# Supramolecular Recognition of Quaternary Phosphonium

## Cations

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## ABSTRACT

9 The modes of supramolecular recognition of quaternary phosphonium cations mediated by 1,1'- bi-2-naphthol (BINOL) are identified and characterized. In contrast to our previous work on ammonium cations, the recognition of the quaternary phosphonium cations via the formation of a 12 PR<sub>4</sub><sup>+</sup>·Br·BINOL ternary complex was found to be mediated by: a hydrogen bond from an  $\alpha$ - carbon center of the phosphonium cation, encapsulation within a continuous hydrogen bond network between the halide–BINOL network, or a combination of these effects working in tandem. The solid state structures of these ternary complexes were analyzed by X-ray crystallography, aided by Hirshfeld surface analysis, to confirm the presence of characteristic intermolecular interactions for the identified modes. In all cases, the quaternary phosphonium cation acts as a

 hydrogen bond donor (HBD) in these supramolecular interactions, and thus is key to the recognition process with BINOL. The characterization of such mechanisms offers insight to the supramolecular and crystal engineering communities in the future design of agents capable of the supramolecular recognition of phosphonium cations and their abstraction from the solution phase.

#### INTRODUCTION

 Understanding the interactions between molecular entities is key to the design of new supramolecular systems. The solution phase recognition of cations has led to significant achievements in the field of supramolecular chemistry. Such recognition has been commonly achieved through hydrogen bond formation between the recognition unit and the target cation. The most commonly studied species are ammonium cations, which have been observed to undergo 28 supramolecular recognition by crown ethers,  $1-4$  cyclodextrins,  $5$  cavitands,  $6,7$  cucurbiturils,  $8,9$ 29 pillar[n]arenes,<sup>10,11</sup> and calix[n]arenes.<sup>12-14</sup> Generally these examples employ protonated 30 ammonium cations where the polarized  $N^+$ -H bond allows strong hydrogen bonding in the 31 host:guest complex (for examples, hydrogen bond donor parameter,  $\alpha = 4.5$  for NEt<sub>3</sub>HBPh<sub>4</sub>).<sup>15</sup> Similarly, for the small number of examples of supramolecular recognition of phosphorus centers, phosphine oxides have garnered the most study due to the strong hydrogen bond acceptor (HBA) 34 properties of the oxide motif (hydrogen bond acceptor parameter,  $\beta = 10.7$ , for Bu<sub>3</sub>P=O).<sup>16,17</sup> BINOL was first identified as a means to recognize and resolve phosphine oxides by direct  $\delta$  hydrogen bonding by Toda *et al*<sup>18</sup> and was recently expanded to include α,α,α',α'-tetraaryl-2,2- disubstituted-1,3-dioxolane-4,5-dimethanol (TADDOL) and dibenzoyltartaric acid (DBTA) as 38 competent recognition units by Bagi *et al.*<sup>19,20</sup> In terms of quaternary phosphonium salts, much less attention has been dedicated to its supramolecular recognition of these species. The 40 importance of such interactions has been recently highlighted by Phipps and coworkers.  $21$  In their

 work, Phipps *et al.* exploit an ion pairing mechanism to perform a supramolecular assembly of a phosphonium substrate and a sulfonate containing iridium catalyst to mediate a regioselective borylation. Due to our group's recent work highlighting the enantioselective recognition of quaternary ammonium cations using enantiopure BINOL via a proposed hydrogen bonding event 45 to the  $\alpha$ -carbon center of the ammonium cation, we sought to explore the possibility of this same 46 behavior in quaternary phosphonium cations. $^{22}$ 

 Based on our previous recognition of ammonium cations, we foresaw that multiple interactions may be possible within the supramolecular recognition of a phosphonium cation. The first possibility was the formation of a continuous hydrogen bond network *via* BINOL-halogen hydrogen bonding [\(Figure 1,](#page-3-0) *Type A*). Such a hydrogen bond network would encapsulate the phosphonium cation, which is bound to the halogen counterion by electrostatic interactions. A second possibility is the expected hydrogen bonding from the BINOL hydroxyl to the halogen counterion, while the α-centre of the phosphonium cation also acts as a HBD to the halogen, 54 forming the ternary complex  $(PR_4^+ \cdot X \cdot BINOL)$  [\(Figure 1,](#page-3-0) *Type B*). In this case, the continuous hydrogen bond network between the BINOL hosts is not strictly necessary to recognize the phosphonium cation, instead it is indirectly recognized using the halogen as a mediator species. The third case would be the hydroxyl groups on the BINOL species participating as a HDB to the 58 halogen counterion, while also acting as a HBA for the  $\alpha$ -centre of the phosphonium cation (Figure [1,](#page-3-0) *Type C*). In this manner, the phosphonium cation would be directly recognized by the BINOL host within the ternary complex. The final possibility would be a combination of the above effects; where the mixture of encapsulation, indirect, and direct recognition mechanisms determine how the phosphonium cation is recognized in the presence of BINOL.



<span id="page-3-0"></span>Figure 1 - Modes of supramolecular recognition

RESULTS

 *Synthesis.* The synthesis of the quaternary phosphonium salts was achieved through simple alkylation of commonly available achiral phosphines with either allyl or benzyl bromide 81 (see [Figure 2\)](#page-4-0). Alkylation of triphenylphosphine yielded  $1 (R_2 = CH = CH_2, 96\%)$  and  $2 (R_2 = Ph,$ 82 91%), and alkylation of tributylphosphine yielded  $3 (R_2 = CH = CH_2, 99%)$ . Upon their synthesis and isolation, single crystals of both **1** and **2** were subsequently obtained and analyzed by X-ray crystallography (see Supplementary Information).

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<span id="page-4-0"></span>Figure 2 - Synthesis of the quaternary phosphonium salts and their respective ternary complexes with (*R*)*-*BINOL

 *Recognition.* The recognition of these quaternary phosphonium salts was then attempted. In concentrated CHCl<sup>3</sup> solutions **1** and **3** formed solid ternary complexes with (*R*)-BINOL to give **4** (74%) and **6** (80%) respectively and were isolated by filtration. **2** failed to yield the desired ternary complex in CHCl3, even upon concentration. When the solvent was changed to ethanol however, the complex **5** formed in lower yield (39%) compared to the previously mentioned 93 ternary complexes. <sup>1</sup>H NMR and melting points of the isolated solids indicated transformation to the desired ternary complexes. However, only crystallographic analysis of the solid-state structure could give further information of the supramolecular recognition.

 *Crystallography.* Single crystals of each ternary complex were grown. **4** readily crystallised in CD3OD to give large clear prisms. Crystals of **5** and **6** were grown by gradual cooling and

 evaporation in ethanol to give large clear prisms and plates respectively. Acquisition of the single crystal diffraction data for each sample gave unambiguous evidence for the formation of the ternary complexes in the solid state. The X-ray crystallographic data for each ternary complex is given in [Table 1.](#page-8-0)

 In the crystal structure of complex **4**, there is clear hydrogen bonding between the phenolic 103 hydroxyl group of the BINOL and the Br counterion (O–H···Br;  $d = 2.349$  Å,  $\theta = 174.11^{\circ}$ ). In addition, another hydrogen bond is also present from the α-carbon center of the phosphonium salt 105 to the bromide counterion (C–H···Br;  $d = 2.826$  Å,  $\theta = 162.72^{\circ}$ ). Such interactions are consistent with supramolecular ternary complex formation between the BINOL, halide and the target phosphonium cation. Interestingly, unlike the previous examples in quaternary ammonium complexation, no continuous hydrogen bonding network between the BINOL and bromide counterions appears to be present, with each BINOL bound to only a single bromide anion (as displayed in [Figure 3](#page-7-0) a). For this reason, complex **4** is designated a *Type B* recognition, where 111 there is no continuous  $BINOL··halogen$  network and the phosphonium is acting as a HBD to the halogen counterion via an acidic α-centre of the phosphonium cation. Minor aryl C–H*···*O interactions (*d* = 2.693 Å, *θ* = 120.68°; *d* = 2.622, *θ* = 124.05° & *d* = 2.630 Å, *θ* = 142.10°) from the electronically deficient phenyl rings of the phosphonium cation to the hydroxyl groups of the 115 BINOL are present.<sup>23</sup> These minor interactions were not classed as forms of recognition (see *Hirshfeld Surface Analysis*, below), however, their presence in the crystal structure should not be discounted as the full intricacies of this phenomenon is not fully understood.

 The crystal structure of complex **5** presented also contained strong interactions between the BINOL and bromide counterion. A strong hydrogen bond between the bromide counterion and 120 BINOL was immediately evident (O–H···Br;  $d = 2.442 \text{ Å}, \theta = 177.77^{\circ}$ ). However, in contrast to

 complex **4**, the packing in this crystal structure revealed a continuous hydrogen bonding network linking each BINOL moiety to two adjacent bromide anions (see [Figure 3](#page-7-0) b). A hydrogen bond 123 from the acidic α-carbon center of the phosphonium cation is also observed (C–H···Br;  $d = 2.652$ ) 124 Å,  $\theta = 172.00^{\circ}$ ), allowing for the formation of the ternary complex. There is also a noteworthy aryl 125 C–H···O hydrogen bond ( $d = 2.409$  Å,  $\theta = 164.81^{\circ}$ ), from the phenyl ring of the phosphonium to an adjacent hydroxyl group of the BINOL species. In this way, it is shown that the recognition of this phosphonium salt is mediated by all three types of hypothesized recognition modes: a *Type A* encapsulation mechanism (*via* the continuous BINOL-halogen hydrogen bond network), a *Type B* indirect recognition of the α-center of the phosphonium cation acting as a HBD to the halogen counterion and a *Type C* direct recognition via aryl C–H*···*O hydrogen bonding to the hydroxyl HBA on the BINOL species. This demonstrates that that an acidic α-centre on the phosphonium is not the only possible HBD motif in this recognition process and that electron deficient aryl rings are also sufficient for hydrogen bonding to the BINOL species. It also highlights the variety in recognition modes that can be responsible for the formation of one ternary complex.

 Analysis of complex **6** revealed the expected hydrogen bonding between the BINOL and bromide 136 counterion (O–H···Br;  $d = 2.349$  Å,  $\theta = 174.11^{\circ}$ ) (see [Figure 3](#page-7-0) c). This crystal structure also had a continuous hydrogen bonding network between the BINOL and bromide counterions, which appear to encapsulate the phosphonium cation, consistent with a *Type A* recognition. However, no close contacts between the phosphonium cation and the bromide counterion could be classed as 140 hydrogen bonding. Instead, C–H···O hydrogen bonds (C–H···O;  $d = 2.517 \text{ Å}, \theta = 161.39^{\circ}, d =$ 141 2.575 Å,  $\theta$  = 126.28°) from the α-carbon centre of the phosphonium cation acting as the HBD and the BINOL hydroxyl group acting as a HBA were present. This demonstrates that both *Type A* and *Type C* modes of recognition are responsible for complex formation.



- <span id="page-7-0"></span>
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<span id="page-8-0"></span>

	4	5	6
Empirical formula	$C_{41}H_{34}BrO_2P$	$C_{45}H_{36}BrO_2P$	$C_{35}H_{46}BrO_2P$
Formula weight	669.56	719.62	609.60
Temperature/K	120.0	120.0	120.0
Crystal system	monoclinic	monoclinic	orthorhombic
Space group	P2 <sub>1</sub>	P2 <sub>1</sub>	$P2_12_12_1$
$a/\text{\AA}$	10.9509(5)	8.8952(15)	12.5247(5)
$b/\text{\AA}$	14.5495(7)	15.823(3)	15.2027(6)
$c/\text{\AA}$	11.0440(6)	12.673(2)	16.4528(7)
$\alpha$ /°	90	90	90
$\beta$ /°	113.207(2)	96.325(7)	90
$\gamma$ <sup>o</sup>	90	90	90
Volume/Å <sup>3</sup>	1617.26(14)	1772.9(5)	3132.8(2)
Z	$\overline{2}$	$\overline{2}$	$\overline{4}$
$\rho_{calc}g/cm^3$	1.375	1.348	1.292
$\mu$ /mm <sup>-1</sup>	1.356	1.243	1.393
F(000)	692.0	744.0	1288.0
Crystal size/mm <sup>3</sup>	$0.273 \times 0.266 \times 0.162$	$0.281 \times 0.222 \times 0.096$	$0.276 \times 0.234 \times 0.216$
Radiation	MoKα ( $\lambda$ = 0.71073)	MoKa ( $\lambda$ = 0.71073)	MoKα ( $λ = 0.71073$ )
20 range for data collection/°	4.012 to 63.166	4.608 to 54.158	4.088 to 64.27
Index ranges	$-16 \le h \le 16, -21 \le k \le$ $21, -16 \le l \le 16$	$-11 \le h \le 11, -20 \le k \le$ 20, $-16 \le l \le 16$	$-18 \le h \le 18$ , $-22 \le k \le$ 22, $-24 \le l \le 24$
Reflections collected	37669	30239	77675
Independent reflections	10748 [ $R_{int} = 0.0555$ , $R_{\text{sigma}} = 0.0726$	7774 $[R_{int} = 0.1175,$ $R_{sigma} = 0.1416$	10995 [ $R_{int} = 0.0744$ , $R_{\text{sigma}} = 0.0625$
Data/restraints/parameters	10748/1/408	7774/1/444	10995/0/403
Goodness-of-fit on $F^2$	0.993	0.930	0.985
Final R indexes $[I>=2\sigma(I)]$	$R_1 = 0.0352$ , w $R_2 =$ 0.0711	$R_1 = 0.0524$ , w $R_2 =$ 0.1029	$R_1 = 0.0323$ , w $R_2 =$ 0.0618
Final R indexes [all data]	$R_1 = 0.0490$ , w $R_2 =$ 0.0740	$R_1 = 0.0968$ , w $R_2 =$ 0.1174	$R_1 = 0.0489$ , w $R_2 =$ 0.0655
Largest diff. peak/hole / e $\AA$ <sup>-3</sup>	$0.52/-0.49$	$0.51/-0.85$	$0.45/-0.38$
Flack parameter	0.041(4)	0.012(9)	0.016(3)

166 Table 1 – Crystallographic data for the ternary complexes

 *Hirshfeld Surface analysis.* Further characterization of the recognition modes within the crystal structures was achieved using Hirshfeld surface analysis [\(Figure 4\)](#page-10-0). <sup>24</sup> Highlighting the C–H*···*Br- interactions in each plot qualitatively displays the presence and strength of this interaction in each crystal structure (blue traces). In complexes **4** and **5** [\(Figure 4](#page-10-0) a & b) a sharp characteristic fingerprint is revealed indicative of a hydrogen bond present in these crystals, further confirming the observation of both crystals having a *Type B* component in their recognition modes. Complex **6** however, has a much weaker and more diffuse fingerprint for this interaction [\(Figure 4](#page-10-0) c). While this interaction is present in the crystal, it is unlikely to be a defined hydrogen bond as its overall contribution to the entire crystal packing is minimal representing just 2.6% of the surface area of the plot. This is likely indicative to the fact that this crystal has been identified to have both *Type A* and *Type C* recognition, which does not involve strong C–H···Br interactions.

 The C–H*···*O interactions with the hydroxyl group of the BINOL acting as a HBA could also be clearly examined (purple traces). In complex **4**, noticeably more diffuse, and therefore weaker, C– H*···*O interactions are present, consistent with the absence of *Type C* direct recognition. Instead the 182 plot for complex 4 is dominated by the C–H···Br interactions indicative of *Type B* recognition. In 183 complex 5, equally sharp traces for C–H···Br, and C–H···O interactions are visible, confirming the present of both *Type B* and *Type C* recognition modes in this ternary complex. Finally, the Hirshfeld plot for complex **6** displays the overwhelming contribution of the C–H*···*O hydrogen bonding observed in this crystal (5.8% of the plot), showing strong evidence for a *Type C* recognition in this complex, with little to no *Type B* character.



<span id="page-10-0"></span>Figure 4 – Expanded unit cells and Hirshfeld fingerprint plots for (a) complex **4**, (b) complex **5**, and (c) complex 6; highlighting the  $P^{\dagger}$ -CH···Br interactions (blue) and  $P^{\dagger}$ -CH···O (purple) observed in each ternary complex

*BINOL<sup>\*</sup>··X<sup><i>-</sup>* packing. To view the impact of the different modes of recognition on the packing of</sup> each complex, both the bromide and BINOL species are represented as their Van der Waals surface. The phosphonium cations could then be more easily viewed within the architecture of the 192 BINOL<sup>\*</sup>···Br scaffold [\(Figure 5\)](#page-12-0). Complex **4**, which displays *Type B* recognition, can be seen in [Figure 5](#page-12-0) a. In this rendering, the non-continuous BINOL network is apparent. Instead, the packing is much closer to a standard co-crystallization between the BINOL and the phosphonium salt where the halide counter-ion is more accessible to the phosphonium cation.

 Complexes **5** [\(Figure 5](#page-12-0) b) and **6** [\(Figure 5](#page-12-0) c) both show much more compact packing structure around the phosphonium cation – indicative of a *Type A* recognition – in which the cation is 198 encapsulated within a continuous BINOL ··· Br hydrogen bond network. The phosphonium cation is accommodated in visible cavities within this hydrogen bond network. Interestingly, this is despite the differences in geometry between the respective phosphonium cations within the two crystal structures. This demonstrates that the encapsulation of the cation is a flexible process – rather than the BINOL only able to accommodate one specific shape, the cavities found within these crystal structures are not of a fixed volume or geometry.

 We speculate that initial recognition of the phosphonium cation in complexes **5** and **6** involves either indirect (*via* the halogen counterion, *Type B*) or direct (*Type C*) hydrogen bond interactions between a BINOL species and the phosphonium cation in solution. The BINOL then propagates the growth of the complex through a second hydrogen bond to an adjacent halogen counterion, thus 'recruiting' a second phosphonium cation. These supramolecular interactions greatly increase the rate of nucleation until a critical mass is reached, when the whole recognition network has greatly reduced solubility and the complex is pulled from solution as a microcrystalline solid.



<span id="page-12-0"></span>Figure 5 – The resulting packing in the crystal structures of the recognition complexes, where (a) is complex **4**, (b) is complex **5**, and (c) is complex **6**. The van der Waals radii for each BINOL (teal) and bromide (brown) species is shown, with the phosphonium cations represented as stick representations within the crystal structure

### CONCLUSION

 This work characterizes the modes of supramolecular recognition of quaternary phosphonium cations in the solid phase by BINOL. Three modes have been hypothesized and subsequently identified within the recognition complexes presented here. Complex **4** arises from *Type B* interactions, while complex **5** has a mixture of *Type A, Type B* and *Type C* character as identified by single crystal X-ray crystal structure and Hirshfeld fingerprint plots. Complex **6** contains *Type A & C* recognition motifs in its solid state structure. The crystal packing of each recognition complex was influenced by these modes of interaction and presents new crystal engineering opportunities in the co-crystallisation of BINOL with ionic species.

#### EXPERIMENTAL

 Reagents and solvents used in this work were commercially available (*Sigma Aldrich*; tributylphosphine, triphenylphosphine. *Fluorochem*; (*R*)*-*BINOL, allyl bromide, benzyl bromide) and were used without further purification. NMR spectra were recorded on either a *Bruker* Avance 225 III HD-400 spectrometer with operating frequencies of 400.07 MHz for  $\rm{^1H}$ ; 100.60 MHz for  $\rm{^{13}C}$ 226 and; 161.95 MHz for  ${}^{31}P$  at 298 K. Melting points are uncorrected. X-ray single crystal diffraction data was collected using a *Bruker* D8 Venture (Photon100 CMOS detector, ImSmicrosource, focusing mirrors). The diffractometer was equipped with an *Oxford Cryosystems* cryostream, with open-flow nitrogen cryostats set at a temperature of 120.0 K. All structures were solved by direct 230 methods and refined by full-matrix least squares on F2 for all data using Olex2 (V 13.0)<sup>25</sup> and 231 SHELXTL<sup>26,27</sup> software. All non-disordered non-hydrogen atoms were refined anisotropically and 232 hydrogen atoms were placed in the calculated positions and refined in riding mode. Crystallographic data and related CIFs for the structures related to the submitted publication have

 been deposited with the *Cambridge Crystallographic Data Centre* as supplementary publications: CCDC 2048561–2048565. Hirshfeld surfaces and fingerprint plots were generated using *Crystal Explorer* (V  $17.5)^{24}$  and full unedited fingerprint plots are available in the Supplementary Information. Full descriptions for the synthesis of all compounds and characterization data are described in the Supplementary Information.

#### **Typical alkylation procedure.** *Synthesis of Allyltributylphosphonium bromide* (**3**).

 To a neat solution of tributylphosphine (2.47 mL, 10.0 mmol) was added allyl bromide (0.95 mL, 11.0 mmol) dropwise over 15 mins with stirring. Caution: the reaction is highly exothermic. The solution was allowed to react at room temperature for 30 minutes. The viscous reaction mixture cooled upon completion of the alkylation and crystallised as a colourless solid. The solid was washed with diethyl ether (3 x 15 mL) and dried in vacuo to afford 3 as a colourless white solid (3.18 g, 99% yield).

1 H NMR (400 MHz, CDCl3) δ 5.80 – 5.60 (m, 1H), 5.49 (ddd, *J* = 16.8, 3.8, 1.0 Hz, 1H), 5.38 (ddd, *J* = 10.0, 4.4, 1.1 Hz, 1H), 3.49 (dd, *J* = 15.7, 7.5 Hz, 2H), 2.52 – 2.29 (m, 6H), 1.76 – 1.24 (m, 12H), 0.90 (q, *J* = 7.1 Hz, 9H). <sup>13</sup>C NMR (101 MHz, CDCl<sub>3</sub>) δ 124.3 (d, *J* = 11.7 Hz), 124.0 (d, *J* = 9.9 Hz), 25.1 (d, *J* = 47.0 Hz), 23.8 (d, *J* = 15.3 Hz), 23.6 (d, *J* = 4.9 Hz), 18.8 (d, *J* = 47.0 251 Hz), 13.3. <sup>31</sup>P NMR (202 MHz, CDCl<sub>3</sub>)  $\delta$  31.32. HRMS (ESI-TOF) m/z: [M]<sup>+</sup> Calculated for  $252 \text{ C}_{15} \text{H}_{23} \text{P}^{\text{+}}$ : 243.2242, found 243.2269. mp: 47–50 °C. IR (max/cm<sup>-1</sup>): 2961s, 2871s, 1634m, 1464m, 1232m, 1093m, 929m, 719m, 597m.

 **Typical complexation procedure.** *Allyltributylphosphonium bromide · (R)-1,1′-bi-2-naphthol*  (**6**).

 The quaternary phosphonium bromide salt **3** (0.969 g, 3.00 mmol) was dissolved in CHCl<sup>3</sup> (1.5 mL, 2.0 M) in a 10 mL vial. Solid (*R*)-BINOL (0.858 g, 1.0 equiv) was then added with stirring to the solution, resulting in a pale-yellow homogeneous solution. This solution was allowed to stir at room temperature overnight, which produced the desired complexed product **6** as a white precipitate. The resulting solid complex was isolated by vacuum filtration (1.47 g, 80% yield).

1 H NMR (400 MHz, DMSO-d6) δ 9.26 (s, 2H), 7.96 – 7.74 (m, 4H), 7.39 (d, *J* = 8.8 Hz, 2H), 7.22 (ddd, *J* = 8.1, 6.7, 1.3 Hz, 2H), 7.16 (ddd, *J* = 8.3, 6.7, 1.4 Hz, 2H), 6.96 (dd, *J* = 8.4, 1.2 Hz, 2H), 5.81 (dddd, *J* = 17.4, 10.0, 7.5, 4.8 Hz, 1H), 5.51 – 5.39 (m, 1H), 5.35 (ddd, *J* = 10.0, 12 4.0, 1.6 Hz, 1H), 3.26 (dd, *J* = 14.7, 7.3 Hz, 2H), 2.33 – 2.13 (m, 6H), 1.49 (tdd, *J* = 10.9, 8.2, 5.8 Hz, 6H), 1.39 (h, *J* = 7.0 Hz, 6H), 0.90 (t, *J* = 7.2 Hz, 9H). <sup>13</sup>C NMR (101 MHz, DMSO-d<sub>6</sub>) δ 153.0, 134.1, 267 128.5, 128.1, 127.8, 125.7, 125.5 (d, *J* = 9.4 Hz), 124.4, 123.0 (d, *J* = 11.9 Hz), 122.2, 118.5, 115.4, 268 23.8, 23.3 (d, *J* = 15.7 Hz), 22.6 (d, *J* = 4.5 Hz), 17.4 (d, *J* = 47.2 Hz), 13.2. <sup>31</sup>P NMR (202 MHz, 269 DMSO-d<sub>6</sub>)  $\delta$  32.18. HRMS (ESI-TOF) m/z: [M]<sup>+</sup> calculated C<sub>15</sub>H<sub>32</sub>P<sup>+</sup>: 243.2242, found 243.2263. [M-H]<sup>-</sup> calculated C<sub>20</sub>H<sub>13</sub>O<sub>2</sub><sup>-</sup>: 285.0921, found 285.0929. mp: 140–141 °C (MeOH). IR (max/cm<sup>-</sup> 271 <sup>1</sup>): 3165br, 2954m, 1622m, 1504m, 1324m, 1269s, 964m, 818s, 683m. XRD: Sample was crystallised in ethanol, to give clear colorless prisms. Crystal data: orthorhombic, space group 273 *P*2<sub>1</sub>2<sub>1</sub>2<sub>1</sub> (no. 19).

### ASSOCIATED CONTENT

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The manuscript was written by M.O.K. and M.P.W. All authors have given approval to the final

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M.P.W.

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- ABBREVIATIONS
- HBD hydrogen bond donor, HBA hydrogen bond acceptor, BINOL 1,1′-bi-2-naphthol, *d* –
- interatomic distance, *θ*  bond angle
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