



Experimental and computational study of the crystal packing of isoelectronic 4,11-diaza[5]helicene and 4,7,8,11-tetraaza[5]helicene

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Abstract

The crystal structures of isoelectronic 4,11-diaza[5]helicene (DNH) and 4,7,8,11-tetraaza[5]helicene (TNH) are accurately determined at room temperature by single crystal X-ray diffraction. Analysis of crystal packing with computational chemistry tools, including the evaluation of intermolecular and cohesive energies with force field and charge density (PIXEL) methods, highlights the effect of skeletal replacement of two aromatic C-H with nitrogen atoms. Both crystals are centrosymmetric racemates, with asymmetric units (ASUs) exploiting the same conformation. The ASU of TNH bears two symmetry-independent enantiomers, and the different settings of symmetry operators mirror in different packing motifs with respect to DNH. Both structures exhibit columnar stacking arrangements, which differ in their mutual orientation, distance and strength. Stacking interactions are more frequent and stronger in the tetraaza compound, owing to the greater intrinsic polarization of its ground-state electron density that favors antiparallel arrangements of neighboring molecular dipole moments. Finally, the possible existence of thermodynamically stable chiral forms of the title compounds is investigated by symmetry-constrained Monte Carlo (SC-MC) calculations.

Keywords Azahelicenes · X-ray crystallography · Supramolecular chemistry · Chirality · Crystal structure prediction

Introduction

[n]Helicenes consist of n (>3) ortho-condensed aromatic rings that are forced to adopt a helical configuration due to steric hindrance of the self-folded molecular backbone.

The out-of-plane distortion produces a helical motif, which eliminates all the second order symmetry elements characterizing the flat presentation of the molecules. The corresponding enantiomers are designated as P (“plus” or Δ , right-handed) and M (“minus” or Λ , left-handed). The handedness is given by the folding direction of the helicoidal structure, as viewed down the ring closest to an observer that looks along the helix axis (Scheme 1) [1].

Owing to the presence of an intrinsically chiral chromophore and the easy functionalization, [2] [n]helicene derivatives are attractive for a wide range of chemical and materials science applications, including OLEDs [3], metal sensing, [4] photovoltaics, [5] chiroptical switches, [6] asymmetric catalysis [7] and supramolecular chemistry [8]. Even when racemic mixtures are employed, helicene-based materials are employed in semiconductor devices and humidity sensors, [9] along with molecular springs [10] and Covalent Organic Frameworks (COFs) with photoconductive and photo-emitting activity [11]. Incorporation of heteroatoms – particularly nitrogen – in the aromatic backbone allows fine-tuning of electron transport properties and luminescence, [12] even in response to the acid–base environment.

Contribution submitted for the special issue of Structural Chemistry honoring the career of Angelo Gavezzotti.

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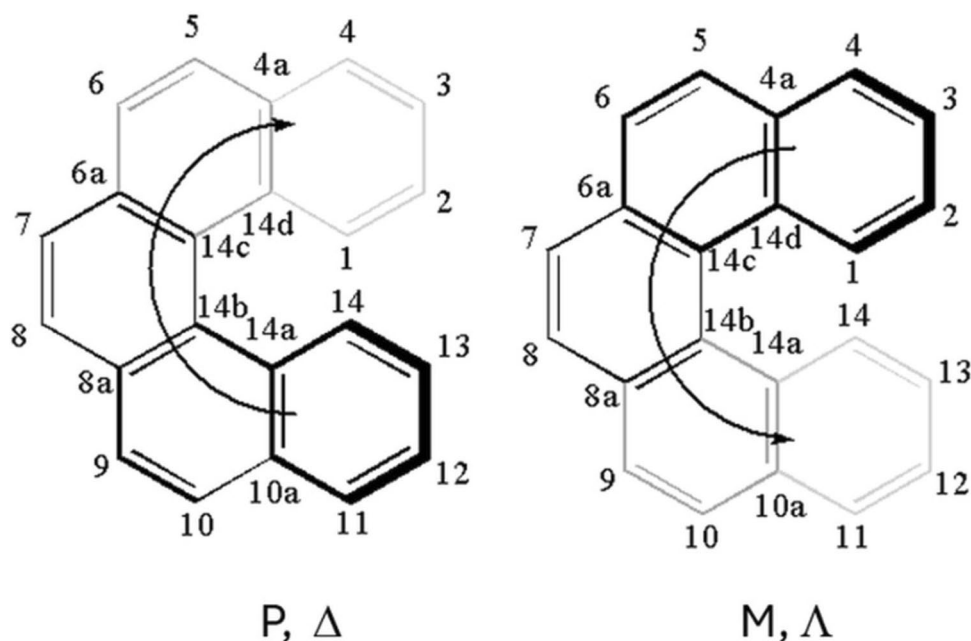
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Scheme 1. Chirality descriptors and IUPAc numbering for the [5] helicene scaffold



[13, 14] Recent studies show indeed that azahelicenes are promising circularly polarized luminescence (CPL) emitters and might pave the way toward next gen chiroptical materials. [15, 16]

The solid-state structure of helicene-based systems provides extra degrees of freedom to control their properties. Crystal packing determines not only the optical response (*e.g.* refraction index and birefringence) but also the dynamics of charge carriers and electronic transitions. [17] Moreover, intrinsic spring-like helicity of their molecular backbone is associated with potentially stimuli-responsive mechanical properties. [18, 19] However, experimental investigations specifically focused on the crystallography of these compounds are scanty. To the best of our knowledge, only a few works address the solid-state structures of aza- and diaza-helicenes [20–22], and no X-ray reports even exist on their poly-nitrogenated analogues.

In this contribution, we investigate how and why the substitution of two skeletal C-H in positions 7 and 8 of 4,11-diaza [5]helicene (DNH) with nitrogen impacts on crystal packing. From the viewpoint of the molecular structure, the change is apparently minimal, as 4,7,8,11-tetraaza [5]helicene (TNH) is isoelectronic with DNH and bears exactly the same solid-state conformation (*vide infra*). However, despite a negligible 0.7% difference in their molecular weights, the TNH crystal is ~5% denser than DNH (Table 1), reflecting the setup of packing motifs that are more efficient in the Kitaigorodskii's sense. [23] We here exploit X-ray crystallography and computational crystal chemistry tools to explore and rationalize how supramolecular motifs, and particularly stacking interactions, are affected by the symmetric nitrogenation of the diaza[5]helicene scaffold.

Methods

Synthesis

All solvents were distilled and dried before use. 6-nitroquinoline, glucose, triphenylphosphine are commercial products and were used without further purification. ^1H and ^{15}N NMR spectra were recorded on Bruker DRX300 or DRX500 spectrometers. The products were dissolved in CDCl_3 or in $\text{DMF-}d_7$. The synthesis of 4,11-diaza[5]helicene was described in [20].

Synthesis of 4,7,8,11-tetraaza[5]helicene (TNH). TNH was prepared following published procedures [24–27]. In particular, 13,2 mmol of 6-nitroquinoline were suspended in 10 mL of 20% aqueous NaOH and heated at 90 °C. To this mixture 12,8 mmol of solid glucose were added portion wise during 30 min. At the end the mixture is diluted with water and extracted three times with 150 mL of CH_2Cl_2 , then once with ethyl acetate. Overall, 839 mg of dark green solid, mainly N-oxide derivative with minor amounts of TNH and of N,N-dioxide, are obtained after drying and evaporation of the solvent.

Then, 533 mg of the crude product are dissolved in 20 mL of toluene together with 500 mg of Ph_3P and refluxed for 27 h. After cooling the toluene is removed by evaporation, the solid residue is washed several times with hexane obtaining 490 mg of TNH product. ^1H NMR (CDCl_3 , 298 K, 7.05 T): δ 9.13 (1H, CH(3) dd, $J=4.2$ Hz, 1.6 Hz), 9.03 (1H, CH(1) dd, $J=8.5$ Hz, 1.6 Hz), 8.87 (1H, CH(6) d, $J=9.0$ Hz), 8.48 (1H, CH(5) d, $J=9.0$ Hz), 7.45 (1H, CH(2) dd, $J=8.5$ Hz, 4.2 Hz) ppm. ^{15}N NMR ($\text{DMF-}d_7$, 298 K, 11.7 T): δ -68 (2N, N(4) and N(11)), +40 (2N, N(7) and N(8)) ppm.

Table 1 Relevant crystallographic results for DNH and TNH crystals. Full crystallographic information (.cif) files, including PLATON [29] validation reports, are deposited as Supplementary Information to this article

	DNH	TNH
Sample information		
Empirical formula	C ₂₀ H ₁₂ N ₂	C ₁₈ H ₁₀ N ₄
Crystal size, mm ³	0.15 × 0.07 × 0.07	0.36 × 0.09 × 0.09
Formula weight, g·mol ⁻¹	280.32	282.30
Crystal system	Orthorhombic	Monoclinic
Space group	<i>Pbca</i> , no. 61	<i>P2₁/c</i> , no. 14
Z	8	8
Z'	1	2
T, K	293.2(1)	293.2(1)
a, Å	15.8246(2)	14.1140(1)
b, Å	8.76360(10)	17.5038(1)
c, Å	19.9202(3)	11.2428(1)
β°	90.0	107.444(1)
Volume, Å ³	2762.54(6)	2649.78(4)
Density (calculated), g·cm ⁻³	1.348	1.415
F(000)	1168	1168
μ, mm ⁻¹	0.623	0.699
Data collection		
λ, Å	1.54184	1.54184
(sinθ/λ) _{max} , Å ⁻¹	0.641	0.641
Scan type °	ω-scan, variable speed	ω-scan, variable speed
	0.5	0.5
No. collected reflections	91,605	95,803
No. unique reflections	3003	5772
No. observed reflections, I > 2σ(I)	2876	5229
Refinement results		
Gof	1.121	1.052
Parameters	211	417
R(F), all data/observed data	0.0425/0.0410	0.0477/0.0443
wR(F ²), all data/observed data	0.1072/0.1058	0.1297/0.1261
Δρ _{MAX/MIN} , e·Å ⁻³	+0.125/-0.184	+0.271/-0.194

X-ray diffraction

CCDC 2495693 (DNH) and 2495694 (TNH) contain the supplementary crystallographic data for this paper. These data can be obtained free of charge from the Cambridge Crystallographic Data Centre via www.ccdc.cam.ac.uk/structures.

DNH was crystallized from a 1:3 mixture of methanol and acetonitrile, while TNH was crystallized from ethanol; in both cases crystallization was obtained by slow evaporation at room temperature (≈ 24–48 h). X-ray quality crystals were carefully selected by visual inspection on a stereomicroscope. X-ray data collections were carried out at room temperature on a Rigaku Synergy-S four-circle diffractometer mounting a HyPix hybrid photon counter in conjunction

with microfocus X-ray sealed tubes. The temperature was chosen to provide an easy comparison with Literature data (vide infra).

Mirror-monochromated Cu Kα radiation (λ = 1.54184 Å) was used throughout, exploiting nominal source power of 50 kV × 1 mA. For both substances, a > 98% complete sphere of reflections was collected up to a resolution of 0.78 Å. Data integration and reduction was carried out with the CrysAlisPro 1.171.43.141a (Rigaku OD, 2024) suite of programs, while the structure was solved with direct methods and refined with nonlinear least-squares procedures implemented in Shelx (Fig. 1) [28]. Table 1 summarizes the relevant crystallographic details of data collections and structural models.

Computational methods

Hirshfeld surface analysis was carried out with CrystalExplorer [30], while Conquest [31] and Mercury [32] were employed to analyze other crystal structures with the [5] helicene scaffold.

The MiCMoS v2.4 suite of programs [33–35] was used throughout to perform both static and Molecular Dynamics (MD) simulations, exploiting experimentally determined X-ray structures as starting points. In all calculations, the C-H covalent distances were normalized to make them comparable with neutron bond lengths [35, 36]. Static lattice energies were computed with either the Lennard–Jones–Coulomb (LJC) force field [37] through the *Crysa* module of MiCMoS [35] or the PIXEL method, [38, 39] which relies on perturbative MP2/6-31G(p,d) molecular charge densities. Relevant topologies and steering parameters are deposited in the Supporting Information (Section S1 SI). Both CHelpG atomic charges [40] and wavefunction-derived charge densities were obtained by single-point calculations of Gaussian16 [41] on the H-corrected experimental structures. Finally, VMD 1.9 [42] was used to plot MP2/6-31G(p,d)-level molecular electrostatic potential onto the corresponding charge isosurfaces at 0.001 atomic units.

500 ps-long NpT MD simulations [34, 37] were carried out on both DNH and TNH crystal structures at 298 K and 1 atm, using a standard leap-frog integrator with time step of 1 fs. Approximately cubic periodic simulation boxes with edges of ≈ 55–60 Å were employed and a general cutoff of 16 Å was applied for long-range interactions. Thermodynamic boundaries were controlled by a CSVR thermostat with time constant τ = 0.6 ps, [43] and by an anisotropic Parrinello–Rahman barostat with scaling constant ww = 4.0 kg. [44] Results from MD were obtained by averaging the structures over the last 200 ps of the trajectory, when the system is fully equilibrated (see Figure S1 SI). The MD estimates for lattice parameters and density are in good quantitative

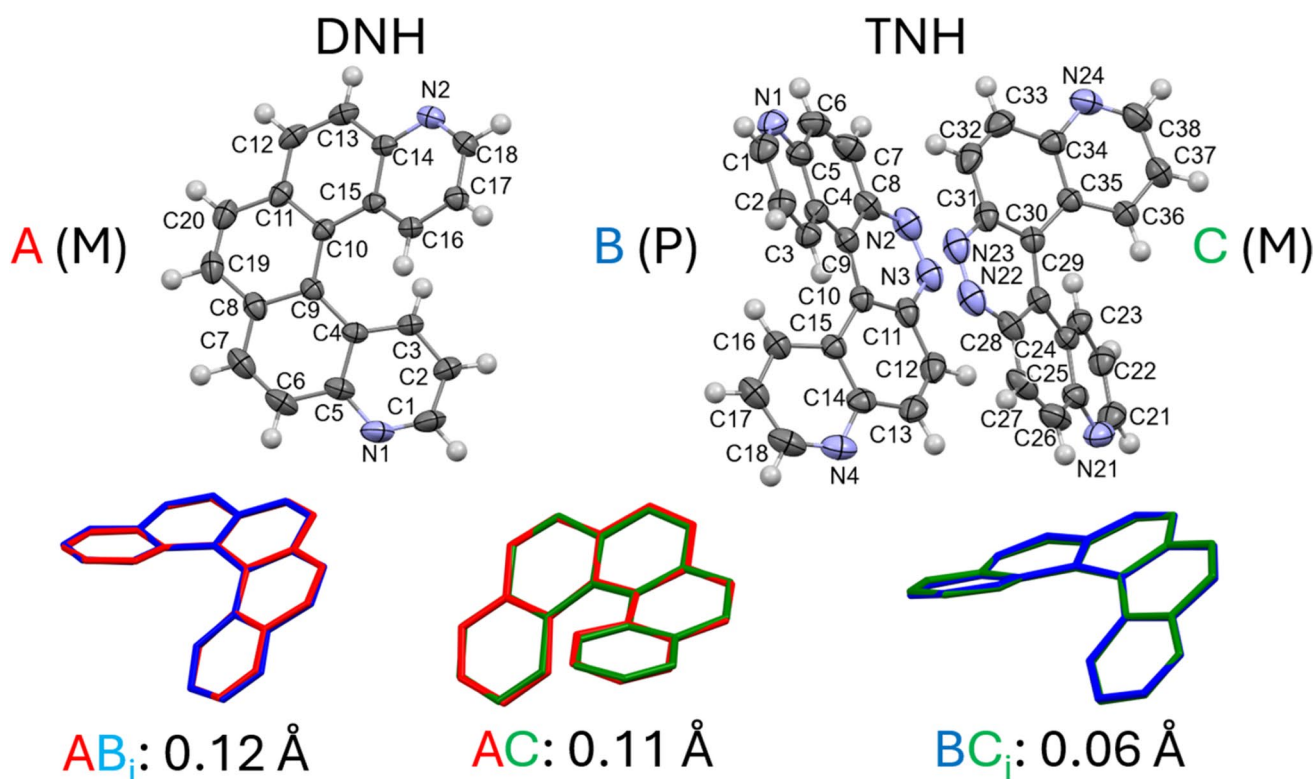


Fig. 1 Up: experimental structure of the asymmetric unit (ASU) of DNH ($Z' = 1$) and TNH ($Z' = 2$) crystals, with the convenience numbering employed the X-ray structure determination (see also Tables S6–S7 SI). Thermal ellipsoids at RT are drawn at the 40% probability level. Molecules are labeled as A, B and C and highlighted in red, blue and

green. The handedness descriptor P or M is also given (see Scheme 1). Down: coordinate RMSD (Å) of different molecular pairings, shown as capped stick representations without hydrogens. An “i” subscript indicates whether that molecule was inverted before calculating the RMSD

agreement with the experiment (Table S4 SI), with relative deviations no greater than +3.6/–3.4%. This is in line with the expected performance of MicMoS. [34, 45]

Crystal structure prediction

The recently developed symmetry-constrained Monte Carlo (SC-MC) method [46] was exploited to generate plausible enantiomerically pure crystal structures of DNH and TNH. The purpose is to predict in what aspects the probable crystal packing of pure P/M enantiomers might differ from the racemates of the title compounds. SC-MC generates guess structures by random choices of lattice parameters and mutual molecular orientations. Any guess structure is tested by some hundreds or thousands of steps of symmetry-constrained Monte Carlo, which means that any change affecting any molecular degree of freedom (conformation, rigid-body translations and rotations) is applied to the whole lattice according to user-selected symmetry operators and Bravais translations. If explorative SC-MC produces a potentially stable structure (*i.e.* with negative cohesive energy), the trajectory is further evolved, until convergence

is achieved. Predicted crystal structures are ranked according to their cohesive energy; duplicates are found and discarded [47].

As for the title compounds, NpT calculations at 298 K and 1 atm with the same LJC force field [37] were carried out, with a 15 Å cutoff for long-range interactions. No internal degrees of freedom were activated, as we assumed that the [5]helicene scaffold is a rigid body. Predictions were restricted to the most common Sohncke groups $P2_1$ (n. 4) and $P2_12_12_1$ (n. 19), up to a maximum of 200 successful structures. Each of these implied 10^3 explorative SC-MC steps, followed by further 10^4 steps for equilibration. Relevant input files are deposited in the Supporting Information (Section S4 SI).

The ten most promising structures in each space group underwent 200 ps-long Molecular Dynamics simulations within the same boundary conditions used for the DNH and TNH experimental structures (298 K and 1 atm, cutoff 16 Å, CSV thermostat and PR barostat) to check their mechanical stability in terms of RMSD of center of mass and relative molecular rotations with respect to the starting predicted (perfect) structure [46].

Reproducibility

Quantum calculations were carried out with the commercial public Gaussian16 software (<https://gaussian.com/>). Classical force field simulations were performed with MiC-MoS v2.4, while crystal structure predictions were carried out with the related SC-MC 1.0 program. These can be both downloaded free of charge under the specified citation and use conditions, upon registration, from https://sites.unimi.it/xtal_chem_group/. Relevant steering parameters and procedures for both programs are deposited in the Supporting Information (Sections S1 and S4 SI). Main output files, including MiC-MoS trajectories and SC-MC predictions, can be downloaded freely from the public repository Zenodo (<https://doi.org/10.5281/zenodo.17407385>). This ensures full reproducibility of all the calculations here described.

Results and discussion

Molecular conformation

The crystal structure of DNH was determined some 20 years ago as part of a systematic investigation of several diaza[5]helicenes (CCDC BOLMEV, structure **1** in ref. [21]). The present results are in fair quantitative agreement with former ones (Table 1 and Table S1 SI), but the current least squares model is significantly more accurate and precise, as can be appreciated by better intensity statistics and lower crystallographic agreement factor (this work: 0.0410; BOLMEV: 0.0806). To the best of our knowledge, the crystal structure of TNH has never been reported before, but the TNH molecule was present in a co-crystal with perhalooctane (CCDC ECASAE). [48] The most striking difference between DNH and TNH resides in their ASUs, which consist of one formula unit in DNH ($Z'=1$) and two formula units in TNH ($Z'=2$).

A survey of the current release of Cambridge Structural Database (v5.46, last update: February 2025) [49] looking for the [5]helicene structure with possible heteroatom and/or substituents in positions 4,7,8 and 11 (Scheme 1) gives only 29 matches (0.002%) under the constraints “only organics” and “single crystal structures”. The full list of CCDC refcodes is deposited in the Supporting Information (Table S5 SI). The dihedral angle between the least squares planes passing through the terminal rings can be taken as a measure of the steepness of the helix. Such dihedrals are symmetrically distributed around an average value of $\sim 50^\circ$ with first 25% quantiles ranging from 48° and 52° , and extreme values at 38° (UFENOM) and 60° (XINLIS). The distance between extremal atoms in positions 1 and 14 (Scheme 1) should measure the extension of the molecular spring, also

including information on how much the parallel “arms” of the loop are open, and correlates well with the above dihedrals (Figure S2 SI). 1–14 distances also exploit a symmetric distribution, ranging from 2.823 (AVUKUA) to 3.058 Å (XINLIS), with average 2.950 Å.

Figure 1 shows the ASUs of DNH and TNH. The helices are slightly compressed, with dihedral descriptors of 48.11° (DNH) and 45.66° (N1-C18, molecule B) or 47.90° (N21-C38, molecule C) (TNH), that is, below the CCDC average but in line with the expected distribution. Interestingly, 1–14 distances are slightly longer in TNH (DNH: 2.929 Å; TNH: 2.961 and 2.967 Å), as nitrogeneration at 7,8 shortens the C-N (8a-8), N-N (8-7) and N-C (7-6a) covalent distances in the central ring by some 0.05 Å (Tables S6-S7 SI). This distortion slightly pushes away the two terminal phenyl groups and increases the 1–14 distances by some 0.03 Å. To the sake of comparison, when TNH is crystallized with perhalogenated n-octane in ECASAE [48], the dihedral is significantly smaller (42.68°) and the 1–14 distance is more like DNH (2.937 Å). This proves that the spatial extension of these compounds can be slightly adjusted, following the requirements of the crystal field, but the structure remains essentially the same, as expected owing to the rigidity of the helix.

Net of these minor differences, all molecules in DNH and TNH bear the same conformation. In fact, DNH and its centrosymmetric image match both molecules in the ASU of TNH, with RMSD of 0.11 Å or 0.12 Å (Fig. 1, bottom). Within the same TNH structure, the two symmetry-independent units are also quantitatively superimposable (RMSD=0.06 Å) upon application of the inversion operator (Fig. 1, bottom). Otherwise, they score an RMSD of 1.06 Å due to the opposite P/M handedness. At variance with other acentric racemates, the breaking of the inversion center in the ASU of TNH is due to small rigid displacements of the two independent moieties with respect to each other, not to coexistence of different conformers [50].

Crystal packing

DNH and TNH crystallize in centrosymmetric space groups (Table 1) meaning that they are centrosymmetric racemates. They both host a total of 8 molecules in cell, equally divided into right-handed and left-handed enantiomers (Scheme 1). The main reason is that the synthesis procedure was not enantioselective (see Methods). Moreover, it is known from DFT simulations that racemization of [n]helicenes is as much easier as smaller is the number [n] of condensed aromatic rings [51]. As for [5]helicenes, the kinetic barrier was estimated to be $24.4 \text{ kcal} \cdot \text{mol}^{-1}$, [51] which is low enough to allow racemization at temperatures close to RT. We did not perform analogue quantum simulations for nitrogenated [5]

helicenes, but we have no reasons to believe that the kinetic barrier is greatly different for the title compounds.

Figure 2a-d compares the main features of crystal packing of DNH and TNH through the corresponding Hirshfeld surface fingerprint plots (HSFP). For each molecule, HSFP displays the frequency of internal and external distances (d_i , d_e) in Å of any pair of atoms from the corresponding Hirshfeld surface. [52] Accordingly, color and shape of HSFP depend only on how a certain molecular shape fits into a given packing, allowing one to highlight qualitatively the main features of molecular recognition in that structure. As molecular conformation is identical in DNH and TNH (see above), HSFP here compares directly packing differences on the same grounds.

Parallel sharp spikes at $\min[d_i + d_e] \approx 2.4\text{--}2.5$ Å are common to both compounds and are due to weak CH \cdots N hydrogen bonded contacts. In molecule A, two lateral “wings” at greater $d_i + d_e$ (≈ 3 Å) are visible, which can be explained by proximity of CH groups with aromatic rings interactions; these are present also in molecules B and C of TNH, although being less appreciable. The most evident difference is due to the bright region along the main diagonal of the TNH fingerprint around 3.8–4.0 Å, which is barely visible – but still present – in DNH. This is the signature of possible

$\pi\cdots\pi$ stacking interactions, which come with more frequent H \cdots H contacts in TNH. H \cdots H interactions populate the long spike at the centre of the diagram and, being ubiquitous in molecular crystals, do not bear any specific structural meaning, apart implying that, on average, molecules are packed closer in this structure.

Inspection of real-space crystal packing (Fig. 2e-f) complements and explains the findings from HSFP. Stacking interactions are explored with the aromatic analyser of Mercury (Table S8 SI) [32]. In brief, this is a geometrical analysis that ranks such interactions as “strong”, “moderate” or “weak” by comparing the mutual distances and orientations of aromatic rings in each experimental structure with those of benzene pairs whose interaction energies are known at the DFT level in the gas phase. The analysis is qualitative, but results are consistent with both the outcomes from HSFP and Molecular Dynamics (vide infra).

In DNH, some stacking interactions are set up between close-packed molecules with closest rings at ~ 5 Å apart from each other (Fig. 2e). These form columns running along the a cell axis, with only one potentially “strong” interaction according to Mercury (red dotted line in Fig. 2e). [32] Along c , parallel columns of stacked molecules are connected each other by weak CH \cdots N hydrogen bonds that involve only the

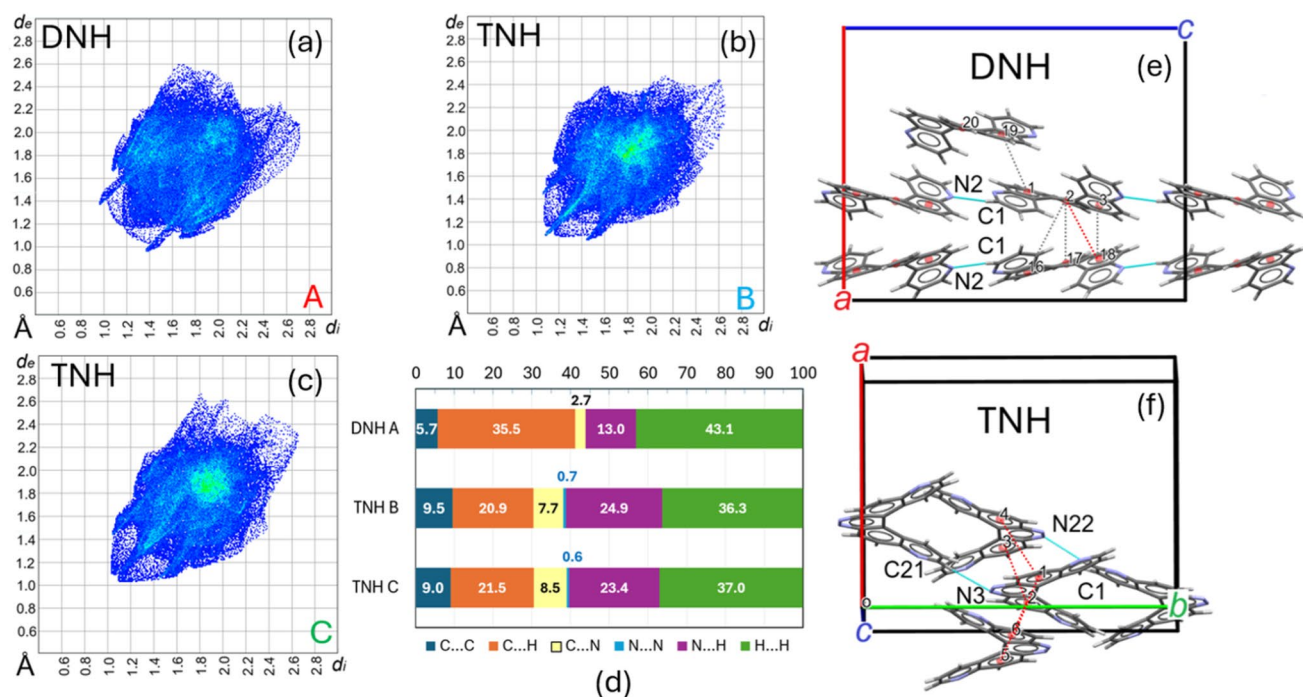


Fig. 2 (a-c) Hirshfeld surface fingerprint plot (HSFP) for symmetry-independent molecules A (DNH) and B, C (TNH) (see Fig. 1). (d) Percent coverage of the HSFP from the various types of atom-atom contacts. (e-f) Relevant intermolecular interactions in the title compounds. Aromatic centroids of coordinated are shown in red, with arbitrary numbering (see Table 2). CH \cdots N hydrogen bonds (Table S9 SI) are coloured in cyan, with labels of donor and acceptor atoms highlighted according to Fig. 1. Dashed lines refer to possible stacking

interactions, with colour referring to their expected strength according to the aromatic analyser utility of Mercury (red: “strong”; grey: “moderate”) [32]. Aromatics numbering is arbitrary and refers to data in Table S8 SI. (e) DNH, as seen along the monoclinic axis. (f) TNH, as seen roughly along the c axis. Only “strong” stacking contacts are shown, as the “moderate” ones cross-link rings also exhibiting “strong” interactions

N2 quinoline nitrogen in position 4 (Fig. 2e, Table S9 SI). Crystallographic glide planes relate such parallel stacked columns along the orthogonal *b* direction (not shown in Fig. 2e).

In TNH, neighbouring molecules lie closer, with nearest rings at ~4 Å apart, and form columnar stackings running along the *a* cell axis (Fig. 2f), where alternating enantiomeric pairs are related by inversion symmetry and produce a “zig-zag” pattern. Stacking interactions are possibly more favourable in TNH than in DNH (red dotted lines in Fig. 2e–f). Along the monoclinic axis *b* of TNH, adjacent columns are held together by weak CH⋯N hydrogen bonds (Fig. 2f, Table S9 SI). At variance with DNH, quinoline nitrogen atoms do not accept hydrogen bonded contacts, which rather involve just one pyridazine nitrogen (either N3 or N22, Figs. 1–2).

Packing energies

Quantitative energy analysis is necessary to understand what interactions prevail in determining the structure. In general, TNH has more negative lattice energy than DNH (Table 2). Both the force fields available in MicMoS predict a difference of ~12–16 kJ·mol⁻¹ (~2.9–3.8 kcal·mol⁻¹) per molecule in favor of TNH, regardless of whether the calculation is carried out on the static X-ray structures, or whether the estimate comes as an average over the last 200 ps of the MD simulations at 298 K.

The PIXEL energies, whose accuracy is more comparable with quantum mechanical models [53], are even closer, with difference not exceeding 4 kJ·mol⁻¹ (~1 kcal·mol⁻¹) per molecule (Table 2). Dissection into individual energy contributions shows that more negative cohesive energies always come with a significant gain in electrostatic terms

E_{el} (Coulomb+polarization), paralleled also by slightly more favorable Lennard–Jones interactions E_{LJ} (dispersion+repulsion). Both static CLP and PIXEL calculations ascribe this gain to electrostatics, as E_{el} terms are more negative in TNH by some kJ·mol⁻¹ per molecule, while E_{LJ} remain almost equal within ~1 kJ·mol⁻¹ per molecule (Table 2). The LJC force field agrees with the other methods in estimating a significantly more negative E_{el} in TNH (by up to 5–8 kJ·mol⁻¹ per molecule) but predicts a considerable gain of dispersive energy as well (up to 8–9 kJ·mol⁻¹ per molecule, Table 2). The reason for this apparent discrepancy resides in how the various energy terms are partitioned in the available functionals of the force field (see the first note to Table 2). At variance with both CLP and PIXEL, LJC does not explicitly account for polarization terms, which instead are implicitly included in the Lennard–Jones functional [37]. In other words, any energy gain due to atomic polarization effects, as implied for example by favorable $\pi\cdots\pi$ stacking interactions, is necessarily included in the E_{LJ} functional by the LJC force field. This picture is consistent with possibly stronger stacking interactions in TNH, as highlighted by the qualitative geometric analysis (see above).

The key role of electrostatics can be traced back to the distribution of electrostatic potential in the title compounds (Fig. 3a). Skeletal substitution of CH groups in 7,8 with nitrogens strongly polarizes the electron density distribution, accumulating negative charges in the region of the pyridazine ring (Fig. 3a, Table S10 SI). Accordingly, MP2/6-31G(p,d) calculations on the isolated molecules at their X-ray geometries predict that DHA is essentially apolar, with a dipole moment of ~0.5 D, while TNH exhibits a roughly tenfold increase of the dipole moment (~4.7 D, Figure S3 SI). The enhanced polarity of TNH was indeed

Table 2 Cohesive energies (E_{coh}) (kJ·mol⁻¹ per molecule) of DNH and TNH, from both static and dynamic calculations, with dissection in terms of individual Coulomb (E_{C}), dispersive (E_{dis}), repulsive (E_{rep}) contributions

Method ^a	E_{C}	E_{pol}	E_{el}^b	E_{dis}	E_{rep}	E_{LJ}^c	E_{coh}^d
DNH	Static, LJC [37]	-14.3	-	-14.3	-193.8	77.0	-116.8
	Static, CLP [34]	-15.6	-15.1	-30.7	-157.8	45.3	-112.5
	Static, PIXEL [39]	-44.5	-20.0	-64.5	-176.5	99.7	-76.8
	MD, LJC ^e	-13.2(1)	-	-13.2(1)	-	-	-107.7(2)
TNH	Static, LJC [37]	-18.9	-	-18.9	-205.2	80.5	-124.7
	Static, CLP [34]	-21.2	-19.2	-40.4	-159.1	45.7	-113.4
	Static, PIXEL [39]	-44.8	-22.4	-67.2	-169.1	91.0	-78.1
	MD, LJC ^e	-20.7(1)	-	-20.7(2)	-	-	-116.4(3)

^aAtom-atom energies, E_{ij} , read $E_{ij}(\text{CLP}) = \frac{q_i q_j}{R_{ij}} + \frac{A_{12}}{R_{ij}^{12}} - \frac{A_6}{R_{ij}^6} - \frac{A_4}{R_{ij}^4}$ and $E_{ij}(\text{LJC}) = \frac{q_i q_j}{R_{ij}} + \frac{A_{12}}{R_{ij}^{12}} - \frac{A_6}{R_{ij}^6}$, being q_i , q_j point atomic charges, A_n empirical 4–6–12 coefficients and R_{ij} the atom–atom distances. PIXEL employs an expression analogue to CLP but exploits interactions among charge density “pixels”, rather than individual atoms. Atomic polarization energy (the R^{-4} term) is not explicitly modelled in LJC. See MicMoS users’ manual [35] and Literature [37, 39] for more details

^bTotal electrostatic energy, $E_{\text{el}} = E_{\text{C}} + E_{\text{pol}}$ (if applicable, see above)

^cTotal Lennard–Jones potential, $E_{\text{LJ}} = E_{\text{dis}} + E_{\text{rep}}$

^dTotal cohesive energy, $E_{\text{coh}} = E_{\text{C}} + E_{\text{pol}} + E_{\text{dis}} + E_{\text{rep}} = E_{\text{el}} + E_{\text{LJ}}$

^eAverage from last 200 ps of 500 ps-long *NpT* Molecular Dynamics trajectories at room temperature. Dispersion and repulsion energies are aggregated. Values in parentheses correspond to 1 estimated standard deviation

exploited already to drive specific interactions with a perhalogenated alkane [48].

Figure 3b shows the PIXEL in-crystal molecule–molecule interaction energies, E_{ij} , as a function of their center of mass (COM) distance. Each point on the graph matches a symmetry-related molecular pair, which can be either homochiral (full dots) or heterochiral (*i.e.* a pair of enantiomers, empty dots). Some points are worth noting. First, molecular pairs with COM distance up to 10 Å provide the most stabilizing contributions to the cohesive energy in both systems. These in fact correspond to the most relevant peaks at short distance of the COM radial distribution functions, $g(r)$, as highlighted by MD simulations (Figs. 3c-d).

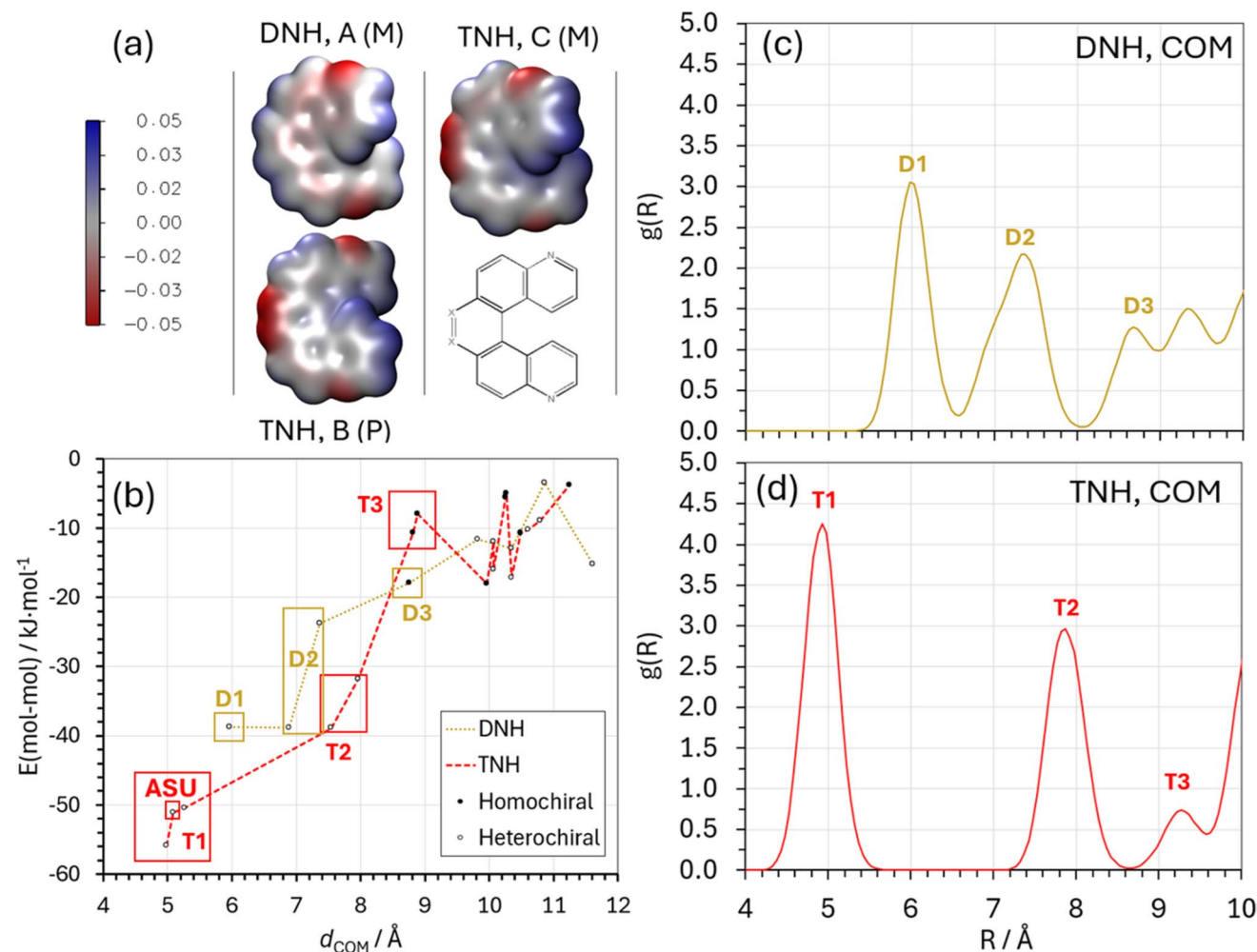
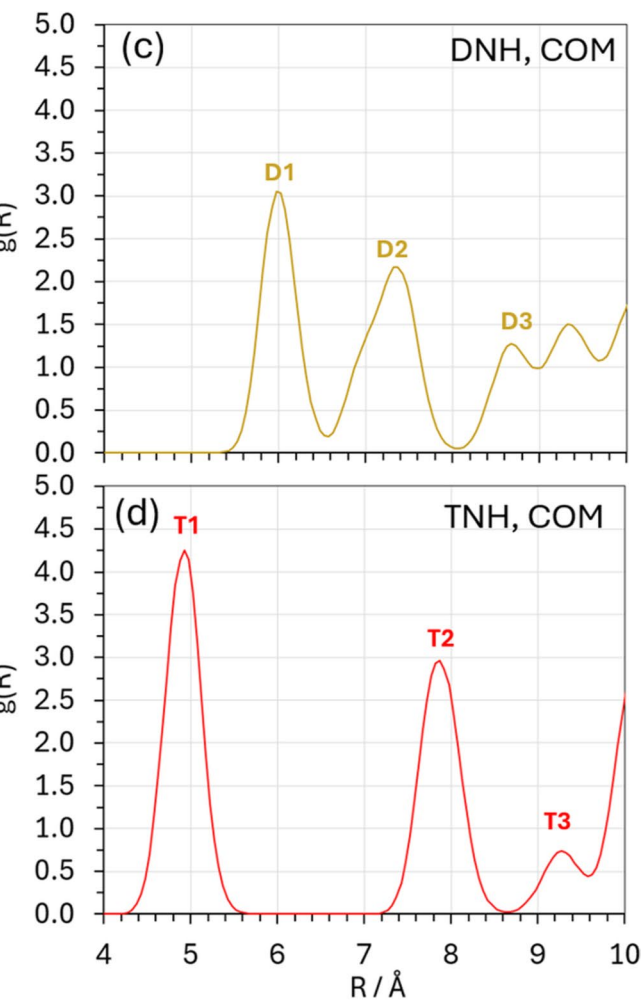


Fig. 3 Electrostatic potential mapped onto the 0.001 charge density isosurfaces of symmetry-independent molecules of DNH and TNH. Both scalar fields were evaluated at the MP2/6-31G(p,d) level of theory at the experimental X-ray geometries and expressed in atomic units (au), where 1 au = 6.7483 e·Å⁻³ for charge density and 1 au = 2625.5 kJ·mol⁻¹ for electrostatic potential. Red (blue) colors highlight regions of negative (positive) electrostatic potential. In the molecular scheme, X=C-H for DNH and X=N for TNH. (b) Total molecule–molecule energies (kJ·mol⁻¹) from PIXEL calculations against the X-ray structures as a function of the molecular center of

mass distance. Numerical entries are shown in Tables S11–S13 SI and in Figure S4 SI. Dashed lines serve only as guides for the eye. Homochiral (heterochiral) pairs are shown as full (empty) dots. Yellow (DNH) and red (TNH) boxes highlight molecular pairs that fall into specific peaks (see labels) of the corresponding radial distribution functions, as retrieved from MD simulations. (c-d) Radial distribution functions, $g(r)$, for molecular center of mass (COM), as a function of the center of mass distance. Labels D1-3 and T1-3 serve just to identify peaks and correspond to the boxes shown in Fig. 3b



mass distance. Numerical entries are shown in Tables S11–S13 SI and in Figure S4 SI. Dashed lines serve only as guides for the eye. Homochiral (heterochiral) pairs are shown as full (empty) dots. Yellow (DNH) and red (TNH) boxes highlight molecular pairs that fall into specific peaks (see labels) of the corresponding radial distribution functions, as retrieved from MD simulations. (c-d) Radial distribution functions, $g(r)$, for molecular center of mass (COM), as a function of the center of mass distance. Labels D1-3 and T1-3 serve just to identify peaks and correspond to the boxes shown in Fig. 3b

generally smaller and not monotonic (compare for example purple and green lines in Figure S4 SI). Interestingly, the curves of electrostatic energies (Figure S4 SI, orange and blue lines) cross at $\sim -6 \text{ kJ}\cdot\text{mol}^{-1}$ at around 9 \AA , meaning that the closest homochiral pairs are equally (and weakly) stabilized by electrostatics in both crystals.

A take-home message from these results is that the significantly greater polarity of TNH dictates its crystal packing. Antiparallel arrangements of polar molecules are favored by second-kind symmetry elements [55], which establish strongly bonded enantiomeric pairs. The electrostatic-driven closer proximity of adjacent molecules in TNH facilitates the setup of stacking interactions between facing aromatic rings. On the contrary, the very low polarity of DNH makes dispersive contributions prevail, and in fact these are almost invariably more negative than the electrostatic ones (Figure S4 SI). The packing of DNH results from the best tradeoff between shape effects, which call for glide operators to fit left-handed with right-handed structures, and attractive van der Waals contributions, possibly enhanced by (weaker) stacking interactions. Finally, weak $\text{CH}\cdots\text{N}$ hydrogen bonds connect glide-related molecules in both crystals (Tables S9 SI and S11-S13 SI) but their contribution is barely significant. In fact, hydrogen-bonded pairs are systematically overwhelmed by closer pairs without hydrogen bonds, which provide significantly more negative molecular contributions to cohesive energy. This evidence raises a warning against relying on pure geometrical criteria, including analysis of HSFP, to determine what interactions are significant to explain cohesion in crystals. [56]

Chiral forms of the title compounds: A CSP study

It is often reported that racemic crystallization is largely preferred over formation of enantiomerically pure conglomerates [57, 58]. Azahelicenes are usually consistent with this general trend [59, 60]. In fact, to the best of our knowledge, homochiral crystals of DNH and TNH were never characterized to date. However, it is known that racemic crystals are not necessarily more stable than their homochiral counterpart. [61] Understanding how energetic and packing factors [62–64] contribute to homochiral crystallization is mandatory to design the best synthetic strategy toward the desired optically active materials. [65] In this context, we generate enantiopure models for the title compounds, with the purpose of providing a tentative – and motivated – answer to the question of whether the crystallization of homochiral DNH and TNH is even experimentally plausible or not. We restricted our screening to $P2_1$ and $P2_12_12_1$ space groups, which host most frequently enantiomerically pure substances, [66] and to $Z'=1$ structures, to deal with predictions comparable on the same grounds. Tables S15 and S16 SI list the 20 top-ranking structures for each prediction.

Figure 4a shows the structural landscape obtained through the SC-MC method [46]. A general linear correlation is appreciable between cohesive energy and packing efficiency, as expected. $P2_12_12_1$ structures (squares in Fig. 4a) tend to achieve greater absolute cohesive energies and C_{pack} than $P2_1$ ones, likely mirroring the greater availability of accessible packing modes in the higher-order symmetry. However, as usual, [61] this is just a general trend more than a rule. In analogy with experimental findings, TNH crystals (blue dots) often display greater cohesion than DNH (red dots), most likely due to more favorable electrostatics (see above). In any case, the title compounds follow Wallach rule, [67] meaning that experimental racemic crystals are more closely packed (and have higher density and greater, or comparable, $|E_{\text{coh}}|$) than homochiral ones. In most $P2_1$ top-ranking predictions, both substances form parallel stacks of aromatic rings piled up along either a or c axes, allegedly exploiting favorable aromatic $\pi\cdots\pi$ interactions. In $P2_12_12_1$ the packing is more complex, but there is always at least one crystallographic direction where patent stacking interactions are set up. Figure S8 SI provides some examples.

An important point is that the space group $P2_1$ is polar and admits nonzero dipole moment in the unit cell, in principle paving the way to the design of TNH-based pyroelectric or ferroelectric materials. Unfortunately, this seems not so likely. Most top-ranking SC-MC predictions for TNH in $P2_1$ consistently bear immaterial ($\ll 1 \text{ D}$) cell dipole moments, since individual molecular dipoles (Figure S9 SI) preferentially lie orthogonal to the unique screw operator along the monoclinic cell axis. Interestingly, this ensures an energetically stable antiparallel arrangement of dipoles (see for example Figure S9a SI), analogue to that imposed by second-kind symmetry operators in the experimental $Pbca$ and $P2_1/c$ lattices. Though, there are some exceptions. For example, the 5th-rank $P2_1$ structure (Figure S9b SI) has nonzero cell dipole ($\sim 7 \text{ D}$), which contributes by $\sim 4 \text{ kJ}\cdot\text{mol}^{-1}$ per molecule to the total cohesive energy.

In general, results from brute-force approaches, including those shown in Fig. 4a, should be considered with caution. To get rid of possible inadequacies of the SC-MC algorithm, particularly of those due to the imposed rigidity of the molecular building blocks, the inclusion of thermal effects is mandatory. [46] Actually, we expect that stable structures lie on reasonably deep potential energy wells. MD allows to check the mechanical stability of the predicted structures by probing the curvature of the potential around the Monte Carlo minimum, with the pros and cons of the same classical force field. Thus, each prediction within the first 10 top-ranked structures underwent 200 ps of MD relaxation (see Methods). Relevant results, together with the new ranking, are shown in Fig. 4b. The first indication of the role of

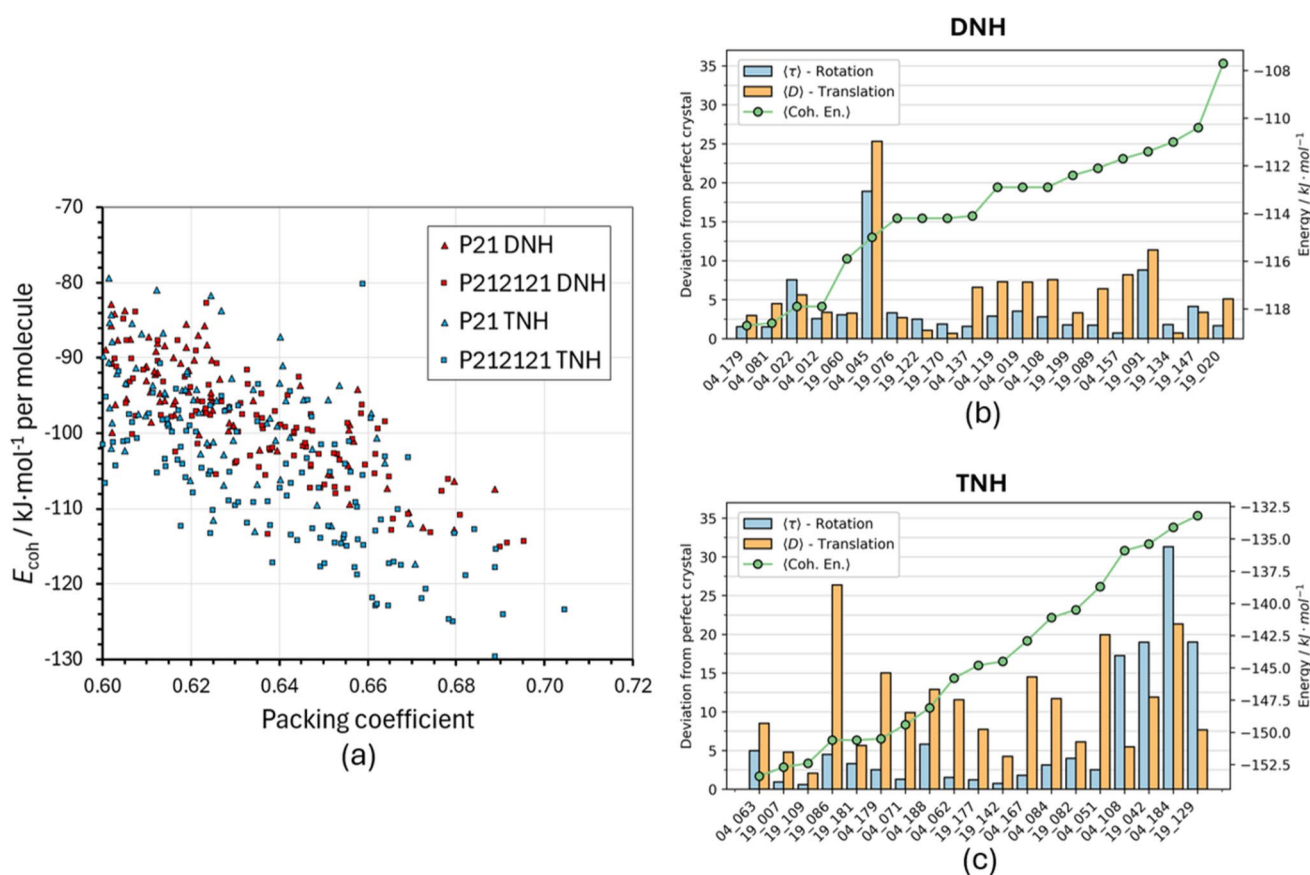


Fig. 4 Structural landscape for possible homochiral structure of DNH and TNH, as estimated by the Symmetry-Constrained Monte Carlo (SC-MC) algorithm. [46] (a) Cohesive energies ($\text{kJ}\cdot\text{mol}^{-1}$ per molecule) vs. packing coefficient (C_{pack}) for all the unique SC-MC predictions with $C_{\text{pack}} \geq 0.6$ (duplicates removed). Red (blue) dots refer to DNH (TNH), while triangles (squares) represent the $P2_1$ ($P2_12_12_1$) space group. All predictions are based on static X-ray geometries, *i.e.* no relaxation of the molecular conformation was allowed. To the sake of comparison, $C_{\text{pack}} = 0.73$ for experimental DNH in $Pbca$ and 0.74 for experimental TNH in $P2_1/c$. (b-c) Final energy ranking for the most

dynamic effects emerges from the structure relaxation, evident in the absolute energy scale. For DNH, the minimum remains essentially unchanged (-119 kJ mol^{-1} , in line with SC-MC values of $\sim -115 \text{ kJ mol}^{-1}$). In contrast, TNH undergoes a marked stabilization, with the most stable structure shifting from -130 to $-152.5 \text{ kJ mol}^{-1}$ per molecule, corresponding to a gain of more than 20 kJ mol^{-1} per molecule.

For DNH, two pairs of energetically equivalent structures were identified, all belonging to the $P2_1$ space group. The 1st, 2nd, and 4th-ranked structures showed reasonable mechanical stability, with rotational correlations below $\sim 2.5\%$ and translational deviations under 5 \AA , consistent with good crystallinity. The 3rd-ranked structure displayed larger displacements (7.5% rotational and 5.0 \AA translational) and was discarded. The packing modes of these 4 structures are similar: the helices pack along the a -axis, with the formation of π - π interactions of translation-related molecules

promising DNH (panel b) and TNH (panel c) structures after applying Molecular Dynamics to top-ranked predictions in (a). Structures are labelled as XX_YYY, where XX is the space group number, and YYY is the cardinal number of the generated structure. The different structures are ranked in terms of energy from left to right. The primary y-axis indicates the difference in terms of rotational ($\langle \tau \rangle$, %) and translational ($\langle D \rangle$, \AA) coefficients from the perfect crystal; the secondary y-axis reports the average cohesive energy ($\langle E_{\text{coh}} \rangle$). The molecular vectors used to calculate $\langle \tau \rangle$ are reported in Figure S7 SI

with different tilting angles. Weak but ubiquitous $\text{C-H}\cdots\text{N}$ and $\text{C-H}\cdots\pi$ interactions are set up along the other directions, connecting different pillars. The packing motifs are reported in Figures S10 and S11, Supporting Information. As expected, the top-ranking $P2_1$ structures after thermalization are all essentially apolar (Table S17 SI), with cell dipole moments never exceeding 1 D. The cohesive energy of the highest-ranked structure after MD relaxation is approximately $-119 \text{ kJ}\cdot\text{mol}^{-1}$, which is comparable to that of the racemate ($\sim -121 \text{ kJ}\cdot\text{mol}^{-1}$; Table 2). The similarity in energy arises from the low polarity of DNH molecules, with dispersion interactions providing the dominant contribution to the total cohesive energy. The small energy difference, which lies within the intrinsic accuracy of the chosen force field ($\approx 5 \text{ kJ}\cdot\text{mol}^{-1}$) [34], does not allow any definitive conclusion to be drawn regarding the relative energetic stability of the enantiomerically pure and racemic structures.

Similar conclusions can be drawn for TNH. Figure 4b shows 3 nearly isoenergetic structures, with an energy gap of $\sim 1 \text{ kJ mol}^{-1}$ with the 4th one. The lowest-energy structure exhibits poor crystallinity, with rotational and translational discrepancies of $\sim 5\%$ and 10 \AA , respectively. Between the second and third ranked structures, the latter proves mechanically more stable. Both belong to the $P2_12_12_1$ space group, which enforces a null dipole moment, aside from minor numerical deviations introduced by molecular dynamics. The polar ($\mu_{\text{cell}} \sim 7 \text{ D}$) structure mentioned above and ranking as 5th in the static SC-MC prediction, shifts to the 18th position once dynamic effects are considered, and shows strong mechanical instabilities (Fig. 4c, structure 04_184). This outcome highlights once more the importance of including molecular dynamics in CSP workflows, as dynamic effects are essential for a reliable assessment of both the energetic ranking and the mechanical stability of the predicted crystal forms. From a structural viewpoint, the top-ranked $P2_1$ form has a similar packing detected for the DNH compound and exploits a cell dipole moment of $\sim 1.6 \text{ D}$, while in the next four $P2_12_12_1$ structures π -stacking motifs are built between C_2 -related molecules (see Figures S10 and S11 SI). As for TNH, the cohesive energy of the highest-ranked structure after MD relaxation ($-153.4 \text{ kJ}\cdot\text{mol}^{-1}$) differs significantly from that of the racemic crystal ($-137.1 \text{ kJ}\cdot\text{mol}^{-1}$, Table 2). Thus, the enantiomerically pure polymorph is predicted to be thermodynamically more stable than the corresponding racemate. A decomposition of the cohesive energy reveals that this stabilization is mainly driven by the Lennard–Jones contribution, which amounts to $-151.5 \text{ kJ}\cdot\text{mol}^{-1}$, substantially larger than the corresponding value observed for the racemic form ($E_{\text{LJ}} = -116.3 \text{ kJ}\cdot\text{mol}^{-1}$, Table 2). In contrast, the Coulombic contribution in the enantiopure crystal is nearly negligible ($-1.9 \text{ kJ}\cdot\text{mol}^{-1}$), whereas a significantly larger electrostatic contribution is found in the racemate ($-20.7 \text{ kJ}\cdot\text{mol}^{-1}$). These results indicate that the enantiomerically pure crystal adopts a more efficient packing motif, which does not maximize electrostatic interactions associated with the relatively large molecular dipole moment of TNH, but instead favors dispersion-dominated stabilization. As a result, enhanced packing efficiency leads to a lower overall cohesive energy. The case of TNH provides a further example of how the crystal packing of an organic substance depends, often in a non-trivial way, on a subtle interplay of competitive intermolecular interactions [68]. However, it must be stressed that, despite this predicted thermodynamic preference, the enantiomerically pure form of TNH has not been experimentally observed to date.

In conclusion, $P2_1$ structures seem to be favored for both compounds, even though some $P2_12_12_1$ lattices cannot be

excluded for TNH. In any case, no unquestionable evidence of pyroelectric behavior was found in energetically and mechanically stable crystal forms of enantiomerically pure TNH and DNH. Such predictions wait for experimental confirmation and possibly call for a full study of the crystal structure landscape to be confirmed.

Conclusions

In this work, a thorough crystal chemical study was carried out on two symmetrically nitrogenated [5]helicenes, namely 4,11-diaza[5]helicene (DNH) and 4,7,8,11-tetraaza[5]helicene (TNH). Their crystal structure was accurately determined by X-ray diffraction at room temperature. Experiments were complemented by both quantum mechanical calculations at the MP2/6-31G(p,d) level and classical force-field based simulations, to retrieve as much accurate intermolecular interaction energies.

It was found that the conformation of the [5]helicene scaffold does not depend on nitrogen substitution in positions 7 and 8. Therefore, differences in the crystal packing can be genuinely attributed to the skeletal replacement of two CH groups in 7,8. with nitrogen atoms. Both crystals are centrosymmetric racemates and contain an equimolar amount of P and M enantiomers, as expected owing to the non-enantioselective synthesis procedure. Both crystals exploit stacked columnar arrangements of aromatic rings to achieve efficient packing and differ in how such stacking motifs are mutually oriented, as well as in the relative distance between facing rings. TNH exhibits closer packing and greater cohesion. Weak $\text{CH}\cdots\text{N}$ hydrogen bonds are also present, but they are overwhelmed by stronger contributions to cohesive energies and barely bear any structural significance.

In TNH, electrons are accumulated in the central pyridazine ring, polarizing the static density so that a tenfold increase in the molecular dipole moment is observed. Accordingly, the TNH crystal is dominated by antiparallel packing of adjacent dipole moments. The attractive dipole–dipole interactions present in TNH, which are entirely absent in DNH, likely draw neighboring molecules closer together. This leads to a significant $\sim 5\%$ increase in crystal density compared to DNH.

Finally, the crystal structure landscape of both compounds was explored by the recently developed SC-MC algorithm, to look for plausible homochiral forms in common $P2_1$ and $P2_12_12_1$ space groups. Based on energy ranking and Molecular Dynamics simulations, enantiomerically pure DNH and TNH might indeed crystallize in some $P2_1$ forms, although patent polar forms are improbable. TNH can in principle also have $P2_12_12_1$ forms that are favored

both thermodynamically and mechanically. We are aware that the present screening is necessarily partial, as real-life scenarios might require to consider other Sohncke groups. For sure, our predictions suggest that homochiral forms of the title compounds will likely imply significant stacking motifs, while TNH should exploit first-kind symmetry elements to maximize the antiparallel pairing of molecular dipole moments. This information could be of some use in future crystallization screening experiments, for example to select solvents or solvent mixtures with the proper polarity.

The problem of why $Z'=2$ in TNH and $Z'=1$ in DNH remains open. As suggested by one Reviewer, $P2_1/c$ (TNH) is a maximal non-isomorphic subgroup of $Pbca$ (DNH). This implies that more efficient packing is related to the loss of a coset of symmetry elements, which allows to lose the symmetry of first-neighbor molecular pairs. Extensive experimental screening of the crystal structure landscape would be useful to gain further insights on this matter.

Supplementary Information The online version contains supplementary material available at <https://doi.org/10.1007/s11224-026-02724-6>.

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Data availability Relevant steering parameters and procedures for computer simulations described in this paper are deposited in the Supporting Information. Molecular dynamics trajectories and outputs of quantum chemical simulations can be downloaded freely from the public repository Zenodo (<https://doi.org/10.5281/zenodo.17407385>).

Declarations

Ethics approval Not applicable, as no human participants or animals were involved in this research.

Competing interests The authors declare no competing interests.

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