

An Experimental and Computational Investigation of the Elusive Anhydrous form of Oxyma-B

**Rafel Prohens,^{*a} Rafael Barbas,^a Beatriz G. de la Torre,^d Fernando Albericio^{*b,c,e}
and Antonio Frontera^{*f}**

^a Unitat de Polimorfisme i Calorimetria, Centres Científics i Tecnològics, Universitat de Barcelona, Baldri Reixac 10, 08028 Barcelona, Spain.

^b CIBER-BBN, Networking Centre on Bioengineering, Biomaterials and Nanomedicine, and Department of Organic Chemistry, University of Barcelona, Martí i Franqués 1-11, 08028 Barcelona, Spain.

^c Institute for Advanced Chemistry of Catalonia (IQAC-CSIC), 08034 Barcelona, Spain

^d KwaZulu-Natal Research Innovation and Sequencing Platform (KRISP), School of Laboratory Medicine and Medical Sciences, College of Health Sciences, University of KwaZulu-Natal, Durban, KwaZulu-Natal 4041, South Africa.

^e Peptide Science Laboratory, School of Chemistry and Physics, University of KwaZulu-Natal, Westville, KwaZulu-Natal 4000, South Africa

^f Departament de Química, Universitat de les Illes Balears, Crta. de Valldemossa km 7.5, 07122 Palma, Spain.

Electronic Supplementary Information

1. Characterization of the solids	2
2. List of 30 solvents used in the crystallization battery at 25 °C	10
3. Structure Determination from Direct Space Methods	11
4. CSD Survey of supramolecular synthon	14

1.- Characterization of the synthesis product of Oxyma-B

Figure S1: DSC of Oxyma-B (synthesis product) at a heating rate of 10 °C/min.

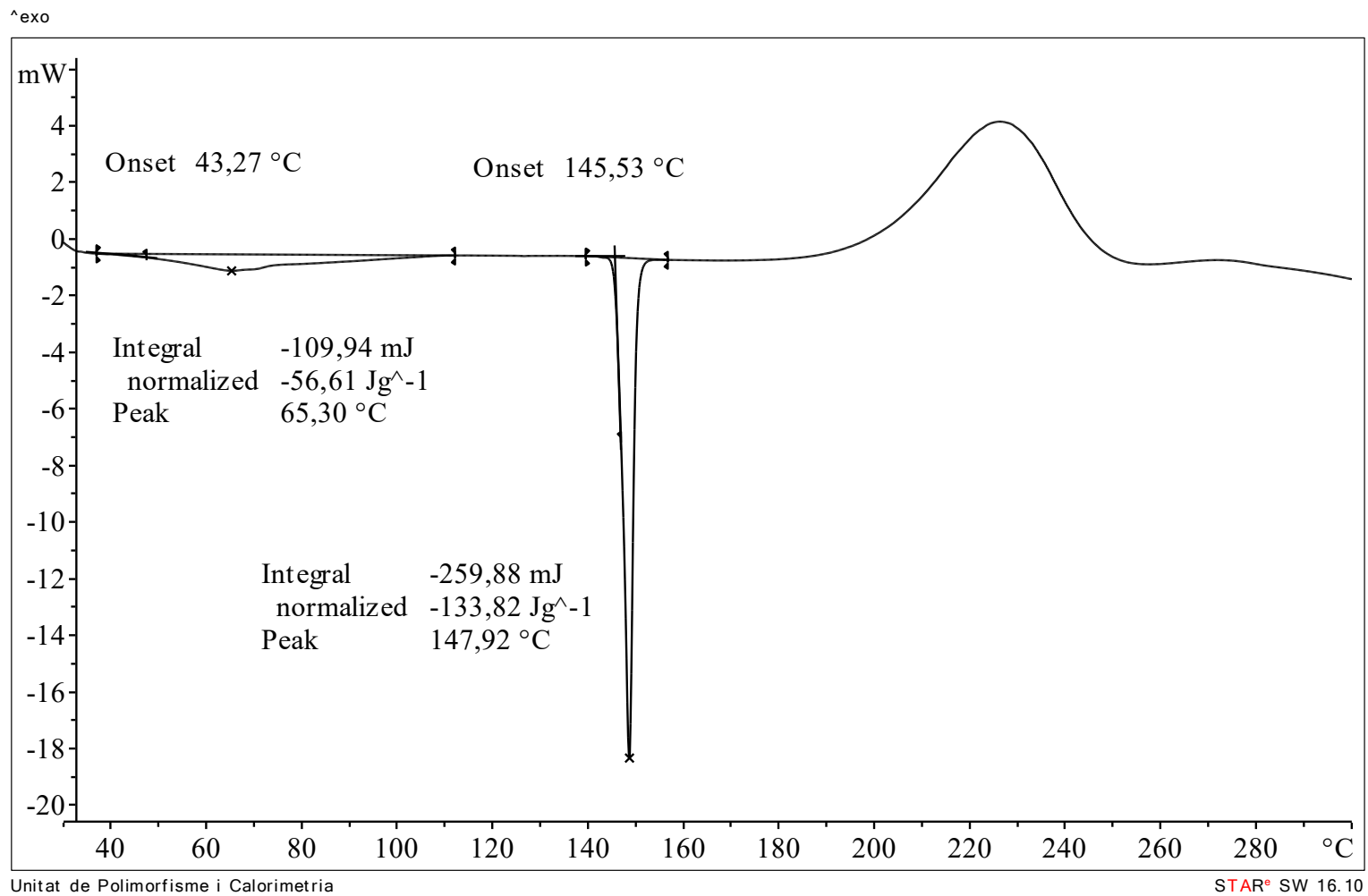


Figure S2: TGA of Oxyma-B (synthesis product).

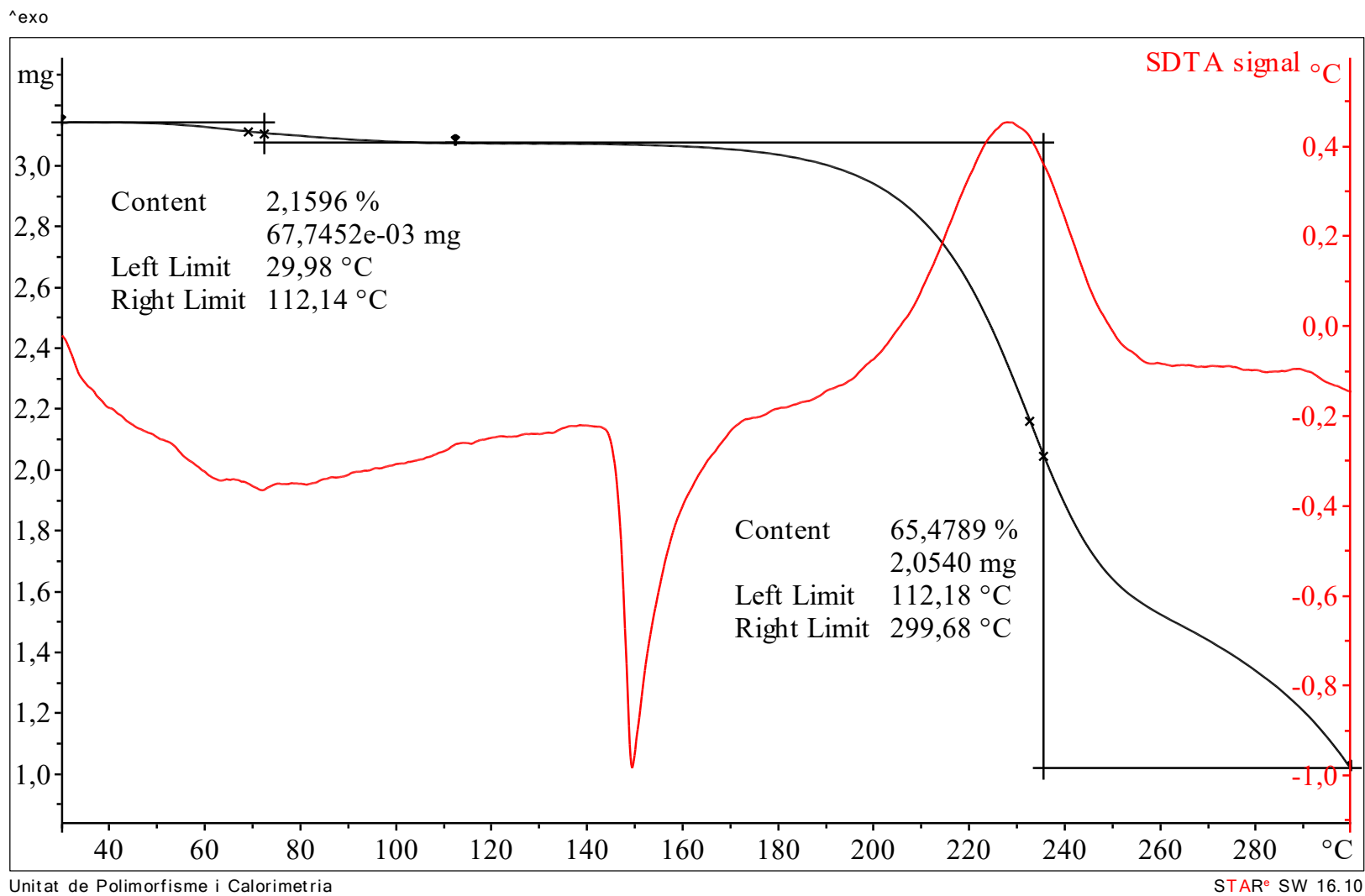


Figure S3: XRPD of Oxyma-B (synthesis product).

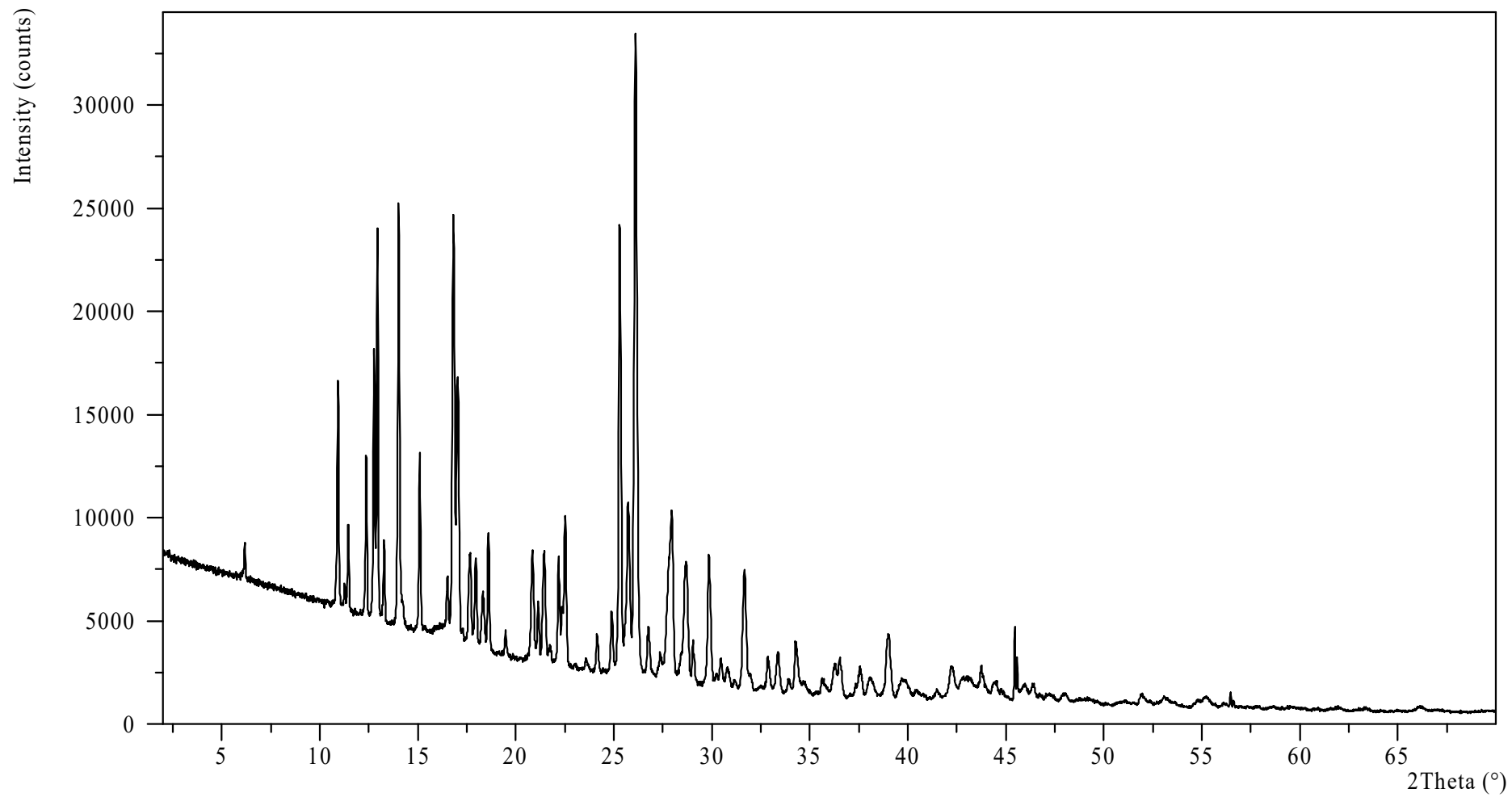


Figure S4: Comparative XRPD diffractograms of Oxyma-B (synthesis product) bulk powder (black) and monohydrate Oxyma-B (CCDC refcode WOWFUL) simulated from the cif file (green).

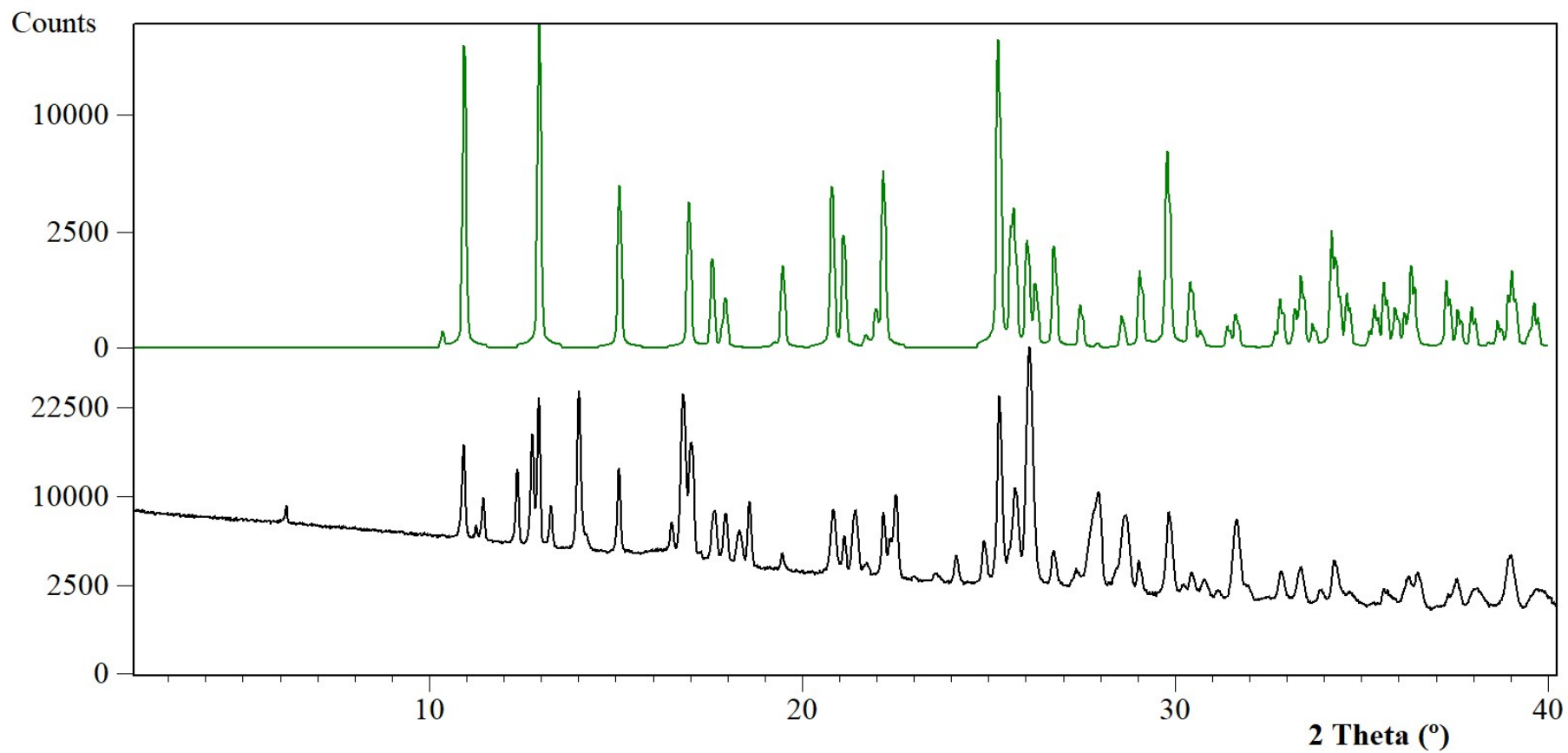


Figure S5: DSC of Oxyma-B (synthesis product) at a heating rates of 10 °C/min and 1 °C/min.

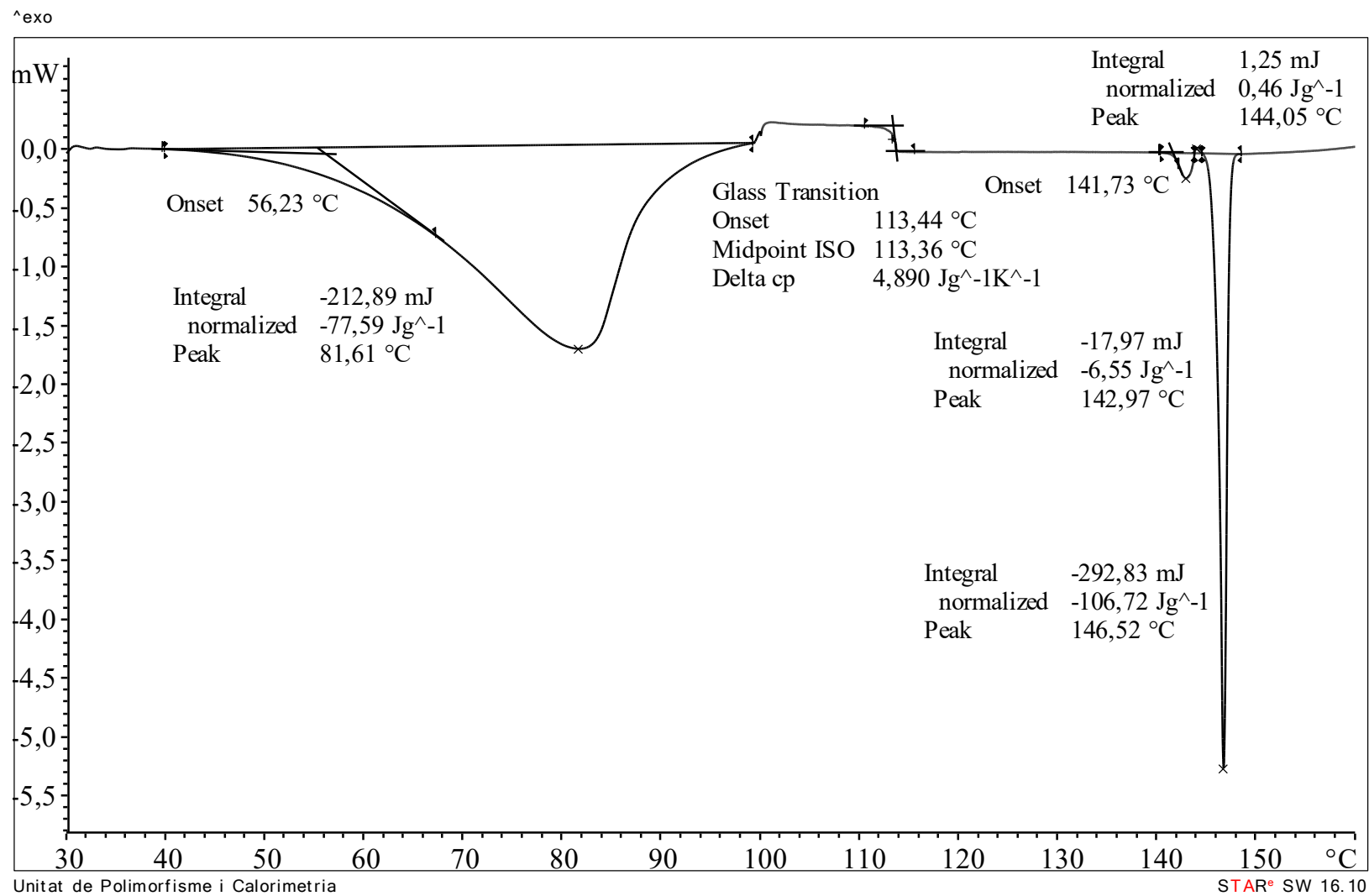


Figure S6: ADSC modulated heat flow curve of Oxyma-B (synthesis product) at a heating rate of 1 °C/min with a modulation period and an amplitude of 60 s and 1 °C, respectively.

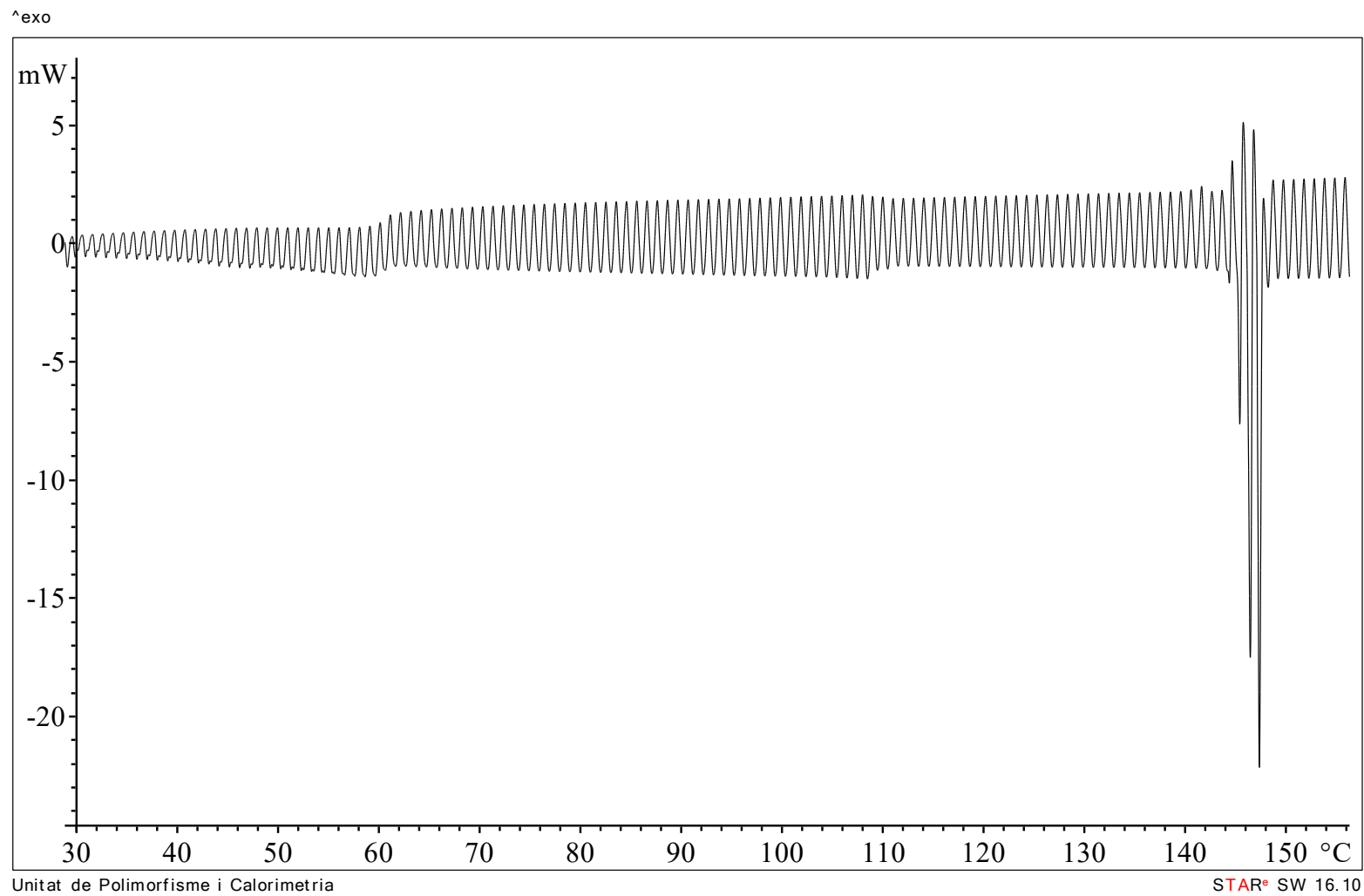
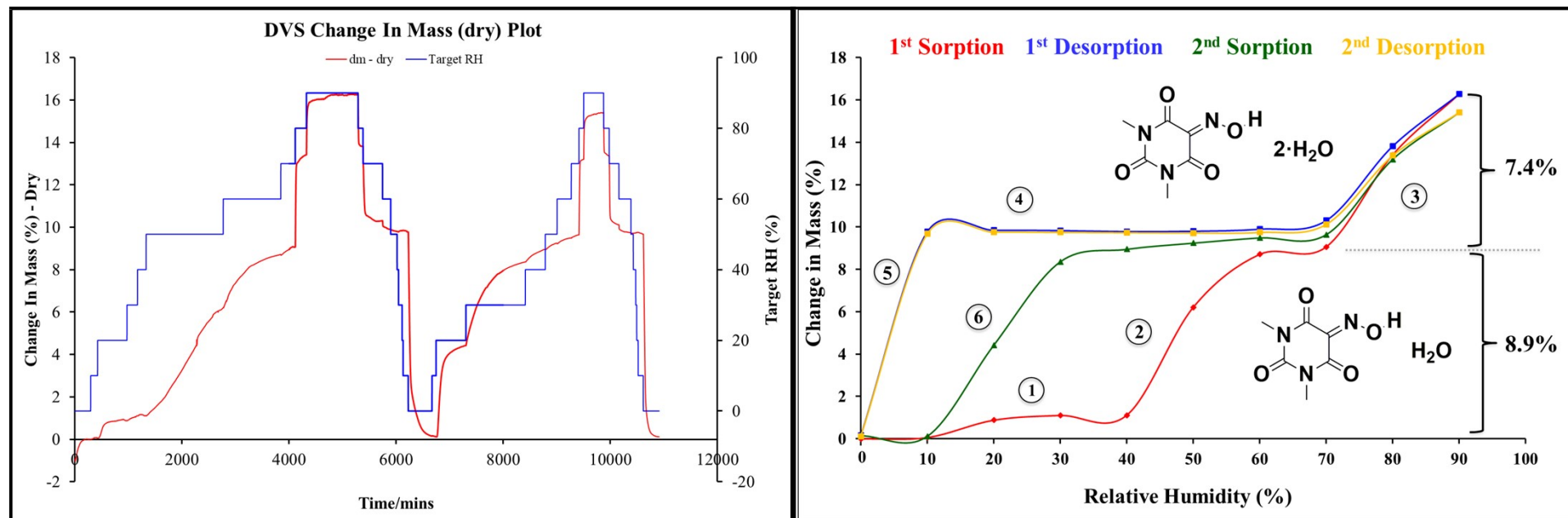


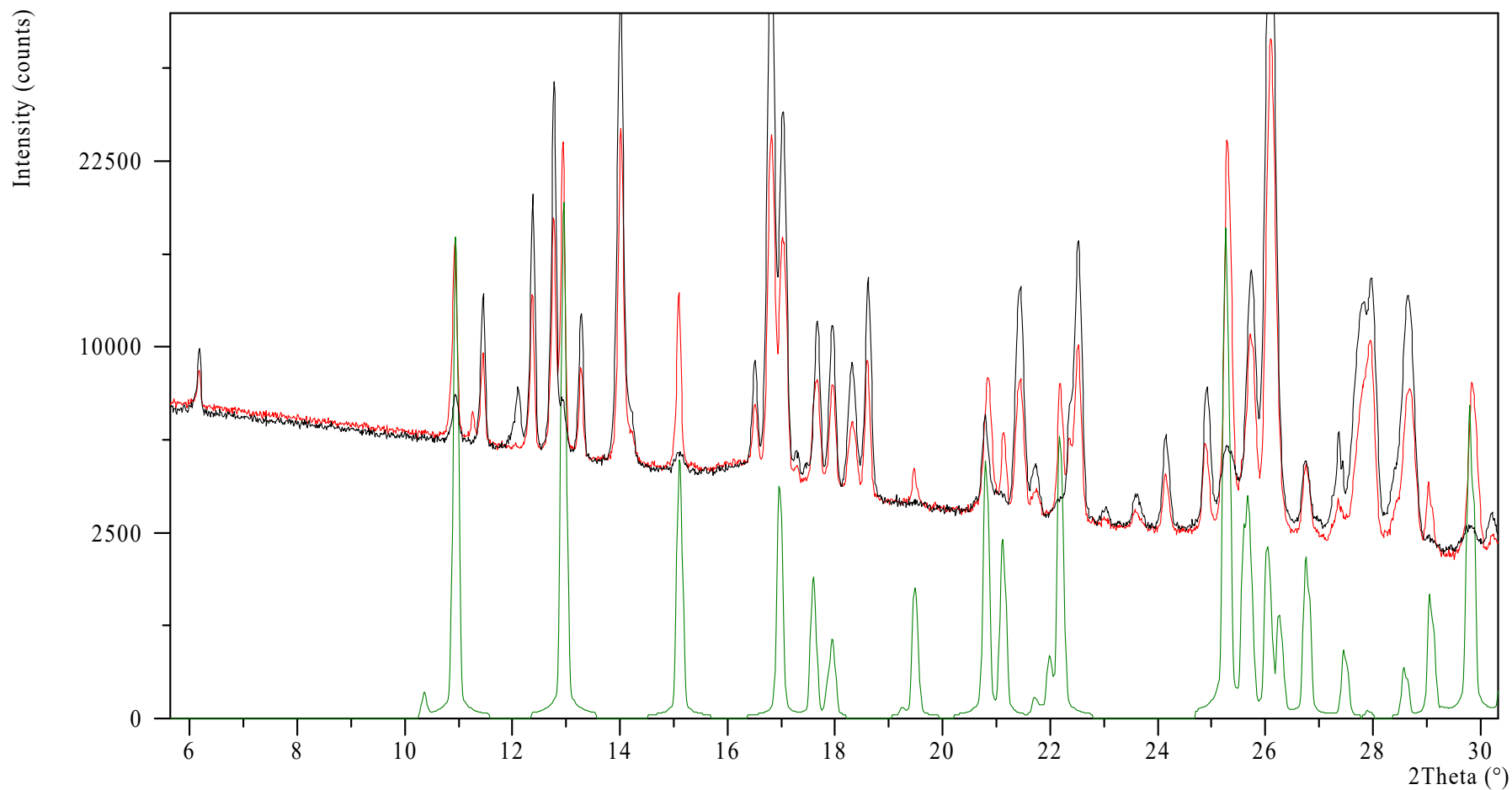
Figure S7: DVS sorption curves (a) and DVS isotherms (b) of Oxyma-B (synthesis product).



(a)

(b)

Figure S8: Comparative XRPD diffractograms of Oxyma-B (synthesis product) bulk powder (red), Oxyma-B (heated until 125 °C during 10 minutes) powder (black) and monohydrate Oxyma-B (CCDC refcode WOWFUL) simulated from the cif file (green).



2.- List of 30 solvents used in the crystallization battery at 25 °C

Methanol, ethanol, IPA, butanol, formic acid, ethylene glycol, dimethyl ethylene glycol, benzyl alcohol, ACN, MEK, acetone, MiBK, water, DMF, DMSO, AcOEt, THF, dioxane, dichloromethane, chloroform, acetic acid, NH₃ (32 %) in water, diethylamine, toluene, xylene, heptane, cyclohexane, pentane, diethyl ether and diisopropyl ether.

3.- Structure Determination from Direct Space Methods

Figure S9: PRF plot showing the Le Bail fit of the XRPD pattern of anhydrous Oxyma-B. The plot shows the experimental powder XRD profile (red marks), the calculated powder XRD profile (black solid line), the difference profile (blue lower line) and Bragg positions (green lines).

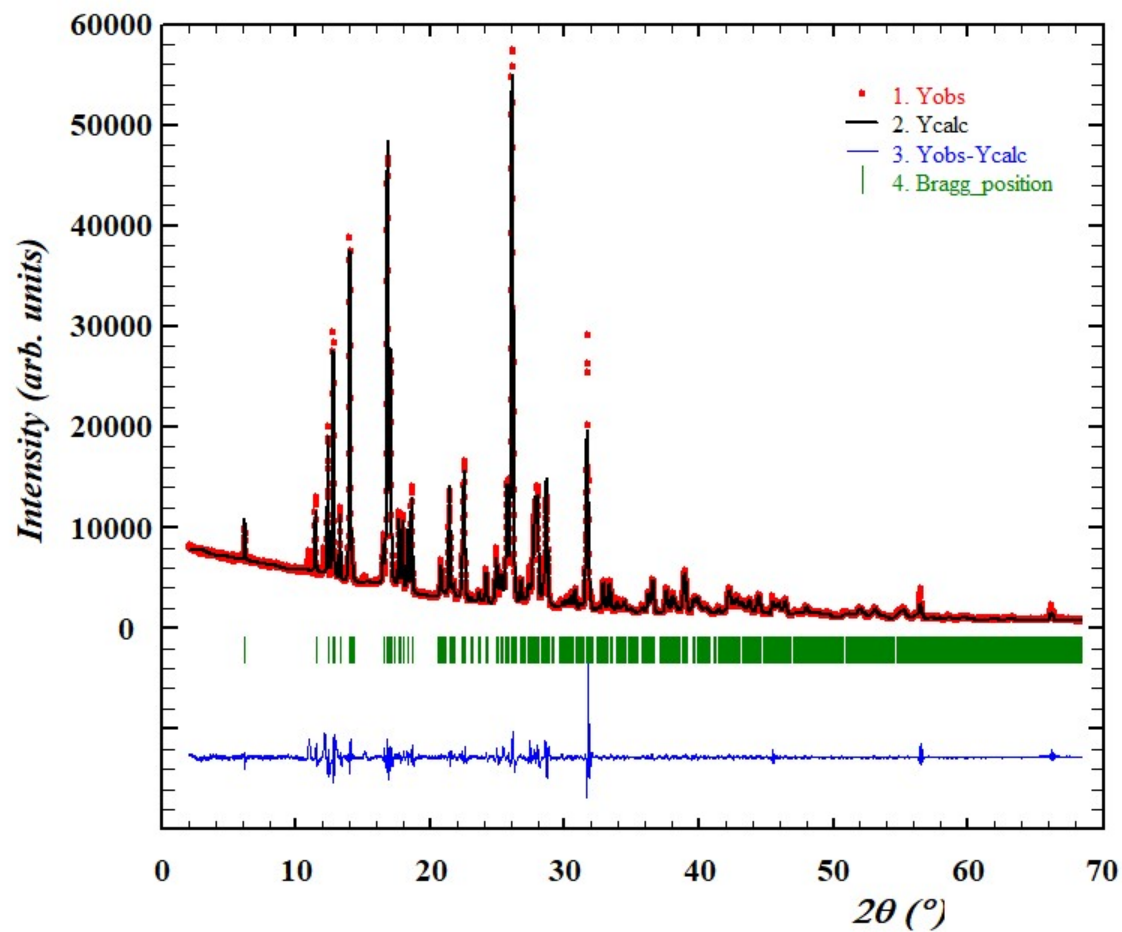
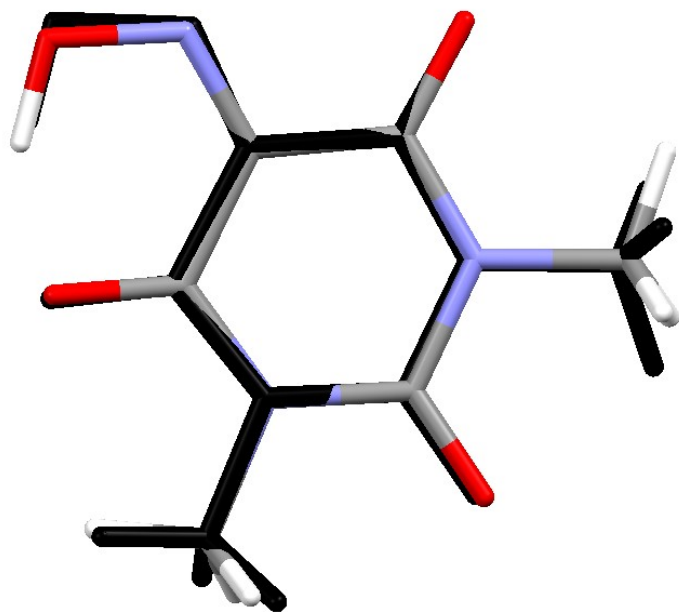
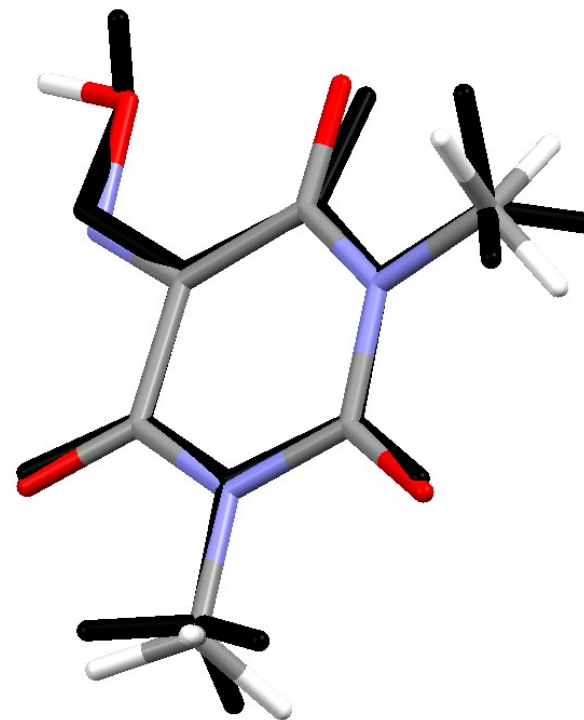


Figure S10: Overlay of the asymmetric unit molecules between the best FOX solution (black) and DFT-D with fixed cell parameters (colour by Element)

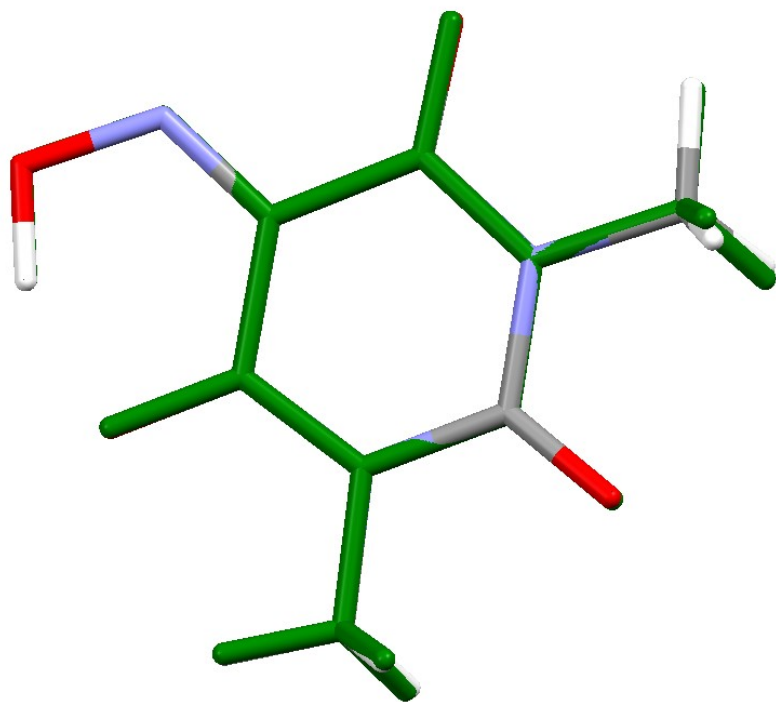


Molecule 1

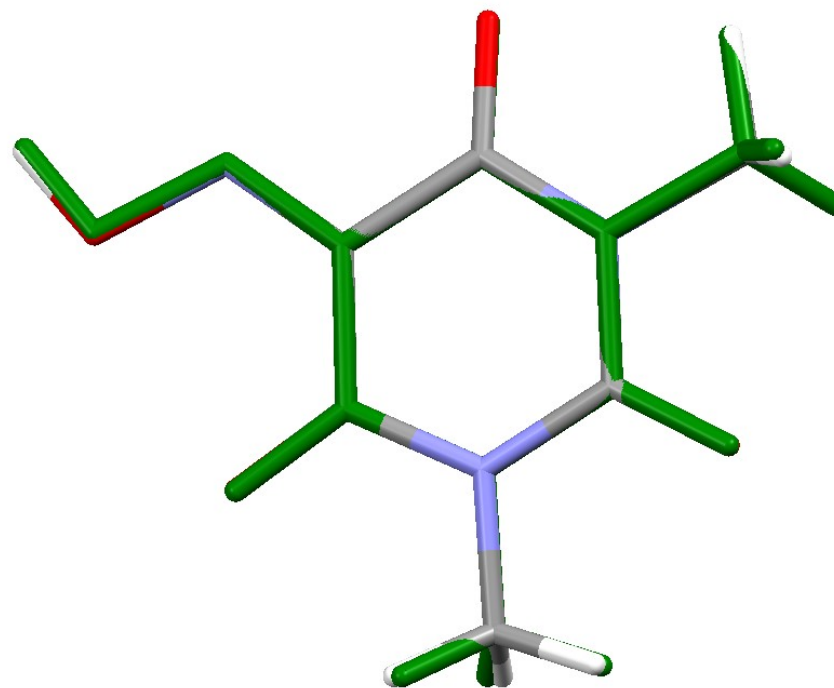


Molecule 2

Figure S11: Comparative of asymmetric unit molecules: final cif deposited (colour by Element) and after validation procedure (green) with DFT geometry optimization



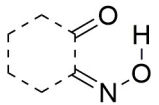
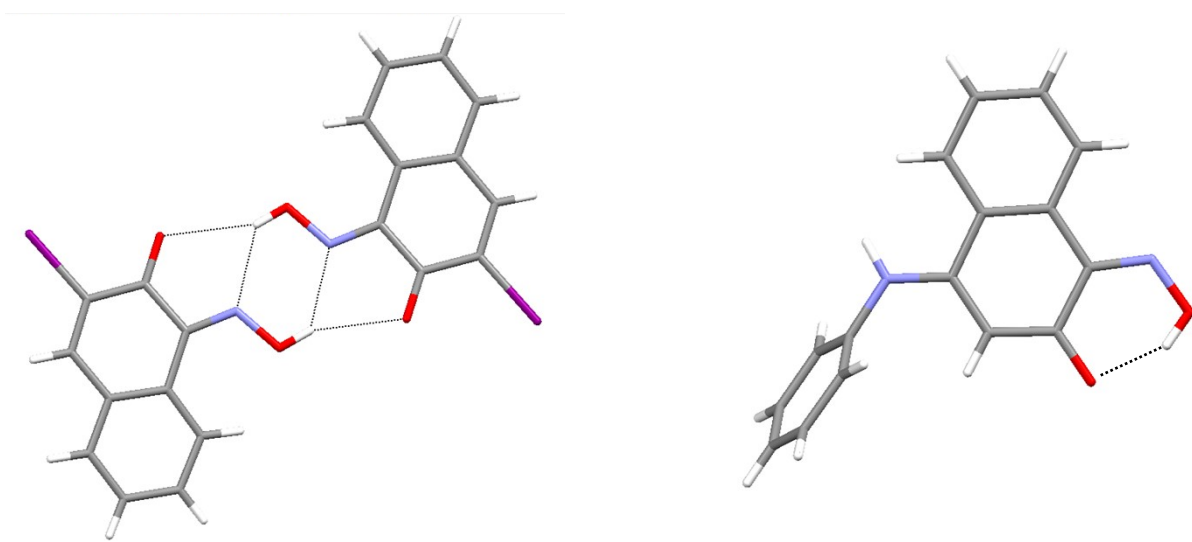
Molecule 1 (RMSD: 0.0391)



Molecule 2 (RMSD: 0.0267)

4.- CSD survey of supramolecular synthons

Table S1. List of CSD refcodes showing the supramolecular synthons searched. Search filters of structures containing only determined atomic coordinates and without organometallic compounds have been applied. Supramolecular synthon $R_1^2(5)$ (blue), intramolecular $R_1^1(6)$ interaction motif (red) and molecules with conformation as molecule 2 but showing different intermolecular interactions (black).

Search	
 --- means any bond	
CCDC	
<p> CAGFOJ BNPQOX CLEBQI DOXROX02 DOXROY FAGYUL GEXQOR KNQXSO MNTRPH MURJEQ NANGEO NQOXSX NQXSLA NQXSNK </p>	<p> NTSHQS PBZQOX SNQUOX SNQXSO UJESEH VIRDEH XOHMUC XOHNAJ YIBGIZ YIBGOF ZAMKIH ZATKUA ZIVJAP ZIVJET </p>
 <div style="display: flex; justify-content: space-around; width: 100%;"> CAGFOJ GEXQOR </div>	