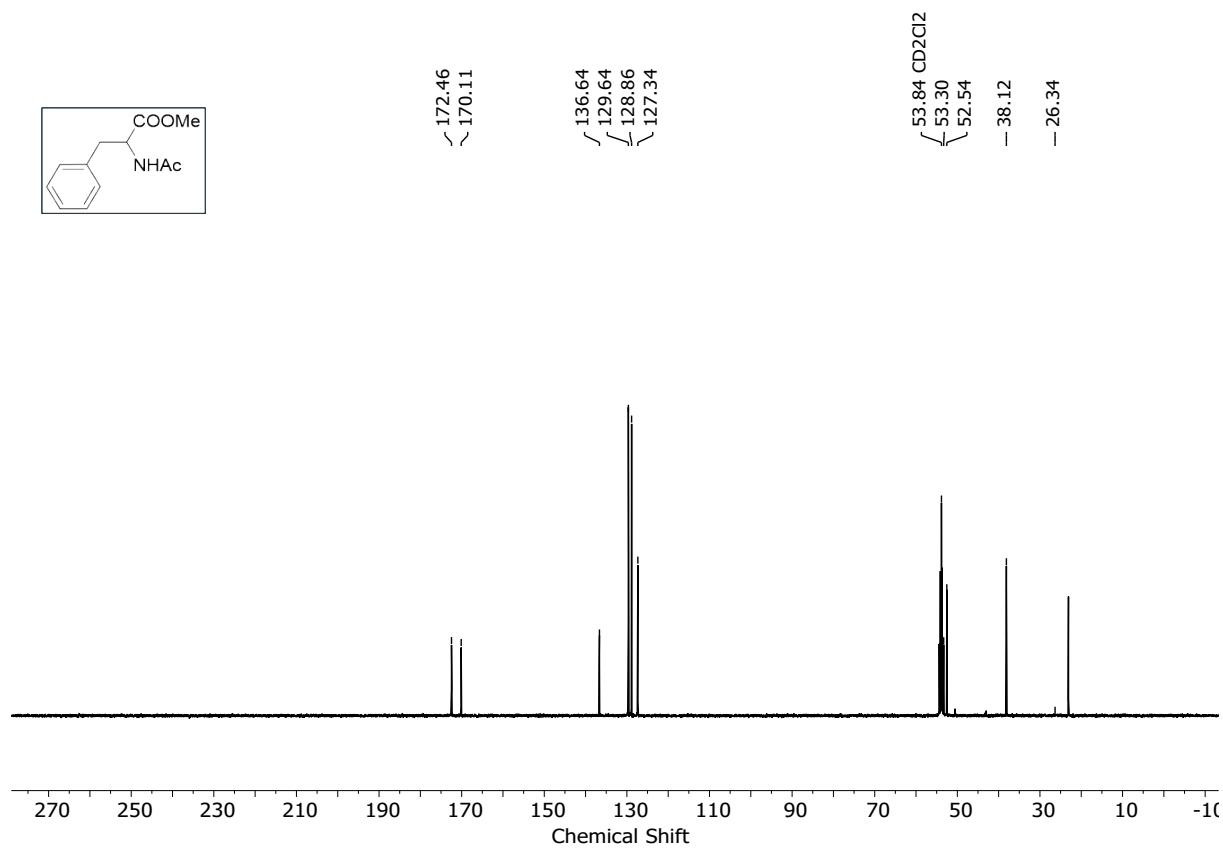


Figure S 28 ¹³C NMR of **Ic**

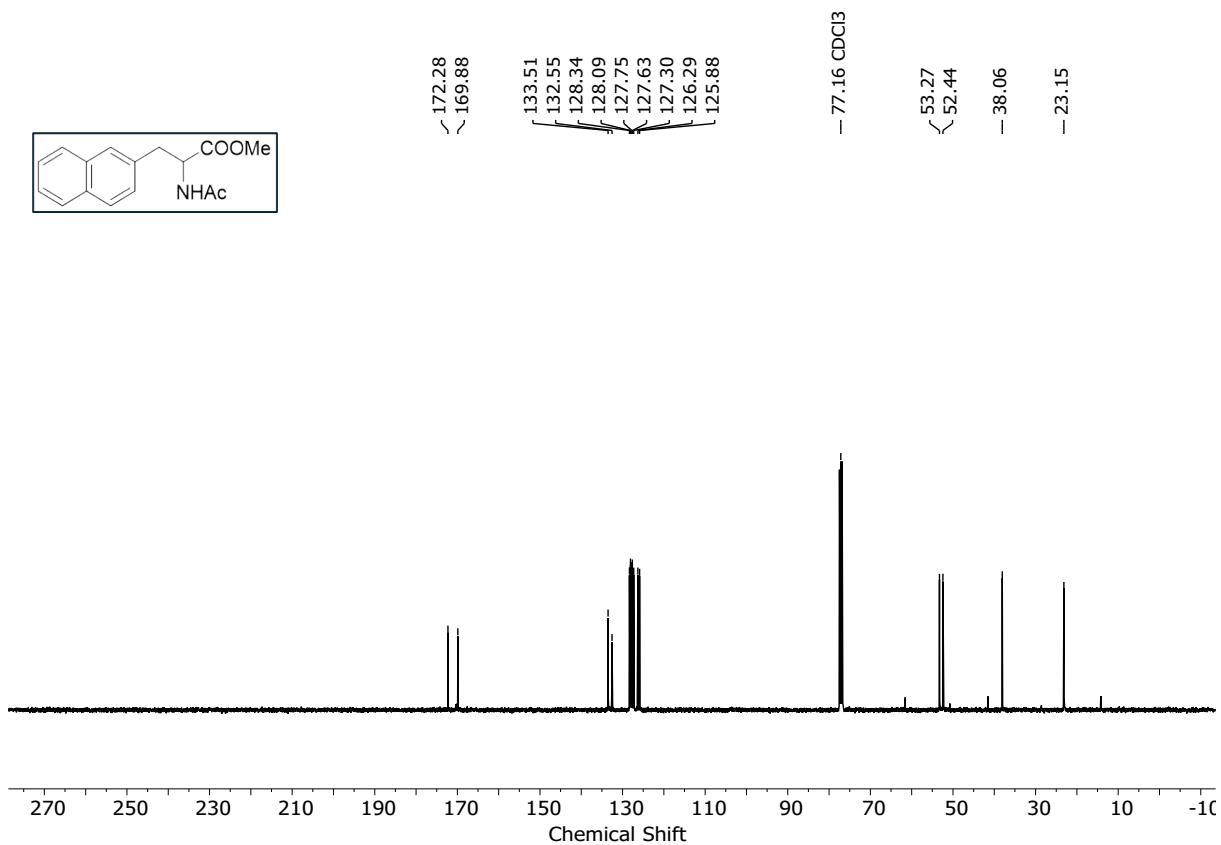
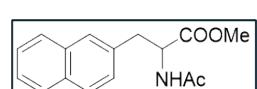
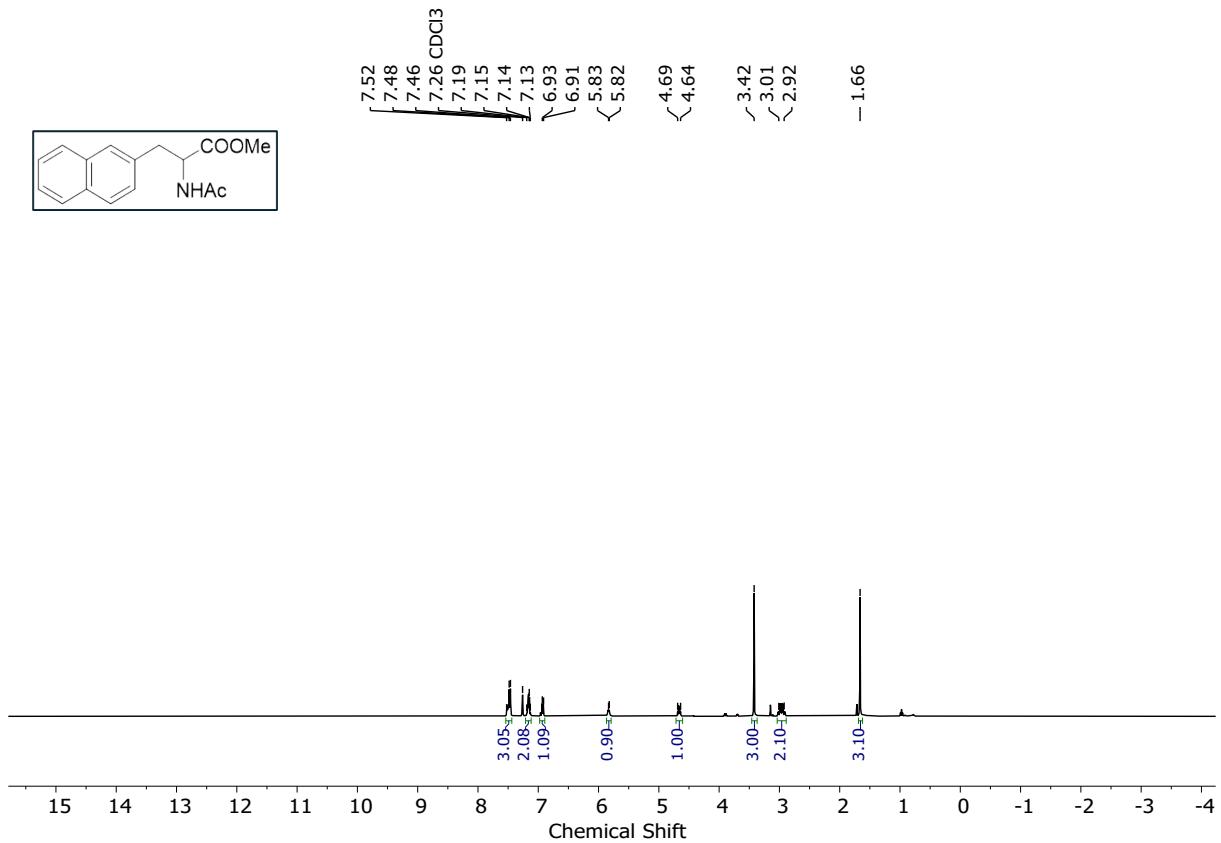
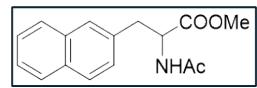


Figure S 30 ^{13}C NMR of **1d**

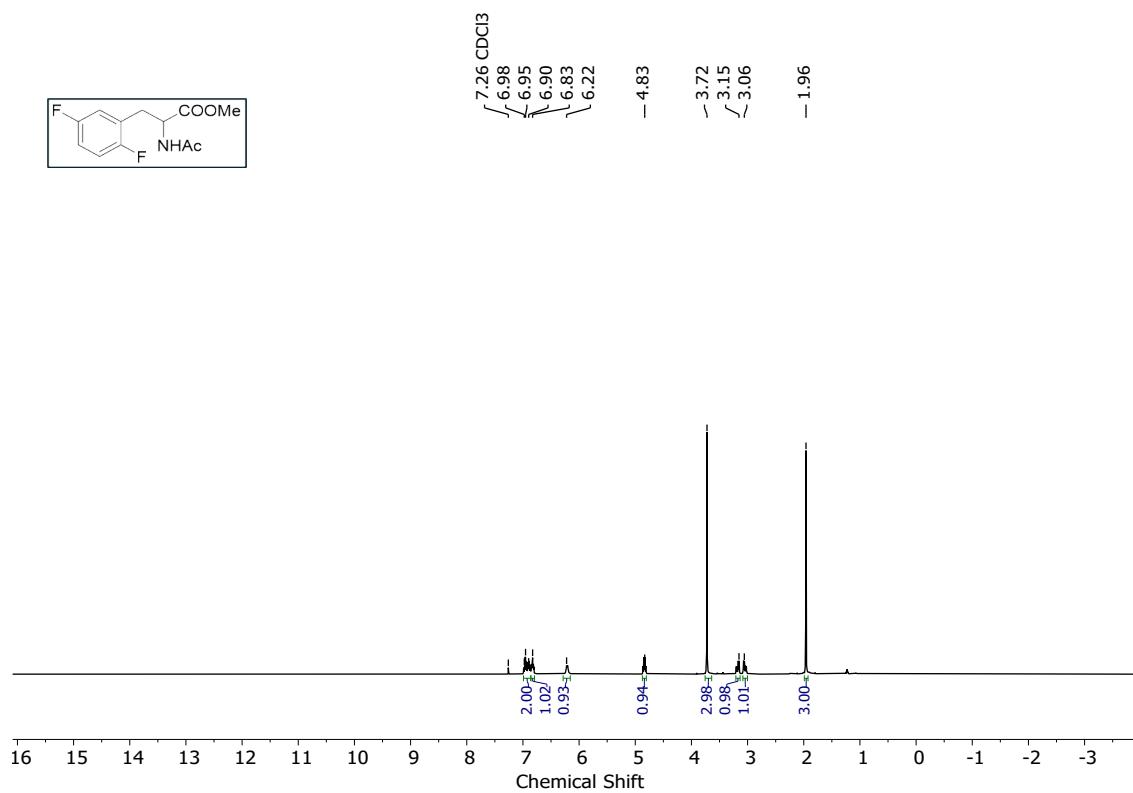
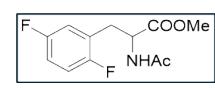


Figure S 31 ^1H NMR of **1e**

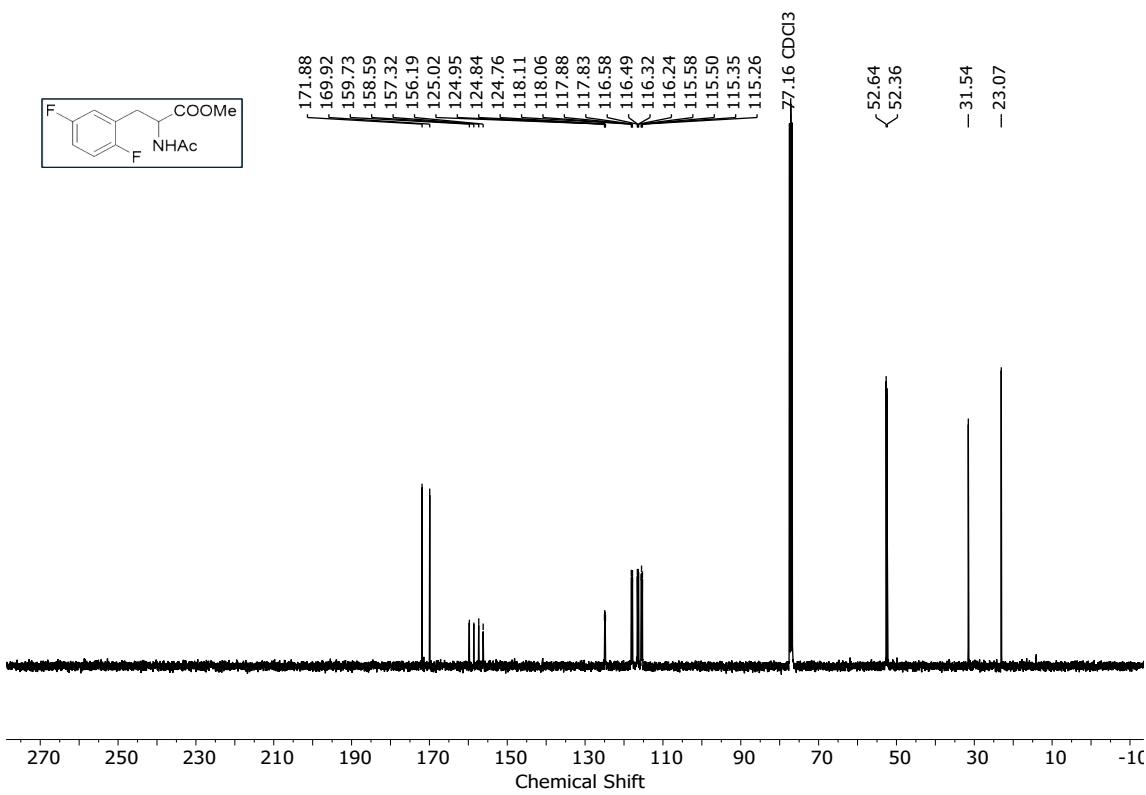


Figure S 32 ^{13}C NMR of **1e**

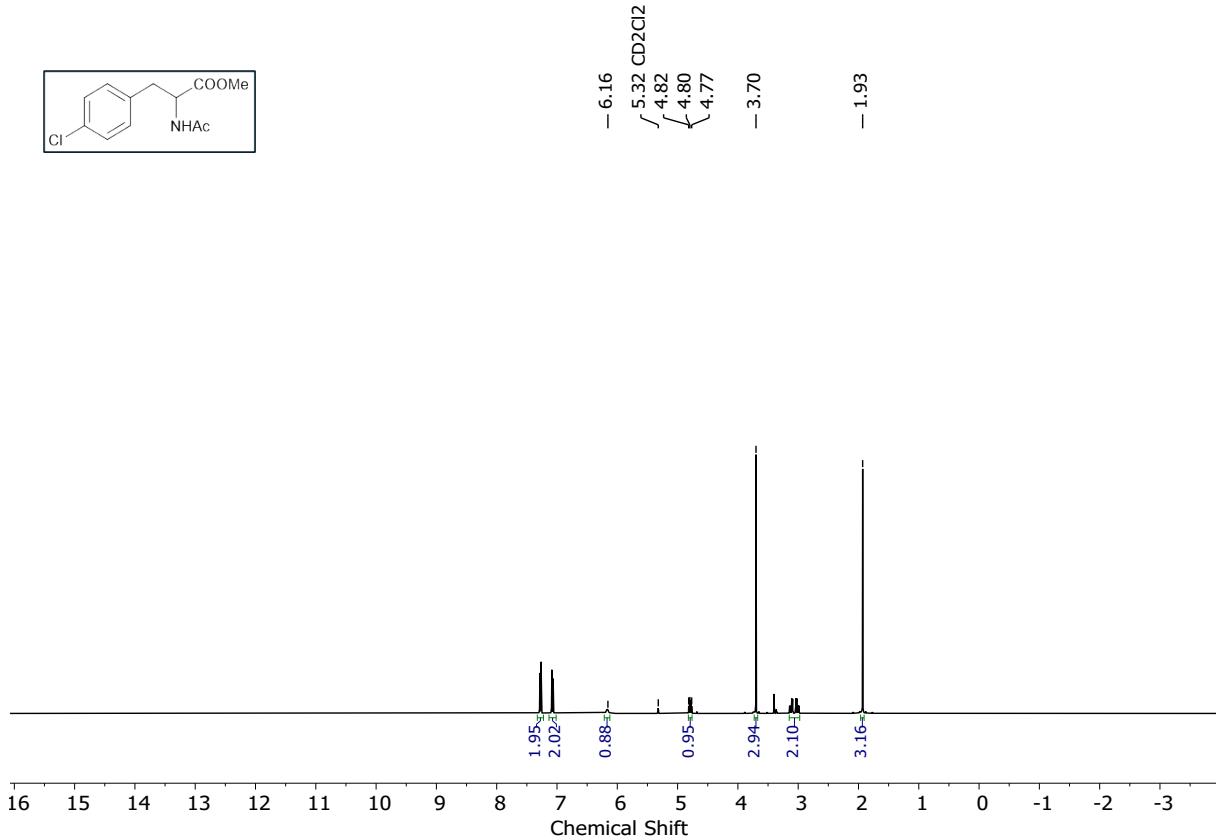


Figure S 33 ^1H NMR of **If**

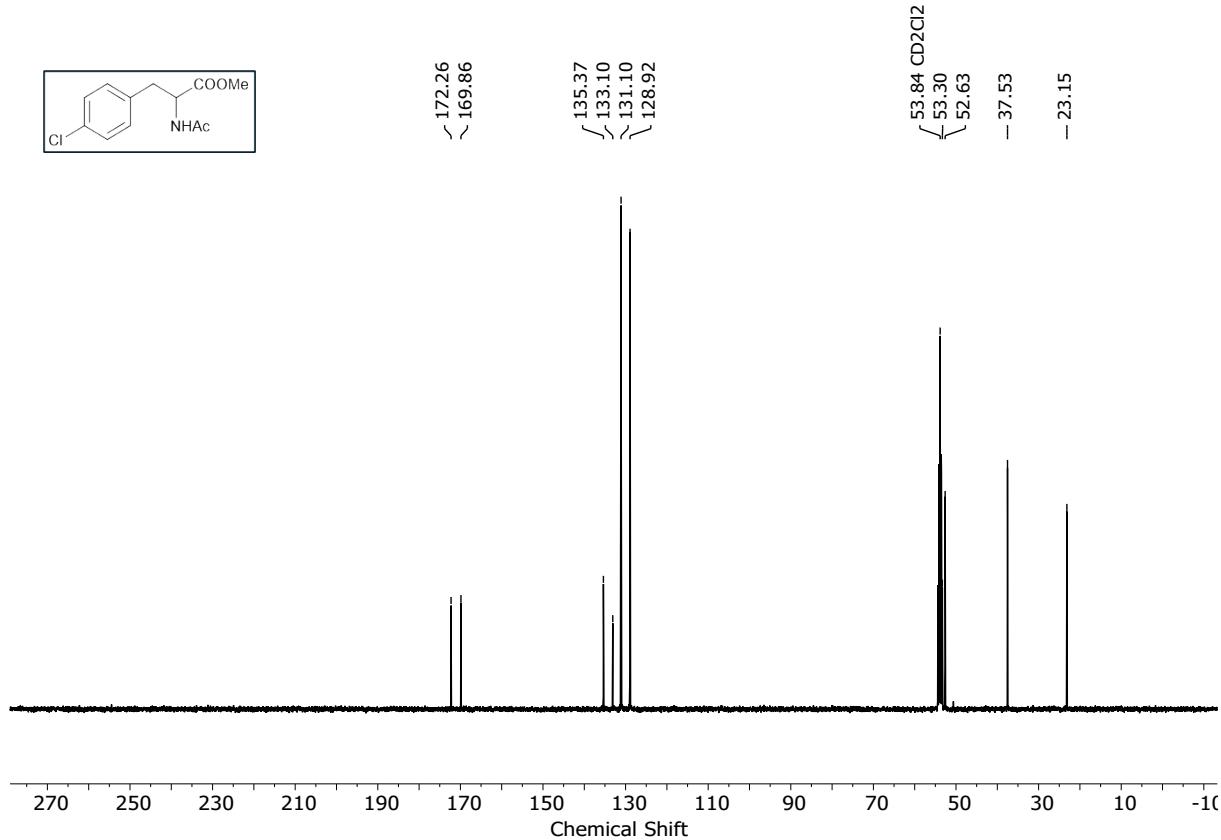


Figure S 34 ^{13}C NMR of **If**

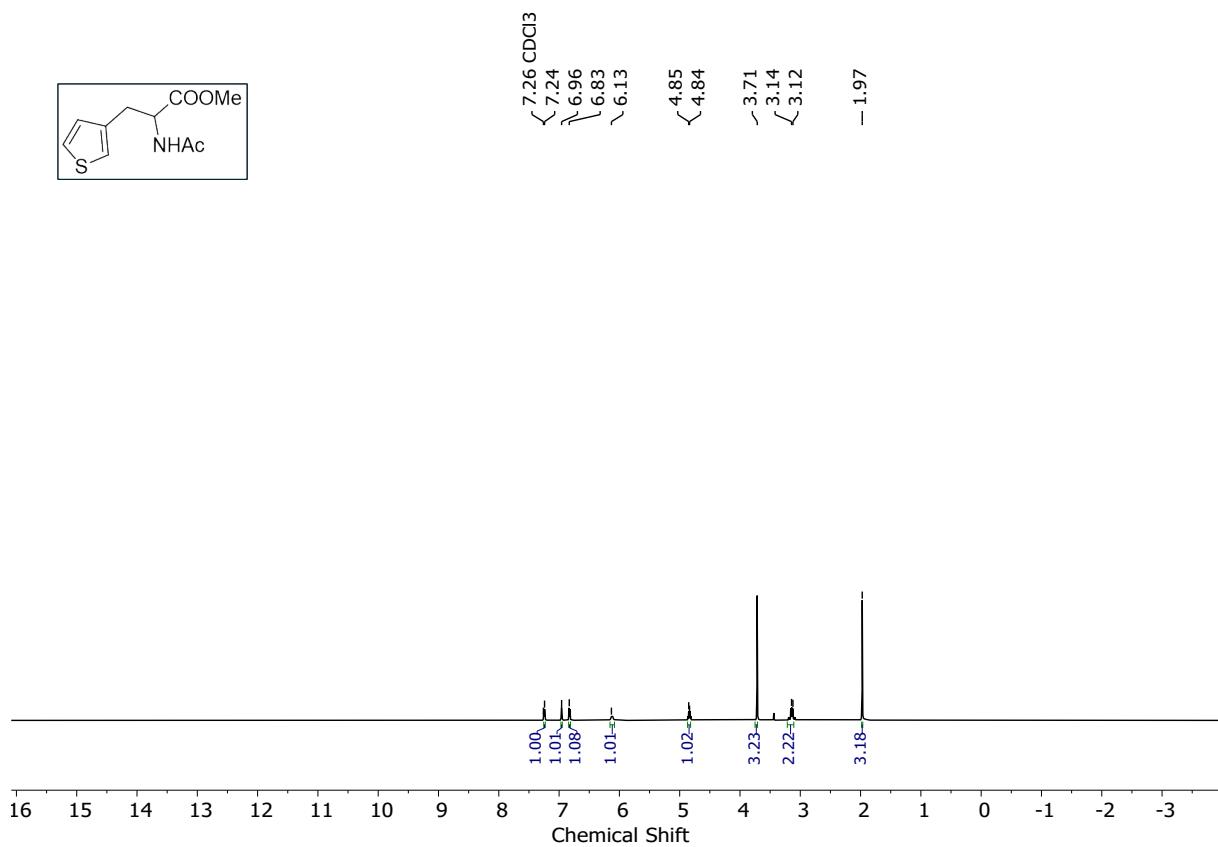


Figure S 35 ¹H NMR of **Ig**

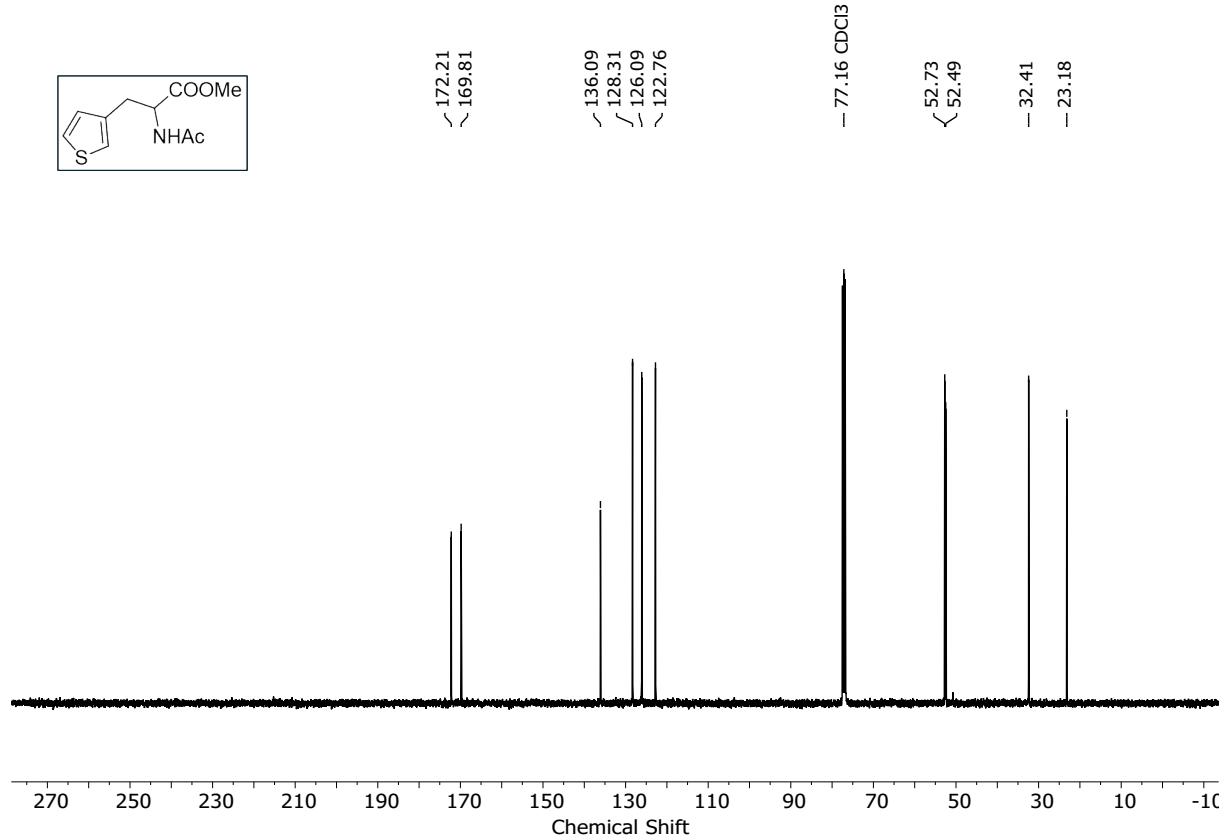


Figure S 36 ¹³C NMR of **Ig**

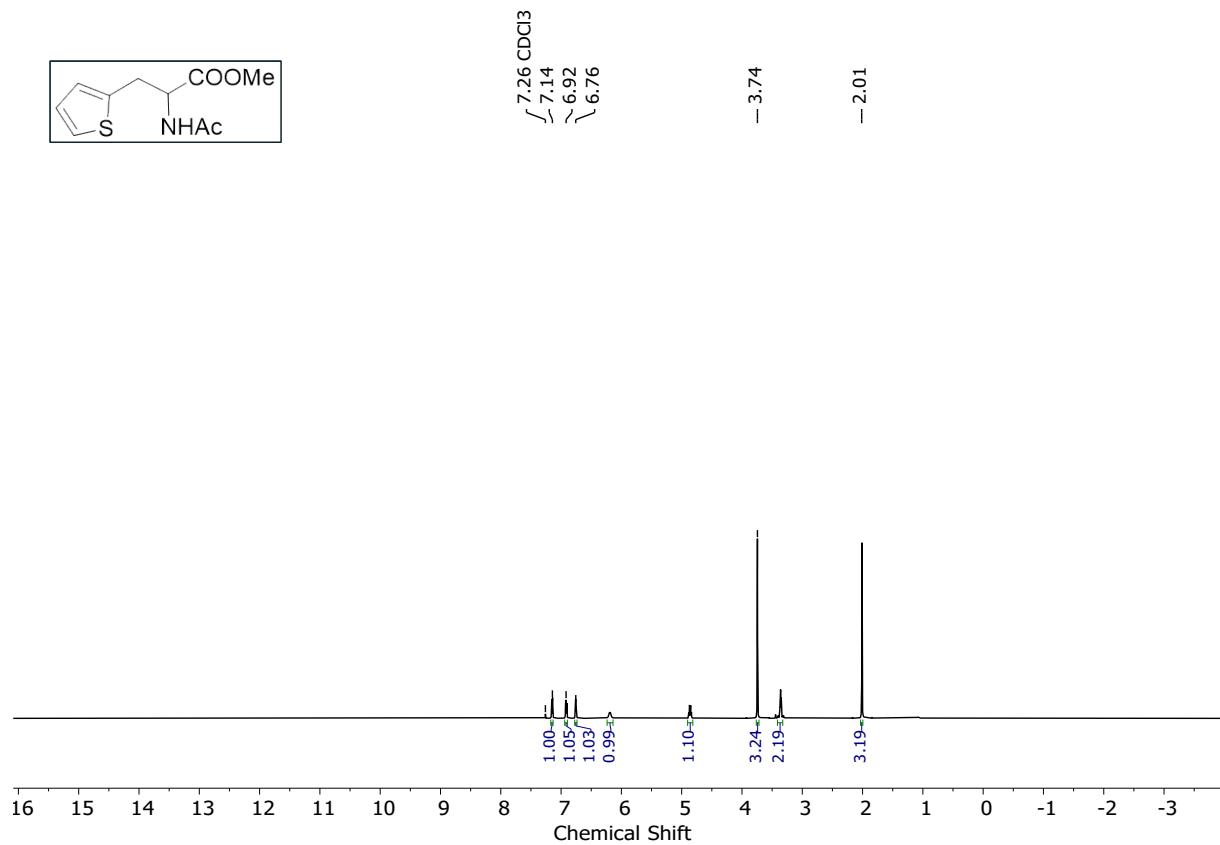
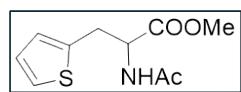


Figure S 37 ^1H NMR of Ih

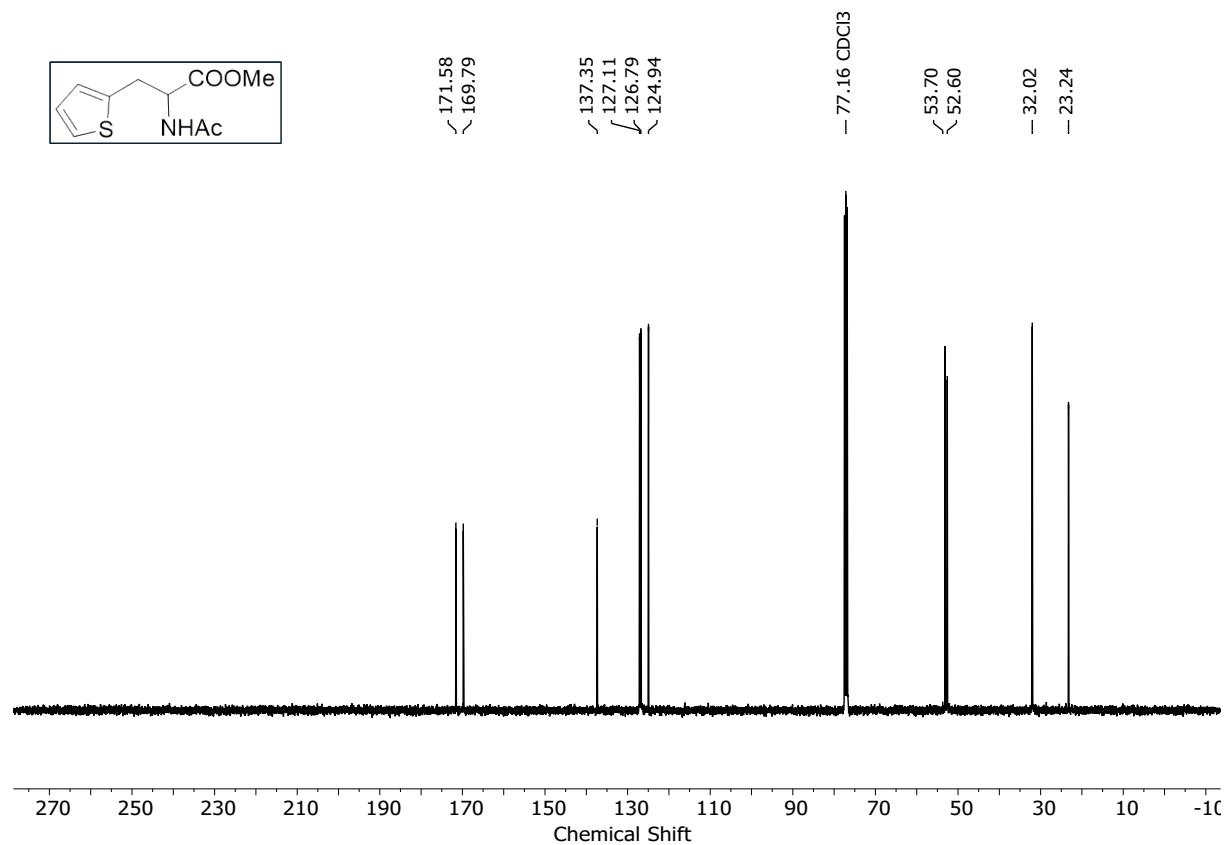
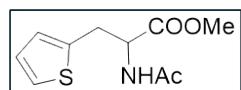
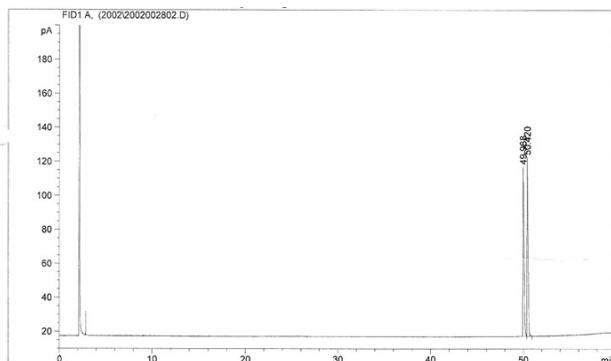
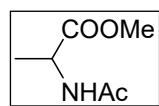


Figure S 38 ^{13}C NMR of Ih

▪ **HPLC Traces for enantiomeric excess determination of the product**
Method Information for HPLC

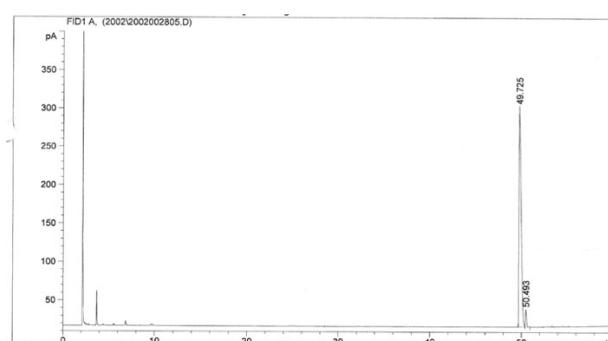
Substrate	Column
1b	25m Lipodex E 90/40-6-200/10
1a, 1c-1h	Chiralcel OJ-H Heptan/EtOH 95:5
2a	AD-H, Heptan/EtOH 90:10

• **Asymmetric Hydrogenation**



Peak #	RetTime [min]	Type	Width [min]	Area [pA*s]	Height [pA]	Area %
1	49.968	BV	0.1062	839.65417	99.74786	50.08221
2	50.420	VB	0.1073	836.89764	105.11221	49.91779

Figure S 39 HPLC traces of racemic **1b**



Peak #	RetTime [min]	Type	Width [min]	Area [pA*s]	Height [pA]	Area %
1	49.725	BV	0.2083	4816.83594	287.48532	95.93251
2	50.493	VB	0.1139	204.23126	22.84838	4.06749

Figure S 40 HPLC traces of enantioenriched product **1b** using *Rh/S,S,R_P,S,R_P-L2* catalytic system

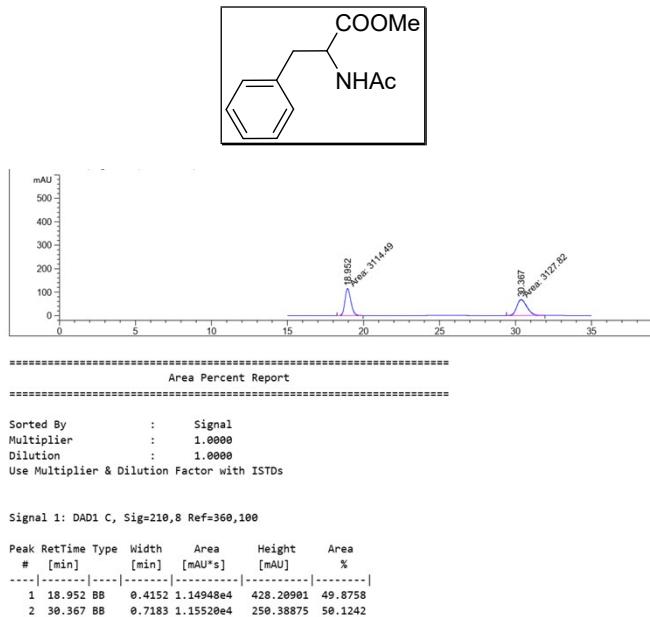


Figure S 41 HPLC traces of racemic **1c**

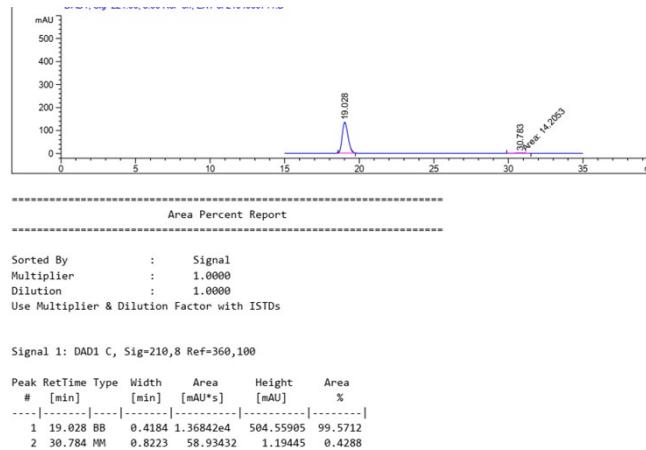


Figure S 42 HPLC traces of enantioenriched product **1c** using *Rh/S,S,R_P,S,R_P-L2* catalytic system

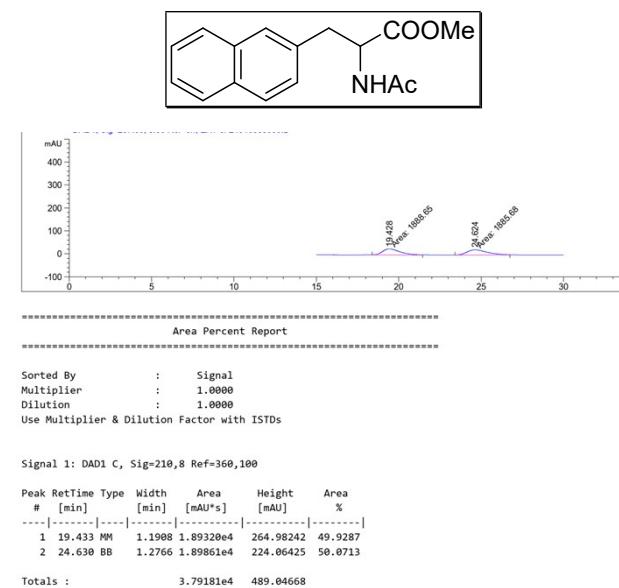


Figure S 43 HPLC traces of racemic **1d**

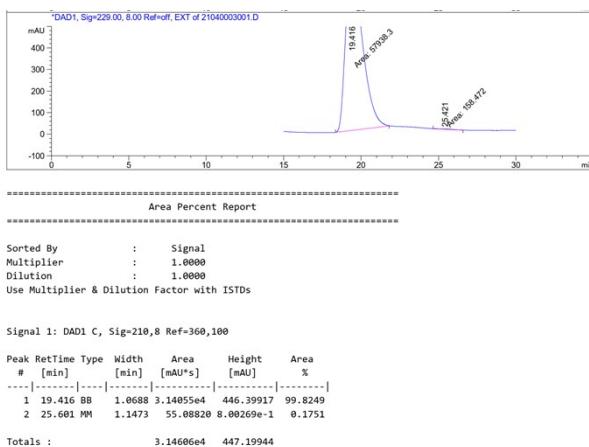


Figure S 44 HPLC traces of enantioenriched product **1d** using *Rh/S,S,R_P,S,R_P-L2* catalytic system

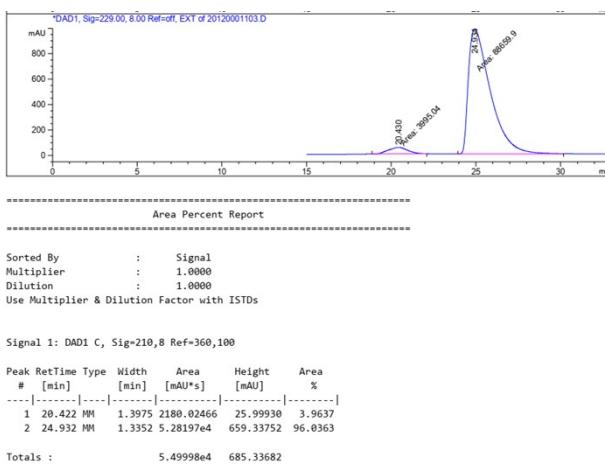


Figure S 45 HPLC traces of enantioenriched product **1d** using *Rh/R,R,S_P,R,S_P-L3* catalytic system.

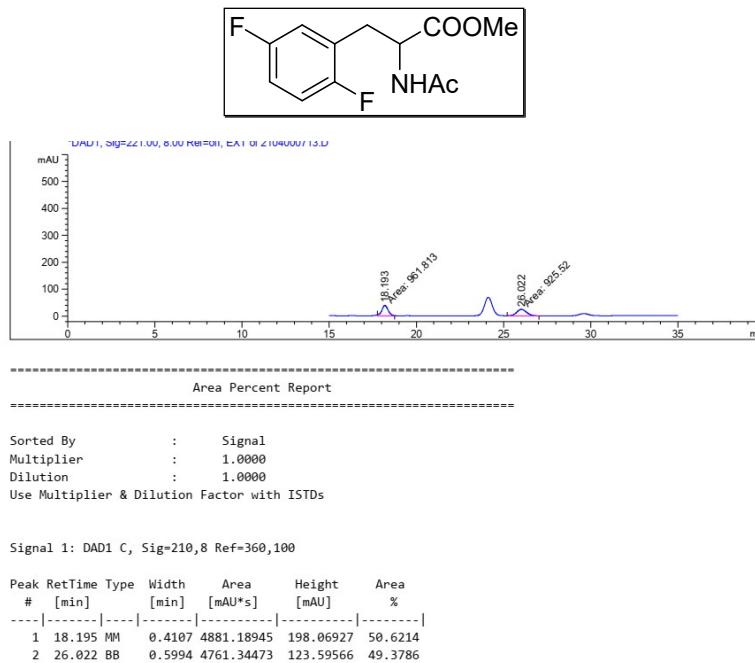


Figure S 46 HPLC traces of racemic **1e**

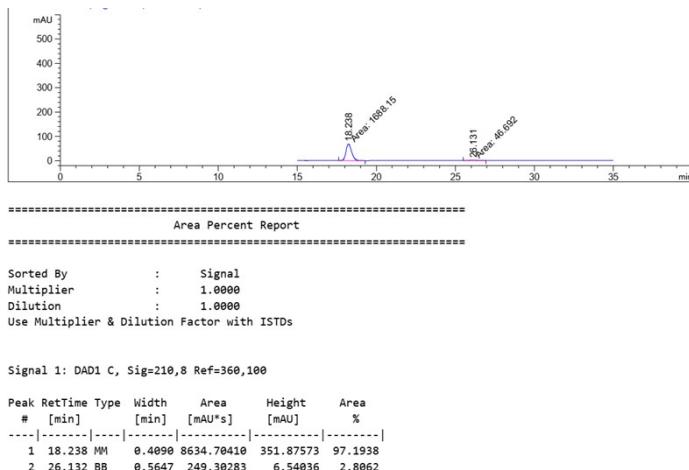


Figure S 47 HPLC traces of enantioenriched product **1e** using *Rh/S,S,R_P,S,R_P-L2* catalytic system

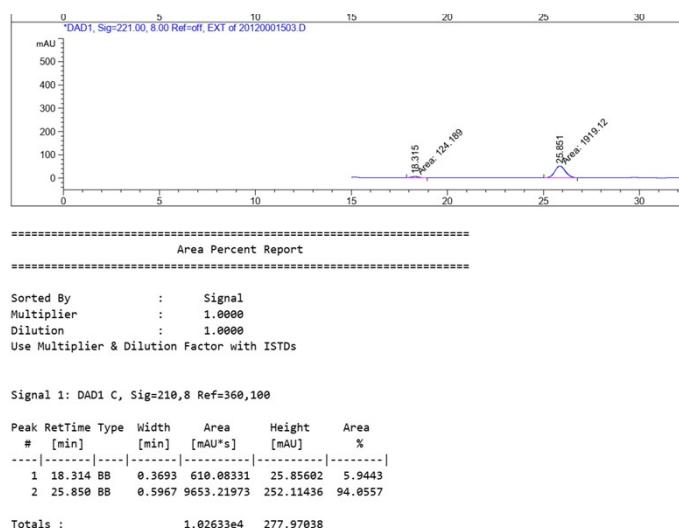


Figure S 48 HPLC traces of enantioenriched product **1e** using *Rh/R,R,S_P,R,S_P-L3* catalytic system.

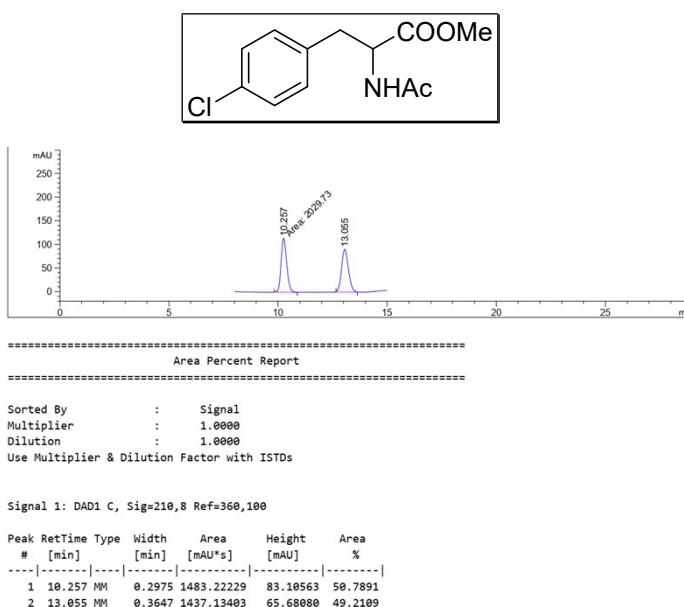
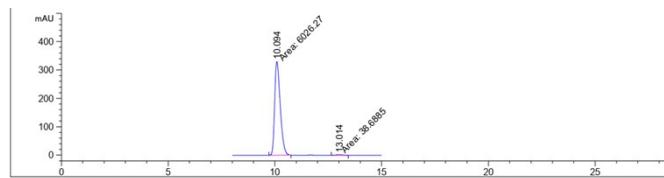


Figure S 49 HPLC traces of racemic **1f**



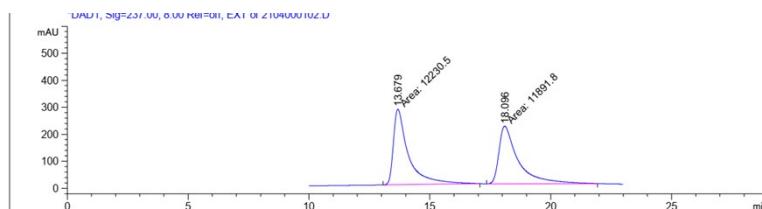
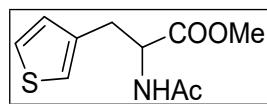
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Area Percent Report
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Sorted By : Signal
Multiplier : 1.0000
Dilution : 1.0000
Use Multiplier & Dilution Factor with ISTDs

Signal 1: DAD1 C, Sig=210,8 Ref=360,100

Peak RetTime	Type	Width	Area	Height	Area %
# [min]		[min]	[mAU*s]	[mAU]	
1 10.094	MM	0.3084	7780.92529	431.75073	99.4528
2 13.009	BB	0.3065	42.81082	2.07715	0.5472

Figure S 50 HPLC traces of enantioenriched product **1f** using *Rh/S,S,R_P,S,R_P-L2* catalytic system



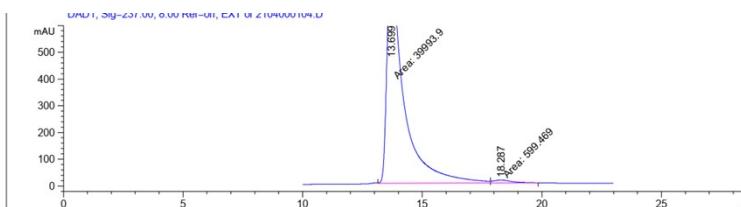
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Area Percent Report
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Sorted By : Signal
Multiplier : 1.0000
Dilution : 1.0000
Use Multiplier & Dilution Factor with ISTDs

Signal 1: DAD1 C, Sig=210,8 Ref=360,100

Peak RetTime	Type	Width	Area	Height	Area %
# [min]		[min]	[mAU*s]	[mAU]	
1 13.678	MM	0.7358	8787.10645	199.05054	50.9399
2 18.096	MM	0.9297	8462.84668	151.71397	49.0601

Figure S 51 HPLC traces of racemic **1g**



```
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Area Percent Report
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Sorted By : Signal
Multiplier : 1.0000
Dilution : 1.0000
Use Multiplier & Dilution Factor with ISTDs

Signal 1: DAD1 C, Sig=210,8 Ref=360,100

Peak RetTime	Type	Width	Area	Height	Area %
# [min]		[min]	[mAU*s]	[mAU]	
1 13.698	MF	0.8000	2.86061e4	595.94580	98.3492
2 18.287	FM	0.9872	480.16110	8.10677	1.6508

Figure S 52 HPLC traces of enantioenriched product **1g** using *Rh/S,S,R_P,S,R_P-L2* catalytic system

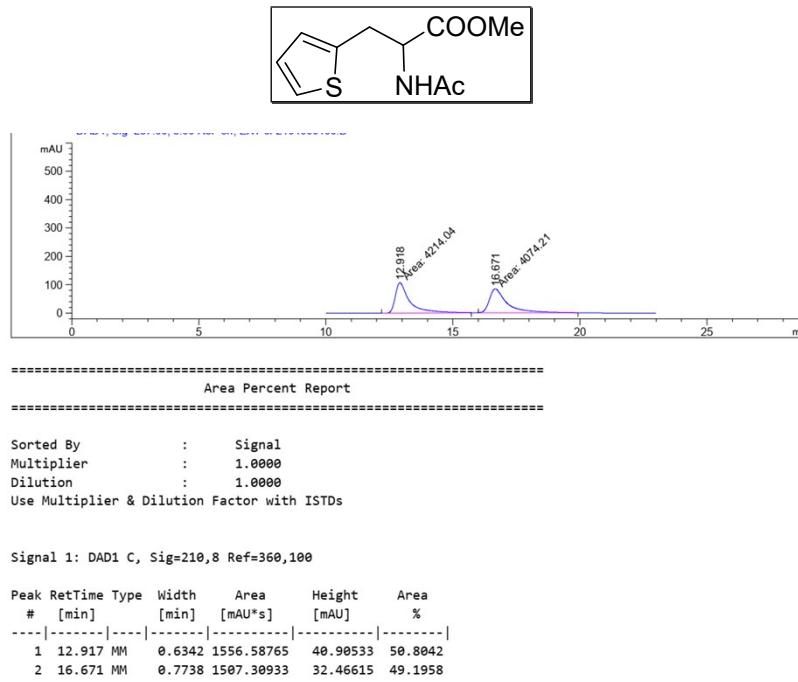


Figure S 53 HPLC traces of racemic **1h**

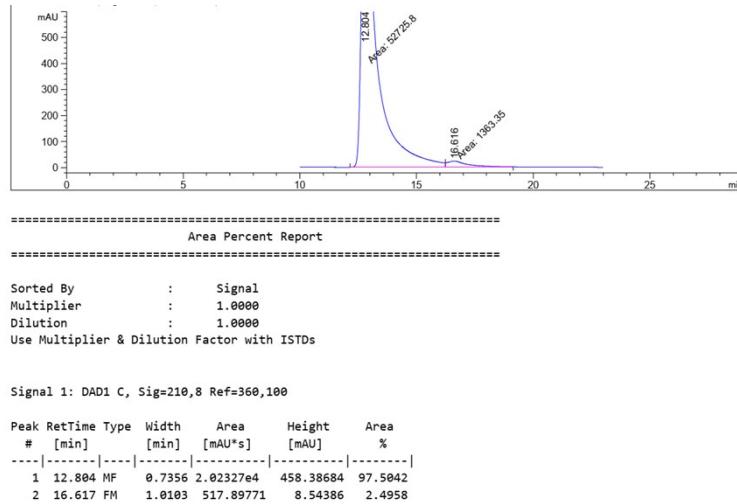
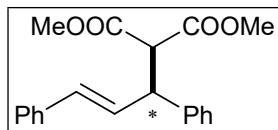


Figure S 54 HPLC traces of enantioenriched product **1h** using *Rh/S,S,R_P,S,R_P-L2* catalytic system

- Asymmetric Allylic substitution



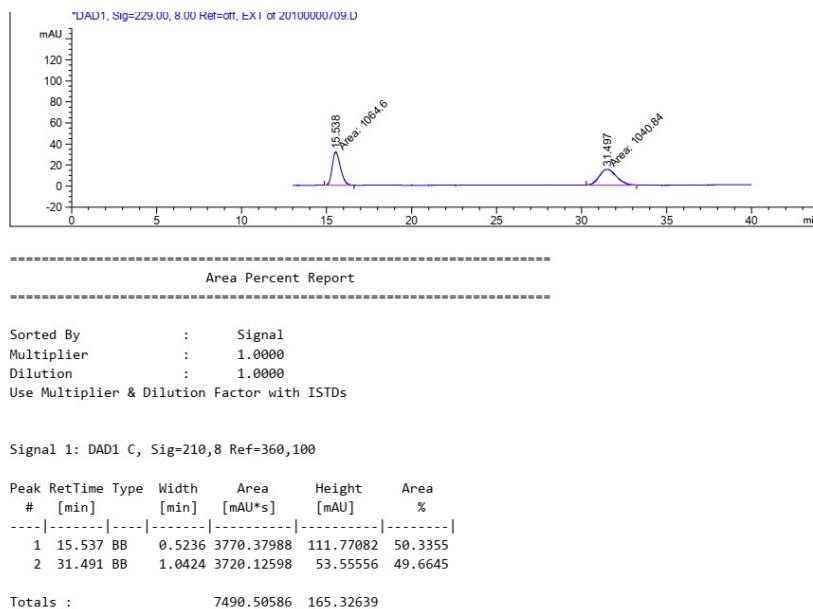


Figure S 55 HPLC traces of racemic **2a**

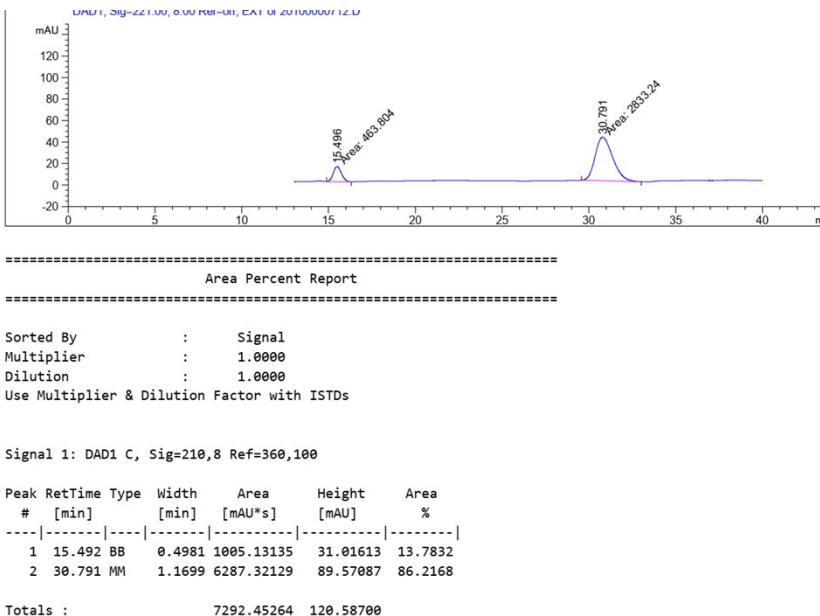


Figure S 56 HPLC traces of enantioenriched product **2a** using *Pd/R,R,S_P,S,R_P-L4* catalytic system.

■ X-ray Crystallographic data

Single crystals were prepared under argon and measured under a continuous flow of nitrogen on a Bruker Kappa APEX II Duo diffractometer. The structure was solved by direct methods (SHELXS-97: Sheldrick, G. M. Acta Cryst. 2008, A64, 112.) and refined by full-matrix least-squares procedures on F²

(SHELXL-2018: Sheldrick, G. M. *Acta Cryst.* 2015, C71, 3.). XP (Bruker AXS) was used for graphical representations.

CCDC 2091899 contains the supplementary crystallographic data for this paper. These data are provided free of charge by the joint Cambridge Crystallographic Data Centre and Fachinformationszentrum Karlsruhe Access Structures service www.ccdc.cam.ac.uk/structures.

Crystal data of *S,S,R_P,R,S_P-L1*: C₄₃H₄₄N₂OP₂, $M = 666.74$, orthorhombic, space group $P2_12_12_1$, $a = 10.0784(3)$, $b = 11.7543(3)$, $c = 30.5885(7)$ Å, $V = 3623.65(16)$ Å³, $T = 150(2)$ K, $Z = 4$, 30024 reflections measured, 6377 independent reflections ($R_{\text{int}} = 0.0251$), final R values ($I > 2\sigma(I)$): $R_1 = 0.0246$, $wR_2 = 0.0645$, final R values (all data): $R_1 = 0.0247$, $wR_2 = 0.0646$, 438 parameters, largest diff. peak/hole: 0.38/-0.25 eÅ⁻³, Flack parameter x = -0.002(4).

■ Reference

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