

Organocatalytic Enantioselective Hydrophosphonylation of Aldehydes

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ELECTRONIC SUPPLEMENTARY INFORMATION (ESI)

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General Experimental Methods. Purification of reaction products was carried out by flash chromatography using silical-gel (0.063-0.200 mm). Analytical thin layer chromatography was performed on 0.25 mm silical gel 60-F plates. ESI ionization method and mass analyzer type MicroTof-Q were used for the HRMS measurements. ¹H-NMR spectra were recorded at 300 MHz and 400 MHz; ¹³C-NMR spectra were recorded at 75 MHz and 100 MHz; CDCl₃ as the solvent. Chemical shifts were reported in the δ scale relative to residual CHCl₃ (7.26 ppm) for ¹H-NMR and to the central line of CDCl₃ (77 ppm) for ¹³C-NMR.

Materials. All commercially available solvents and reagents were used as received. The ¹H and ¹³C NMR spectra for compounds **4eb**,^[1] **4ed**^[2] **4ef**,^[2] are consistent with values previously reported in the literature.

Representative procedure for squaramide-organocatalyzed hydrophosphonylation reaction of aldehydes

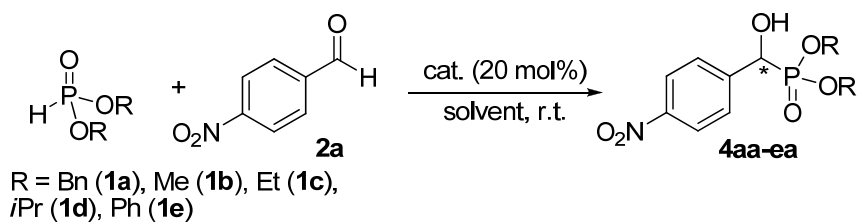
To a mixture of catalyst **3c** (0.02 mmol), and aldehyde **2a-k** (0.1 mmol) in CH₃CN (0.5 mL), phosphite **1e** (0.2 mmol) was further added in a test tube at -38 °C. After the reaction time (see Table 3), adducts **4** were isolated by flash chromatography or medium pressure liquid chromatography (SiO₂, hexane/EtOAc 7:3). Yields and enantioselectivities are reported in Table 3.

¹ A. R. Katritzky, B. V. Rogovoy, A. Y. Mitrokhin, *Arkivoc* **2002**, (xiii), 17.

² X. Zhu, S. Wang, S. Zhou, Y. Wei, L. Zhang, F. Wang, Z. Feng, L. Guo, X. Mu, *Inorg. Chem.* **2012**, 51, 7134.

Screening of Catalysts and Study of the Reaction Conditions

Table S1. Catalytic enantioselective hydrophosphonylation using aldehydes.^[a]



Entry	Catalyst	Solvent	Phosphite	Time (days)	Yield (%) ^[b]	ee (%) ^[c]
1		Toluene (0.5)	1a	2	n.r. ^[d]	n.d. ^[e]
2		Toluene (0.5)	1a	2	n.r. ^[d]	n.d. ^[e]
3		Toluene (0.5)	1a	2	n.r. ^[d]	n.d. ^[e]
4		Toluene (0.5)	1a	2	n.r. ^[d]	n.d. ^[e]
5		Toluene (0.5)	1a	2	n.d. ^[e]	Racemic
6		CH ₃ CN (0.5)	1e	3	52	37

Figure S1. ^1H and ^{13}C NMR spectra of compound **4ea**

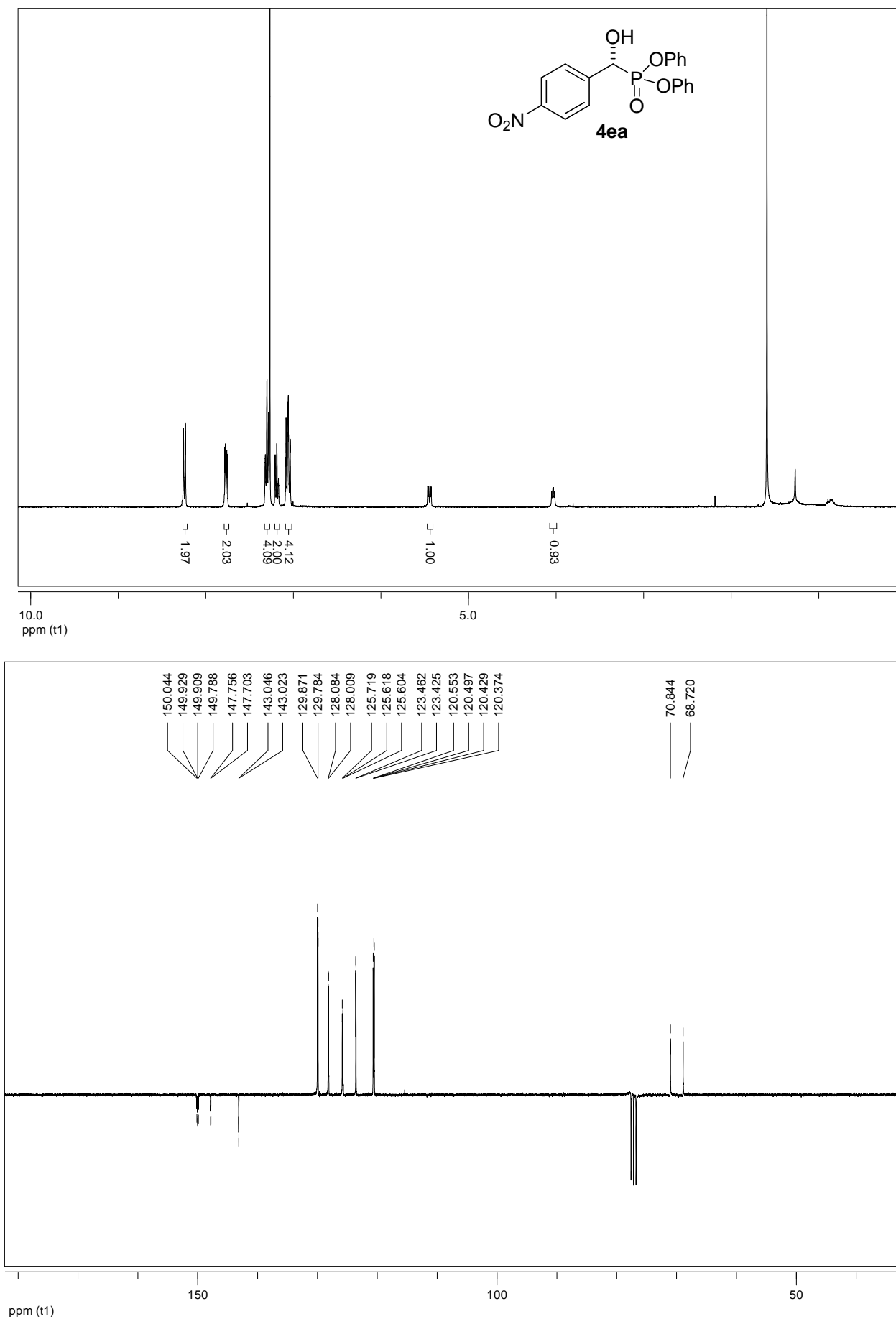


Figure S2. ^1H and ^{13}C NMR spectra of compound **4ec**

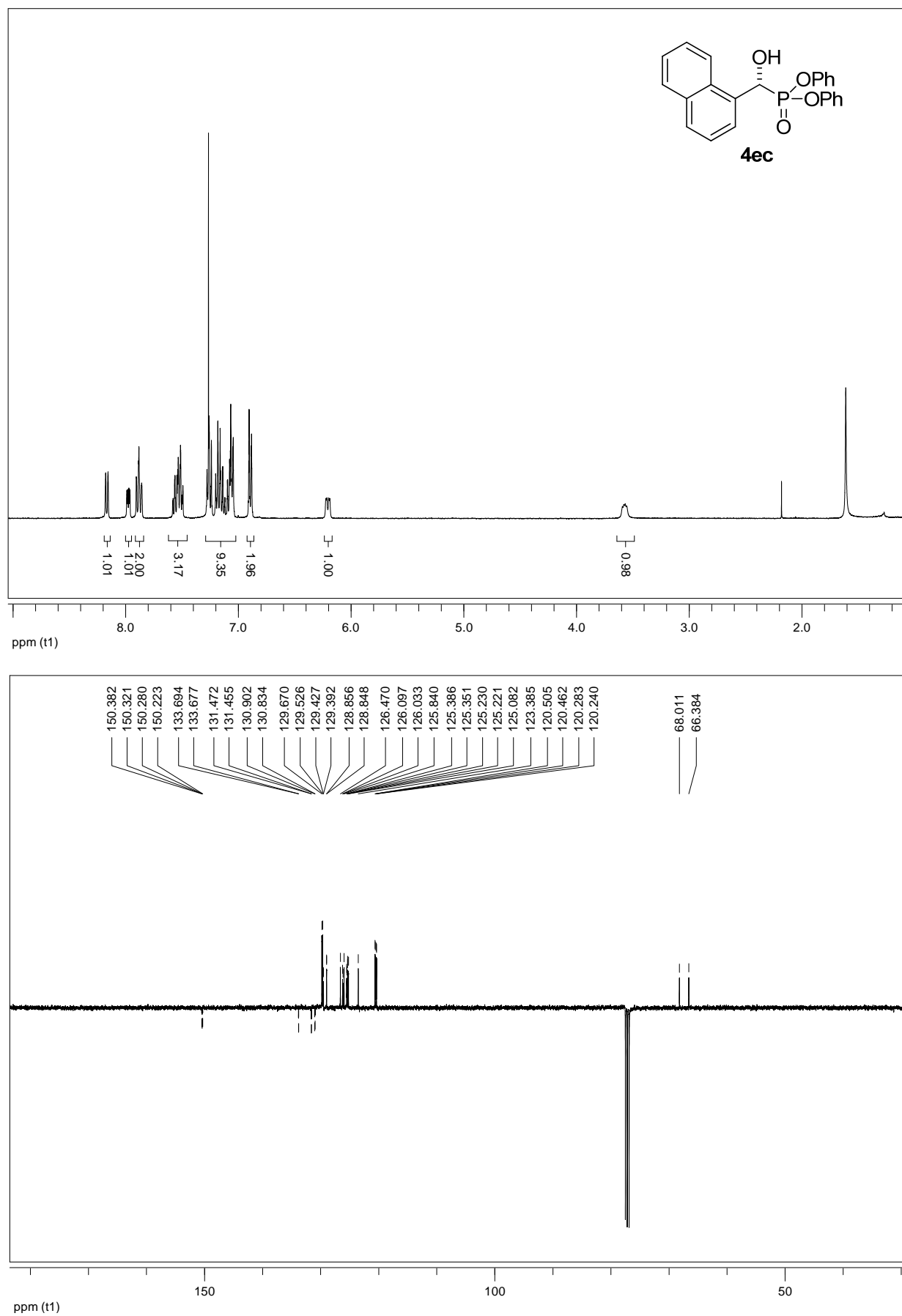


Figure S3. ^1H and ^{13}C NMR spectra of compound **4ee**

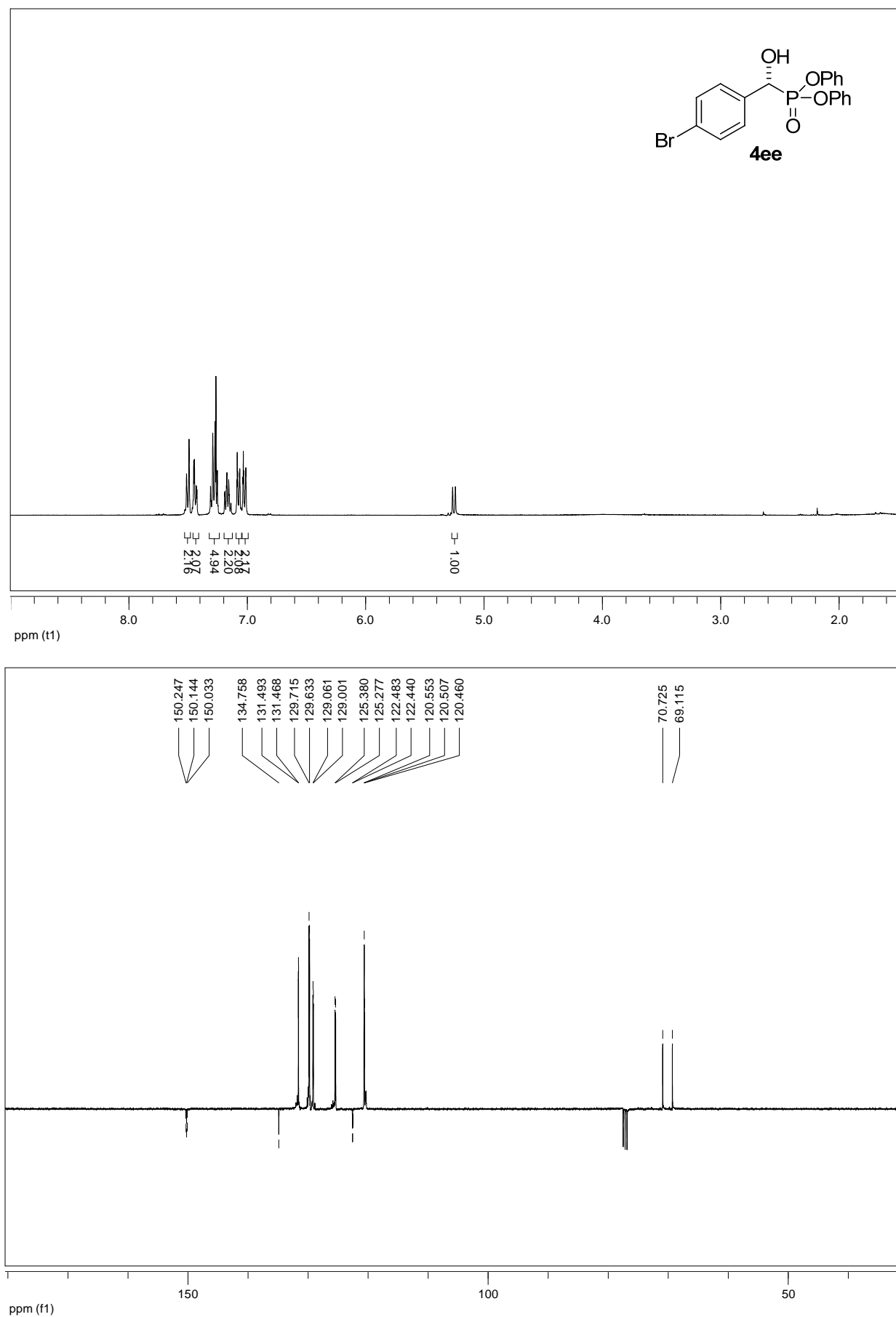


Figure S4. ^1H and ^{13}C NMR spectra of compound **4eg**

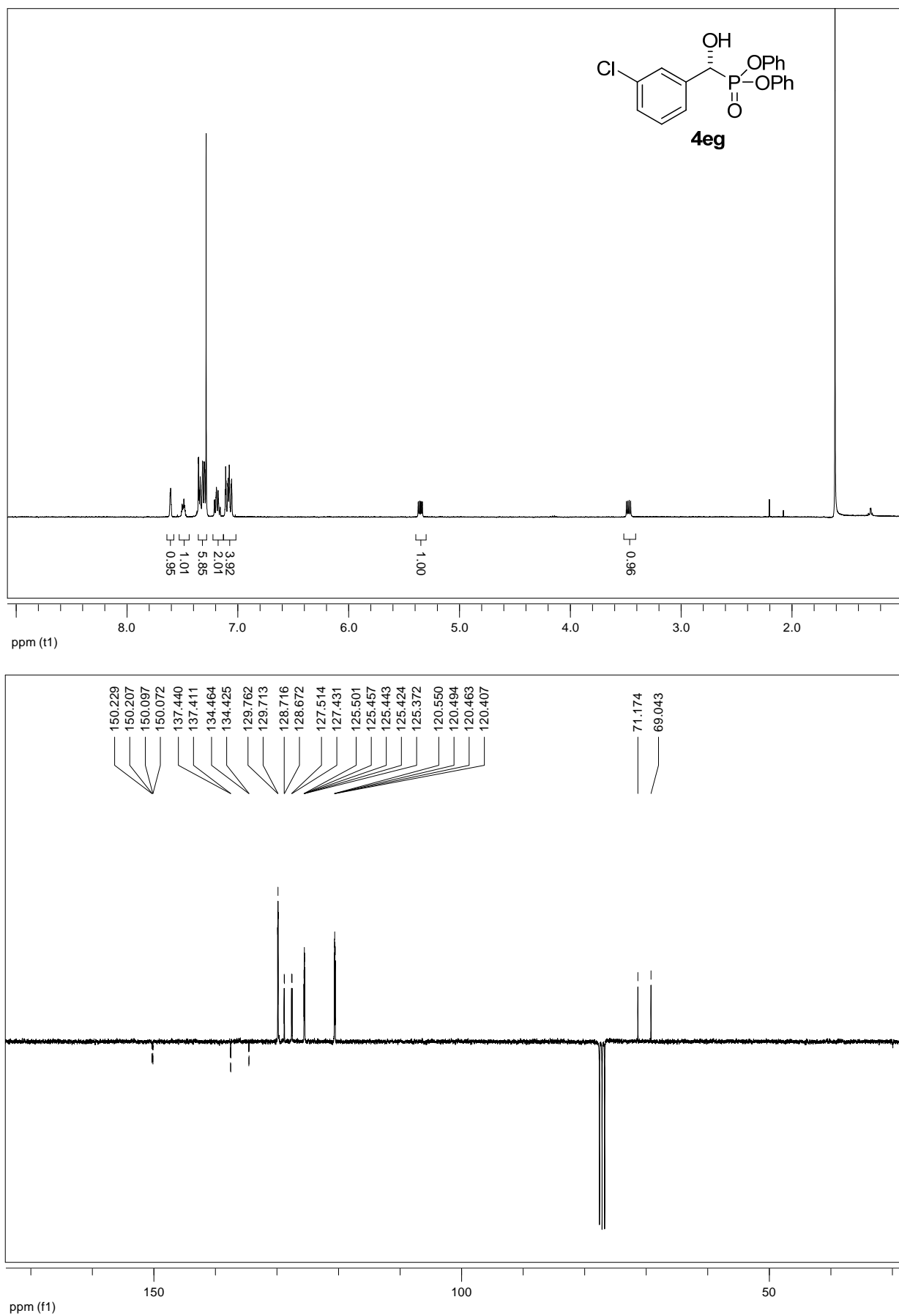


Figure S5. ^1H and ^{13}C NMR spectra of compound **4eh**

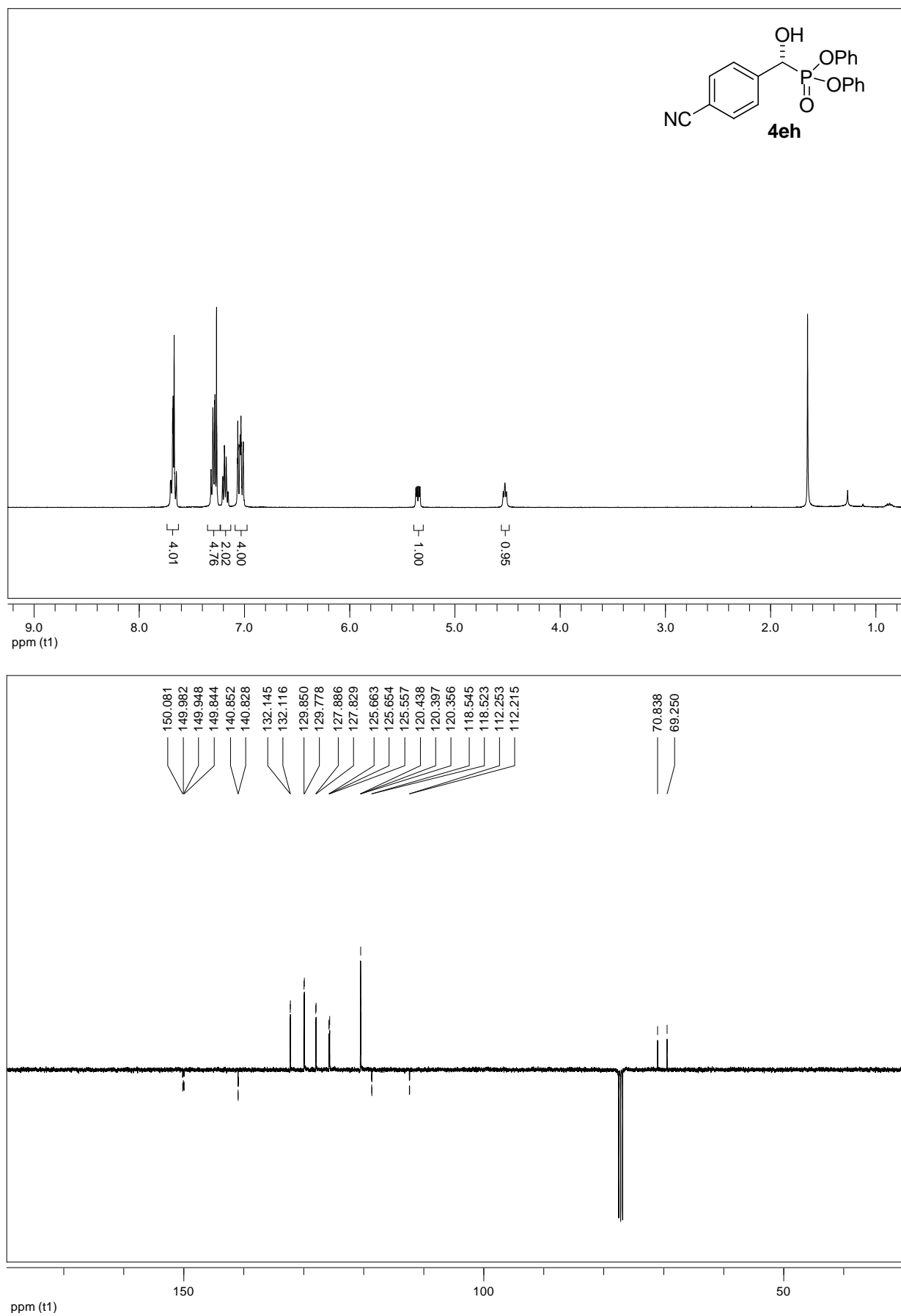


Figure S6. ^1H and ^{13}C NMR spectra of compound **4ei**

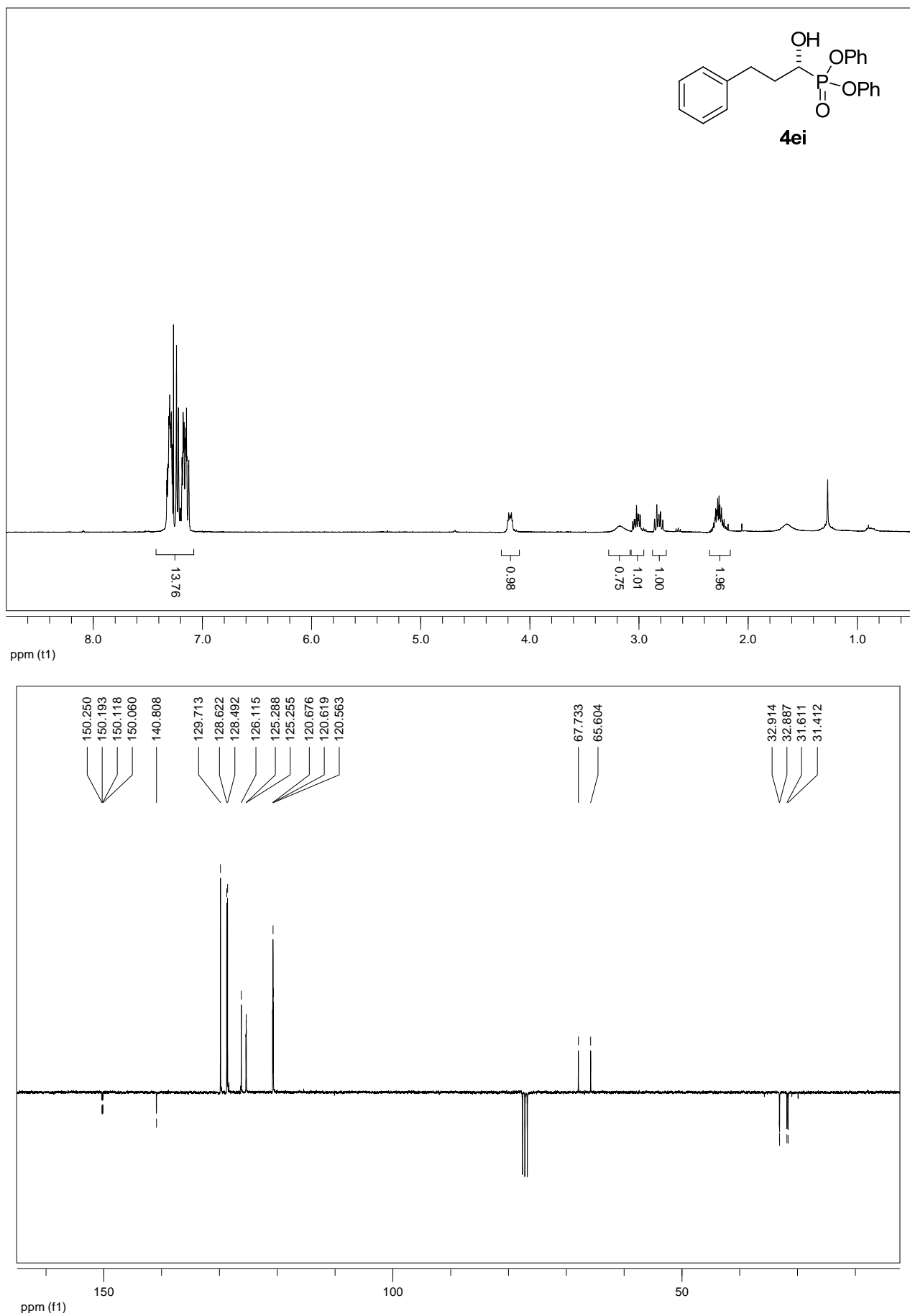


Figure S7. ^1H and ^{13}C NMR spectra of compound **4ej**

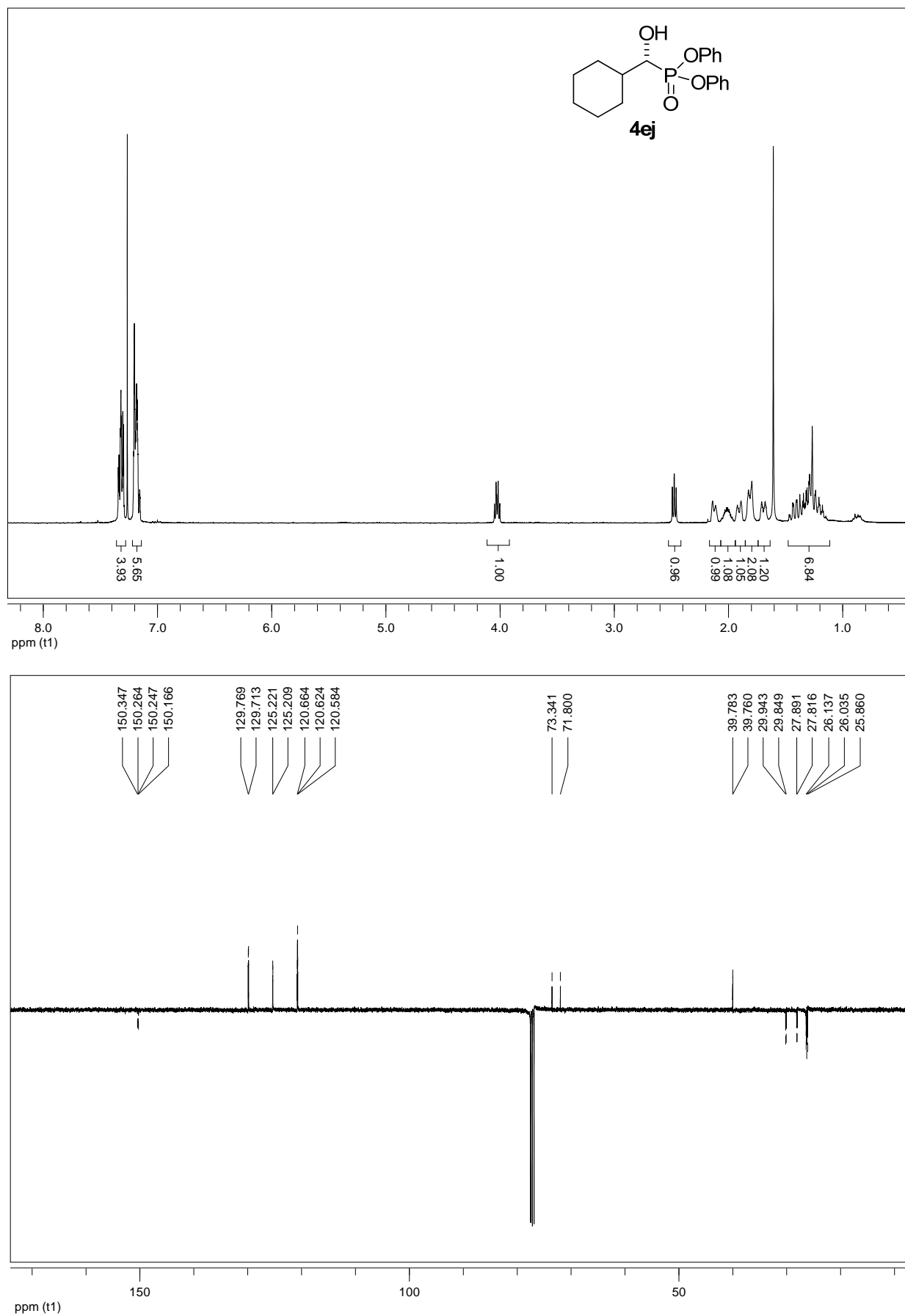
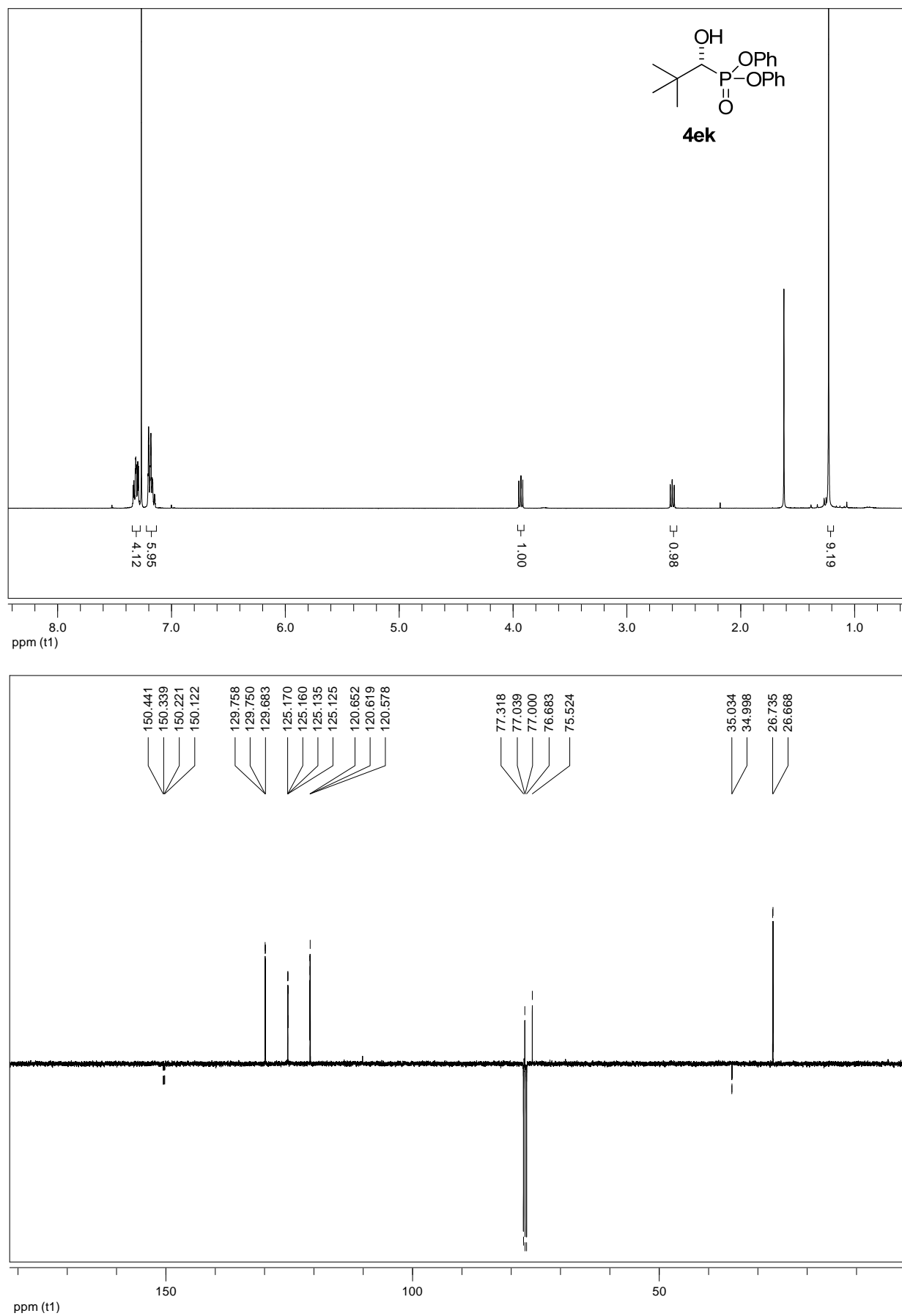
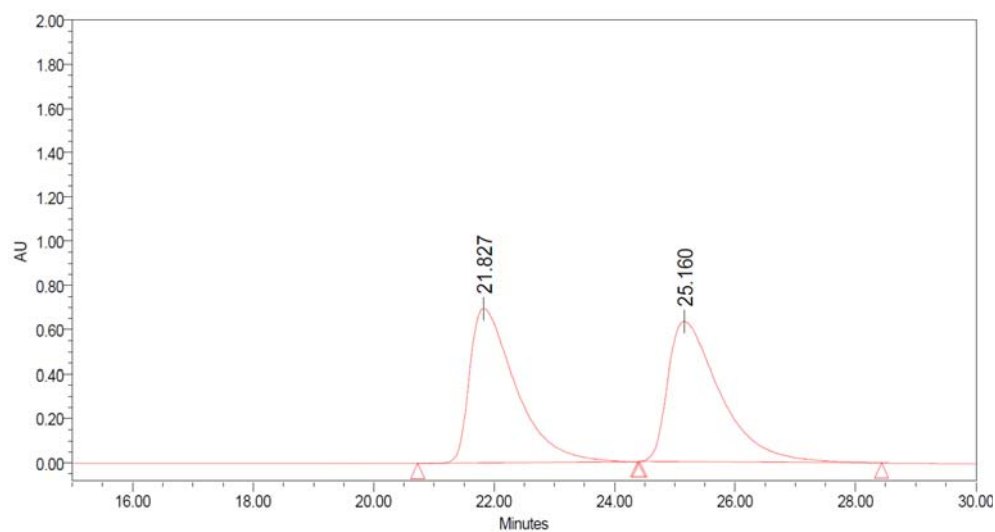


Figure S8. ^1H and ^{13}C NMR spectra of compound **4ek**



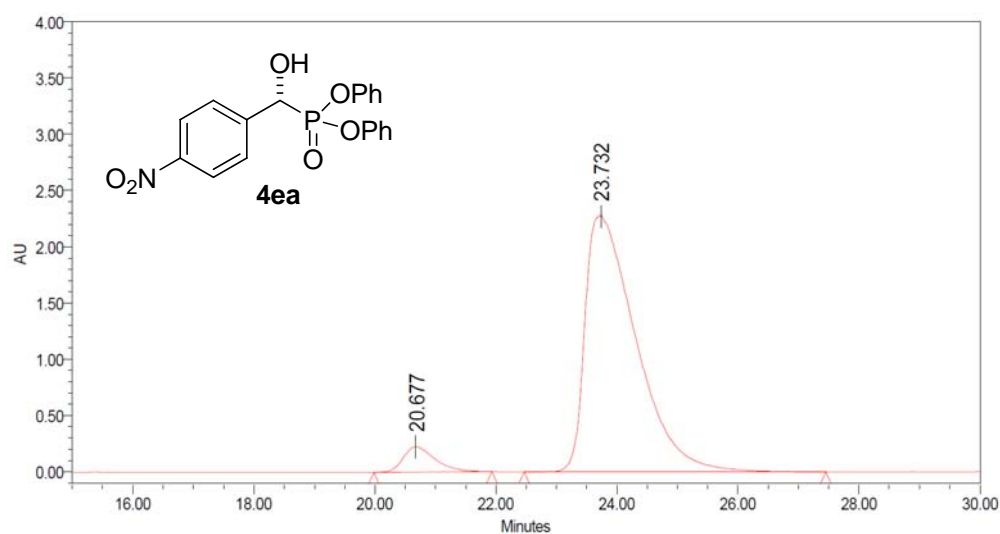
HPLC CROMATOGRAMS OF COMPOUNDS 4



Processed Channel: PDA 268.3 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 268.3 nm	21.827	36965003	49.20	695744
2	PDA 268.3 nm	25.160	38163293	50.80	632692

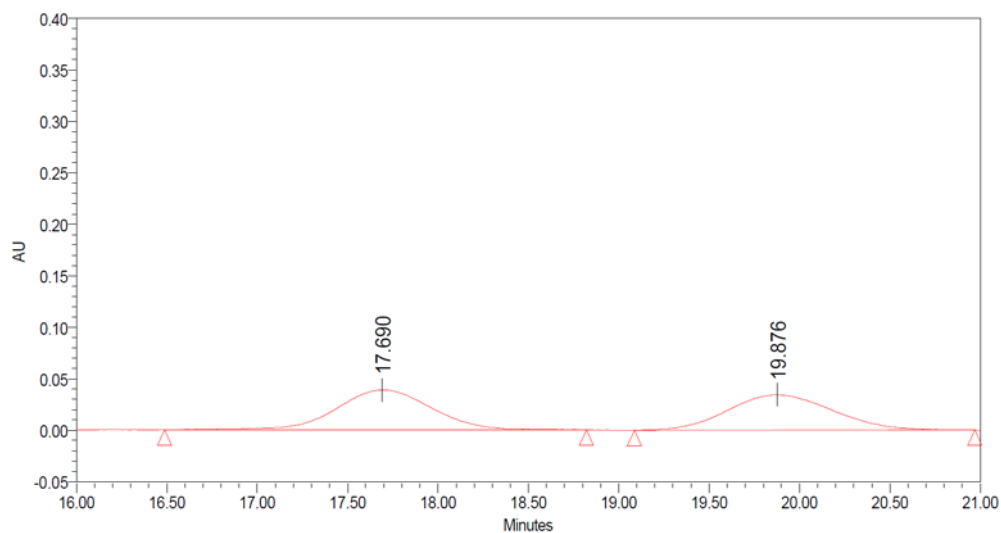
Figure S9. Racemic mixture of **4ea**. Daicel Chiralpak IA column (*n*-hexane/AcOEt = 70:30, flow rate 1 mL/min).



Processed Channel: PDA 268.3 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 268.3 nm	20.677	8332458	5.88	225378
2	PDA 268.3 nm	23.732	133424521	94.12	2280242

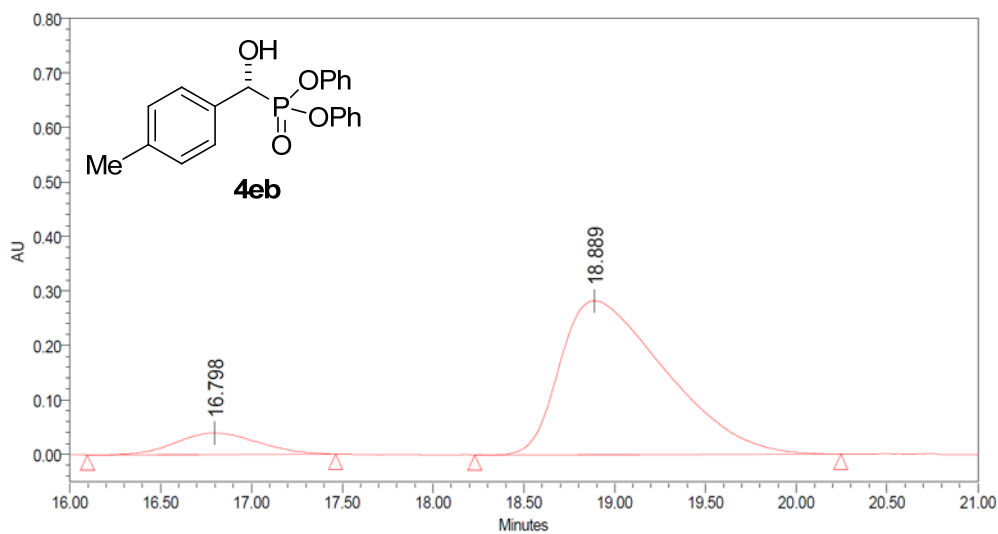
Figure S10. Chiral sample of (*R*)-diphenyl hydroxy(4-nitrophenyl)methylphosphonate (**4ea**)



Processed Channel: PDA 262.0 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 262.0 nm	17.690	1424339	50.89	38732
2	PDA 262.0 nm	19.876	1374490	49.11	34164

Figure S11. Racemic mixture of **4eb**. Daicel Chiralpak IC column (*n*-hexane/*i*-PrOH = 90:10, flow rate 1 mL/min).



Processed Channel: PDA 262.0 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 262.0 nm	16.798	1239109	9.80	39547
2	PDA 262.0 nm	18.889	11407210	90.20	282613

Figure S12. Chiral sample of (*R*)-diphenyl hydroxy(*p*-tolyl)methylphosphonate (**4eb**)

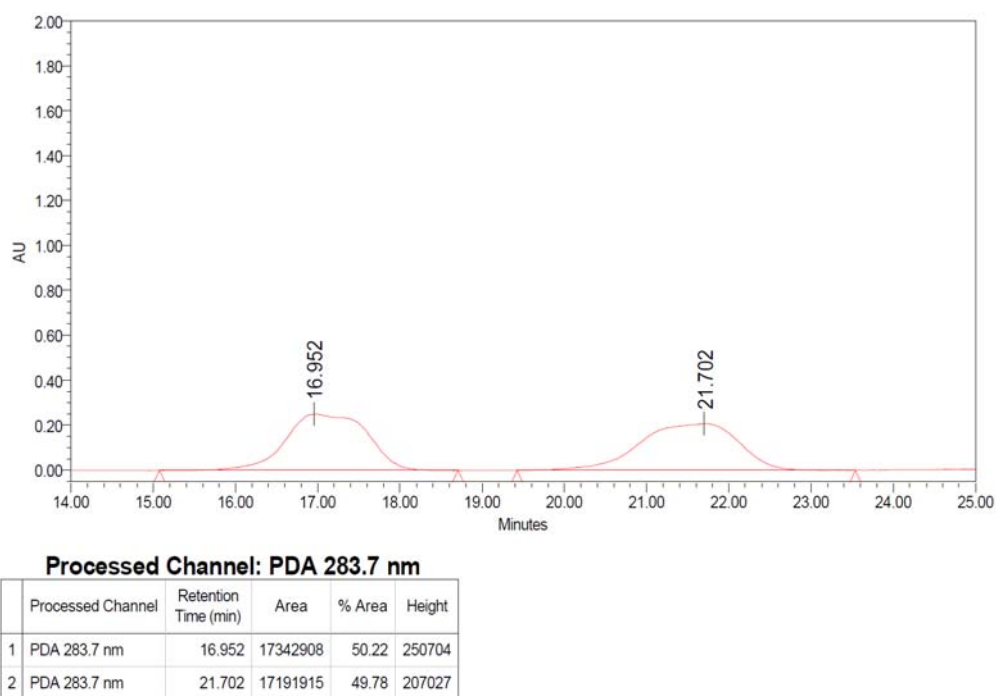


Figure S13. Racemic mixture of **4ec**. Daicel Chiralpak IC column (*n*-hexane/*i*-PrOH = 90:10, flow rate 1 mL/min).

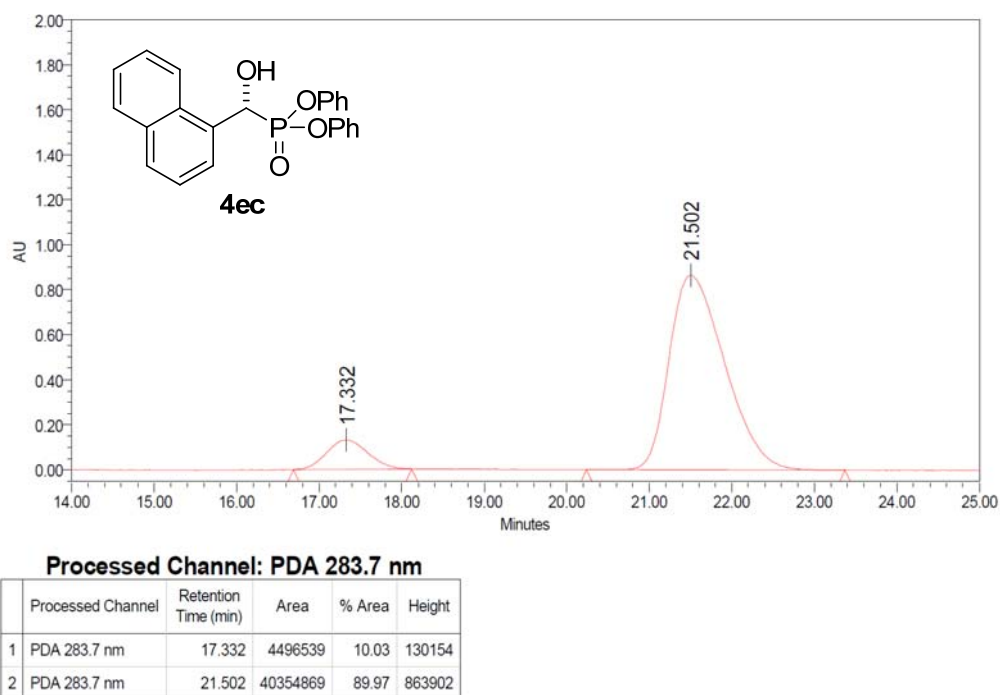
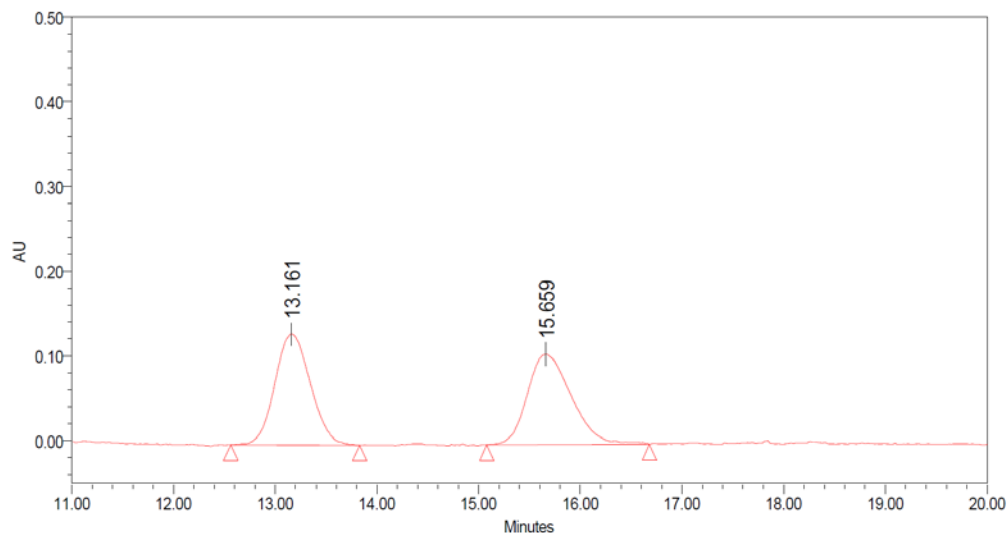


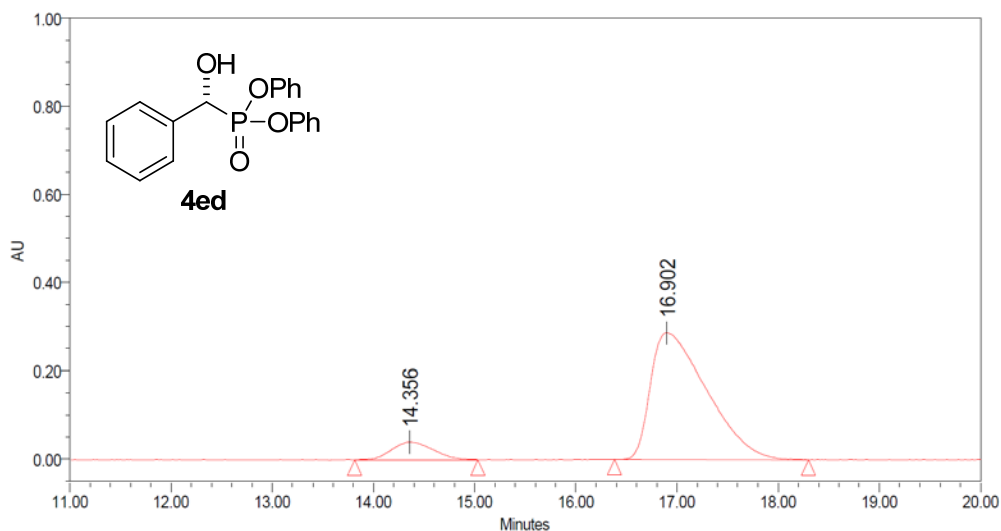
Figure S14. Chiral sample of (*R*)-diphenyl hydroxy(naphthalen-1-yl)methylphosphonate (**4ec**)



Processed Channel: PDA 261.2 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 261.2 nm	13.161	3181004	49.68	131025
2	PDA 261.2 nm	15.659	3222248	50.32	106750

Figure S15. Racemic mixture of **4ed**. Daicel Chiralpak IC column (*n*-hexane/*i*-PrOH = 90:10, flow rate 1 mL/min).



Processed Channel: PDA 261.2 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 261.2 nm	14.356	1133451	9.26	39930
2	PDA 261.2 nm	16.902	11101272	90.74	286940

Figure S16. Chiral sample of (*R*)-diphenyl hydroxy(phenyl)methylphosphonate (**4ed**).

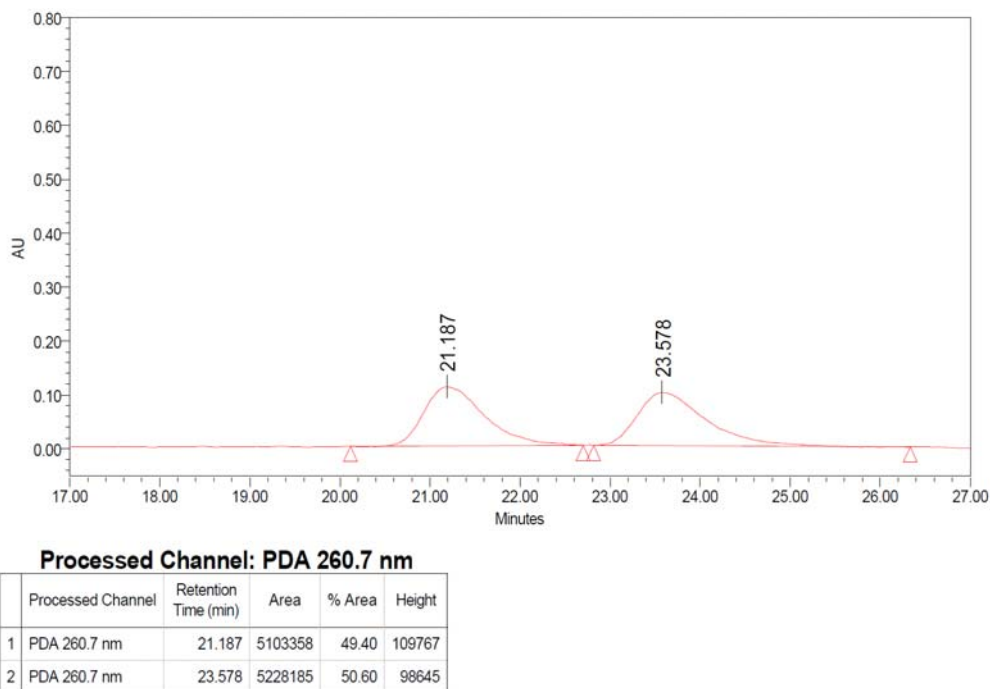


Figure S17. Racemic mixture of **4ee**. Daicel Chiralpak IA column (*n*-hexane/*i*-PrOH = 90:10, flow rate 1 mL/min).

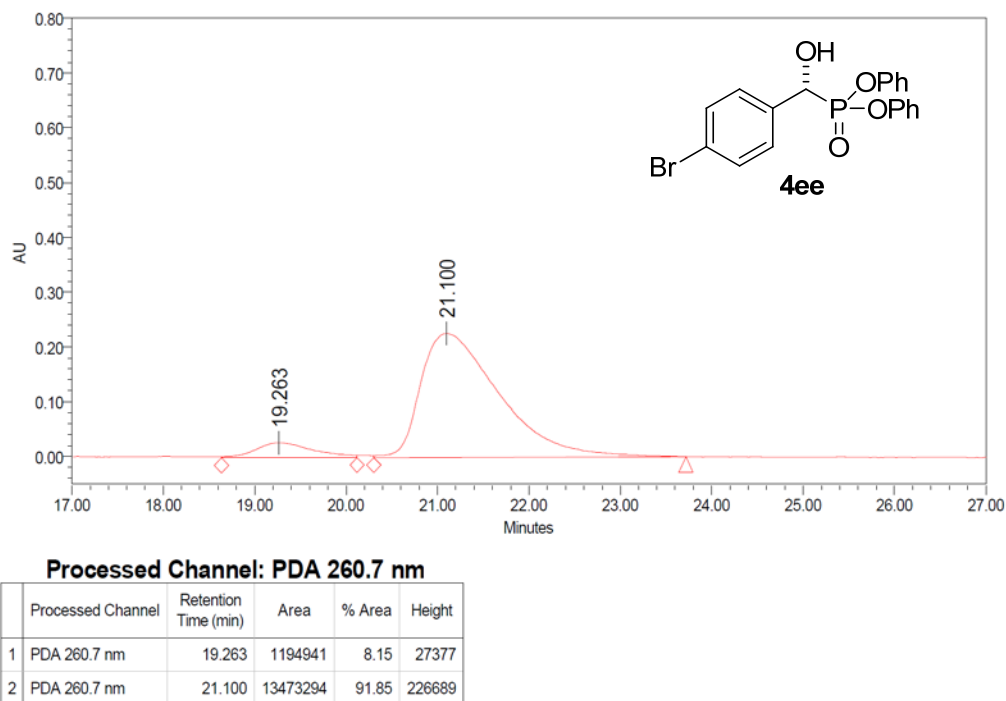
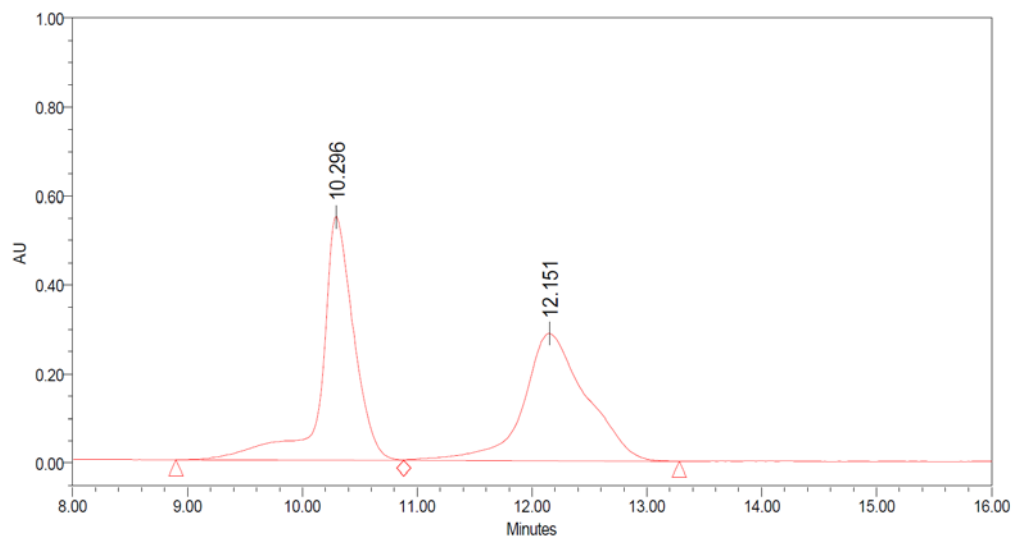


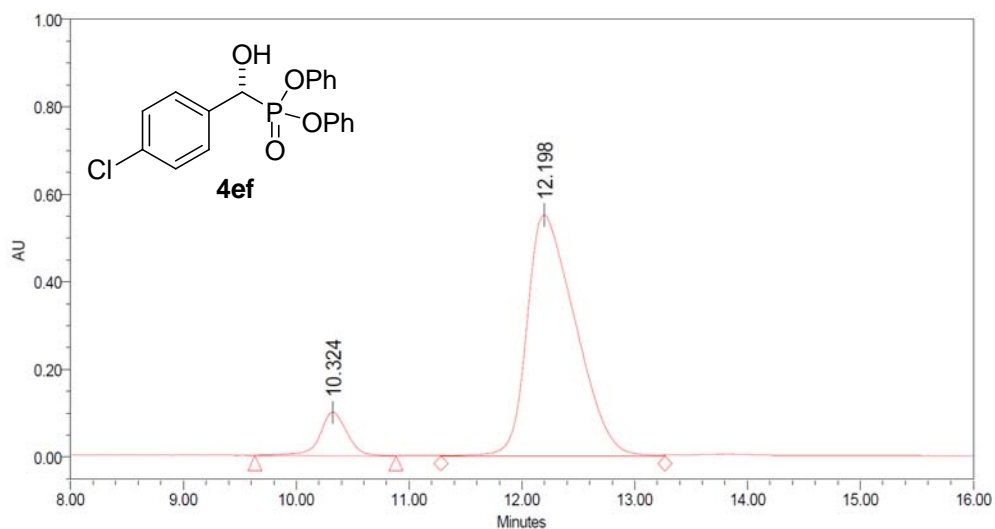
Figure S18. Chiral sample of (*R*)-diphenyl (4-bromophenyl)(hydroxy)methylphosphonate (**4ee**).



Processed Channel: PDA 261.3 nm

Processed Channel	Retention Time (min)	Area	% Area	Height
1 PDA 261.3 nm	10.296	10904399	50.07	548564
2 PDA 261.3 nm	12.151	10874242	49.93	286768

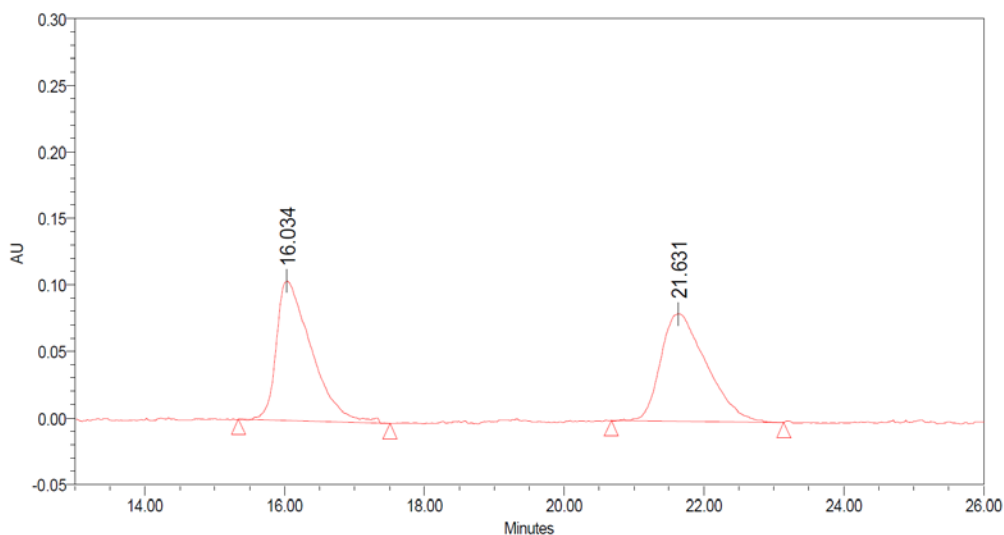
Figure S19. Racemic mixture of **4ef**. Daicel Chiralpak IC column (*n*-hexane/*i*-PrOH = 98:2, flow rate 1 mL/min).



Processed Channel: PDA 261.3 nm

Processed Channel	Retention Time (min)	Area	% Area	Height
1 PDA 261.3 nm	10.324	1640691	9.13	99015
2 PDA 261.3 nm	12.198	16333188	90.87	550764

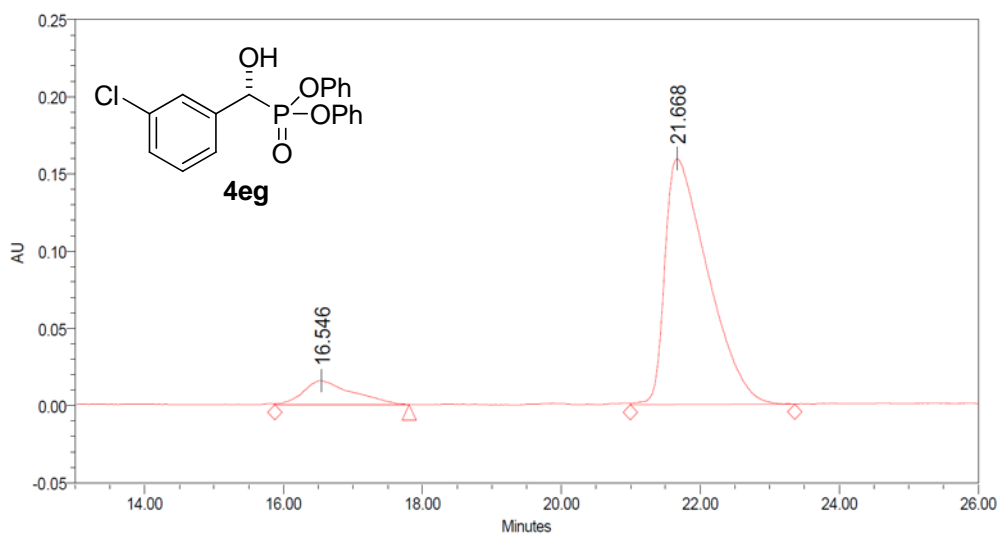
Figure S20. Chiral sample of (*R*)-diphenyl (4-chlorophenyl)(hydroxy)methylphosphonate (**4ef**).



Processed Channel: PDA 272.6 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 272.6 nm	16.034	3639007	49.75	104602
2	PDA 272.6 nm	21.631	3675027	50.25	80550

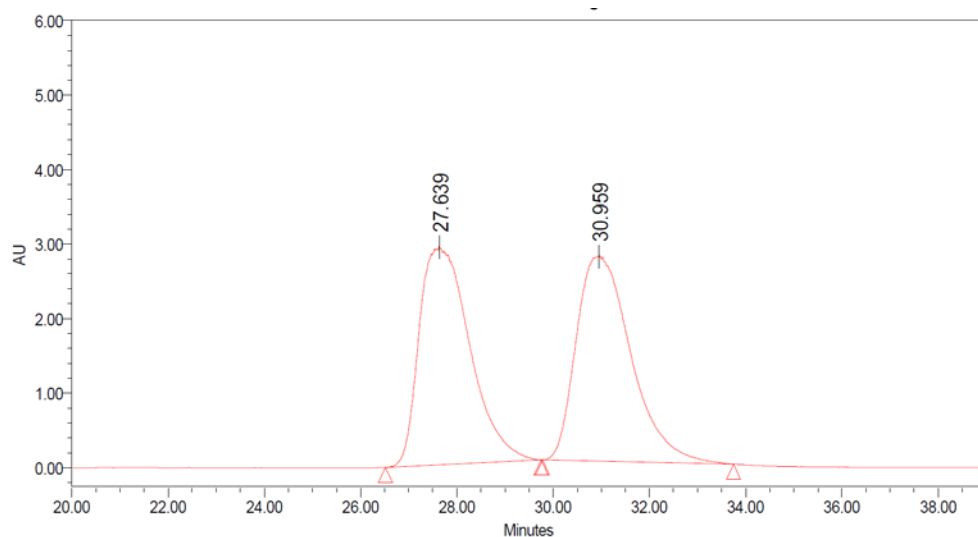
Figure S21. Racemic mixture of **4eg**. Daicel Chiralpak IC column (*n*-hexane/*i*-PrOH = 95:5, flow rate 1 mL/min).



Processed Channel: PDA 272.6 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 272.6 nm	16.546	757006	9.96	15531
2	PDA 272.6 nm	21.668	6843497	90.04	159285

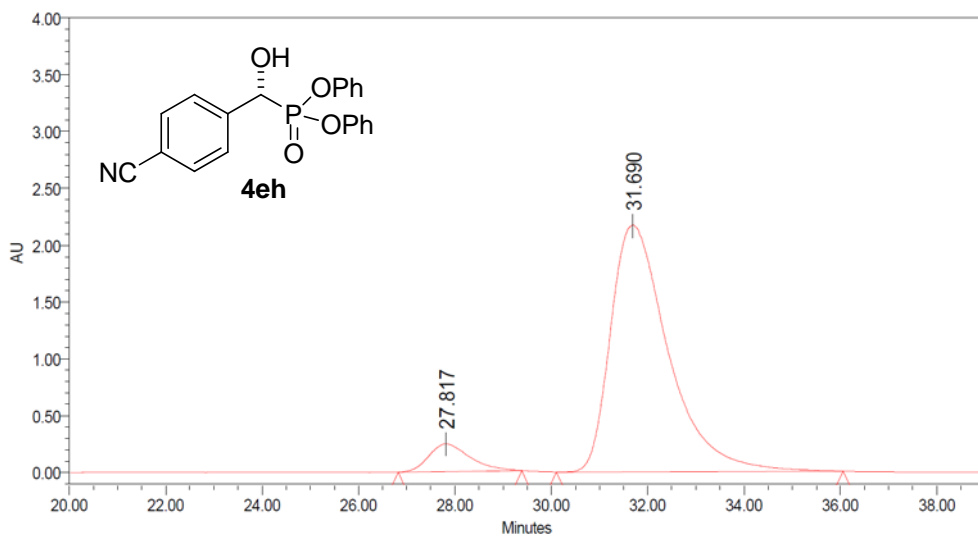
Figure S22. Chiral sample of (*R*)-diphenyl (3-chlorophenyl)(hydroxy)methylphosphonate (**4eg**).



Processed Channel: PDA 236.4 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 236.4 nm	27.639	213162102	49.55	2910449
2	PDA 236.4 nm	30.959	216998274	50.45	2746862

Figure S23. Racemic mixture of **4eh**. Daicel Chiralpak IA column (*n*-hexane/*i*-PrOH = 90:10, flow rate 1 mL/min).



Processed Channel: PDA 236.4 nm

	Processed Channel	Retention Time (min)	Area	% Area	Height
1	PDA 236.4 nm	27.817	14410196	7.39	243508
2	PDA 236.4 nm	31.690	180593876	92.61	2166462

Figure S24. Chiral sample of (*R*)-diphenyl (4-cyanophenyl)(hydroxy)methylphosphonate (**4eh**)

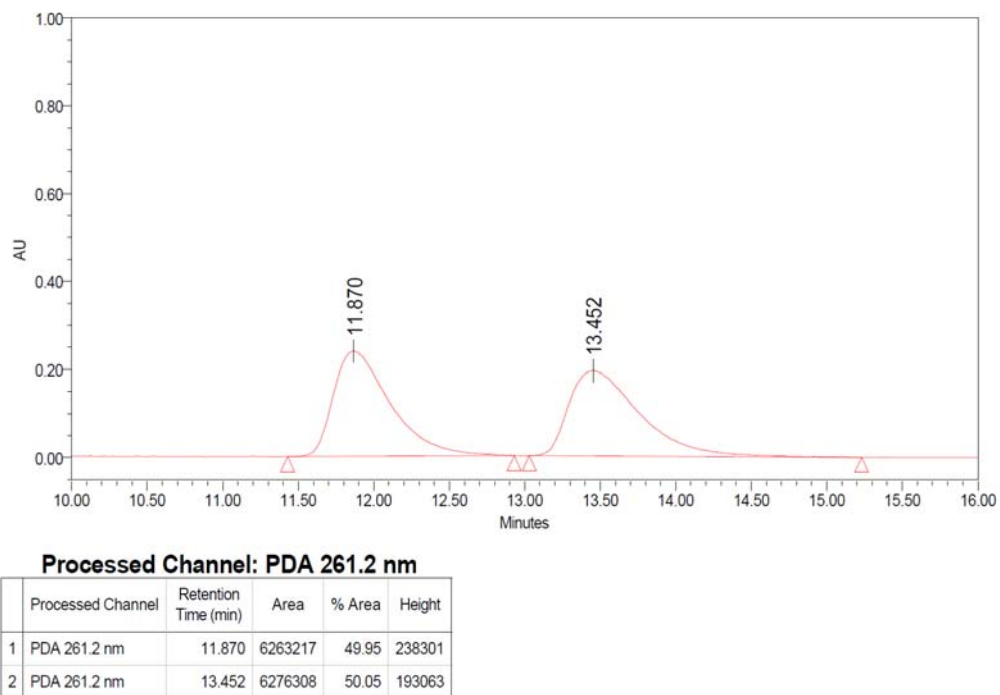


Figure S25. Racemic mixture of **4ei**. Daicel Chiralpak IA column (*n*-hexane/*i*-PrOH = 90:10, flow rate 1 mL/min).

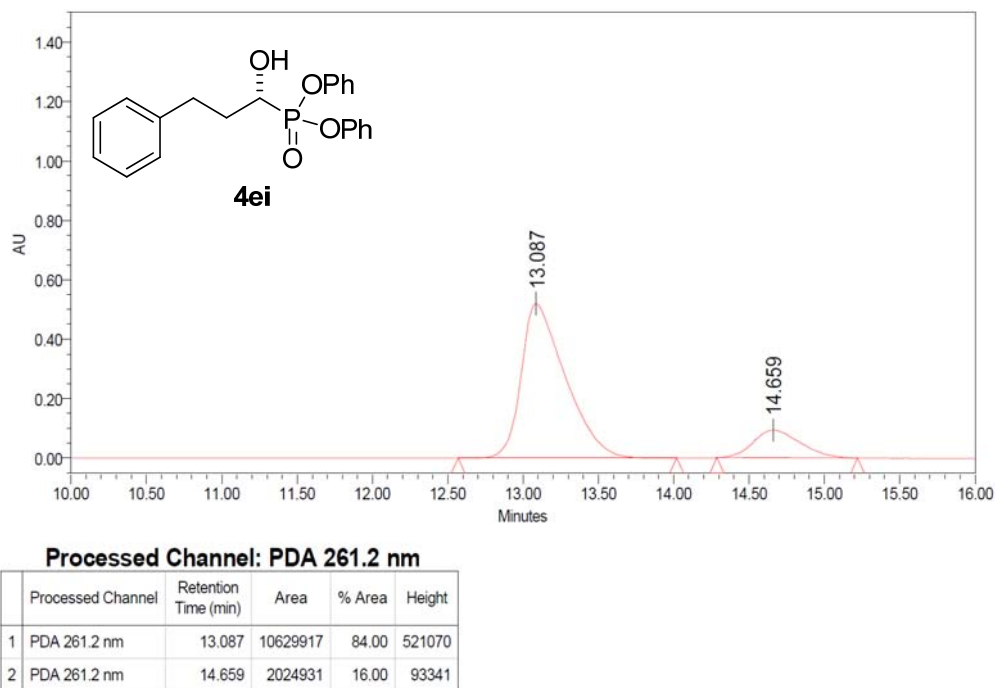
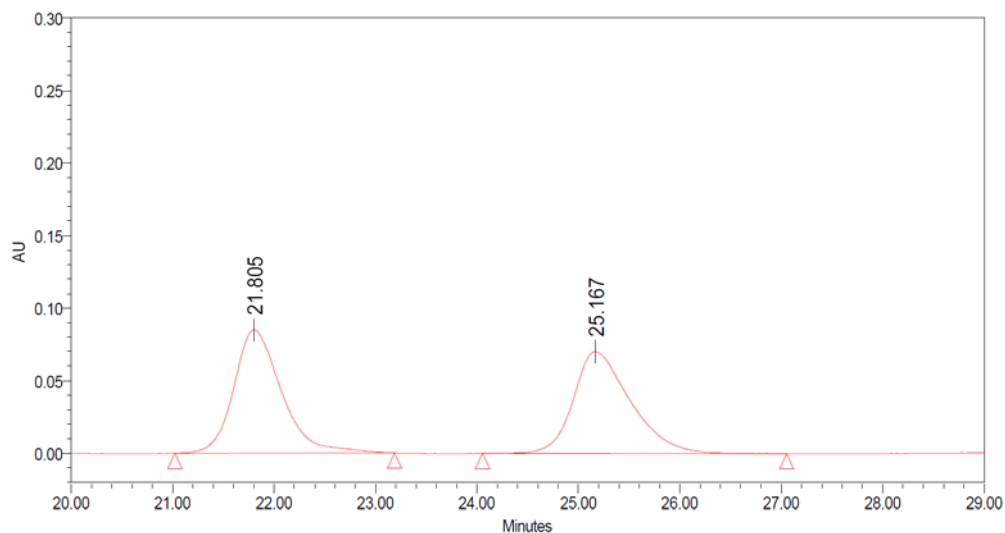


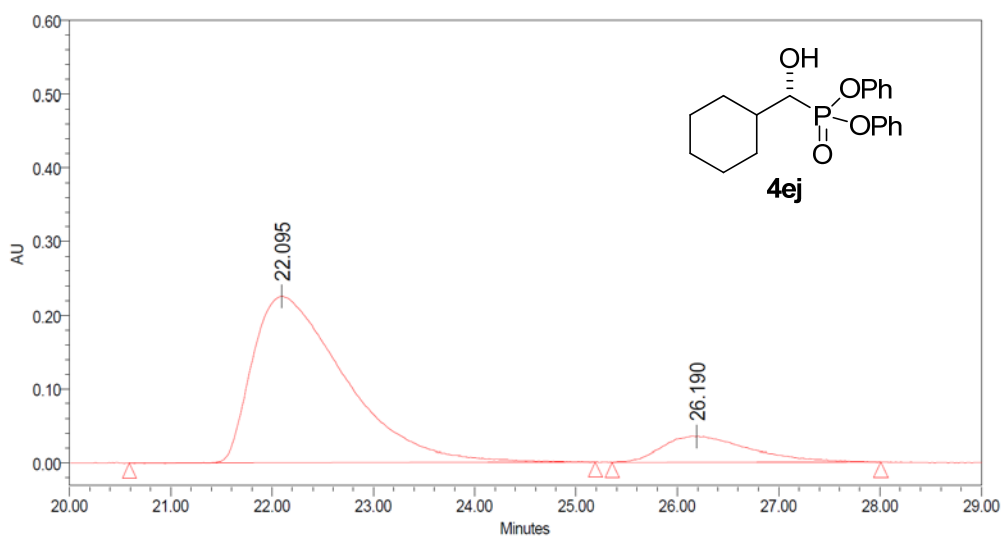
Figure S26. Chiral sample of (*R*)-diphenyl 1-hydroxy-3-phenylpropylphosphonate (**4ei**)



Processed Channel: PDA 262.4 nm

Processed Channel	Retention Time (min)	Area	% Area	Height
1 PDA 262.4 nm	21.805	2784600	50.16	84848
2 PDA 262.4 nm	25.167	2766741	49.84	70156

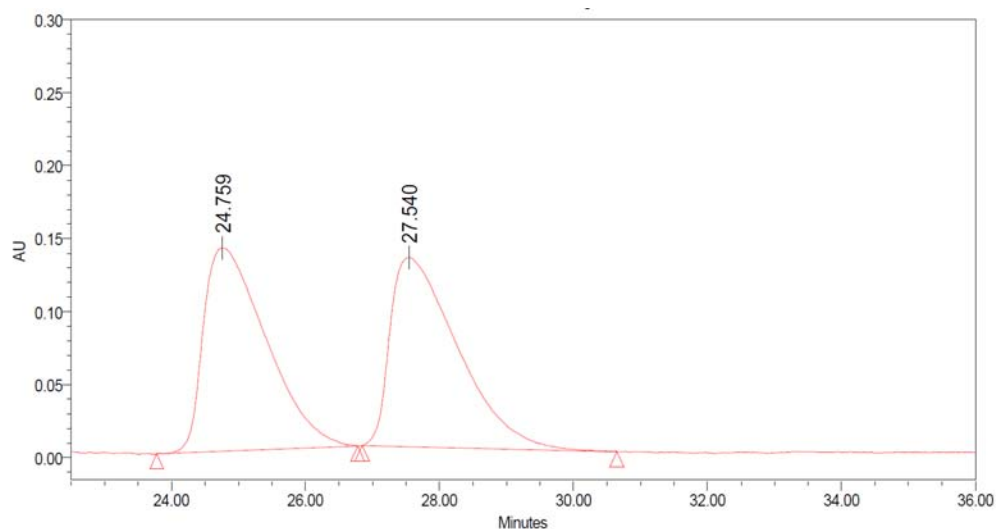
Figure S27. Racemic mixture of **4ej**. Daicel Chiralpak IA column (*n*-hexane/*i*-PrOH = 95:5, flow rate 1 mL/min).



Processed Channel: PDA 262.4 nm

Processed Channel	Retention Time (min)	Area	% Area	Height
1 PDA 262.4 nm	22.095	14249877	87.35	225719
2 PDA 262.4 nm	26.190	2062997	12.65	34386

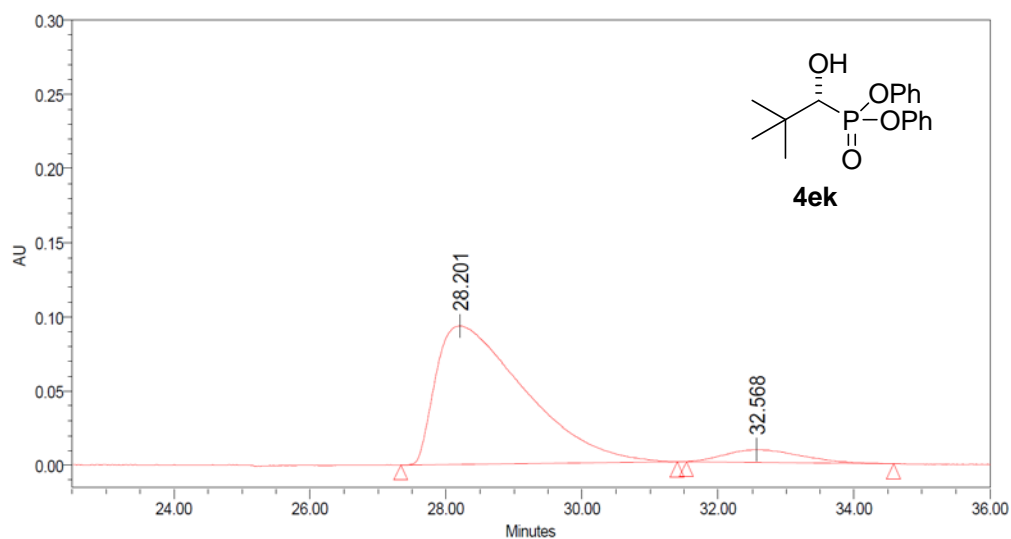
Figure S28. Chiral sample of (*R*)-diphenyl cyclohexyl(hydroxy)methylphosphonate (**4ej**)



Processed Channel: PDA 262.4 nm

Processed Channel	Retention Time (min)	Area	% Area	Height
1 PDA 262.4 nm	24.759	9060405	49.86	139257
2 PDA 262.4 nm	27.540	9110016	50.14	129584

Figure S29. Racemic mixture of **4ek**. Daicel Chiralpak IA column (*n*-hexane/*i*-PrOH = 98:2, flow rate 1 mL/min).



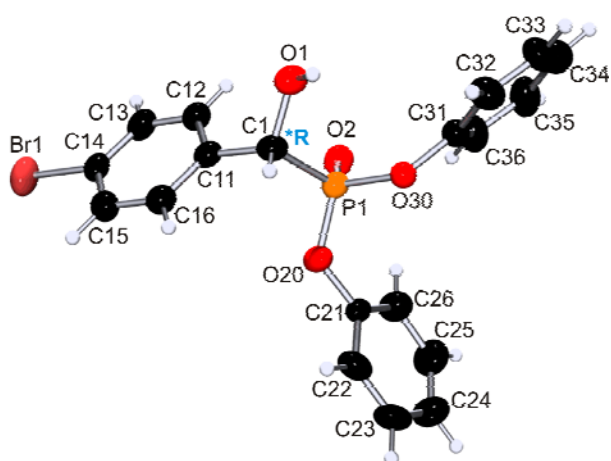
Processed Channel: PDA 262.4 nm

Processed Channel	Retention Time (min)	Area	% Area	Height
1 PDA 262.4 nm	28.201	8470790	92.72	93208
2 PDA 262.4 nm	32.568	665509	7.28	8616

Figure S30. Chiral sample of (*R*)-diphenyl-1-hydroxy-2,2-dimethylpropylphosphonate (**4ek**)

X-RAY CRYSTALLOGRAPHY

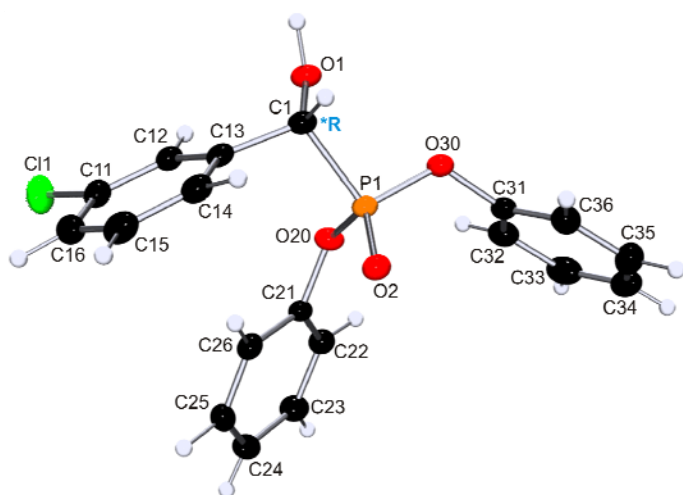
Crystal data, data collection and refinement parameters for complexes **1** and **2** were recorded with a Bruker Kappa APEX2 diffractometer equipped with an area detector and graphite monochromated Mo K α radiation (0.71073 Å). Data reduction was done with the APEX2 software.[3] The structure was solved by direct methods and refined by full-matrix least-squares methods based on F^2 using SHELXL-97 and WinGX programs.[4] Non-hydrogen atoms were refined anisotropically, while hydrogen atoms were positioned geometrically and refined with isotropic displacement parameters according to the riding model. Distance and angle calculations were performed using the SHELXL-97 and WinGX programs.[4]



Crystal data of (**R**)-**4ee**: [C₁₉H₁₆BrO₄P], orthorhombic, $P21212$, $a = 9.2741(10)$ Å, $b = 38.324(4)$ Å, $c = 5.0810(5)$ Å, $Z = 4$, $M_r = 419.20$ g mol⁻¹, $V = 1805.9(3)$ Å³, $D_{\text{calcd}} = 1.542$ g cm⁻³, $\lambda(\text{Mo K}\alpha) = 0.71073$ Å, $T = 100$ K, $\mu = 2.385$ mm⁻¹, 28933 reflections collected, 4788 unique ($R_{\text{int}} = 0.0931$), $R1(F_o) = 0.0599$ [$I > 2\sigma(I)$], $wR2(F_o^2) = 0.1583$ (all data), GOF = 1.038. CCDC 953498.

³ APEX2 Bruker AXS Inc., Madison, Wisconsin, USA, 2011.

⁴ (a) G. M. Sheldrick, SHELXS-97 and SHELXL-97; University of Göttingen, Germany, 1997; (b) Farrugia, L. J. WinGX; University of Glasgow, Great Britain, 1998.



Crystal data of **(R)-4eg**: [C₁₉H₁₆ClO₄P], orthorhombic, *P212121*, *a* = 6.2873(11) Å, *b* = 12.402(2) Å, *c* = 22.054(4) Å, *Z* = 4, *M_r* = 374.74 g mol⁻¹, *V* = 1719.7(5) Å³, *D_{calcd}* = 1.447 g cm⁻³, λ(Mo Kα) = 0.71073 Å, *T* = 100 K, μ = 0.336 mm⁻¹, 19889 reflections collected, 4095 unique (*R_{int}* = 0.0352), *R1*(*F_o*) = 0.0315 [*I* > 2σ(*I*)], *wR2* (*F_o*²) = 0.0883 (all data), GOF = 1.047. CCDC 953499.