

Cite this: *CrystEngComm*, 2012, 14, 7848–7850

www.rsc.org/crystengcomm

COMMUNICATION

Investigating the influence of the sulfur oxidation state on solid state conformation†

Carla A. Daly,^a Kevin S. Eccles,^a Lorraine M. Bateman,^a Noel M. O'Boyle,^a Simon E. Lawrence^a and Anita R. Maguire^{*b}

Received 14th August 2012, Accepted 12th September 2012

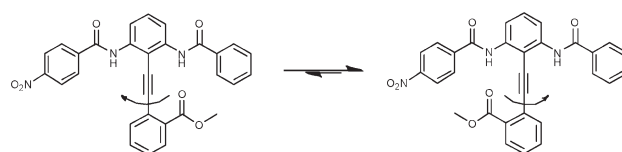
DOI: 10.1039/c2ce26298c

Design, synthesis and structural characterization of a series of diphenylacetylene derivatives bearing organosulfur, amide and amine moieties has been achieved in which the molecular conformation is controlled through variation of the hydrogen bond properties on alteration of the oxidation level of sulfur.

The ability to understand and rationally predict the conformation adopted by solid state structures has been actively pursued for many years.¹ Crystal engineering specifically focuses on intermolecular interactions with the aim to identify supramolecular synthons for the design of materials with specific properties *e.g.* optical, magnetic, electronic.^{2,3} Control of the solid state physical properties of organic and inorganic materials *e.g.* solubility, bioavailability, dissolution rate, hygroscopicity also demands an understanding of the nature of the interactions in the solid state at a fundamental level.^{4–6}

Previous research in our group focused on organosulfur functional groups, specifically sulfides, sulfoxides and sulfones, with the aim to develop an understanding of how the molecular structure of the compounds impacts upon the solid state crystalline structure and, in particular, to probe the relative importance of different inter/intramolecular non-covalent interactions. In particular, our research highlighted the effective use of sulfoxides in supramolecular synthons, due to their nature as strong hydrogen bond acceptors,^{7,8} including with amides as N–H donors.⁹

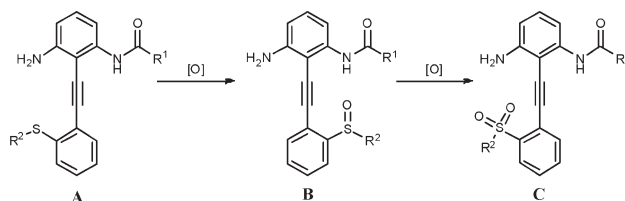
To further expand on this work we aimed to incorporate sulfur and amide functionalities within a single molecule and study the effects of varying the oxidation level of sulfur on the hydrogen bond interactions in the solid state. The diphenylacetylene unit involving ester and amide functionalities recently explored by Hamilton provided us with a suitable scaffold on which to construct this system (Scheme 1).^{10,11} Their success in controlling the conformation of the molecule by varying the acidity of the



Scheme 1 Controlling the conformation of benzamidodiphenylacetylenes by changing the acidity of the hydrogen bond donors.¹¹

amide encouraged us to expand this system by incorporating sulfur functionalities (Scheme 2).

The basic concept involves creating competition between hydrogen bond acceptors for the strongest hydrogen bond donor by altering the oxidation level of the sulfide and exploiting the difference in acidity between amides and amines.¹² At the sulfide level, interaction between the sulfur and amide or amine is not expected based on results from earlier fundamental studies¹² and the dominant solid state interaction predicted is the N–H⋯O=C intermolecular interaction. As a result we would expect the sulfide to lie on the opposite side to the amide as illustrated (A), thereby enabling the intermolecular N–H⋯O=C interaction. On oxidation to the sulfoxide, the strong intramolecular N–H⋯O=S interaction should compete effectively with the intermolecular N–H⋯O=C interaction as sulfoxides are potent hydrogen bond acceptors¹³ and amides are stronger hydrogen bond donors than amines.¹² In this case we expect the sulfoxide to lie on the same side as the amide (B), following Hamilton's model. On further oxidation to the sulfone, which is a weaker hydrogen bond acceptor than the sulfoxide, we anticipated at the outset that the strong N–H⋯O=C intermolecular interaction would once again dominate, resulting in the sulfone lying on the opposite side to the amide (C).

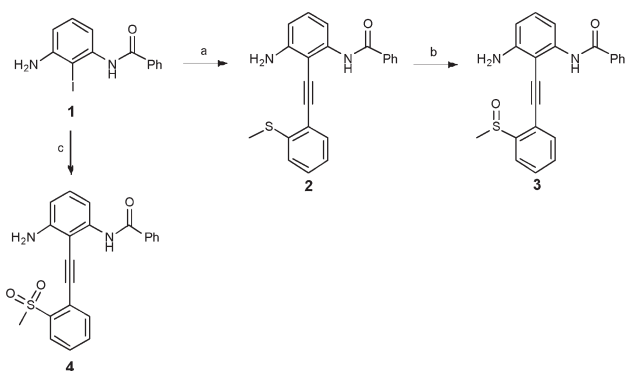


Scheme 2 Predicting the conformation of A, B and C by applying the rationale of differential hydrogen bonding ability of sulfur functionalities.

^aDepartment of Chemistry, Analytical and Biological Chemistry Research Facility, University College Cork, Cork, Ireland

^bDepartment of Chemistry and School of Pharmacy, Analytical and Biological Chemistry Research Facility, University College Cork, Cork, Ireland. E-mail: a.maguire@ucc.ie

† Electronic supplementary information (ESI) available: CCDC 891708–891710. Synthetic procedures for 1–4; computational studies on 3. For ESI and crystallographic data in CIF or other electronic format see 10.1039/c2ce26298c



Scheme 3 The synthesis of **2**, **3** and **4**. Reagents and conditions: a) 1-ethynyl-2-methylthiobenzene, $\text{PdCl}_2(\text{PPh}_3)_2$, CuI , DMF, NEt_3 , b) NaIO_4 , $\text{MeOH}/\text{H}_2\text{O}$. c) 2-methylsulfonyl-1-ethynylbenzene, $\text{PdCl}_2(\text{PPh}_3)_2$, CuI , DMF, NEt_3 .

To explore this concept, *N*-(2-iodo-3-aminophenyl)benzamide **1**, was synthesised following Hamilton's procedure.¹⁰ Then the alkynes, bearing sulfide and sulfone functional groups, were attached *via* Sonogashira coupling to form **2** and **4** (Scheme 3). The sulfoxide, **3**, was readily obtained by oxidation of **2**. These systems with the substituents in the *ortho* position were designed to allow the exploration of intramolecular hydrogen bonding between the key functional groups. The successful Sonogashira coupling to provide the sulfide **2** is particularly interesting in the context of Larock's report involving a related system where the coupling product could not be obtained.¹⁴

Single crystal X-ray diffraction of compounds **2**, **3** and **4**, each recrystallized from the same solvent, CH_2Cl_2 , demonstrated the predicted conformational change as a result of altering the oxidation level of sulfur (Fig. 1).[‡] As expected the sulfide lies on the opposite side to the amide, then switches after oxidation to the sulfoxide and switches back again when the sulfone is formed. For compound **2**, the strong intermolecular $\text{N-H}\cdots\text{O}=\text{C}$ dominates the crystal packing, and the $\text{C}=\text{O}$ of the amide is involved in bifurcated

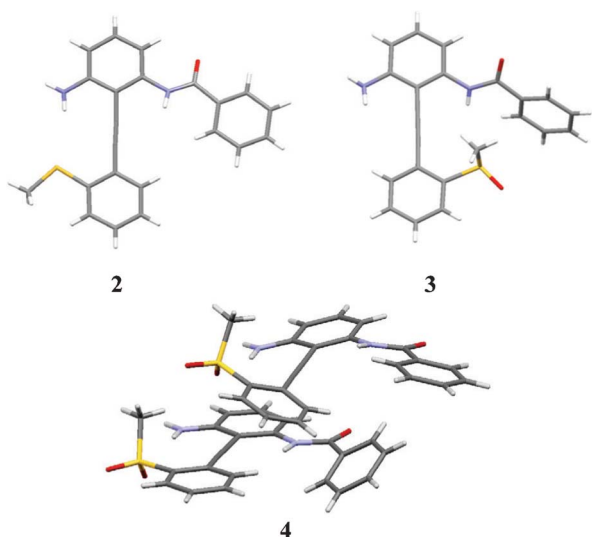


Fig. 1 Single crystal X-ray structures obtained for compounds **2**, **3** and **4**.

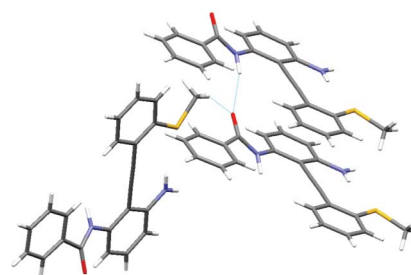


Fig. 2 Hydrogen bond interactions in compound **2**.

hydrogen bonding to both a neighbouring N-H of an amide and C-H of a methyl group (Fig. 2).

Interestingly, although the conformation switches in the sulfoxide, **3**, the key non-covalent interactions observed were not as anticipated (Fig. 3). Instead of an intramolecular $\text{N-H}\cdots\text{O}=\text{S}$ bond occurring between the amide and sulfoxide, an intermolecular $\text{N-H}\cdots\text{O}=\text{S}$ is formed between the sulfoxide and a neighbouring amine. The oxygen from the sulfoxide points away from the amide, with the result that intramolecular hydrogen bonding does not occur. The strong $\text{N-H}\cdots\text{O}=\text{C}$ interaction prevails in the crystal structure and oxidation to the sulfoxide has not disrupted this interaction. Comparison of the structural features of Hamilton's amide-ester system with our amide-sulfoxide system is very interesting. Although the sulfoxide is expected to be a stronger hydrogen bond acceptor than the ester, the planar intramolecular hydrogen bond which we anticipated to form did not occur in practice. Examination of the amide to sulfoxide $\text{N-H}\cdots\text{O}=\text{S}$ intramolecular distance available in **3** (~ 2.05 Å), together with analysis of the Cambridge Structural Database¹⁵ and comparison with the amide-ester $\text{N-H}\cdots\text{O}=\text{C}$ hydrogen bond distance (2.23 Å),¹⁰ suggests that intramolecular hydrogen bonding, while not observed, is feasible in our system.

Overall the solid state structure of the sulfoxide adopts a conformation that enables two structure-defining intermolecular interactions: the amine $\text{N-H}\cdots\text{O}=\text{S}$ and the amide $\text{N-H}\cdots\text{O}=\text{C}$. The key feature that arose was the unanticipated orientation of the sulfoxide out of the plane. While computational studies (see ESI[†]) demonstrate that an intramolecular hydrogen bond is possible, it would require the axial phenyl rings to twist out of planarity, therefore leading to a decrease of extended conjugation and stabilisation. As a result, the observed conformation, which has the

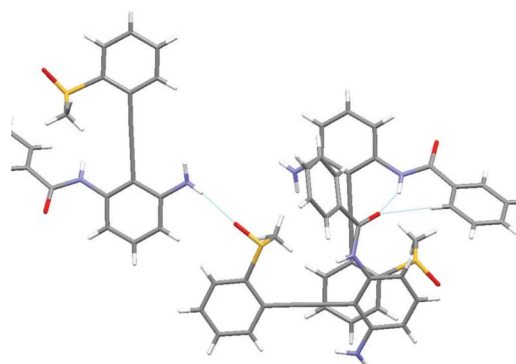


Fig. 3 Hydrogen bond interactions in compound **3**.

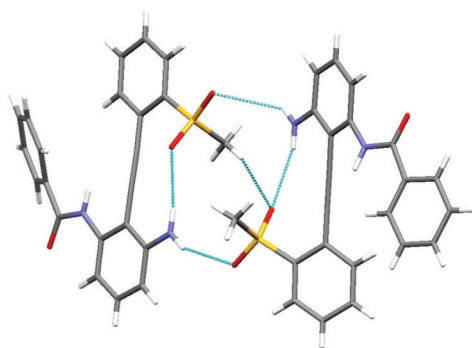


Fig. 4 Hydrogen bond interactions in compound 4.

sulfoxide oxygen pointing away from the amide, is predicted to be slightly lower in energy.

The sulfone, **4**, crystallises with $Z' = 2$, with both molecules adopting the same conformation as seen in the sulfide, *i.e.* the sulfone lies on the opposite side to the amide (Fig. 4). The key interactions involving the two crystallographically independent molecules are intra- and intermolecular $\text{N-H}\cdots\text{O}=\text{S}$ hydrogen-bonds. The combination gives rise to a visually appealing R_4^4 (12) motif at the binary level. Also present within this motif is a $\text{C-H}\cdots\text{O}=\text{S}$ intermolecular interaction between one of the sulfone oxygen atoms and a methyl group. Significantly, the strong intermolecular $\text{N-H}\cdots\text{O}=\text{C}$ between the amides, which was the key structure-defining feature in the sulfide and sulfoxide structures, was disrupted on oxidation to the sulfone, therefore altering very substantially the crystal packing of the molecule.

To investigate the solution properties of compounds **2** and **3** NMR studies were undertaken. Results from NOESY 2D NMR experiments did not result in any substantial correlation between spectroscopic features and the solid state interactions.

In conclusion, the predicted change in molecular conformation of the sulfide **2** to the sulfoxide **3** and sulfone **4** was observed as a direct result of altering the oxidation state of sulfur and therefore impacting on the key hydrogen bonding features in the solid state. This significant result, particularly the observed rotation of the diphenylacetylene unit after oxidation, may lead to future applications in a molecular switching mechanism.

Acknowledgements

This publication has emanated from research conducted with the financial support of Science Foundation Ireland, under Grant Numbers 08/RFP/MTR1664 and 05/PICA/B802/EC07,

and a Health Research Board Career Development Fellowship, PD/2009/13 (N.O'B.). We thank the SFI/HEA Irish Centre for High-End Computing (ICHEC) for the provision of computational facilities.

References

† Single crystal X-ray diffraction data were collected on either a Bruker SMART X2S diffractometer (**2**) or a Bruker APEX II DUO diffractometer (**3** and **4**). All calculations and refinement were made using the APEX software,^{16,17} and diagrams prepared using Mercury.¹⁸ Crystal data for **2**: $\text{C}_{22}\text{H}_{18}\text{N}_2\text{O}_2\text{S}$, $M = 358.44$, $a = 18.140(3)$ Å, $b = 5.0400(9)$ Å, $c = 19.369(3)$ Å, $V = 1770.8(5)$ Å³, $T = 300(2)$ K, orthorhombic, space group $Pna2_1$, $Z = 4$, 13 743 reflections measured, 3012 independent reflections ($R_{\text{int}} = 0.0631$). The final R_1 value was 0.0548 [$I > 2\sigma(I)$] and the final $wR(F^2)$ value was 0.1638 (all data). Crystal data for **3**: $\text{C}_{22}\text{H}_{18}\text{N}_2\text{O}_2\text{S}$, $M = 374.44$, $a = 8.8488(15)$ Å, $b = 21.149(4)$ Å, $c = 10.0801(17)$ Å, $\beta = 98.541(4)^\circ$, $V = 1865.5(5)$ Å³, $T = 296(2)$ K, monoclinic, space group $P2_1/c$, $Z = 4$, 19 006 reflections measured, 3277 independent reflections ($R_{\text{int}} = 0.0763$). The final R_1 value was 0.057 [$I > 2\sigma(I)$] and the final $wR(F^2)$ value was 0.175 (all data). Crystal data for **4**: $\text{C}_{22}\text{H}_{18}\text{N}_2\text{O}_3\text{S}$, $M = 390.44$, $a = 10.511(2)$ Å, $b = 34.171(8)$ Å, $c = 11.778(3)$ Å, $\beta = 113.517(5)^\circ$, $V = 3879.0(15)$ Å³, $T = 296(2)$ K, monoclinic, space group $P2_1/n$, $Z = 8$, 21 664 reflections measured, 7387 independent reflections ($R_{\text{int}} = 0.0505$). The final R_1 value was 0.0504 [$I > 2\sigma(I)$] and the final $wR(F^2)$ value was 0.1279 (all data).

- G. R. Desiraju, *J. Chem. Sci.*, 2010, **122**, 667–675.
- G. R. Desiraju, *Angew. Chem., Int. Ed. Engl.*, 1995, **34**, 2311–2327.
- G. R. Desiraju, *Nature*, 2001, **412**, 397–400.
- L. F. Huang and W. Q. Tong, *Adv. Drug Delivery Rev.*, 2004, **56**, 321–334.
- D. A. Snider, W. Addicks and W. Owens, *Adv. Drug Delivery Rev.*, 2004, **56**, 391–395.
- B. Rodriguez-Spong, C. P. Price, A. Jayasankar, A. J. Matzger and N. Rodriguez-Hornedo, *Adv. Drug Delivery Rev.*, 2004, **56**, 241–274.
- K. N. Lehane, E. J. A. Moynihan, N. Brondel, S. E. Lawrence and A. R. Maguire, *CrystEngComm*, 2007, **9**, 1041–1050.
- N. Brondel, E. J. A. Moynihan, K. N. Lehane, K. S. Eccles, C. J. Elcoate, S. J. Coles, S. E. Lawrence and A. R. Maguire, *CrystEngComm*, 2010, **12**, 2910–2927.
- K. S. Eccles, C. J. Elcoate, A. R. Maguire and S. E. Lawrence, *Cryst. Growth Des.*, 2011, **11**, 4433–4439.
- I. M. Jones and A. D. Hamilton, *Org. Lett.*, 2010, **12**, 3651–3653.
- I. M. Jones and A. D. Hamilton, *Angew. Chem., Int. Ed.*, 2011, **50**, 4597–4600.
- C. A. Hunter, *Angew. Chem., Int. Ed.*, 2004, **43**, 5310–5324.
- A. Nangia and G. R. Desiraju, *Chem. Commun.*, 1999, 605–606.
- S. Mehta, J. P. Waldo and R. C. Larock, *J. Org. Chem.*, 2009, **74**, 1141–1147.
- F. H. Allen, *Acta Crystallogr., Sect. B: Struct. Sci.*, 2002, **58**, 380–388.
- APEX2 v2009.3-0, Bruker AXS, Madison, WI, 2009.
- G. M. Sheldrick, *Acta Crystallogr., Sect. A: Found. Crystallogr.*, 2008, **64**, 112–122.
- C. F. Macrae, I. J. Bruno, J. A. Chrisholm, P. R. Edgington, P. McCabe, E. Pidcock, L. Rodriguez-Monge, R. Taylor, J. van de Streek and P. A. Wood, *J. Appl. Crystallogr.*, 2008, **41**, 466–470.