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Synthesis, X-ray characterization and DFT calculations of a series of 3-substituted 4,5-dichloroisothiazoles†

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This manuscript reports a synthetic protocol to obtain 3-substituted 4,5-dichloroisothiazole derivatives in good yields. This is exemplified by the synthesis of six new compounds, which have been spectroscopically characterized including their solid state structure by single crystal X-ray analysis. As substituents in position 3, both 1,4-dihydropyridinyl and 4*H*-pyranyl moieties have been used, decorated with several groups like cyano, amino, acetyl, ester and methyl, thus providing a variety of possibilities to establish noncovalent interactions. The assemblies have been analysed theoretically, using DFT calculations, molecular electrostatic potential (MEP) surfaces and QTAIM/NCIplot calculations. The π -stacking between the 4,5-dichloroisothiazole rings is a recurrent motif in the solid state of most compounds, in addition to other less conventional interactions like lone pair (LP) $\cdots \pi$ and halogen bonding contacts. Another interesting motif is the formation of centrosymmetric $R_2^2(12)$ motifs where the cyano groups concurrently establish H-bonds and antiparallel CN \cdots CN interactions.

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Introduction

The deep understanding of noncovalent interactions is crucial for the development of supramolecular chemistry and crystal engineering.^{1–3} The analysis of X-ray structures is very useful to keep progressing in the adequate interpretation of the structure directing role of non-covalent interactions.^{4–6} Moreover, the application of theoretical methods is also convenient to investigate the energetic features of interactions and their geometric requirements, allowing the design and

synthesis of new compounds with tailored properties.^{7,8} Furthermore, the utilization of computational tools like the quantum theory of atoms-in-molecules,⁹ noncovalent interaction plots¹⁰ or molecular electrostatic potentials¹¹ is also very convenient to investigate and disclose the physical nature of interactions, charge transfer effects, *etc.*

Substituted isothiazoles and their fused derivatives are important in the pharmacological industry¹² due their analgesic, antipyretic, fungicidal, and herbicidal properties.¹³ For instance, monocyclic isothiazole skeletons such as sulfasomizole and denotivir display antibacterial and antiviral properties.¹⁴ Fused isothiazole derivatives also exhibit interesting effects like cytotoxicity¹⁵ and prostaglandin release regulation from human synovial sarcoma cells. Moreover, the antipsychotic drug zipracidone possessing a [*d*]fused isothiazole ring was proposed to treat schizophrenia and bipolar disorder.¹⁶

It has been also demonstrated that isothiazole scaffolds have emissive properties when they are attached to purine or pyridine nucleobases. Polyfunctional isothiazole derivatives have been also used for the construction of metal complexes of different complexities, like organometallic frameworks and functional materials.¹⁷ It is documented that Co(II) and Cu(II) complexes of 4,5-dichloroisothiazole-3-carboxylic acid and 1,10-phenanthroline as a coligand present interesting cytotoxicity effects.^{18,19}

4*H*-pyrans and 1,4-dihydropyridines are also valuable scaffolds for pharmaceutical chemistry due to their wide

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spectrum of biological activities.^{20,21} Functionally substituted 4*H*-pyrans were found among natural products, compounds with antitumor, antibacterial, antiviral, fungicidal, antispasmodic and diuretic activity.^{20,22} 1,4-Dihydropyridines are one of the most important groups of calcium channel modulators widely used in the treatment of cardiovascular diseases.²¹ These include an extensive group of antihypertensive “dipine” drugs like amlodipine, benidipine, *etc.*

1,4-Dihydropyridine is the main heterocyclic core of compounds displaying antitumor, antioxidant, and gero-, hepato- and neuroprotective effects as well as of derivatives effective in the treatment of diabetes, ischemia, and Alzheimer's disease.²³

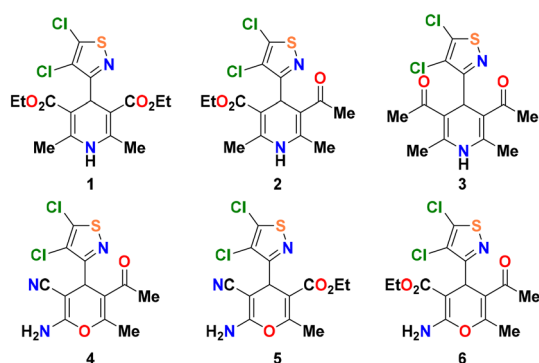
The relevance of 4*H*-pyran and 1,4-dihydropyridine derivatives is emphasized by the abundance of studies on optimizing their synthesis.²⁴ These biologically active 4*H*-pyrans and 1,4-dihydropyridines often bear an aromatic substituent in the 4 position. However, there are only a few examples with a heterocyclic core in this position.^{23–25} Isothiazole containing derivatives of this type are almost unknown.²⁶

In this work, a convenient synthetic protocol to obtain 4*H*-pyrans and 1,4-dihydropyridines containing a 4,5-dichloroisothiazole fragment in the 4 position is reported, as exemplified by the generation of compounds **1–6** shown in Scheme 1. The target compounds are very rich in functional groups that allow the generation of supramolecular synthons in the solid state that are described in this work and analysed theoretically using a variety of computational tools.

Methods

Synthesis and characterization

Synthesis of heterocycles **1–6** was performed according to known procedures for preparation of 4*H*-pyrans and 1,4-dihydropyridines using 4,5-dichloroisothiazole-3-carbaldehyde²⁷ and commercially available methylene-active components (acetylacetone, malononitrile, ethyl 2-cyanoacetate) as starting materials. Single crystals of **1–6** for XRD analysis were obtained by slow evaporation of their saturated solutions in methanol.



Scheme 1 Chemical diagram of compounds **1–6** reported in this work.

In addition to XRD, all obtained compounds were characterized using standard LCMS, FTIR and NMR spectroscopy. The presence of the 4,5-dichloroisothiazole fragment was elucidated from the characteristic position of quaternary carbon atom signals in ¹³C NMR spectra (δ 101.8–115.5, 118.1–123.3 and 146.2–149.1 ppm). In ¹H NMR spectra, chemical shifts of the protons at the tertiary carbon atoms of the pyran and pyridine systems are observed in the area of δ 4.91–5.50 ppm. Dihydropyridine NH group signals are observed at δ 6.24–6.49 ppm. A slight W-splitting of the proton signals at the tertiary carbon atom (H-4) and one of the methyl groups (⁵*J*_{4,Me} ~ 0.7 Hz) was observed in the cases of **5** and **6**.

X-ray analysis

The crystal structure of heterocycles **2**, **3**, **5** and **6** was determined by X-ray structural analysis using an automatic four-circle area-detector diffractometer Bruker KAPPA APEX II with MoK α radiation. The crystal structure **1** and **4** was determined by X-ray structural analysis using an XtaLAB Synergy, Dualflex, HyPix with CuK α radiation. The cell parameters of **2**, **3**, **5** and **6** were refined over the entire data set, together with data reduction using SAINT-Plus software.²⁸ Absorption corrections of **2**, **3**, **5** and **6** were introduced using the SADABS program.²⁹ The structures were solved using the SHELXT-2018/2 program³⁰ and refined by full-matrix least squares on *F*² in the anisotropic approximation for all non-hydrogen atoms (SHELXL-2018/3).³¹ The C–H bonded hydrogen atoms were placed in geometrically calculated positions and refined in an idealized geometry with isotropic temperature factors equal to 1.2*U*_{eq}(C) for CH and CH₂-groups, and 1.5*U*_{eq}(C) for CH₃-groups. The N–H bonded atoms were objectively located from difference Fourier synthesis and refined with isotropic temperature factors equal to 1.2*U*_{eq}(N) for NH and NH₂ groups. The orientation of CH₃-groups was refined. Structures **3** and **6** were refined as inversion twins. Tables and figures for the structures were generated using Olex2.³² Crystal data, data collection, and structure refinement details are summarized in Table 1. All other crystallographic parameters of the structures are indicated in Tables S1–S24 (see the ESI†). The atomic coordinates were deposited at the Cambridge Crystallographic Data Centre (CCDC).³³ The CCDC numbers are 2213501–2213504 for **2**, **3**, **5**, and **6**, correspondingly, and 2243955 and 2243956 for **1** and **4**.

DFT calculations

The calculations reported herein were performed at the PBE0 (ref. 34)–D3 (ref. 35)/def2-TZVP (ref. 36) level of theory using the Turbomole 7.0 program.³⁷ The binding energies were computed as the difference between the energy of the assembly and the sum of the isolated monomers. The energies have been corrected for the basis set superposition error.³⁸ The MEP surfaces were generated using a 0.001 isosurface to emulate the van der Waals envelope. The



Table 1 Crystal data and structure refinement for structures 1–6

Identification code	1	2	3	4	5	6
CCDC number	2243955	2236983	2236984	2243956	2236985	2236986
Empirical formula	C ₁₆ H ₁₈ N ₂ O ₄ SCl ₂	C ₁₅ H ₁₆ N ₂ O ₃ SCl ₂	C ₁₄ H ₁₄ N ₂ O ₂ SCl ₂	C ₁₂ H ₉ N ₃ O ₂ SCl ₂	C ₁₃ H ₁₁ N ₃ O ₃ SCl ₂	C ₁₄ H ₁₄ N ₂ O ₄ SCl ₂
Formula weight	405.28	375.26	345.23	330.18	360.21	377.23
Temperature/K	100(2)	100(2)	100(2)	100(2)	100(2)	100(2)
Crystal system	Triclinic	Triclinic	Monoclinic	Triclinic	Triclinic	Monoclinic
Space group	<i>P</i> $\bar{1}$	<i>P</i> $\bar{1}$	<i>P</i> 2 ₁	<i>P</i> $\bar{1}$	<i>P</i> $\bar{1}$	<i>P</i> 2 ₁
<i>a</i> /Å	8.7279(3)	7.3058(11)	7.2276(5)	8.1335(2)	7.5420(14)	12.2812(6)
<i>b</i> /Å	9.6603(3)	10.1615(14)	26.5464(19)	9.2637(2)	8.8365(16)	7.3500(4)
<i>c</i> /Å	11.4482(4)	12.4124(18)	8.5613(6)	10.0924(3)	11.384(2)	17.4340(9)
α /°	86.524(3)	105.388(6)	90	72.817(2)	83.566(6)	90
β /°	77.743(3)	106.574(6)	112.990(3)	72.328(3)	83.963(6)	97.212(2)
γ /°	71.031(3)	100.321(6)	90	76.935(2)	88.682(6)	90
Volume/Å ³	891.98(6)	818.5(2)	1512.16(19)	684.39(3)	749.7(2)	1561.26(14)
<i>Z</i>	2	2	4	2	2	4
ρ_{calc} g cm ^{−3}	1.509	1.523	1.516	1.602	1.596	1.605
μ /mm ^{−1}	4.588	0.539	0.572	5.746	0.587	0.571
<i>F</i> (000)	420.0	388.0	712.0	336.0	368.0	776.0
Crystal size/mm ³	0.25 × 0.2 × 0.18	0.4 × 0.05 × 0.04	0.32 × 0.2 × 0.16	0.24 × 0.21 × 0.15	0.26 × 0.14 × 0.06	0.5 × 0.4 × 0.32
Radiation	CuK α	MoK α	MoK α	CuK α	MoK α	MoK α
$(\lambda = 1.54184)$	$(\lambda = 1.54184)$	$(\lambda = 0.71073)$	$(\lambda = 0.71073)$	$(\lambda = 1.54184)$	$(\lambda = 0.71073)$	$(\lambda = 0.71073)$
2 θ range for data collection/°	7.904 to 158.912	8.186 to 59.998	8.674 to 59.998	9.484 to 159.122	8.162 to 59.986	8.192 to 59.998
Index ranges	−10 ≤ <i>h</i> ≤ 11, −12 ≤ <i>k</i> ≤ 12, −14 ≤ <i>l</i> ≤ 14	−8 ≤ <i>h</i> ≤ 10, −14 ≤ <i>k</i> ≤ 14, −17 ≤ <i>l</i> ≤ 17	−10 ≤ <i>h</i> ≤ 10, −37 ≤ <i>k</i> ≤ 37, −12 ≤ <i>l</i> ≤ 11	−9 ≤ <i>h</i> ≤ 10, −11 ≤ <i>k</i> ≤ 11, −12 ≤ <i>l</i> ≤ 12	−10 ≤ <i>h</i> ≤ 10, −12 ≤ <i>k</i> ≤ 12, −15 ≤ <i>l</i> ≤ 16	−17 ≤ <i>h</i> ≤ 17, −10 ≤ <i>k</i> ≤ 9, −24 ≤ <i>l</i> ≤ 24
Reflections collected	19 809	18 356	25 913	14 966	13 476	26 683
Independent reflections	3836 [<i>R</i> _{int} = 0.0591, <i>R</i> _{sigma} = 0.0317]	4650 [<i>R</i> _{int} = 0.1086, <i>R</i> _{sigma} = 0.1363]	8579 [<i>R</i> _{int} = 0.0655, <i>R</i> _{sigma} = 0.0936]	2945 [<i>R</i> _{int} = 0.0511, <i>R</i> _{sigma} = 0.0293]	4251 [<i>R</i> _{int} = 0.0823, <i>R</i> _{sigma} = 0.1054]	8322 [<i>R</i> _{int} = 0.0379, <i>R</i> _{sigma} = 0.0527]
Data/restraints/parameters	3836/0/234	4650/0/215	8579/2/393	2945/0/190	4251/0/207	8322/1/434
Goodness-of-fit on <i>F</i> ²	1.061	0.965	1.016	1.031	0.993	1.037
Final <i>R</i> indices [<i>I</i> ≥ 2 σ (<i>I</i>)]	<i>R</i> ₁ = 0.0415, <i>wR</i> ₂ = 0.1139	<i>R</i> ₁ = 0.0633, <i>wR</i> ₂ = 0.1243	<i>R</i> ₁ = 0.0507, <i>wR</i> ₂ = 0.0867	<i>R</i> ₁ = 0.0358, <i>wR</i> ₂ = 0.0985	<i>R</i> ₁ = 0.0527, <i>wR</i> ₂ = 0.1013	<i>R</i> ₁ = 0.0362, <i>wR</i> ₂ = 0.0697
Final <i>R</i> indices [all data]	<i>R</i> ₁ = 0.0421, <i>wR</i> ₂ = 0.1147	<i>R</i> ₁ = 0.1396, <i>wR</i> ₂ = 0.1520	<i>R</i> ₁ = 0.0840, <i>wR</i> ₂ = 0.0982	<i>R</i> ₁ = 0.0363, <i>wR</i> ₂ = 0.0992	<i>R</i> ₁ = 0.1077, <i>wR</i> ₂ = 0.1207	<i>R</i> ₁ = 0.0446, <i>wR</i> ₂ = 0.0727
Largest diff. peak/hole/e Å ^{−3}	0.54/−0.61	0.47/−0.68	0.45/−0.42	0.64/−0.50	0.58/−0.64	0.38/−0.28
Flack parameter	—	—	0.04(4)	—	—	0.30(5)

QTAIM (ref. 9) and NCIPLOT (ref. 39) analyses were performed at the same level using the MultiWFN program⁴⁰ and represented using the VMD software.⁴¹ The NCIPLOT method is convenient to reveal interactions in real space. It uses reduced density gradient isosurfaces and a colour code (based on the sign of the second eigenvalue of ρ , λ_2) to identify the attractive or repulsive nature of the interactions. The following settings were used in this work: RDG = 0.5, density cut-off = 0.04 a.u., and colour scale $-0.03 \text{ a.u.} \leq \text{sign}(\lambda_2(\rho)) \leq 0.03 \text{ a.u.}$ Blue and green colours are used here to identify strongly and moderately attractive interactions, respectively.

Results and discussion

Synthesis

Since isothiazole derivatives of 1,4-dihydropyridines and 4*H*-pyrans were not previously described, their synthesis was carried out by known methods,⁴² including ones used for the preparation of other heterocyclic derivatives of this type. 4,5-Dichloroisothiazole-3-carbaldehyde **7** (ref. 27) and its condensation products with acetoacetic ester, acetylacetone,

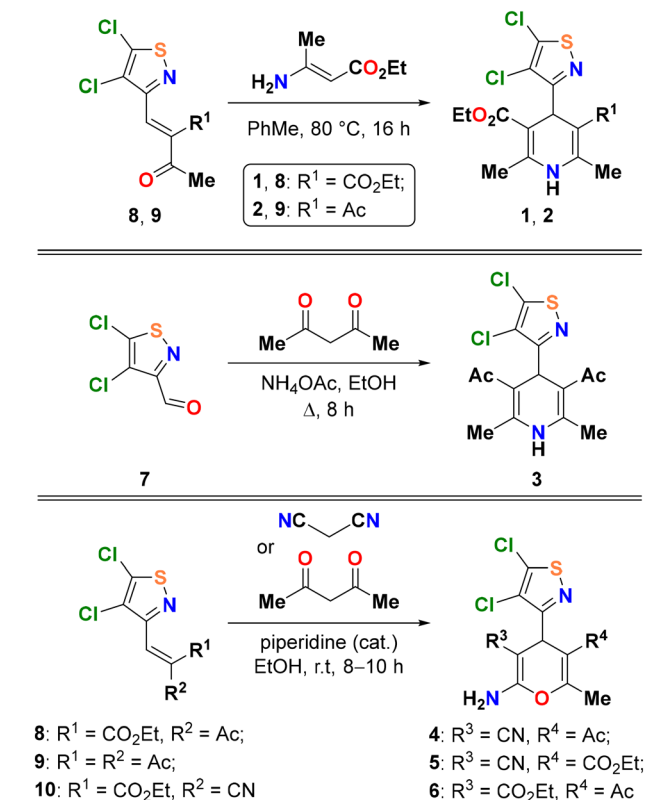
and cyanoacetic ester **8–10** were chosen to be starting compounds (Scheme 2).

Dihydropyridines **1** and **2** were obtained by the interaction of ethyl 3-aminocrotonate with α,β -unsaturated heterocyclic derivatives **8** and **9** (Scheme 2). Ethanol is a common solvent for such processes; however, the reaction in ethanol was not selective. The possible reason for this is side processes involving the nucleophilic substitution of the active chlorine atom in the 5th position of the isothiazole cycle. The same reactions in toluene proceeded smoothly; therefore, compounds **1** and **2** were isolated in 56% and 59% yields, respectively.

The classical Hantzsch synthesis was chosen to obtain symmetric 4-isothiazole-1,4-dihydropyridine **3** from aldehyde **7** (Scheme 2). The yield of product **3** (58%) was similar to those of compounds **1** and **2**. This way seems to be more efficient, since it does not require the preliminary synthesis of the intermediate α,β -unsaturated isothiazole from the corresponding aldehyde **7** and acetylacetone, which proceeds with a high but not quantitative yield.

Finally, isothiazole-containing 4*H*-pyrans **4** and **5** were prepared by the condensation of α,β -unsaturated compounds





8 and 9 with malononitrile; compound 6 was prepared from ethyl 2-cyanocrotonate 10 and acetylacetone. These reactions were catalyzed by piperidine as a base catalyst and yielded the target products with acceptable to good yields (56–76%) even in ethanol.

Structural description of the X-ray structures

The molecular structures of the obtained compounds 1–6 are presented in Fig. 1. Structures 2 and 4 contain one independent molecule of the complex; structures 3 and 6 contain two crystallographically independent molecules. At the same time, 3 contains two molecules with the same conformation (RMSD without inversion is 0.408 Å), and 6 contains right and left independent molecules (Fig. 2), which, however, practically coincide upon inversion (RMSD with inversion is 0.654 Å).

The chlorine atoms are in the plane of the isothiazole rings (the maximum deviation from the plane is 0.02 Å for one of the molecules in 6). The S–N distances in all compounds are close and vary from 1.65 to 1.66 Å. It is worth noting that the isothiazole fragments are turned perpendicular to the plane of the central six-membered ring (the angles between the isothiazole and pyridinyl/pyranil fragments are close to 90°). In all compounds except 5, the central six-membered rings containing a nitrogen or oxygen atom have a slightly distorted conformation. Most of all, the C4 carbon atom deviates from the plane of the six-membered

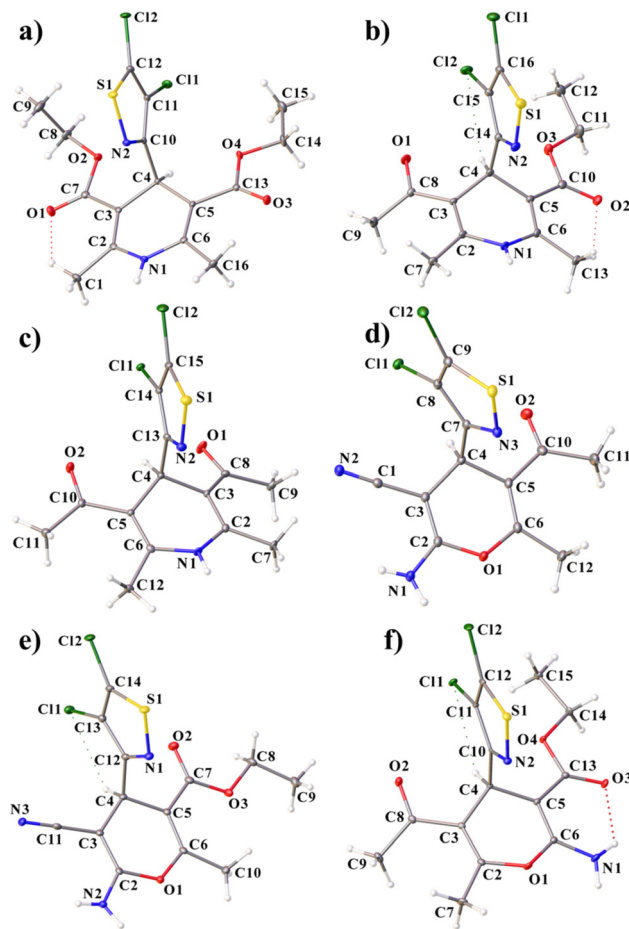


Fig. 1 Molecular structure of 1–6 (a–f), including atom labelling. Only one independent molecule is shown in 3 and 6. Displacement ellipsoids are drawn at the 50% probability level.

ring. In 5, distortion of the ring conformation is practically not observed (RMSD is 0.025 Å).

When searching in the CCDC, only two compounds were found containing isothiazole and pyridinyl fragments,⁴³ but with other bulky substituents in the five-membered ring; therefore, it is difficult to compare them with those obtained in this work. However, it should be noted that in the described analogues, the pyridinyl ring was not substituted

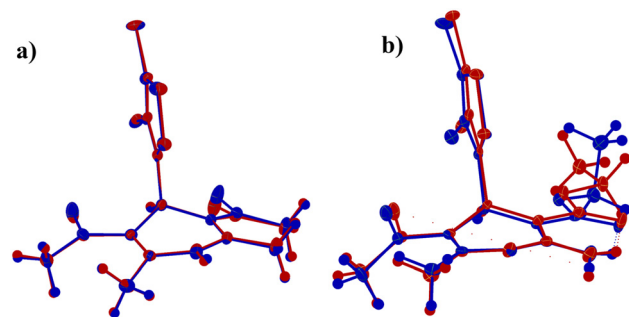


Fig. 2 Superposition of two crystallographically independent molecules in 3 (a) and 6 (b).



and had a planar geometry, while the isothiazole ring was rotated differently.

In all the structures, intramolecular hydrogen bonds of the C–H \cdots Cl type are present. In **2**, an additional weak hydrogen bond of the C–H \cdots O type appears. In **3**, a stronger N–H \cdots O bond compared to the rest is present.

In compounds **2**, **3**, and **6**, the distances between the center of the six-membered ring and the nitrogen atom of the isothiazole fragment vary from 2.71 Å to 2.78 Å, which are shorter than the sum of the van der Waals radii of C and N, and the angle α is close to 70° (Fig. 3). There is an increase in this growth to 2.93–3.11 Å and a decrease in the angle α to 53–61° in **1**, **4** and **5**. All this allows us to assume the presence of an intramolecular N \cdots π interaction in the molecules.⁴⁴

All the compounds present good H-bond donor and acceptor groups and, consequently, exhibit H-bonded assemblies in the solid state as shown in Fig. 4. In most of them, the H-bonds propagate the molecules into 1D supramolecular polymers, as observed in compounds **2**, **3**, **4** and **6**, where either the NH of the 1,4-dihydropyridinyl ring (**2** and **3**) or the exocyclic NH₂ group of the 4*H*-pyranyl ring (**4**

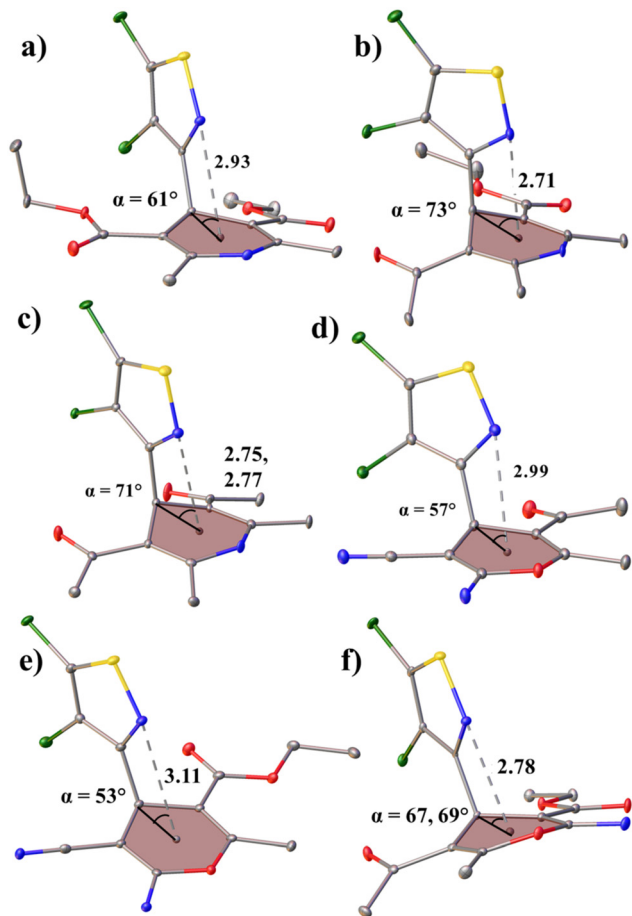


Fig. 3 View showing N \cdots π interactions in structures **1**–**6** (a–f). Distances in Å. H-atoms are omitted for clarity. Only one independent molecule for **3** and **6** is shown.

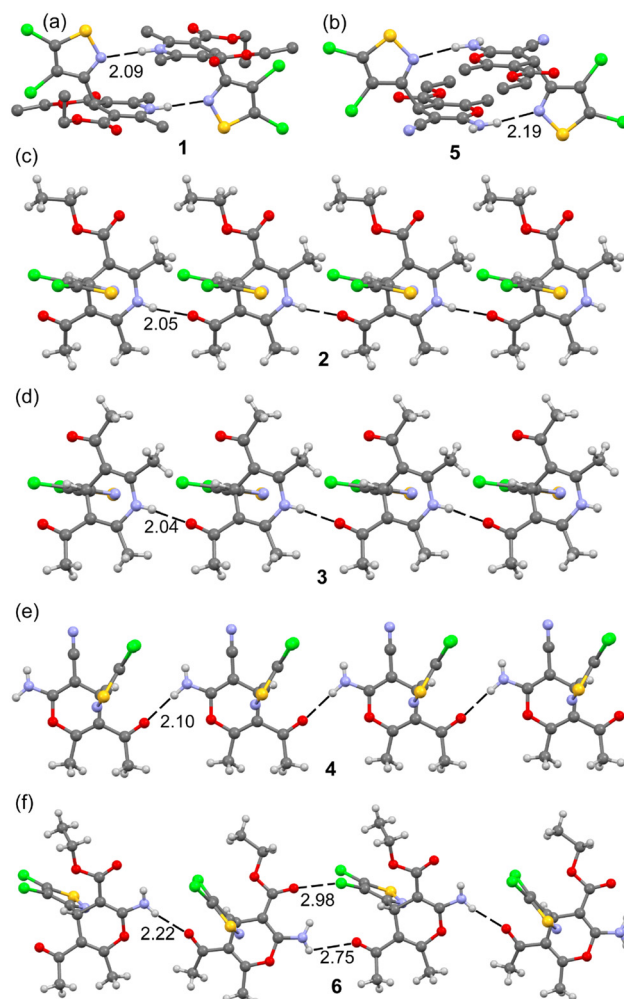


Fig. 4 H-bonded assemblies observed in the solid state X-ray structures of compounds **1** (a), **5** (b), **2** (c), **3** (d), **4** (e) and **6** (f). Distances in Å.

and **6**) acts as a H-bond donor and the O-atom of the carbonyl group (from either keto or ester groups) acts as a H-bond acceptor. It is interesting to comment on compound **6** (Fig. 4f) where two different and alternate types of interactions connect the monomers. On the one hand, there is a directional NH \cdots O H-bond with a typical distance (2.22 Å) and on the other hand, there is a longer and less directional NH \cdots O contact (2.75 Å) in combination with a C–Cl \cdots O halogen bond (2.98 Å) involving one of the Cl-atoms of the 4,5-dichloroisothiazole moiety and the carbonyl O-atom of the ester group. Compounds **1** and **5** (see Fig. 4a and b) form discrete self-assembled dimers instead of 1D polymers, where the H-bond acceptor is the N-atom of the 4,5-dichloroisothiazole moiety. The NH \cdots O distances range from 2.05 to 2.22 Å (apart from the special XB/HB assembly in **6**), which are similar to the NH \cdots H distances in **1** and **5** (2.09 Å and 2.19 Å, respectively). Further analysis of these H-bonds is provided in the theoretical section below.

Additional H-bonded assemblies were also observed in the solid state of compounds **4** and **5** that contain the cyano and



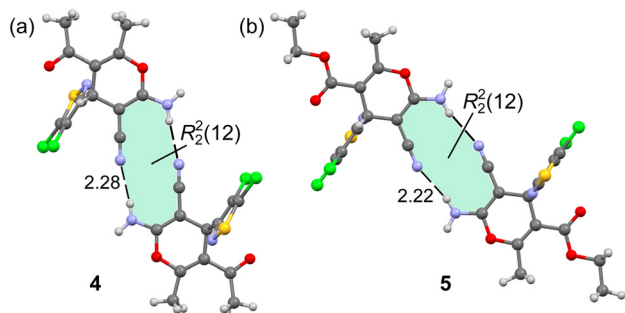


Fig. 5 H-bonded centrosymmetric dimer observed in the solid state X-ray structures of compounds 4 (a) and 5 (b). Distances in Å.

amino substituents attached to the 4*H*-pyranyl ring. Actually, the exocyclic amino group has two protons suitable to form H-bonds, the ones shown in Fig. 4 and another one with the N-atom of the CN group as an acceptor. The ability of the cyano group to establish noncovalent interactions including HBs has been analyzed before in the literature.⁴⁵ Both compounds form centrosymmetric $R_2^2(12)$ supramolecular rings in the solid state, as highlighted in Fig. 5. The NH...N distances are 2.28 and 2.22 Å for 4 and 5, respectively. The conjugation of the electron lone pair of the amino group to the cyano group likely increases the electron acceptor ability of the sp-hybridized N-atom, as studied below.

It is also worthy to comment on the different π -stacked assemblies observed in some of the compounds reported herein. That is, three different binding modes are observed for the π - π interactions between the 4,5-dichloroisothiazole rings, which are detailed in Fig. 6. In the first one, only one carbon atom of each ring are in contact (van der Waals distance), as observed in compound 1 (see Fig. 6a).

This binding mode leaves one of the Cl-atoms over the centre of the aromatic ring of the adjacent molecule and *vice versa* (see blue dashed lines in Fig. 6a), thus establishing LP... π interactions. In the second binding mode, two atoms

of each ring are in contact, one C-atom of one ring interacts with the S-atom of the other ring and *vice versa*. This binding mode is observed in compounds 2 and 5 with distances of 3.53 Å and 3.63 Å, respectively. Finally, in the third binding mode, the N-atom of one ring interacts with the S-atom of the other ring and *vice versa*. This binding mode is observed in compound 4 with a distance of 3.52 Å. The energetic differences between these three binding modes are small, as discussed in the following section.

DFT analysis

The molecular electrostatic potential surfaces of compounds 2, as a representative example for the 1,4-dihydropyridinyl series, and 4 for the 4*H*-pyranyl series are presented in Fig. 7. It can be observed that the MEP maximum is located at the NH group in 2 (51.5 kcal mol⁻¹) and at the NH₂ group in 4. The MEP minimum in 2 is located at the keto O-atom (−42.7 kcal mol⁻¹) which is a better H-bond acceptor than the carbonyl O-atom of the ester group (−35.1 kcal mol⁻¹). In compound 4, it is interesting to highlight that the MEP minimum is located at the sp-hybridized N atom of the cyano group, confirming that the conjugated LPs of the N and O-atoms bonded at the other end of the double bond increase the electron density at the N-atom of the cyano group. The MEP at the O-atom of the keto group is also large and negative (−35.7 kcal mol⁻¹); thus it is also adequate to participate in H-bonds. The MEP also reveals the existence of a σ -hole opposite to the C–Cl bond that is more intense in the Cl-atom bonded at position 5. A detail of the MEP surface around these Cl-atoms is presented at the bottom part of the figure for both compounds, revealing the anisotropy and a modest MEP value at the σ -hole (+12.0 and +13.2 kcal mol⁻¹ for 2 and 4, respectively).

H-bonded dimers extracted from the 1D assemblies described above (see Fig. 4) have been further analysed

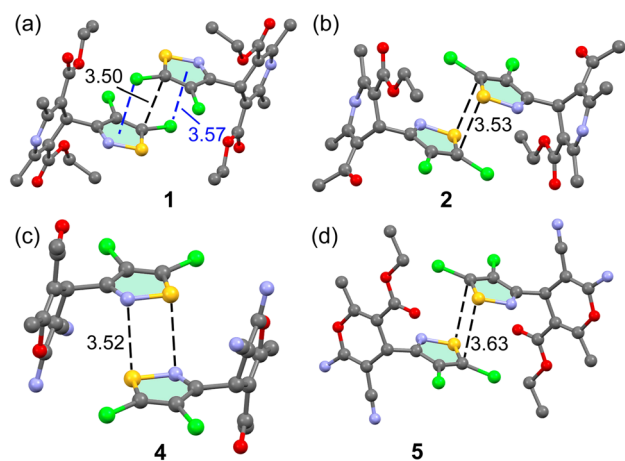


Fig. 6 π -Stacked assemblies observed in the solid state X-ray structures of compounds 1 (a), 2 (b), 4 (c) and 5 (d). Distances in Å.

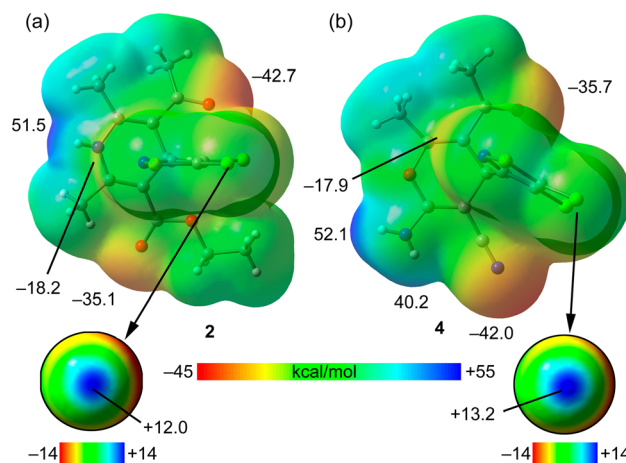


Fig. 7 MEP surfaces of compounds 2 (a) and 4 (b). The values at selected points are given in kcal mol⁻¹. At the bottom, a detail of the MEP surface around the Cl-atom in position 5 is given for both compounds using a reduced scale (± 14 kcal mol⁻¹).



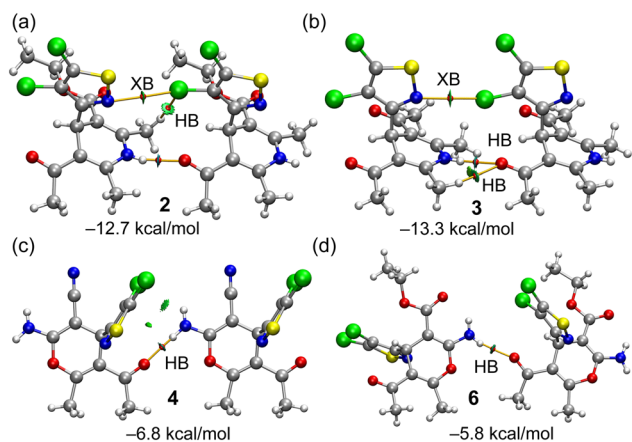


Fig. 8 QTAIM (bond CPs in red and bond paths as orange lines) and RDG isosurfaces (see computational methods for NCIPLOT settings) of compounds 2 (a), 3 (b), 4 (c) and 6 (d). Only intermolecular interactions are presented. The dimerization energies are also given.

energetically and characterized using the NCIPLOT and QTAIM methods. The results are gathered in Fig. 8, evidencing that in all dimers, the $\text{NH}\cdots\text{O}$ interaction is characterized by a bond critical point (CP, small red sphere) and bond path (orange line) and a blue reduced density gradient (RDG) isosurface, thus evidencing the strong nature of these contacts. In the dimers of compounds 2 and 3, the QTAIM analysis evidences the existence of a weaker halogen bond (XB) contact established between one of the Cl-atoms of the 4,5-dichloroisothiazole ring of one monomer and the N-atom of the isothiazole ring of the other monomer, also characterized by the corresponding bond CPs, bond paths and blue RDG isosurfaces. Moreover, additional $\text{CH}\cdots\text{Cl}$ and $\text{CH}\cdots\text{O}$ contacts are revealed in compounds 2 and 3, respectively, which are weaker according to the green colour of the RDG isosurfaces.

In the dimers of compounds 4 and 6, only one bond CP and bond path are observed corresponding to the strong $\text{NH}\cdots\text{O}$ contact. As expected, taking into consideration the QTAIM/NCIPLOT analysis, the dimerization energies are larger in the dimers 2 and 3 than in 4 and 6, due to the presence of the additional $\text{CH}\cdots\text{Cl}$, O contacts, XBs and shorter $\text{NH}\cdots\text{O}$ distances ($-12.7 \text{ kcal mol}^{-1}$ and $-13.3 \text{ kcal mol}^{-1}$, respectively). The dimerization energies of 4 ($-6.8 \text{ kcal mol}^{-1}$) and 6 ($-5.8 \text{ kcal mol}^{-1}$) are moderately strong, and basically correspond to the H-bond strength. The contribution of the XBs has been estimated by using the potential energy density (V_r) value at the bond CP that characterizes the XB. This QTAIM parameter can be used as an energy predictor *via* the equation proposed by Bartashevich & Tsirelson (*i.e.*, $E = 0.49 \times V_r$).⁴⁶ The $\text{Cl}\cdots\text{N}$ XB energies are very small in compounds 2 and 3 (-0.6 and $-0.8 \text{ kcal mol}^{-1}$, respectively). Such small energies disclose that these dimers are dominated by the $\text{NH}\cdots\text{O}$ HBs in line with the dark blue colour of the RDG isosurfaces. Finally, we have also estimated the $\text{Cl}\cdots\text{O}$ XB energy of the XB/HB dimer of compound 6 presented at the bottom of Fig. 4 (middle) and commented above. The XB

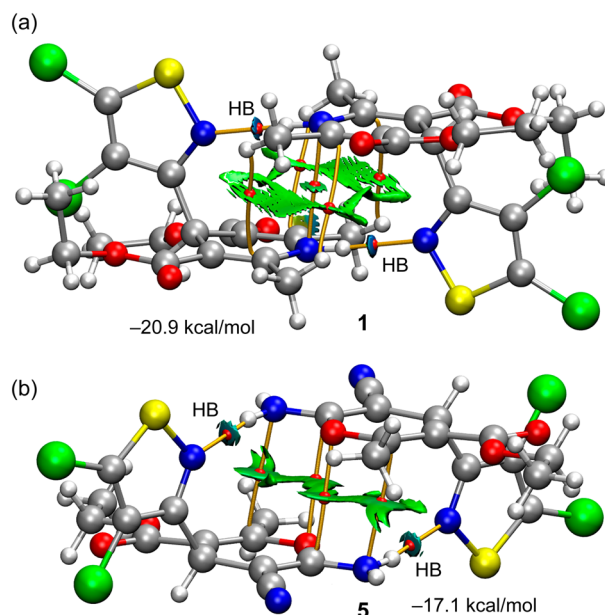


Fig. 9 QTAIM (bond CPs in red and bond paths as orange lines) and RDG isosurfaces (see computational methods for NCIPLOT settings) of compounds 1 (a) and 5 (b). Only intermolecular interactions are presented. The dimerization energies are also given.

energy is $-1.9 \text{ kcal mol}^{-1}$, thus evidencing that the $\text{Cl}\cdots\text{O}$ XB in compound 6 is stronger than the $\text{Cl}\cdots\text{N}$ XBs in compounds 2 and 3.

The QTAIM/NCIPLOT analyses of the H-bonded self-assembled dimers of compounds 1 and 5 are shown in Fig. 9. In both assemblies, the two symmetrically equivalent $\text{NH}\cdots\text{N}$ H-bonds are characterized by bond CPs and bond paths interconnecting the H and N-atoms. An intricate combination of bond CPs and bond paths emerges upon formation of the two strong $\text{NH}\cdots\text{N}$ H-bonds due to the close proximity of the 1,4-dihydropyridinyl rings in 1 and the 4H-pyranil rings in 5. In particular, five bond CPs and bond paths interconnect the 1,4-dihydropyridinyl rings (including the substituents) in compound 1 and four bond CPs and bond paths interconnect the 4H-pyranil rings in compound 5. Moreover, extended green RDG isosurfaces are located between the rings, as typical in van der Waals interactions. The dimerization energies are large and negative ($-20.9 \text{ kcal mol}^{-1}$ and $-17.1 \text{ kcal mol}^{-1}$ in 1 and 5 respectively) due to the contribution of both $\text{NH}\cdots\text{N}$ H-bonds and the van der Waals interactions.

The energetic features of the centrosymmetric $R_2^{2(12)}$ dimers have been also studied. Fig. 10 presents the combined QTAIM/NCIPLOT analyses of the dimers of compounds 4 and 5, where the $\text{NH}\cdots\text{NC}$ H-bonds are characterized by the corresponding bond CPs, bond paths and blue RDG isosurfaces. It is interesting to highlight the existence of a bond CP interconnecting the CN groups. Moreover, an extended green RDG isosurface is located between the CN groups, thus disclosing the presence of antiparallel $\text{CN}\cdots\text{CN}$ interactions. The dimerization energies are $-10.4 \text{ kcal mol}^{-1}$ and $-11.9 \text{ kcal mol}^{-1}$ for 4 and 5, respectively, thus



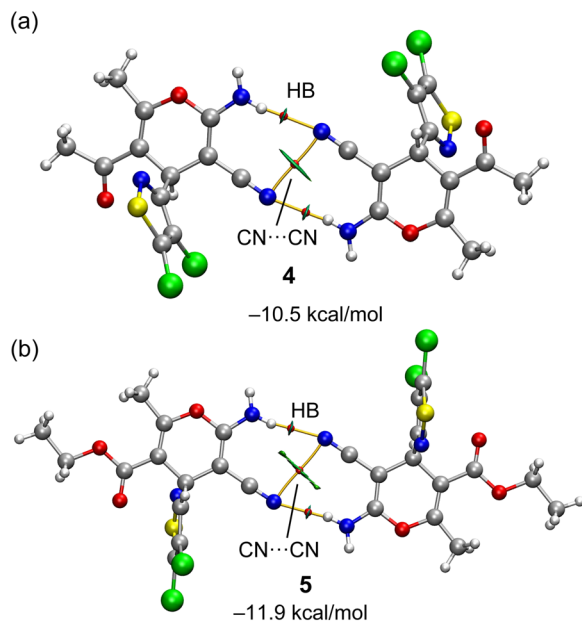


Fig. 10 QTAIM (bond CPs in red and bond paths as orange lines) and RDG isosurfaces (see computational methods for NCIPLOT settings) of the $R_2(12)$ synthons of compounds 4 (a) and 5 (b). Only intermolecular interactions are presented. The dimerization energies are also given.

confirming the relevance of these $R_2(12)$ synthons in the solid state.

Finally, the different π -stacking modes between the 4,5-dichloroisothiazole rings observed in several compounds and described in Fig. 6 have been also studied energetically and analysed *via* QTAIM/NCIPLOT. The results are summarized in Fig. 11, evidencing that all π -stacked dimers exhibit similar interaction energies, ranging from -9.6 kcal mol $^{-1}$ in 2 to -10.5 kcal mol $^{-1}$ in 5, thus explaining the observation of different

stacking modes in the solid state. The $C \cdots C$ π -stacking is characterized only by one bond CP and bond path (in 1).

The double $C \cdots S$ stacking is characterized by two bond CPs and bond paths (compounds 2 and 5) and the double $S \cdots N$ stacking is characterized by three bond CPs and bond paths (two interconnecting the N and S-atoms and one extra CP connecting both S-atoms (see Fig. 11c)). In compounds 1, 2 and 5, the interaction is further complemented by $Cl \cdots \pi$ interactions, as evidenced by both the QTAIM and NCIPLOT methods. In all systems, additional van der Waals interactions exist between the 4,5-dichloroisothiazole rings and either the 1,4-dihydropyridinyl (in 1 and 2) or 4*H*-pyranil rings (4 and 5), characterized by several bond CPs, bond paths and extended green RDG isosurfaces. It is worthy to emphasize that the interaction energies of the π -stacked assemblies are similar to those of the $R_2(12)$ synthons shown in Fig. 11, evidencing that these π -stacked assemblies are energetically very relevant and have an important role in the crystal packing.

Concluding remarks

A convenient synthetic protocol to obtain 3-substituted 4,5-dichloroisothiazole derivatives is reported herein. Moreover, the X-ray structures of six new derivatives are described, showing some recurrent motifs like $R_2(12)$ synthons and HB/XB 1D supramolecular polymers. The interactions have been characterized using a combination of QTAIM and NCIPLOT computational tools and rationalized using MEP surface analysis. The latter is useful to reveal the existence of σ -holes opposite to the C–Cl bonds of the 4,5-dichloroisothiazole ring and rationalize the formation of $Cl \cdots N$ and $Cl \cdots O$ interactions. It is also important to demonstrate the strong H-bond acceptor ability of the sp-hybridized N-atom in compounds 4 and 5, thus explaining the formation of the $R_2(12)$ synthons. Finally, the energetic analysis discloses that both H-bonds and π -stacking interactions are equally important, dictating the X-ray packing of the compounds.

Author contributions

I. A. K., A. G. P. and V. I. P.: synthesis and spectral analysis; V. V. B. and R. M. G.: investigation and methodology; M. S. G. and A. P. N.: X-ray experiments; A. F. and F. I. Z.: international collaboration, conceptualization, supervision, validation, project administration, writing – original draft, and writing – review & editing.

Conflicts of interest

There are no conflicts to declare.

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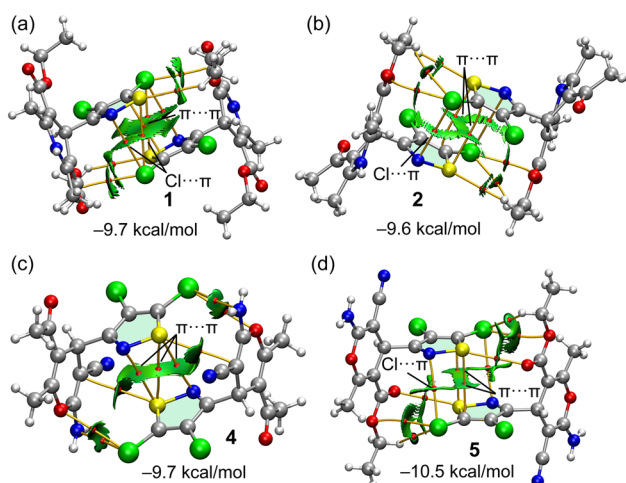


Fig. 11 QTAIM (bond CPs in red and bond paths as orange lines) and RDG isosurfaces (see computational methods for NCIPLOT settings) of the π -stacked dimers of compounds 1 (a), 2 (b), 4 (c) and 5 (d). Only intermolecular interactions are presented. The dimerization energies are also given.



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