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Edge chlorination of hexa-*peri*-hexabenzocoronene investigated by density functional theory and vibrational spectroscopy[†]

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We investigate the molecular structure and vibrational properties of perchlorinated hexa-*peri*-hexabenzocoronene (HBC-CI) by Density Functional Theory (DFT) calculations and IR and Raman spectroscopy, in comparison to the parent HBC. The theoretical and experimental IR and Raman spectra demonstrated very good agreement, elucidating a number of vibrational modes corresponding to the observed peaks. Compared with the parent HBC, the edge chlorination significantly alters the planarity of the molecule. Nevertheless, the results indicated that such structural distortion does not significantly impair the π -conjugation of such polycyclic aromatic hydrocarbons.

Keywords: PAH; Density Functional Theory (DFT) Calculations; Raman spectroscopy; IR spectroscopy

1 Introduction

Perchlorination of extended polycyclic aromatic hydrocarbons (PAHs) has recently been developed as a convenient way not only to modulate the optical properties¹, or the formation of self-assembled monolayers², but also to improve the solubility of such disk-shaped molecules³, which can also be considered as molecularly defined model systems of graphene, *i.e.*, graphene molecules^{4,5}. Solubility is a technology-enabling key concept in





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[†] Electronic Supplementary Information (ESI) available: description and animations of the vibrational normal modes of HBC and HBC-Cl discussed in the text. See DOI: 10.1039/b000000x/



Fig. 1 Representation of a π stacked dimer of HBC-Cl obtained from geometry optimization with DFT methods including Grimme's dispersion (B3LYP/6-31G(d,p) GD3BJ)¹⁰. This structure is very similar to the one determined by X-ray diffraction³.



Fig. 2 Left panel: the molecular structure of HBC-CI. Right panel: the notation scheme for the possible out-of-plane conformations at the chlorinated edge (see text for details).

of the expected range found in other graphenic molecules with planar molecular shapes⁸. This shows that perchlorination and the non-planarity of the molecule are not suppressing the charge transport properties: by proper tuning of the relative molecular disposition and design of the chlorination pattern one could possibly enhance the value of t, which is known to dramatically depend on even small changes of the relative orientation angle and relative sliding of the aromatic cores^{8,9}.

Intrigued by the appealing properties of HBC-Cl and its nonplanar molecular structure we have carried out a joint experimental and theoretical study of the vibrational properties and the structure of HBC-Cl, adopting IR, Raman and DFT methods.

2 Results and Discussion

2.1 Molecular structure of perchlorinated HBC

The chemical structure of perchlorinated HBC (HBC-Cl) is shown in Figure 2. While the parent HBC is a planar π -conjugated system¹¹, substitution with chlorine atoms at the edge introduces significant steric hindrance at the bay positions such as 1-18 (see Figure 2). For this reason the equilibrium structure of HBC-Cl significantly deviates from planarity in order to increase the Cl-Cl distances at 1-18 and other bay positions (*e.g.*, 3-4). On the other hand, because of the larger Cl-Cl distances involving the substitutions at positions 1, 2 and 3, we may consider the chlorinated aryl moieties at the edge carbons as locally planar. This is a first

#	description	short form	S	energy	group
1	(ududud)	$(ud)_3$	12	0.0	D_{3d}
2	$(u\alpha du\alpha d)$	$(u\alpha d)_2$	8	4.6	D_2
3	$(udu\alpha d\beta)$	-	8	6.5	C_s
4	$(\alpha du du \alpha)$	$\alpha_2(du)_2$	8	8.9	C_2
5	$(\alpha \alpha \alpha \alpha d u)$	$(\alpha_4 du)$	4	17.3	C_2
6	$(\alpha \alpha \alpha d\beta u)$	$(\alpha_3 d\beta u)$	4	19.1	C_2
7	$(\alpha \alpha d\beta \beta u)$	$(\alpha_2 d\beta_2 u)$	4	22.8	C_{2h}
8	(αααααα)	(α_6)	0	28.2	\overline{D}_6

 Table 1
 The description of the eight conformers of HBC-Cl and their relative energies (kcal/mol) as determined by DFT calculations.

approximation useful to simplify the notation for describing the possible out-of-plane conformations. Depending on the position of a given edge aryl moiety with respect to the average molecular plane, we may have four possible conditions, which are exemplified in Figure 2:

- up (or down) if the aryl moiety lies above (or below) the plane;
- α (or β) if the aryl moiety lies in a propeller blade fashion, with α being related with *P*-helicity.

Because of steric hindrance (see Figure 2) the conformation sequence can not contain pairs such uu, dd, $\alpha\beta$, $\beta\alpha$, $u\beta$, $d\alpha$, αu , and βd . The exhaustive enumeration of all possible combinations satisfying this prescription, considering the equivalence between enantiomeric pairs, leads to the eight conformers listed in Table 1. These conformers have been considered for geometry optimization with the DFT method in order to obtain information on their relative energies, which are also reported in Table 1. When α or β symbols are present in the structure, it is possible to have enantiomeric pairs. For the sake of compactness, for each enantiomeric pair in Table 1 we report the representative with a number of α symbols greater than the number of β symbols (this does not apply to conformers #3 and #7 which have the same number of α and β symbols). Obviously the conformation symbols can be cyclically permuted without changing the nature of the conformation. They can be also subjected to mirror symmetry with respect to the average molecular plane, *i.e.* $\alpha \rightarrow \beta$, $u \rightarrow d$, and so on. This operation exchanges with one another the members of one enantiomeric pair, which of course possess the same relative energy. Hence the conformation shown in Figure 2, namely $(\alpha dud\beta u)$, by cyclic permutation can be transformed to $(dud\beta u\alpha)$ and by mirror symmetry is transformed to $(udu\alpha d\beta)$, which corresponds to conformer #3 in Table 1.

Interestingly, the most stable conformation of HBC-Cl corresponds to the only one non-chiral conformation. This is $(ududud) \equiv (ud)_3$, which belongs to point group symmetry D_{3d} (see Figure 3). On the contrary (α_6), the conformation with more extensive chirality (point group symmetry D_6), is predicted to be the one with highest relative energy. Inspection of the relative energies of HBC-Cl conformations reported in Table 1 reveals an approximate correlation of their stability with the molecular structure at the edge, as described by the conformation symbol. We



Fig. 3 (a) Three-dimensional representation of $(ud)_3$, the most stable conformer of HBC-CI. The model represents the equilibrium structure obtained from DFT calculations. (b) Orthogonal projection of the eight conformations of HBC-CI where the out-of-plane *z*-coordinate of the middle point of each bond is coded in shades of red (z > 0) and blue (z < 0). In-plane bonds are coded with light gray shade.

	HBC	HBC^\dagger	HBC*	HBC-Cl $(ud)_3$
ΔE (kcal/mol)	0	53	54	-
HOMO (hartree)	-0.1929	-0.1898	-0.1897	-0.2316
LUMO (hartree)	-0.061	-0.0618	-0.0626	-0.115
Δ_{HL} (hartree)	0.132	0.128	0.127	0.117
λ_{max} (nm)	357	372	375	415
$\Delta\lambda$ (nm)	0	15	18	58

Table 2 Comparison of the relative positions of frontier orbitals (HOMO, LUMO) and of the lowest lying bright excited state (absorption maximum λ_{max}) in planar HBC, distorted HBC*, HBC[†] and HBC-Cl in its more stable conformation. ΔE is the energy difference between the non-planar models HBC* and HBC[†] and planar HBC.

can introduce a score number *S* defined as follows:

$$S = s_1 N_1 + s_2 N_2 + s_3 N_3 \tag{1}$$

where N_1 represents the number of ud (or du) sequencies along the conformation string, considered in a cyclic manner, so that N_1 = 6 for $(ud)_3$. Similarly, N_2 represents the total number of $u\alpha$, αd , $d\beta$, βu sequencies along the string and N_3 is the total number of $\alpha \alpha$, $\beta \beta$ sequencies along the string. s_1 , s_2 , s_3 are suitable numerical coefficients which weight the different stability of the possible local edge conformation. The choice made in Table 1 simply assumes $s_1 = 2$, $s_2 = 1$ and $s_3 = 0$, so that higher scores are associated to higher molecular stabilities. This effectively allows highlighting in Table 1 clusters of conformations characterized by similar energies and the same score number *S*.

After the evaluation of the stable conformations of HBC-Cl by means of the analysis of the DFT results, we can conclude that the most stable structure is fully consistent with experimental observation by X-ray diffraction, which clearly reveals $(ud)_3$ as the unique structure of HBC-Cl in the crystal phase³. This is expected based on the relatively large energy separation (4.6 kcal/mol) of $(ud)_3$ with the second most stable conformer, $(u\alpha d)_2$. We can conclude on this basis that the packing motif observed in the crystal simoultaneusly minimizes the intramolecular (conformational) energy and allows an effective intermolecular packing in the dimer.

Functionalisation of HBC with Cl at the molecular edge drives the molecule out of planarity, with a rich possibility of conformations, even though $(ud)_3$ is markedly more stable than all the others (see Table 1). In principle out-of-plane distortions could negatively affect π -conjugation. Notably this is observed in π conjugated polymers possessing torsional degrees of freedom able to affect the nearest neighbor π -interactions¹². Hence, to dwell more on the effects of distortion from planarity, we have considered a molecular model with the same conformation of the aromatic core as in $(ud)_3$, but with a hydrogen-terminated molecular edge (the hydrogen positions have been fully optimized while keeping the position of the carbon atoms frozen at the positions they have in perchlorinated $(ud)_3$). For simplicity we name this model HBC*. In addition to this model, we consider also HBC[†], which is obtained starting from the structure of HBC* and fully optimizing all internal coordinates except dihedral angles. This effectively allows to maintain the characteristic curved shape

found in $(ud)_3$, while fully relaxing the bond lengths and valence angles. The energy difference ΔE between the total energy of HBC^{*}, HBC[†] and HBC (see Table 2) is a measure of the energy cost associated to the distorsion of the aromatic core from planarity. By considering the number of π -conjugated carbon atoms in the HBC analogues (42) this results in about 1.3 kcal/mol per carbon, a rather low value which explains the good stability of the compound despite its seeming dramatic distorsion from planarity. Turning now to the electronic properties, we observe in Table 2 that along the sequence HBC, HBC*, HBC[†] the position of the frontier orbitals does not change dramatically, which enforces the idea that π -conjugation is not seriously affected by the deviation of HBC from planarity. On the other hand perchlorination causes the decrease of the position of the frontier orbitals, as expected from an electron-withdrawing substitution. We notice that, compared to HBC*, in HBC-Cl the position of the HOMO (decrease by 0.04 ha) is relatively less affected than the position of the LUMO (decrease by 0.05 ha), which explains the slight decrease of the HOMO-LUMO gap in HBC-Cl. Following the trend of the position and spacing between the frontier orbitals, according to TDDFT calculations the low-lying doubly degenerate bright state red shifts from 3.47 eV (357 nm) in HBC to 2.98 eV (415 nm) in HBC-Cl, while the total oscillator strength of the doublet slightly increases from f = 1.44 to f = 1.54, respectively.

The representation of the doubly degenerate HOMO and LUMO orbitals of HBC-Cl and HBC is reported in Figure 4. For both degenerate HOMO and LUMO pairs (a,b) represented in Figure 4 it is possible to observe the close similarity existing between HBC-Cl and HBC. This further supports the conclusion that, despite the marked out-of-plane distorsion, π -conjugation in HBC-Cl extends over the whole aromatic core of the molecule. This is also consistent with a recent report which highlights the remarkable non-linear optical properties of several non-planar chlorinated graphene molecules, including HBC-Cl¹.



Fig. 4 Representation of the HOMO and LUMO orbitals of HBC (left (a,b) panels) and HBC-Cl (right (a,b) panels).

To further characterize the deviation from planarity in HBC-Cl it is useful introducing the parameter Δ , given by the absolute value of the difference of a CCCC dihedral angle τ along the edge of the molecule from the *trans* or *cis* conformation expected in the



Fig. 5 The deviation from planarity Δ of the dihedral angles at the molecular edge of HBC-CI in the $(ud)_3$ conformation obtained from DFT calculations. The definition of Δ is given in the text.

planar case, *i.e.*:

 $\Delta = | au - 180^\circ|$ (trans); $\Delta = | au - 0^\circ|$ (cis).

 Δ approaches zero as the molecular geometry approaches planarity in correspondence of a given CC bond around which the dihedral angle can be defined. In Figure 5 we report the values of Δ as a function of the position of the CC bonds R_i along the molecular edge of HBC-Cl. Through Eq. (2) each Δ_i is obtained from the corresponding dihedral τ_i , which is defined in terms of the sequence of CC bonds (R_{i-1}, R_i, R_{i+1}) . The maximum deviations from planarity, by almost 45°, are found in correspondence of the CC bonds connecting two consecutive chlorinated aryl moieties along the molecular edge. This is expected based on the three dimensional representation of the molecule in $(ud)_3$ conformation (see Figure 3). Furthermore, the low Δ values of the three dihedral angles defined for each chlorinated aryl unit (close to 2°), confirm the validity of our initial assumption of considering them as locally planar in order to simplify the description of the molecular conformations of HBC-Cl.

Interestingly, as shown in Figure 6, perchlorination slightly affects the CC bond lengths, without altering the basic pattern based on the Clar structure formed by seven aromatic sextets 13,14 . In particular, adopting the label scheme proposed in Figure 6, we notice that upon perchlorination the inner bonds d, e, f becomes shorter, while the outer bonds a, b, c become longer. Finally, it is worth mentioning that interesting non-planar structures have been also reported for fluorinated PAHs based on a coronene core 15 . These structures are distorted to a degree

(2)



Fig. 6 Comparison of the equilibrium bond lengths in HBC and HBC-Cl $((ud)_3 \text{ conformer})$ computed with DFT.

similar to the case of HBC-Cl here investigated.

2.2 IR spectroscopy of HBC-Cl

The IR spectra of both HBC and HBC-Cl have been measured in the solid state and compared with results from DFT calculations carried out on HBC and on the lowest energy conformation of HBC-Cl (see Table 1 and the experimental and computational methods for further details). The principal IR features found in the range between 650 and 2000 cm⁻¹ have been labeled from 1 to 9 and assigned to the corresponding IR transitions predicted by DFT calculations (see Figure 7 and Table 3). Compared with HBC-Cl, HBC presents a similar number of IR signatures. We observe a satisfactory agreement between theory and experiments, which allows to propose the following assignment of selected IR features, based on the analysis of the nuclear displacements computed for each normal mode (see Electronic Supplementary information for further details and animations of selected modes).

Band 2 in HBC is assigned to the collective out-of-plane bending of all CH bonds and correlates with the characteristic TRIO features of PAHs^{16–18}. However, the IR feature 2 of HBC-Cl is assigned to a doubly degenerate mode involving the out-of-phase C-Cl stretching of the bonds at 1 and 3 of the chlorinated aryl moieties (see Figure 2).

Band 3 in both molecules is assigned to a collective ringbreathing mode of the aromatic core which mainly involves the six outer Clar rings; this occurs with an alternated pattern in HBC-Cl, but in HBC half of the molecule vibrates out-of-phase with respect to the other half and the mode is degenerate.

Band 6 is assigned to two closely located degenerate modes in both molecules. In both cases the pattern of the nuclear displacements is complex and mainly involves the CC bonds of the aromatic core. In HBC the normal mode is coupled with in-plane CH bending.

Band 9 is assigned in both molecules to a collective doubly degenerate ring stretching vibration, whose pattern is close to that found in discussing the Raman G line of PAHs 19 .



Fig. 7 IR absorption spectra of HBC and HBC-CI. Wavenumbers computed by DFT have been scaled by 0.98.

feature #	wavenumber (DFT)	wavenumber (expt.)	
HBC-Cl	$D_{3d}, (ud)_3$		
1	773 (E_u)	771	
2	836 (<i>E</i> _{<i>u</i>})	836	
3	1121 (A_{2u})	1108	
4	1189 (E_u)	1175	
5	1277 (E_u)	1258	
6	1308 (E_u), 1316 (E_u)	1291	
7	1337 (A _{2u})	1315	
8	1463 (E_u), 1475 (A_{2u})	1437, 1444	
9	1583 (E_u)	1555	
HBC	D_{6h}		
1	742 (A_{2u})	740	
2	784 (A _{2u})	766	
3	1125 (E_{1u})	1094	
4	1259 (E_{1u})	1227	
5	1341 (E_{1u})	1310	
6	1408 (<i>E</i> _{1<i>u</i>})	1377	
7	1435 (E_{1u})	1409	
8	1531 (E_{1u})	1493	
9	1639 (E_{1u}), 1650 (E_{1u})	1585, 1602	

 Table 3
 List of observed and computed (unscaled) IR features in HBC and HBC-CI.







Fig. 8 Raman spectra of HBC and HBC-Cl excited with 458 nm and 325 nm laser lines, respectively. The simulated Raman spectra computed with DFT (off-resonance) are displayed below each experimental spectrum. For HBC-Cl just the $(ud)_3$ conformation has been considered in simulating the Raman spectrum. Wavenumbers computed by DFT have been scaled by 0.98. For HBC-Cl the lower wavenumber side of the simulated spectrum is also displayed with enhanced intensity $(10\times)$ to help the assignment of the experimental features. This intensity mismatch in the simulation is due to limitations in the treatment of resonance effects in the standard implementation of Raman scattering currently available in Gaussian09²⁰.

feature #	wavenumber (DFT)	wavenumber (expt.)	
HBC-Cl	$D_{3d}, (ud)_3$		
A_1	276 (A_{1g})	292	
A_2	415 (A_{1g}, E_g)	412	
B_1	1095 (A_{1g})	1086	
B_2	1173 (E_g)	1157	
D_1	1259 (E_g)	1235	
D_2	1280 (A_{1g})	1259	
D_3	1336 (A_{1g})	1321	
D_4	1399 (A _{1g})	1376	
G_1	1500 (E_g)	1470	
G_2	1544 (A_{1g})	1514	
G_3	1567 (E_g)	1536	
G_4	1621 (E_g)	1589	
HBC	D_{6h}		
A_0	279 (E_{2g})	279	
A_1	352 (A_{1g})	355	
B_1	1022 (A_{1g})	1007	
D_1	1282 (E_{2g})	1255	
D_2	1184 (A_{1g})	1150	
D_3	1325 (A_{1g})	1304	
D_4	1426 (A_{1g})	1402	
D_5	1384 (A_{1g})	1363	
G_1	1499 (E _{2g})	1476	
G_2	1647 (A_{1g})	-	
G_3	1650 (E_{2g})	1606	
G_{4a}	1635 (E_{2g})	-	
G_{Ab}	1578 (E_{2g})	1533	

Table 4 List of observed and computed (unscaled) Raman features in HBC-Cl and HBC. The G and D features of HBC have also discussed previously²¹. Experimentally feature G_2 is unresolved because it is very close to the strong G_3 feature, while feature G_{4a} is too weak to be observed.

2.3 Raman spectroscopy of HBC-Cl

The Raman spectra of HBC and HBC-Cl are reported in Figure 8. The two molecules display a similar spectral pattern, dominated by features which have been attributed to G and D modes in HBC^{19,21}. The analysis of the nuclear displacements computed by DFT for the G and D Raman lines displays typical and recognizable patterns which can be put in correspondence between HBC and HBC-Cl.

Following the labeling scheme adopted in Figure 8, here below we discuss and compare the nuclear displacement patterns of selected modes of HBC and HBC-Cl which are associated to relatively intense experimental Raman lines. The complete list of modes is reported in Table 4 and further details are given in the Electronic Supplementary information.

In both HBC and HBC-Cl the mode A_1 is assigned to the inphase collective breathing of the molecule along a radial direction (Figure 9b). The collective breathing of HBC (feature A_1 , observed at 355 cm⁻¹) is significantly red-shifted in HBC-Cl (feature A_1 , observed at 292 cm⁻¹). This is due to the mass effect of the heavy chlorine atoms at the molecular edge. Interestingly, the A_0 feature observed in HBC (assigned to a doubly degenerate mode which involve mainly the CC stretching of aromatic core – see Figure 9a) is characteristic of HBC: no similar mode is computed or observed in HBC-Cl.



Fig. 9 Representation of the low wavenumber (A region) normal modes of HBC and HBC-CI relevant for Raman spectroscopy according to DFT calculations: (a) the A_0 doubly degenerate mode of HBC; (b) the compared breathing modes of HBC-CI and HBC; (c) the degenerate A_{1g} and E_g modes computed at 415 cm⁻¹. For in-plane modes red lines represent displacement vectors; CC bonds are represented as green (blue) lines of different thickness according to their relative stretching (shrinking). For out-of-plane modes the size of blue/red circles of the molecular sketch are proportional to nuclear displacements in the *z* direction.



Fig. 10 Representation of the normal modes of HBC-Cl relevant for Raman spectroscopy in the B region according to DFT calculations.

In the low wavenumber region we find in the experimental spectrum of HBC-Cl a strong Raman line at 412 cm⁻¹ (A₂) which does not find a counterpart in HBC. Based on DFT calculations, this corresponds to three very close Raman active modes computed at 415 cm^{-1} (see Figure 9c) which arise from the degeneracy of an A_{1g} mode with a doublet of E_g species. These modes are characterized by vibrational displacements which have a sizable contribution both along the *z*-axis and within the (x, y) plane. DFT calculations predict a significantly stronger intensity for the A_{1g} mode than the degenerate E_g doublet. Hence it is reasonable to associate the experimental A2 feature mainly to the totally symmetric mode represented in Figure 9c; looking at the (x,y) representation of this A_{1e} mode, one recognizes the breathing pattern of the 7 inner rings (which could be associated to a coronene moiety) in the center of HBC-Cl. During this vibration the outer part of the molecule breathes out-of-phase with respect to the center. As for the E_g degenerate doublet, the associated nuclear displacements shown in Figure 9c display a collective pattern characterized by alternated out-of-plane displacements along the *z*-direction accompanied by displacements in the (x, y) plane which are mostly localized at the molecular periphery.

At higher wavenumber, in the region located between the Amodes and the D-band modes we find Raman active modes which are described as collective breathing vibrations of the Clar rings which can be identified in the molecular structure (see Figure 10



Fig. 11 Representation of the normal modes of HBC and HBC-CI relevant for Raman spectroscopy in the D region according to DFT calculations.

– further details in ESI). Similar modes have been also identified in a previous study on the graphene molecule $C78^{22}$.

At progressively higher wavenumbers one finds modes which can be related to the pattern expected for D-modes¹⁹. The Raman feature D_2 is assigned to the ring breathing of the central Clar ring, out of phase with respect to the stretching of the radial CC bonds which emanate from the central Clar ring. It partially displays the canonical D-mode pattern¹⁹, the notable exception being the wrong phase of the stretching of the bonds of kind *d* (see Figure 6). In HBC-Cl the D_2 mode is coupled with the inphase C-Cl stretching of the chlorinated aryl moieties. In HBC the D_2 mode is coupled with the collective in-plane bending of the CH bonds at positions (1,3) (see Figure 2). D_2 is notably blue-shifted in HBC-Cl compared to HBC (expt. 109 cm⁻¹).

The strong Raman active line D_3 is assigned to the in-phase ring breathing of all the seven Clar rings coupled with the CC shrinking of the bonds of kind *c* and *e* (see Figure 6). This is the mode which fully displays the expected D-mode pattern¹⁹. In HBC the D_3 mode is coupled with the collective in-plane bending of the CH bonds at positions 1 and 3 (see Figure 2). Interestingly, comparing modes D_2 and D_3 in HBC, one finds that the relative phase between the inner Clar ring breathing and the outer CHbending inverts. Hence the modes D_2 and D_3 of HBC can be approximately described as the doublet arising from the vibrational coupling between the collective breathing coordinate of the seven Clar rings and the collective CH bending of the CH bonds at positions 1 and 3. Mode D_4 is assigned to the breathing of the central Clar ring, out-of-phase with respect to the approximate breathing of the outer six Clar rings. In HBC the D_4 mode is coupled with the collective in-plane bending of the CH bonds at 1 and 3 (see Figure 2) while in HBC-Cl the D_4 mode is coupled with the collective stretching of the C-Cl bonds at 2 (see Figure 2). Finally, feature D_5 is found just in HBC and is assigned to the collective CC stretching at the edge of the molecule, mainly at bonds of kind c and d (Figure 6) coupled with the collective in-plane CH bending at positions 1 and 3 (see Figure 2).

The G-modes appear in the next wavenumber region, above the D-modes. In graphene molecules the G-modes display collective displacement patterns¹⁹ which can be associated to that of the $v_{16} E_{2g}$ ring-stretching mode of benzene²³.

The G_3 Raman feature of HBC is assigned to a doubly degenerate mode which involves CC stretching mainly in outer part of the molecule and it is coupled with collective in-plane CH bending at positions 1 and 3 (see Figure 2). As observed in Figure 12, the G_3 feature of HBC-Cl is assigned to a doubly degenerate mode with a similar pattern as HBC. However, compared to HBC, the G_3 mode of HBC-Cl is remarkably red-shifted (expt. 70 cm⁻¹). Finally, the G_4 feature of HBC-Cl is assigned to a doubly degenerate mode which involves the ring stretching of the three central rings next to each other along a row. In HBC there are two doubly degenerate modes with a similar nuclear displacement pattern which have been named G_{4a} and G_{4b} . They both involve ring stretching vibrations coupled with a collective in-plane CH bending.

3 Conclusions

Out-of-plane distortions in molecular models of graphene are not seriously impairing the π -conjugation: this is supported by DFT calculations on perchlorinated HBC which shows a markedly nonplanar equilibrium structure due to the steric hindrance of Cl atoms at the molecular edge, which is in agreement with the structure obtained from the single-crystal X-ray analysis. Both the computed HOMO-LUMO gap and the vibrational properties observed with Raman and IR spectroscopies show that HBC-Cl possesses a π -conjugation similar to that of HBC. Interestingly, even in the non-planar case of HBC-Cl, the π -stacking, which is a crucial property for charge transport in molecular electronics devices based on HBC^{7,24}, is still possible due to specific steric interactions and interlocking of chlorine hindrances, as depicted in Figure 1. Vibrational spectroscopy complemented with DFT calculations proves to be informative about the chemical structure of HBC and HBC-Cl: distinct markers can be directly associated to specific moieties. For instance, the TRIO marker in the IR of HBC (766 cm⁻¹) is due to the symmetric hydrogen-terminated molecular edge, while the persistence of the strong D_3 peak in



Fig. 12 Representation of the normal modes of HBC and HBC-CI relevant for Raman spectroscopy in the G region according to DFT calculations.

HBC (1304 cm⁻¹) and HBC-Cl (1321 cm⁻¹) proves the similar π -conjugated nature of the two molecules, independently of planarity. A thorough analysis of the different stable conformers of HBC-Cl, carried out by geometry optimization with DFT methods, revealed that a rich variety of structures, including chiral enantiomers, can be obtained by suitable chemical substitution at the edges of HBC. While, in the case under study, the lowest energy conformation with achiral (*ud*)₃ conformation is the structure found in the crystal³, we can infer that the introduction of selected edge substituents and/or synthetic pathways could give rise to novel structures, for instance based on propeller-shaped chiral units. These could exhibit appealing chiroptical properties, similar to the class of helicenes²⁵.

4 Experimental and computational methods

Samples of HBC and HBC-Cl were synthesized as described in²⁶ and³, respectively. Micro FT-IR measurements on all the molecules were carried out with Nicolet Nexus equipment coupled with a Thermo-Nicolet Continuµm infrared microscope and a cooled MCT detector (77 K). The spectra of the samples (as powders) were acquired by using the diamond anvil cell technique with a $15 \times$ infrared objective (64 scans, 1 cm⁻¹) resolution). Compared with the KBr pellet technique, the micro FT-IR setup allows recording spectra with a minimal sample amount. The micro-Raman measurements reported in this work have been carried out with a Jobin-Yvon Labram HR800UV equipment using laser excitations at 458 nm and 325 nm, which have been selected to optimize the Raman signal and keep the fluorescence background as low as possible. All DFT calculations reported in this work have been carried out using Gaussian09²⁰ adopting the B3LYP functional and the 6-31G(d,p) basis set. The computer rendered representations of the molecular models reported in Figure 3 and Figure 1 have been obtained with the program YASARA²⁷. The representation of the molecular orbitals reported in Figure 4 has been obtained with the open source program Avogadro (version 1.1.1)²⁸. A set of post-processing programs developed at Politecnico di Milano has been used to generate the representation of the vibrational normal modes and simulate the Raman and IR spectra from the results of DFT calculations.

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