1 Supramolecular Recognition of Quaternary Phosphonium

2 Cations

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8 ABSTRACT

9 The modes of supramolecular recognition of quaternary phosphonium cations mediated by 1,1'-10 bi-2-naphthol (BINOL) are identified and characterized. In contrast to our previous work on 11 ammonium cations, the recognition of the quaternary phosphonium cations via the formation of a 12 $PR_4^+ \cdot Br^- \cdot BINOL$ ternary complex was found to be mediated by: a hydrogen bond from an α -13 carbon center of the phosphonium cation, encapsulation within a continuous hydrogen bond 14 network between the halide–BINOL network, or a combination of these effects working in tandem. 15 The solid state structures of these ternary complexes were analyzed by X-ray crystallography, 16 aided by Hirshfeld surface analysis, to confirm the presence of characteristic intermolecular 17 interactions for the identified modes. In all cases, the quaternary phosphonium cation acts as a

18 hydrogen bond donor (HBD) in these supramolecular interactions, and thus is key to the 19 recognition process with BINOL. The characterization of such mechanisms offers insight to the 20 supramolecular and crystal engineering communities in the future design of agents capable of the 21 supramolecular recognition of phosphonium cations and their abstraction from the solution phase.

22 INTRODUCTION

23 Understanding the interactions between molecular entities is key to the design of new 24 supramolecular systems. The solution phase recognition of cations has led to significant 25 achievements in the field of supramolecular chemistry. Such recognition has been commonly 26 achieved through hydrogen bond formation between the recognition unit and the target cation. The 27 most commonly studied species are ammonium cations, which have been observed to undergo supramolecular recognition by crown ethers,^{1–4} cyclodextrins,⁵ cavitands,^{6,7} cucurbiturils,^{8,9} 28 pillar[n]arenes,10,11 and calix[n]arenes.12-14 Generally these examples employ protonated 29 30 ammonium cations where the polarized N⁺-H bond allows strong hydrogen bonding in the host:guest complex (for examples, hydrogen bond donor parameter, $\alpha = 4.5$ for NEt₃HBPh₄).¹⁵ 31 32 Similarly, for the small number of examples of supramolecular recognition of phosphorus centers, 33 phosphine oxides have garnered the most study due to the strong hydrogen bond acceptor (HBA) properties of the oxide motif (hydrogen bond acceptor parameter, $\beta = 10.7$, for Bu₃P=O).^{16,17} 34 35 BINOL was first identified as a means to recognize and resolve phosphine oxides by direct hydrogen bonding by Toda *et al*¹⁸ and was recently expanded to include $\alpha, \alpha, \alpha', \alpha'$ -tetraaryl-2,2-36 37 disubstituted-1,3-dioxolane-4,5-dimethanol (TADDOL) and dibenzoyltartaric acid (DBTA) as competent recognition units by Bagi et al.^{19,20} In terms of quaternary phosphonium salts, much 38 39 less attention has been dedicated to its supramolecular recognition of these species. The importance of such interactions has been recently highlighted by Phipps and coworkers.²¹ In their 40

41 work, Phipps *et al.* exploit an ion pairing mechanism to perform a supramolecular assembly of a 42 phosphonium substrate and a sulfonate containing iridium catalyst to mediate a regioselective 43 borylation. Due to our group's recent work highlighting the enantioselective recognition of 44 quaternary ammonium cations using enantiopure BINOL via a proposed hydrogen bonding event 45 to the α -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 47 48 may be possible within the supramolecular recognition of a phosphonium cation. The first 49 possibility was the formation of a continuous hydrogen bond network via BINOL-halogen 50 hydrogen bonding (Figure 1, Type A). Such a hydrogen bond network would encapsulate the 51 phosphonium cation, which is bound to the halogen counterion by electrostatic interactions. A 52 second possibility is the expected hydrogen bonding from the BINOL hydroxyl to the halogen 53 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, *Type B*). In this case, the continuous 55 hydrogen bond network between the BINOL hosts is not strictly necessary to recognize the 56 phosphonium cation, instead it is indirectly recognized using the halogen as a mediator species. 57 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 α-centre of the phosphonium cation (Figure 59 1, Type C). In this manner, the phosphonium cation would be directly recognized by the BINOL 60 host within the ternary complex. The final possibility would be a combination of the above effects; 61 where the mixture of encapsulation, indirect, and direct recognition mechanisms determine how 62 the phosphonium cation is recognized in the presence of BINOL.

63



Figure 1 - Modes of supramolecular recognition

78 RESULTS

79 Synthesis. The synthesis of the quaternary phosphonium salts was achieved through simple alkylation of commonly available achiral phosphines with either allyl or benzyl bromide 80 81 (see Figure 2). Alkylation of triphenylphosphine yielded 1 ($R_2 = CH = CH_2$, 96%) and 2 ($R_2 = Ph$, 91%), and alkylation of tributylphosphine yielded 3 ($R_2 = CH=CH_2$, 99%). Upon their synthesis 82 83 and isolation, single crystals of both 1 and 2 were subsequently obtained and analyzed by X-ray 84 crystallography (see Supplementary Information).

85

 $\begin{array}{c} R^{1}_{P} R^{1} \xrightarrow{Br} R^{2} \xrightarrow{R^{1}_{P} P^{+}} R^{2} \xrightarrow{R^{1}_{P} P^{+}} R^{2} \xrightarrow{(R)-BINOL (1 equiv)} \xrightarrow{(R)-BINOL (1 equ$ 86 (91–99% yield) (39-80% yield) **1** ($R^1 = Ph, R^2 = CH_2 = CH_2$) 4 ($R^1 = Ph, R^2 = CH_2 = CH_2$) 87 **2** ($R^1 = Ph, R^2 = Ph$) 5 (R^1 = Ph, R^2 = Ph) **3** ($R^1 = {}^nBu, R^2 = CH_2 = CH_2$) 6 ($R^1 = {}^nBu, R^2 = CH_2 = CH_2$)

Figure 2 - Synthesis of the quaternary phosphonium salts and their respective ternary complexes with (*R*)-BINOL

88 Recognition. The recognition of these quaternary phosphonium salts was then attempted. 89 In concentrated CHCl₃ solutions 1 and 3 formed solid ternary complexes with (R)-BINOL to give 90 4 (74%) and 6 (80%) respectively and were isolated by filtration. 2 failed to yield the desired 91 ternary complex in CHCl₃, even upon concentration. When the solvent was changed to ethanol 92 however, the complex 5 formed in lower yield (39%) compared to the previously mentioned 93 ternary complexes. ¹H NMR and melting points of the isolated solids indicated transformation to 94 the desired ternary complexes. However, only crystallographic analysis of the solid-state structure 95 could give further information of the supramolecular recognition.

96 Crystallography. Single crystals of each ternary complex were grown. 4 readily crystallised 97 in CD₃OD to give large clear prisms. Crystals of 5 and 6 were grown by gradual cooling and

98 evaporation in ethanol to give large clear prisms and plates respectively. Acquisition of the single 99 crystal diffraction data for each sample gave unambiguous evidence for the formation of the 100 ternary complexes in the solid state. The X-ray crystallographic data for each ternary complex is 101 given in Table 1.

102 In the crystal structure of complex 4, there is clear hydrogen bonding between the phenolic hydroxyl group of the BINOL and the Br⁻ counterion (O–H···Br⁻; d = 2.349 Å, $\theta = 174.11^{\circ}$). In 103 104 addition, another hydrogen bond is also present from the α -carbon center of the phosphonium salt to the bromide counterion (C–H···Br⁻; d = 2.826 Å, $\theta = 162.72^{\circ}$). Such interactions are consistent 105 106 with supramolecular ternary complex formation between the BINOL, halide and the target 107 phosphonium cation. Interestingly, unlike the previous examples in quaternary ammonium 108 complexation, no continuous hydrogen bonding network between the BINOL and bromide 109 counterions appears to be present, with each BINOL bound to only a single bromide anion (as 110 displayed in Figure 3 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 112 halogen counterion via an acidic a-centre of the phosphonium cation. Minor aryl C-H-O interactions (d = 2.693 Å, $\theta = 120.68^{\circ}$; d = 2.622, $\theta = 124.05^{\circ}$ & d = 2.630 Å, $\theta = 142.10^{\circ}$) from 113 114 the electronically deficient phenyl rings of the phosphonium cation to the hydroxyl groups of the BINOL are present.²³ These minor interactions were not classed as forms of recognition (see 115 116 Hirshfeld Surface Analysis, below), however, their presence in the crystal structure should not be 117 discounted as the full intricacies of this phenomenon is not fully understood.

118 The crystal structure of complex **5** presented also contained strong interactions between the 119 BINOL and bromide counterion. A strong hydrogen bond between the bromide counterion and 120 BINOL was immediately evident (O–H…Br⁻; d = 2.442 Å, $\theta = 177.77^{\circ}$). However, in contrast to

121 complex 4, the packing in this crystal structure revealed a continuous hydrogen bonding network 122 linking each BINOL moiety to two adjacent bromide anions (see Figure 3 b). A hydrogen bond from the acidic α -carbon center of the phosphonium cation is also observed (C–H···Br⁻; d = 2.652123 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 126 an adjacent hydroxyl group of the BINOL species. In this way, it is shown that the recognition of 127 this phosphonium salt is mediated by all three types of hypothesized recognition modes: a Type A 128 encapsulation mechanism (via the continuous BINOL-halogen hydrogen bond network), a Type B 129 indirect recognition of the α -center of the phosphonium cation acting as a HBD to the halogen 130 counterion and a Type C direct recognition via aryl C-H...O hydrogen bonding to the hydroxyl 131 HBA on the BINOL species. This demonstrates that that an acidic α -centre on the phosphonium is 132 not the only possible HBD motif in this recognition process and that electron deficient aryl rings 133 are also sufficient for hydrogen bonding to the BINOL species. It also highlights the variety in 134 recognition modes that can be responsible for the formation of one ternary complex.

135 Analysis of complex 6 revealed the expected hydrogen bonding between the BINOL and bromide counterion (O–H···Br; d = 2.349 Å, $\theta = 174.11^{\circ}$) (see Figure 3 c). This crystal structure also had 136 137 a continuous hydrogen bonding network between the BINOL and bromide counterions, which 138 appear to encapsulate the phosphonium cation, consistent with a Type A recognition. However, no 139 close contacts between the phosphonium cation and the bromide counterion could be classed as hydrogen bonding. Instead, C-H···O hydrogen bonds (C-H···O; d = 2.517 Å, $\theta = 161.39^{\circ}$, d =140 141 2.575 Å, $\theta = 126.28^{\circ}$) from the α -carbon centre of the phosphonium cation acting as the HBD and 142 the BINOL hydroxyl group acting as a HBA were present. This demonstrates that both Type A and 143 *Type C* modes of recognition are responsible for complex formation.



	4	5	6
Empirical formula	$C_{41}H_{34}BrO_2P$	$C_{45}H_{36}BrO_2P$	C ₃₅ H ₄₆ BrO ₂ P
Formula weight	669.56	719.62	609.60
Temperature/K	120.0	120.0	120.0
Crystal system	monoclinic	monoclinic	orthorhombic
Space group	<i>P</i> 2 ₁	<i>P</i> 2 ₁	$P2_{1}2_{1}2_{1}$
a/Å	10.9509(5)	8.8952(15)	12.5247(5)
b/Å	14.5495(7)	15.823(3)	15.2027(6)
c/Å	11.0440(6)	12.673(2)	16.4528(7)
$\alpha/^{\circ}$	90	90	90
β/°	113.207(2)	96.325(7)	90
$\gamma/^{\circ}$	90	90	90
Volume/Å ³	1617.26(14)	1772.9(5)	3132.8(2)
Z	2	2	4
$\rho_{calc}g/cm^3$	1.375	1.348	1.292
μ/mm^{-1}	1.356	1.243	1.393
F(000)	692.0	744.0	1288.0
Crystal size/mm ³	$0.273 \times 0.266 \times 0.162$	$0.281 \times 0.222 \times 0.096$	$0.276 \times 0.234 \times 0.216$
Radiation	MoKa ($\lambda = 0.71073$)	MoKa ($\lambda = 0.71073$)	MoKa ($\lambda = 0.71073$)
2Θ 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 < 1 < 16	$-11 \le h \le 11, -20 \le k \le 20, -16 \le 1 \le 16$	$-18 \le h \le 18, -22 \le k \le 22, -24 \le 1 \le 24$
Reflections collected	37669	30239	77675
Independent reflections	10748 [$R_{int} = 0.0555$, $R_{sigma} = 0.0726$]	7774 [$R_{int} = 0.1175$, $R_{sigma} = 0.1416$]	10995 [$R_{int} = 0.0744$, $R_{simms} = 0.0625$]
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>= 2σ (I)]	$R_1 = 0.0352, wR_2 =$	$R_1 = 0.0524, wR_2 =$	$R_1 = 0.0323, wR_2 =$
Final R indexes [all data]	0.0711 $R_1 = 0.0490, wR_2 = 0.0740$	0.1029 $R_1 = 0.0968$, $wR_2 = 0.1174$	0.0618 $R_1 = 0.0489$, $wR_2 = 0.0655$
Largest diff. peak/hole / e Å ⁻³	0.52/-0.49	0.51/-0.85	0.45/-0.38
Flack parameter	0.041(4)	0.012(9)	0.016(3)

Table 1 – Crystallographic data for the ternary complexes

168 Hirshfeld Surface analysis. Further characterization of the recognition modes within the crystal 169 structures was achieved using Hirshfeld surface analysis (Figure 4).²⁴ Highlighting the C-H-Br 170 interactions in each plot qualitatively displays the presence and strength of this interaction in each 171 crystal structure (blue traces). In complexes 4 and 5 (Figure 4 a & b) a sharp characteristic 172 fingerprint is revealed indicative of a hydrogen bond present in these crystals, further confirming 173 the observation of both crystals having a *Type B* component in their recognition modes. Complex 174 6 however, has a much weaker and more diffuse fingerprint for this interaction (Figure 4 c). While 175 this interaction is present in the crystal, it is unlikely to be a defined hydrogen bond as its overall 176 contribution to the entire crystal packing is minimal representing just 2.6% of the surface area of 177 the plot. This is likely indicative to the fact that this crystal has been identified to have both *Type* 178 A and Type C recognition, which does not involve strong C–H…Br interactions.

179 The C-H...O interactions with the hydroxyl group of the BINOL acting as a HBA could also be 180 clearly examined (purple traces). In complex 4, noticeably more diffuse, and therefore weaker, C-181 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 184 present of both Type B and Type C recognition modes in this ternary complex. Finally, the 185 Hirshfeld plot for complex 6 displays the overwhelming contribution of the C-H…O hydrogen 186 bonding observed in this crystal (5.8% of the plot), showing strong evidence for a Type C187 recognition in this complex, with little to no *Type B* character.



Figure 4 – Expanded unit cells and Hirshfeld fingerprint plots for (a) complex 4, (b) complex 5, and (c) complex 6; highlighting the P⁺–CH···Br⁻ interactions (blue) and P⁺–CH···O (purple) observed in each ternary complex

BINOL...*X*[•] *packing.* To view the impact of the different modes of recognition on the packing of 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 BINOL...Br⁻ scaffold (Figure 5). Complex 4, which displays *Type B* recognition, can be seen in Figure 5 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.

196 Complexes 5 (Figure 5 b) and 6 (Figure 5 c) both show much more compact packing structure 197 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 199 is accommodated in visible cavities within this hydrogen bond network. Interestingly, this is 200 despite the differences in geometry between the respective phosphonium cations within the two 201 crystal structures. This demonstrates that the encapsulation of the cation is a flexible process – 202 rather than the BINOL only able to accommodate one specific shape, the cavities found within 203 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.



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

212 CONCLUSION

213 This work characterizes the modes of supramolecular recognition of quaternary phosphonium 214 cations in the solid phase by BINOL. Three modes have been hypothesized and subsequently 215 identified within the recognition complexes presented here. Complex 4 arises from Type B 216 interactions, while complex 5 has a mixture of Type A, Type B and Type C character as identified 217 by single crystal X-ray crystal structure and Hirshfeld fingerprint plots. Complex 6 contains Type 218 A & C recognition motifs in its solid state structure. The crystal packing of each recognition 219 complex was influenced by these modes of interaction and presents new crystal engineering 220 opportunities in the co-crystallisation of BINOL with ionic species.

221 EXPERIMENTAL

222 Reagents and solvents used in this work were commercially available (Sigma Aldrich; 223 tributylphosphine, triphenylphosphine. Fluorochem; (R)-BINOL, allyl bromide, benzyl bromide) 224 and were used without further purification. NMR spectra were recorded on either a Bruker Avance III HD-400 spectrometer with operating frequencies of 400.07 MHz for ¹H; 100.60 MHz for ¹³C 225 and; 161.95 MHz for ³¹P at 298 K. Melting points are uncorrected. X-ray single crystal diffraction 226 227 data was collected using a Bruker D8 Venture (Photon100 CMOS detector, ImSmicrosource, 228 focusing mirrors). The diffractometer was equipped with an Oxford Cryosystems cryostream, with 229 open-flow nitrogen cryostats set at a temperature of 120.0 K. All structures were solved by direct methods and refined by full-matrix least squares on F2 for all data using Olex2 (V 13.0)²⁵ and 230 SHELXTL^{26,27} software. All non-disordered non-hydrogen atoms were refined anisotropically and 231 232 hydrogen atoms were placed in the calculated positions and refined in riding mode. 233 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.

239

240 **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).

¹H 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). ¹³C NMR (101 MHz, CDCl₃) 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 Hz), 13.3. ³¹P NMR (202 MHz, CDCl₃) δ 31.32. HRMS (ESI-TOF) m/z: [M]⁺ Calculated for C₁₅H₂₃P⁺: 243.2242, found 243.2269. mp: 47–50 °C. IR (max/cm⁻¹): 2961s, 2871s, 1634m, 1464m, 1232m, 1093m, 929m, 719m, 597m.

254

255 Typical complexation procedure. Allyltributylphosphonium bromide · (R)-1,1'-bi-2-naphthol
256 (6).

The quaternary phosphonium bromide salt **3** (0.969 g, 3.00 mmol) was dissolved in CHCl₃ (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).

262 ¹H NMR (400 MHz, DMSO-d₆) δ 9.26 (s, 2H), 7.96 – 7.74 (m, 4H), 7.39 (d, J = 8.8 Hz, 2H), 7.22 263 (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), 264 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, 1.6 265 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), 266 1.39 (h, J = 7.0 Hz, 6H), 0.90 (t, J = 7.2 Hz, 9H). ¹³C NMR (101 MHz, DMSO-d₆) 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. ³¹P NMR (202 MHz, 269 DMSO-d₆) δ 32.18. HRMS (ESI-TOF) m/z: [M]⁺ calculated C₁₅H₃₂P⁺: 243.2242, found 243.2263. 270 [M-H]⁻ calculated C₂₀H₁₃O₂⁻ : 285.0921, found 285.0929. mp: 140–141 °C (MeOH). IR (max/cm⁻ 271 ¹): 3165br, 2954m, 1622m, 1504m, 1324m, 1269s, 964m, 818s, 683m. XRD: Sample was 272 crystallised in ethanol, to give clear colorless prisms. Crystal data: orthorhombic, space group 273 *P*2₁2₁2₁ (no. 19).

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275 ASSOCIATED CONTENT

Supporting Information. A listing of the contents of each file supplied as Supporting Information
should be included. For instructions on what should be included in the Supporting Information as
well as how to prepare this material for publications, refer to the journal's Instructions for Authors.
The following files are available free of charge. Supporting Information (PDF) Crystallographic
files (CIF)

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- 295 ABBREVIATIONS
- HBD hydrogen bond donor, HBA hydrogen bond acceptor, BINOL 1,1'-bi-2-naphthol, d –
- 297 interatomic distance, θ bond angle
- 298 REFERENCES
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