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## Various types of non-covalent interactions contributing towards crystal packing of halogenated diphospha-dicarbaborane with an open pentagonal belt<sup>†</sup>

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We have prepared 3-Cl-10-I-*nido*-7,8,9,11- $P_2C_2B_7H_7$  by the reaction of 3-Cl-*nido*-7,8,9,11- $P_2C_2B_7H_8$  with  $I_2$  in the presence of AlCl<sub>3</sub>. The product was obtained in a yield of 49% and was characterized by NMR spectroscopy and X-ray crystallography. Quantum chemical calculations have demonstrated that the crystal structure is stabilized by hydrogen, dihydrogen and pnictogen bonds.

Innerly and/or outerly substituted boron cluster hydrides (heteroboranes) of various architectures are known for various types of non-covalent intermolecular interactions and thus serve as good models for the evaluation and mutual comparison of various types of hydrogen, dihydrogen (diH), halogen (X), chalcogen, pnictogen (Pn) or even tetrel bonds.<sup>1,2</sup> The neutral *nido*- $B_7P_2[C(R)]_2H_7$  cage with an open  $P_2CBC$  pentagonal belt can be formally obtained from  $nido-B_{11}H_{11}^{4-}$  (nido-B<sub>11</sub>H<sub>14</sub> exists)<sup>3</sup> by replacing four BH vertices with two tetrel and Pn vertices. Miscellaneous dicarba dipnictogen nido eleven-vertex heteroboranes have been prepared by reacting arachno-4,5-C<sub>2</sub>B<sub>7</sub>H<sub>13</sub> with PCl<sub>3</sub>, AsCl<sub>3</sub> and SbCl<sub>3</sub>.<sup>4-6</sup> The crystal structures of 3-Cl-nido-7,8,9,11-P<sub>2</sub>C<sub>2</sub>B<sub>7</sub>H<sub>8</sub> (1), 3-I-nido-7,8,9,11-As<sub>2</sub>C<sub>2</sub>B<sub>7</sub>H<sub>8</sub> (2) and nido-7,8,9,11-Sb<sub>2</sub>C<sub>2</sub>B<sub>7</sub>H<sub>9</sub> (3) compounds are already known.<sup>4-6</sup> The crystal packing of **3** is governed by Pn-bonds.<sup>6</sup> In contrast to 3, it has not been possible to record the singlecrystal X-ray data of the parent nido-7,8,9,11-P2C2B7H9 and nido-7,8,9,11-As<sub>2</sub>C<sub>2</sub>B<sub>7</sub>H<sub>9</sub> in sufficient quality. Interestingly, it is feasible to determine the structure of single-crystal halogenated compounds 1 and 2, revealing a supramolecular architecture mainly resulting from extensive Pn-bonding.4-6 In order to

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Fig. 1 A simplified reaction scheme.

proceed in the examination of the non-covalent interactions of *nido*-heteroboranes, we attempted to iodinate 1 (Fig. 1) by the reaction with elemental iodine in the presence of aluminum trichloride (details in ESI†). The resulting 3-Cl-10-I-*nido*-7,8,9,11- $P_2C_2B_7H_7$  (4) was obtained in a moderate yield of 49% and was characterized in solution in terms of multinuclear NMR spectroscopy (Table 1).

In the crystal structure of 4 (Fig. 2), four molecules have been found in the unit cell. The structure of the cage of 4 is essentially the same as found for  $1^4$  except for the very slight shortening of the B(10)–C(9/11) bonds inside the open P<sub>2</sub>CBC pentagonal belt caused by the substitution of the B(10) atom by the heavy iodine. In fact, there are only two examples of a similar structural pattern in the Cambridge Structural Database, *i.e.* dicarba-*closo*-dodecaborane-1,2-diyl-1,2-bis(diisopropylamino)diphosphines,<sup>9</sup> but these are heavily distorted with no option of a serious structural comparison.

Table 1 N	VMR re	esults	for	<b>4</b> <sup>a</sup>
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	B(1)	B(3)	B(2,4)	$B(10)^b$	B(5,6)
GIAO Exp.	$-29.3 \\ -31.2$	$\begin{array}{c} 2.5 \\ -0.7 \end{array}$	$-2.0 \\ -1.9$	$5.9\\-14.7$	3.5 0.3

<sup>*a*</sup> All in ppm with respect to  $BF_3$ ·OEt<sub>2</sub>. Atomic numbering is shown in Fig. 2. <sup>*b*</sup> The discrepancy between the calculated and experimental chemical shift results from the neglect of the effect of spin–orbit (SO) coupling on the overall shielding, which is not taken into account in the GIAO-MP2 scheme. However, it has been shown that due to iodine being terminally bonded to a boron atom within a neutral boron cluster, SO coupling increases the contribution of this B atom towards overall shielding by about 20 ppm,<sup>7,8</sup> which would considerably improve the fit in the present case.

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**Fig. 2** The molecular structure (ORTEP 50% probability level) and atomic numbering of **4**. The selected interatomic distances [Å] are: I1–B10 2.184(2), C11–B3 1.786(2), B2–C11 1.730(3), B2–P7 2.053(2), B3–P8 2.1117(19), B3–P7 2.113(2), B4–C9 1.732(2), B4–P8 2.0516(18), B5–C9 1.715(3), B6–C11 1.715(2), C9–B10 1.608(2), C9–P8 1.8681(17), C9–H9 0.91(3), B10–C11 1.604(3), C11–P7 1.8706(19) and P8–P7 2.1970(6). For all the distances and other structural characteristics, see Table S2 (ESI†).



Fig. 3 The computed electrostatic potential (ESP) surface for compound 4. The color range of the ESP in kcal mol<sup>-1</sup>. The direction of the dipole moment is shown as an arrow derived at the HF/cc-pVTZ level.

To better understand non-covalent interactions of 4, we computed its ESP potential (Fig. 3) and compared it with other recently studied analogous compounds, 1-3. Compound **1** has only one highly positive  $\sigma$ -hole located on the P vertices. The other expected  $\sigma$ -holes of 1 could not be defined on the P vertices due to the more positive neighboring CH vertices. The Sb and As vertices of 2 and 3 have more positive ESP surfaces than the CH vertices. Both molecules thus have three well-defined  $\sigma$ -holes on the Pn atoms. From this point of view, the ESP surface of 4 is thus analogous to 2 and 3. Compound 4 has three  $\sigma$ -holes located on P atoms with the same magnitude  $(V_{\rm S,max})$  of 31.5 kcal mol<sup>-1</sup>. The P vertices of 4 have even more positive ESP values than the CH vertices. The ESP surface on top of the CH vertices has the value of 28.2 kcal mol<sup>-1</sup>. Besides the  $\sigma$ -holes located on P atoms, the 4 compound also has  $\sigma$ -holes on the X atoms. While the I atom has a positive  $\sigma$ -hole

**Table 2** The magnitudes of  $\sigma$ -holes ( $V_{S,max}$ ) on the surfaces of group V and VII atoms and dipole moments ( $\mu$ ) computed at the HF/cc-pVDZ level

$V_{\rm s,max}/{\rm kcal}~{\rm mol}^{-1}$		
Group V atoms	Group VII atoms	
25.4 <sup>6</sup>	$-3.1^{6}$	
$32.2; 2 \times 29.0^{6}$	8.2 <sup>6</sup>	
$1 \times 42.7$ ; $2 \times 25.7^{6}$	_	
$3 \times 31.5^{6}$	6.8 (I); -1.2 (Cl)	
	$\frac{V_{s,max}/kcal mol^{-1}}{Group V atoms}$ 25.4 <sup>6</sup> 32.2; 2 × 29.0 <sup>6</sup> 1 × 42.7; 2 × 25.7 <sup>6</sup> 3 × 31.5 <sup>6</sup>	

with the  $V_{S,max}$  value of 6.8 kcal mol<sup>-1</sup>, the Cl atom only has a relative  $\sigma$ -hole (*i.e.* the top of the Cl atom is less negative than its belt but the  $V_{S,max}$  value is negative). The 4 compound can thus form Pn-, H- and X-bonds, the last of which are, however, less probable due to the lower  $V_{S,max}$  values on the X atoms (Table 2). The most negative ESP surface of 4 is on the negative belt on the I and Cl atoms (the magnitudes of -15.1 and -12.3 kcal mol<sup>-1</sup>, respectively).

Finally, we have studied pairwise interactions in the crystal structure of 4 and compared them with those of 1, 2 and 3. Highly-accurate MP2/CBS interaction energy ( $\Delta E$ ) values are summarized in Table 3 (see also Fig. 4). The A····B motif is the most stable motif of 4. The motif can be described as two symmetrical P<sub>2</sub>···I Pn-bonds. Since the Pn-bond is quite long (3.8 Å), it could also be seen as a nonspecific dispersion contact with an optimal dipole–dipole arrangement. The  $\Delta E$  value of -9.5 kcal mol<sup>-1</sup> is very large, considerably more negative than the most negative bonding motifs of 1 and 2 (the  $\Delta E$  of about -6.0 kcal mol<sup>-1</sup>), which also had two Pn-bonds. However, considering that the A···B motif of 4 has two Pn-bonds, they are weaker than the Pn-bonds of 3. Compound 3 has a binding motif that is exclusively formed by a single Pn-bond and has the  $\Delta E$  value of -6.5 kcal mol<sup>-1</sup>.<sup>6</sup>

The A···C motif, formed by two symmetrical C-H···I-B H-bonds, has  $\Delta E$  of -6.5 kcal mol<sup>-1</sup>. The H-bond is again rather long (3.2 Å). The third most stable motif is A···D with  $\Delta E$ of -6.1 kcal mol<sup>-1</sup>. It is formed by the C-H···H-B dihydrogen bond and the C-H···I-B H-bond. The C-H···H-B diH-bond is the only contact in the crystal structure of 4 with the length below the sum of van der Waals radii (2.9 Å between the H and

	Table 3	Interaction energies ( $\Delta E$ ) in kcal mol <sup>-1</sup>
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	Interaction	MP2.5/CCSD(T)
3-Cl-nido-7,8,9,11-	$P_{2}C_{2}B_{7}H_{8}(1)$	
A···B	2× Pn-bond	$-5.93^{6}/-6.10^{6}$
3-I-nido-7,8,9,11-A	$s_2C_2B_7H_8$ (2)	
A···B	2× Pn-bond	$-5.98^{6}$
$A{\cdot}\cdot{\cdot}C$	$2 \times $ Pn-bond	$-5.67^{6}$
nido-7,8,9,11-Sb <sub>2</sub> C	$_{2}B_{7}H_{9}(3)$	
A···B	$2 \times$ Pn-bond, HH-bond	$-6.80^{6}/-7.05^{6}$
$A{\cdots}C$	Single Pn-bond	$-6.22^{6}/-6.46^{6}$
3-Cl-10-I-nido-7,8,9	$P_{11}-P_{2}C_{2}B_{7}H_{7}$ (4)	
A···B	2× Pn-bond	-9.5
$A{\cdots}C$	$2 \times$ H-bond	-6.5
$A\!\cdot\cdot\cdot D$	diH and H-bonds	-6.1
$A{\cdot}\cdot{\cdot}E$	Stacking	-4.8



Fig. 4 The most significant interaction motifs from the crystal structure of 3-Cl-10-I-*nido*-7,8,9,11-P<sub>2</sub>C<sub>2</sub>B<sub>7</sub>H<sub>7</sub> (4). The distances are in Å. The positions of the H atoms have been optimized at the DFT-D3/BLYP/DZVP level.<sup>10</sup>

the center of the B–H bond;  $H \cdots H$  and  $H \cdots B$  distances of 2.3 and 3.0 Å, respectively). The  $A \cdots E$  motif also has all contacts much higher than the sum of the van der Waals radii (*e.g.* the  $P_2 \cdots Cl$  Pn bond of 3.9 Å).

In summary, compound 4 (3-Cl-10-I-*nido*-7,8,9,11-P<sub>2</sub>C<sub>2</sub>B<sub>7</sub>H<sub>7</sub>) was prepared by reacting 3-Cl-*nido*-7,8,9,11-P<sub>2</sub>C<sub>2</sub>B<sub>7</sub>H<sub>8</sub> with elemental iodine in the presence of aluminum trichloride. Its molecular structure has been characterized by NMR spectroscopy, X-ray crystallography and *ab initio* computations. Apart from B(10) there is a good accord between the experimental and computed <sup>11</sup>B chemical shifts. It is the spin–orbit coupling that

is responsible for such a disagreement related to B(10). The combination of Pn-, diH and H-bonds turned out to be a driving force for the crystal packing as revealed by quantum chemical computations.

### Conflicts of interest

There are no conflicts to declare.

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