

Crystal morphology prediction of structures determined by X-ray powder diffraction

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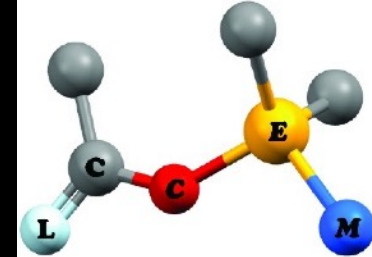
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ICDD Website - www.icdd.com

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- Prof. Dr. Giuseppe M. Lombardo



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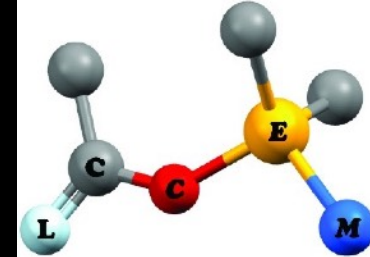
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- Prof. Dr. Fanny N. Costa
- Dr. Letícia Kuplich
- Dr. Amanda L. Ibiapino
- Dr. Laysa P. de Figueiredo



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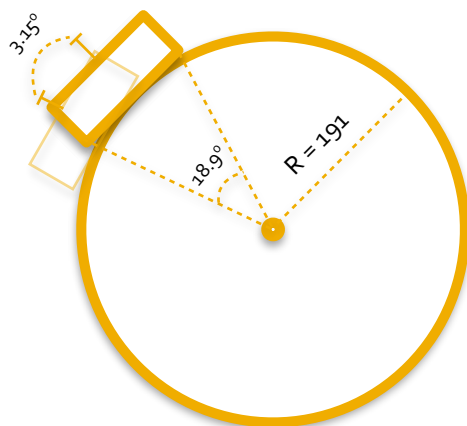
Experimental details



Stoe® STADI-P X-ray Powder Diffractometer (fully functional @ LCCEM)



- Variable counting time (VCT) acquisitions
- Transmission geometry → capillaries or acetate-cellulose foils
- $\text{CuK}\alpha_1$ radiation ($\lambda = 1.54056 \text{ \AA}$)



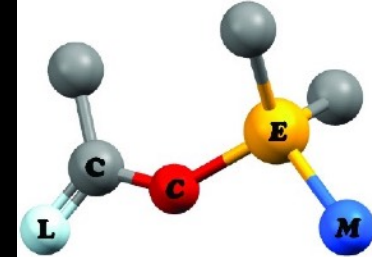
LASSBio-1773/1774		LASSBio-1755	
2θ range (°)	Time (s)	2θ range (°)	Time (s)
5-26.000	100	6-28.035	100
26.015-45.950	200	28.050-47.985	200
45.965-65.900	400	48.000-68.985	400
65.915-90.050	800	69.000-100.485	800



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Discovery of new lead-compounds

Laboratory of Evaluation and Synthesis
of Bioactive Substances (LASSBio®)



instituto nacional
de ciência e tecnologia

de Fármacos e Medicamentos

www.inct-inofar.ccs.ufrj.br

Research program to develop a series of compounds with **anti-inflammatory**, **antinociceptive**, **anticancer**, **antidiabetes** activities by structural modifications of some prototype compounds

(In this presentation) Studies devoted to the discovery of:
new oral **hypoglycemiants** with dual activity **hypoglycemiants and anti-inflammatory**
and **antinociceptive and anti-inflammatory**



LASSBio-1755

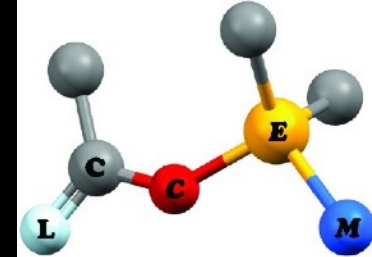
LASSBio-1773

LASSBio-1774

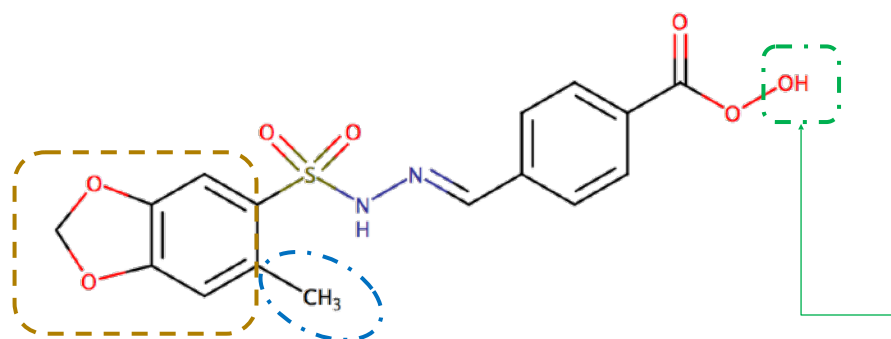


LASSBio-1773 and LASSBio-1774

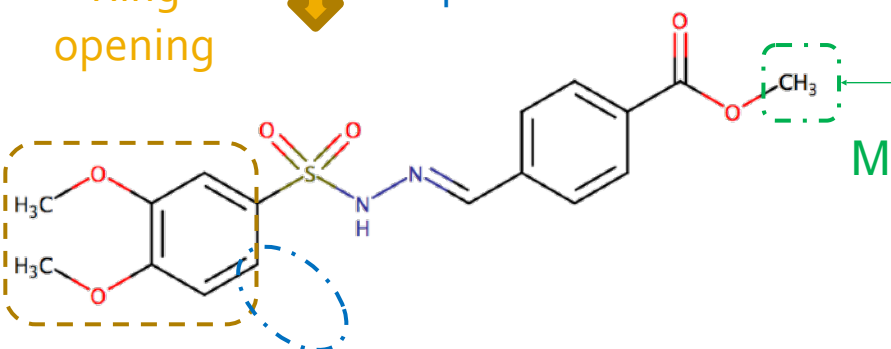
Synthesis procedure



LASSBio-1471



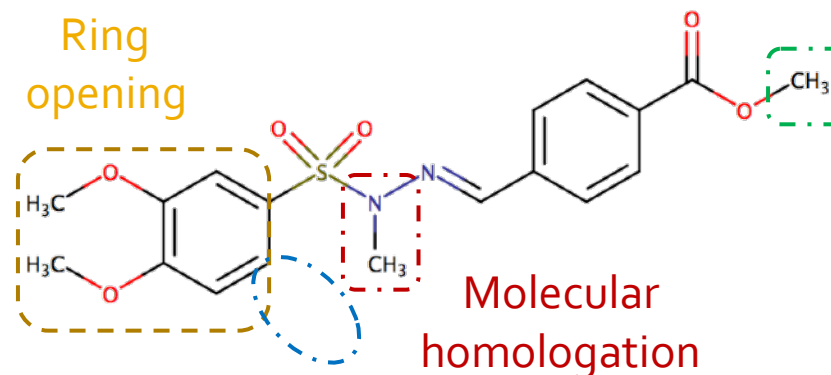
Ring opening
Molecular simplification



LASSBio-1773



LASSBio-1774



Molecular simplification

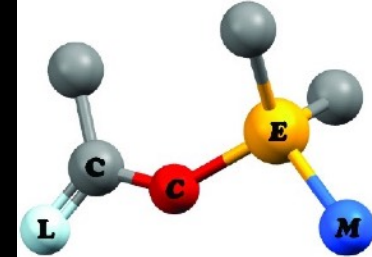
Methylation



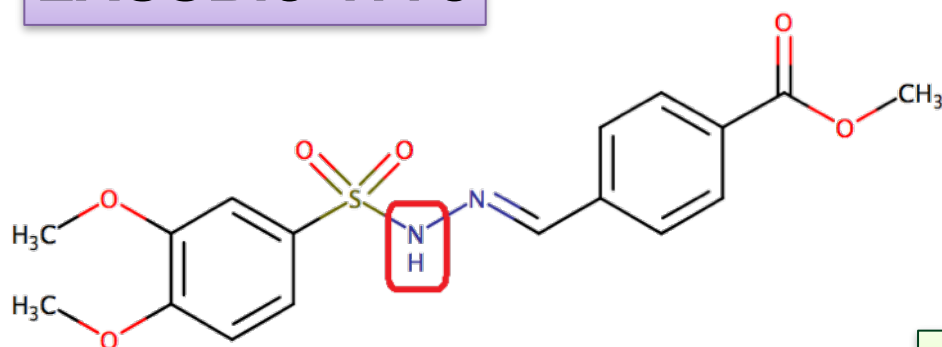
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Structural modification

LASSBio-1773 and LASSBio-1774



LASSBio-1773

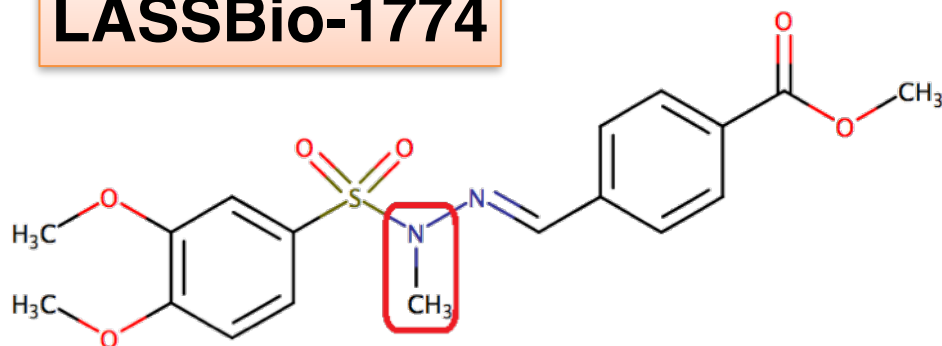


✓ *PPAR γ antagonist*

✓ Hypoglycemic activity

✓ **Samples:** *powders*
✓ **Aspect:** *white crystalline solids*

LASSBio-1774



✓ *PPAR γ agonist*

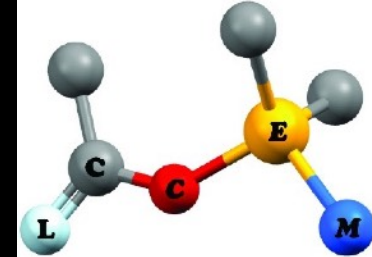
✓ Hypoglycemic and
anti-inflammatory activities



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LASSBio-1773

Structure determination



1) Indexing procedure

✓ Crystal system

Orthorhombic

✓ unit cell parameters

$a = 22.7330(6) \text{ \AA}$;

$b = 10.6880(4) \text{ \AA}$;

$c = 7.2519(2) \text{ \AA}$;

✓ unit cell volume

$V = 1762.32(1) \text{ \AA}^3$

✓ formula unit(s) per unit cell

$Z = 4$

$Z' = 1$

2) Space group determination

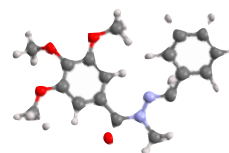
✓ space group

$P2_12_12_1$



3) Structure determination

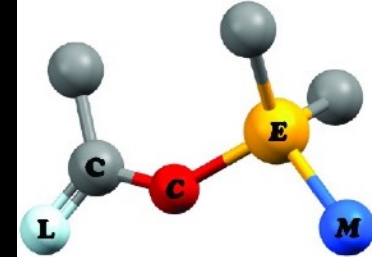
simulated annealing



4) Rietveld refinement

LASSBio-1774

Structure determination



1) Indexing procedure

✓ Crystal system
Orthorhombic (*Pbca*)

✓ unit cell parameters

***a* = 28.9007(11) Å;**

***b* = 15.0348(4) Å;**

***c* = 9.1482(2) Å;**

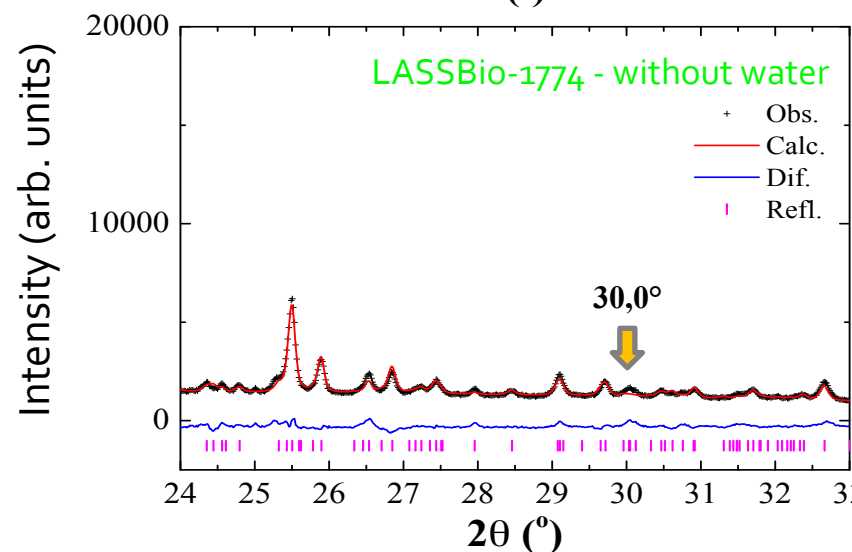
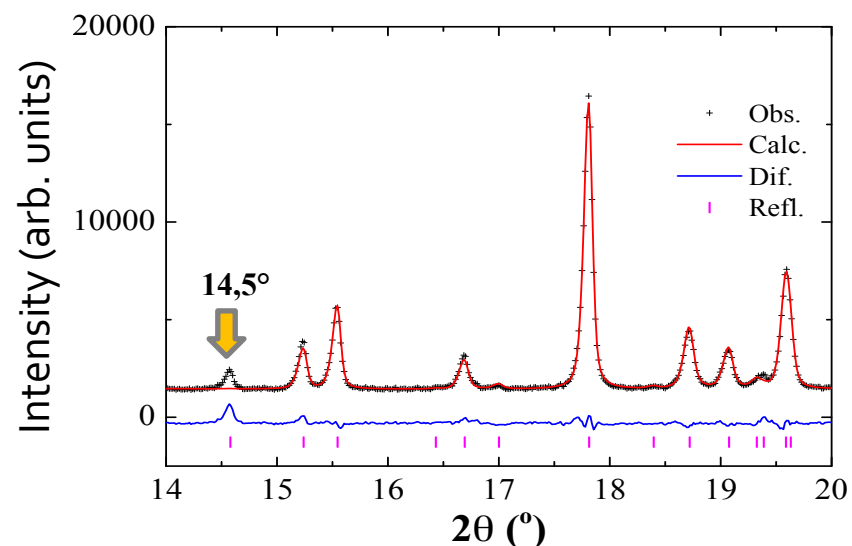
✓ unit cell volume

***V* = 3975.1(2) Å³**

✓ formula unit(s) per unit cell

***Z* = 8**

***Z'* = 1**

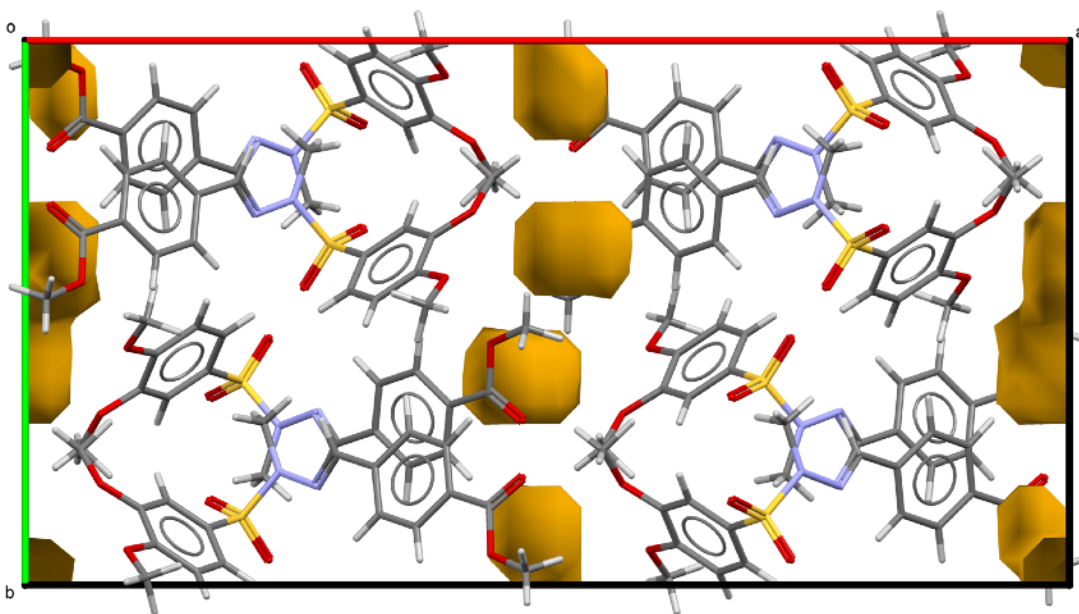
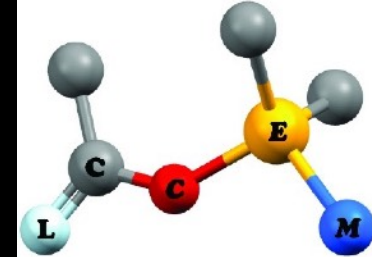




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Voids



Packing along the **c-axis**

✓ “Voids” within the unit cell

Volume: $\sim 179.4 \text{ \AA}^3$



Volume: water
(H_2O)

Volume: ethanol
($\text{C}_2\text{H}_6\text{O}$)

Volume: $\sim 22 \text{ \AA}^3$

Volume: $\sim 70 \text{ \AA}^3$



8 voids: $\sim 176 \text{ \AA}^3$



8 voids: $\sim 560 \text{ \AA}^3$

$$\frac{-H}{v_i} = 5.08 \text{ \AA}^3, \frac{-O}{v_i} = 11.39 \text{ \AA}^3 \text{ and } \frac{-C}{v_i} = 13.87 \text{ \AA}^3$$

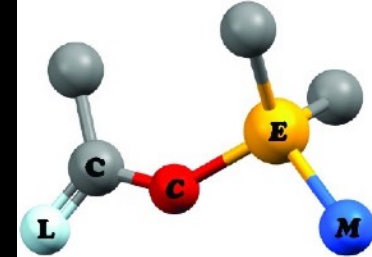
Hoffmann, D. W. M., Acta Cryst. (2002). B57, 489-493



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LASSBio-1774

Structure re-determination



1) Indexing procedure

✓

Crystal system

Orthorhombic

✓

unit cell parameters

***a* = 28.9030(10) Å;**

***b* = 15.0357(3) Å;**

***c* = 9.1476(1) Å;**

✓

unit cell volume

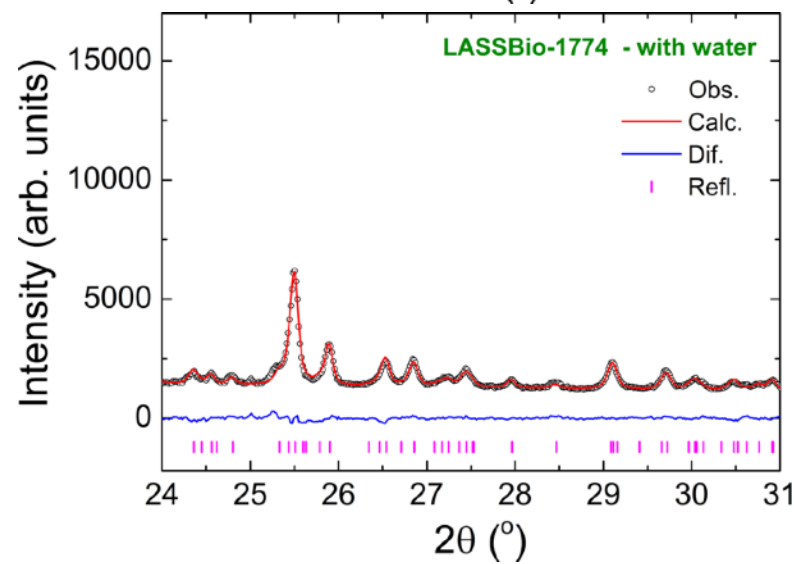
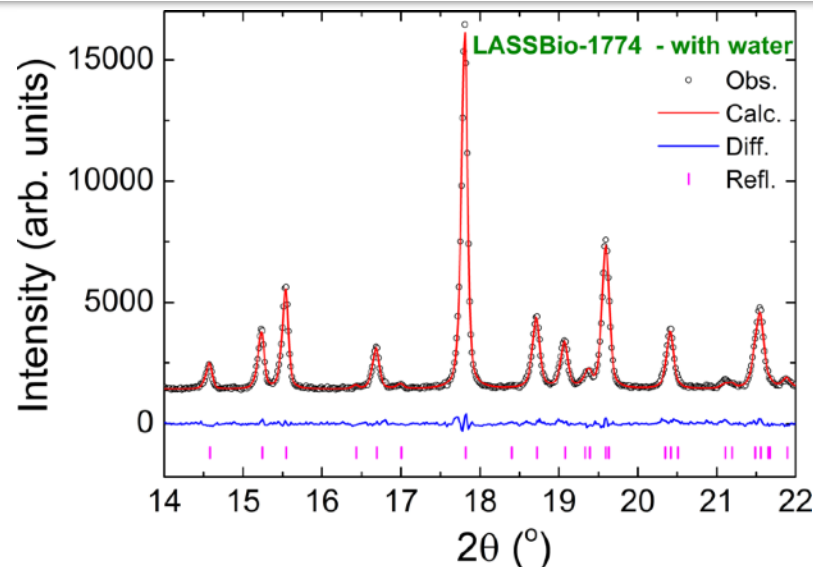
***V* = 3975.40(8) Å³**

✓

formula unit(s) per unit
cell

***Z* = 8**

***Z'* = 1**

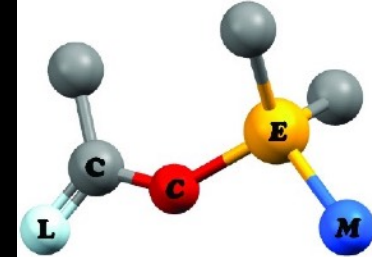




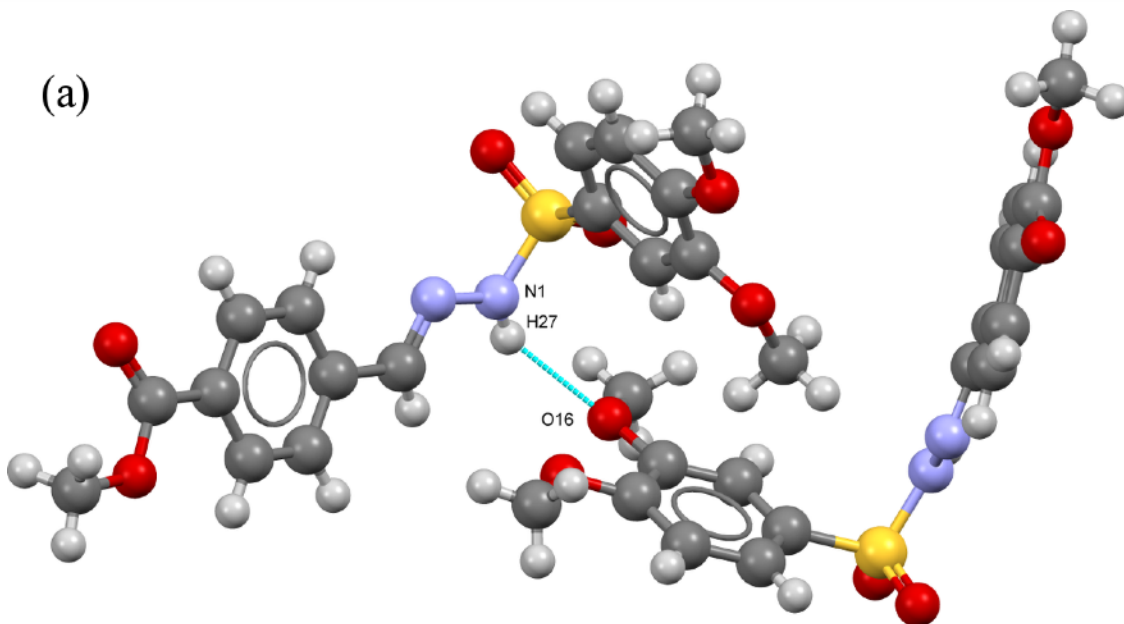
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Hydrogen interactions



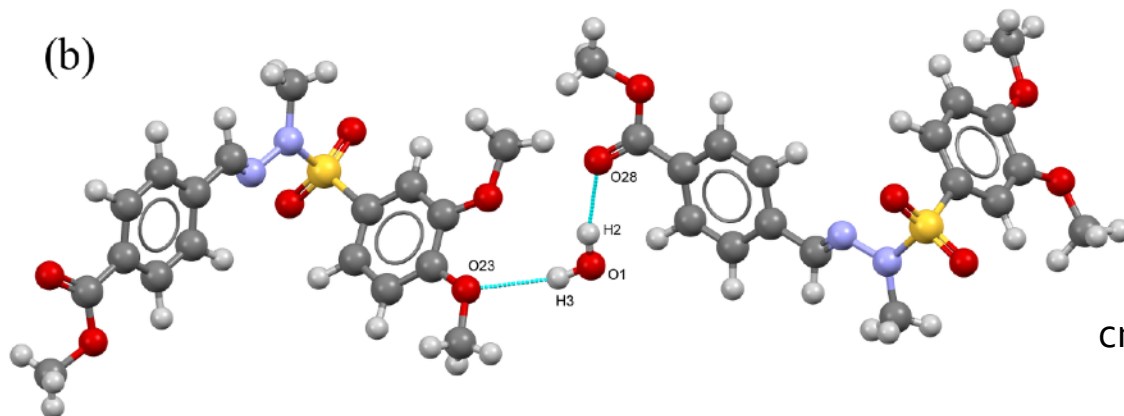
(a)



folded shape with an apical polar part and two mainly apolar wings

crystallized in water

(b)



mainly a non-polar landscape
has been transformed in a
clearly polar one in all the
considered crystal faces

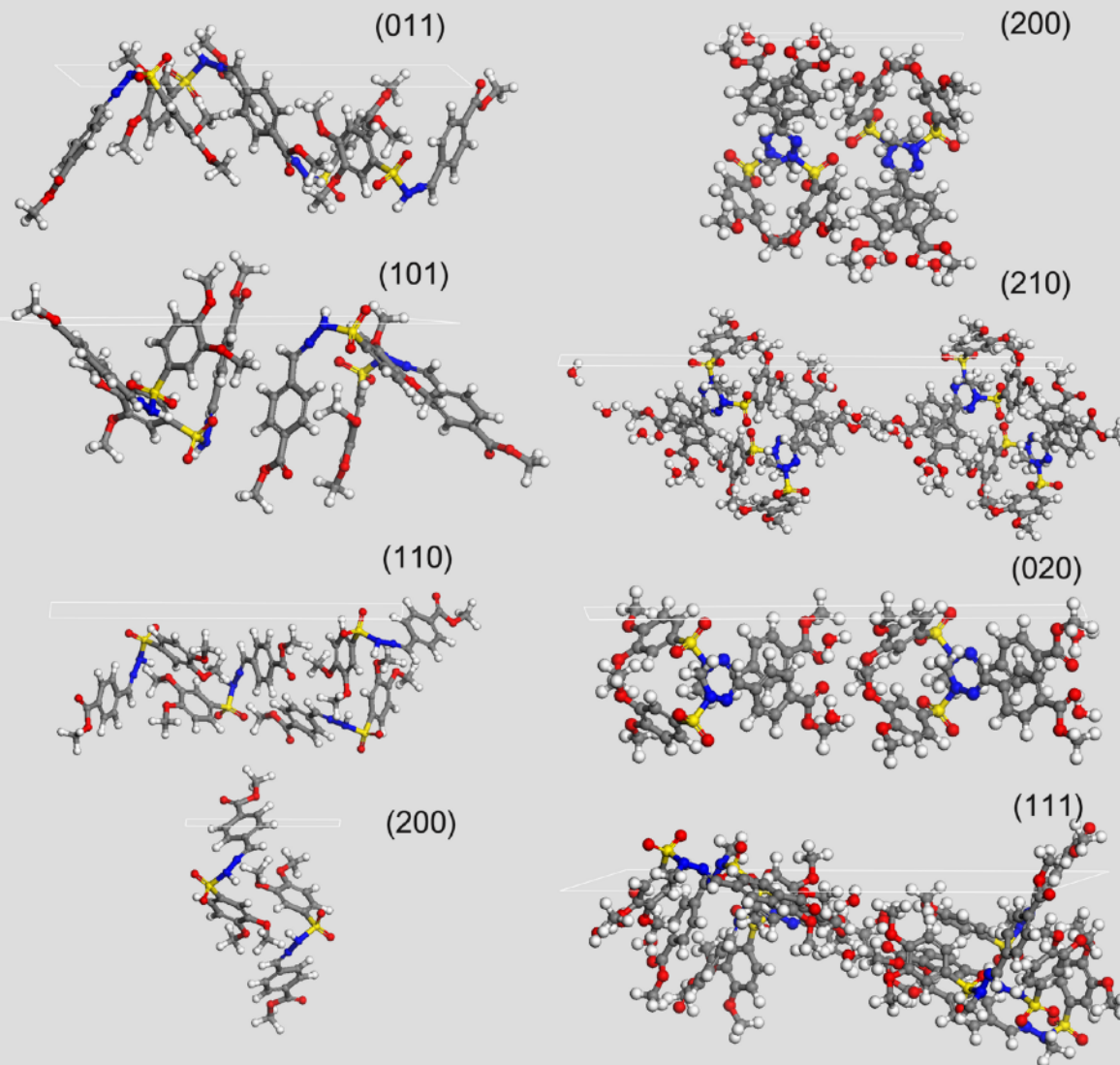
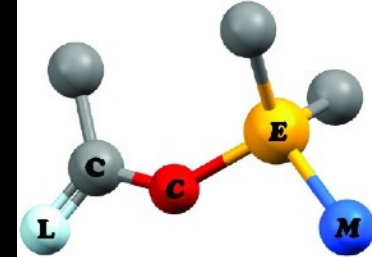
crystallized in a mixture of ethanol/water



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Morphology prediction

LASSBio-1773 and LASSBio-1774



Growth morphology method

The **growth morphology (GM)** method can predict the **shape of a crystal** more accurately than the BFDH method because it takes the **energetics of the system** into account

The **crystallization energy**, E_{cr} , is defined as:

$$E_{cr} = E_{slice} + E_{att}$$

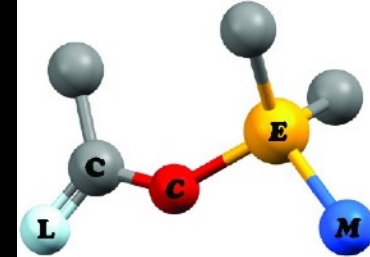
E_{slice} = energy resulting from the **lateral interaction** of each formula unit **within a slice**

E_{att} = energy released as a consequence of the **vertical interaction** of the formula unit with an **underlying slice**

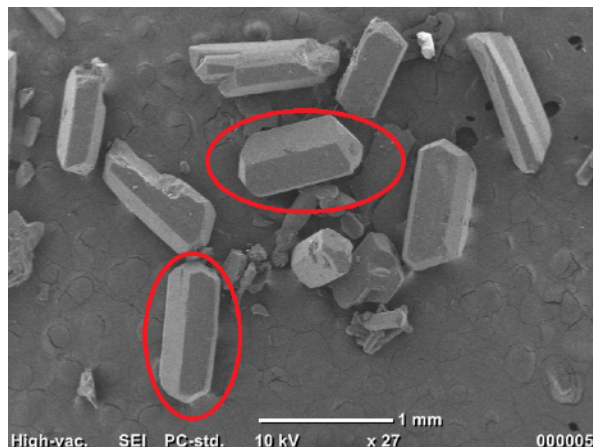
$$G_r \propto E_{att}$$

Morphology prediction

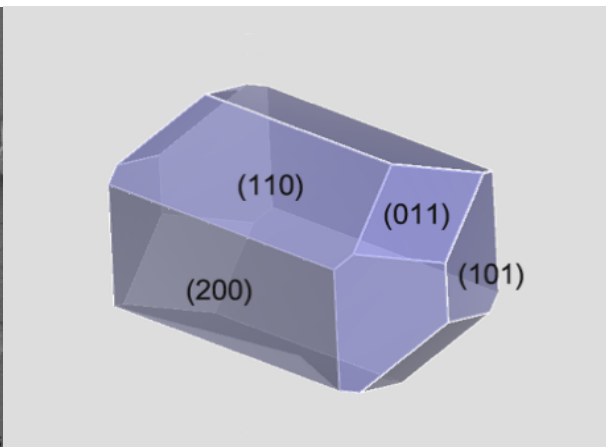
LASSBio-1773



SEM



Simulation (*Growth morphology*)



Potential effect of solvents in the growth mechanism of crystal faces

$$(200) \quad E_{att} = -126.2876 \text{ kcal.mol}^{-1}$$

$$(101) \quad E_{att} = -180.9218 \text{ kcal.mol}^{-1}$$

$$(011) \quad E_{att} = -177.3885 \text{ kcal.mol}^{-1}$$

apolar

$$(110) \quad E_{att} = -120.7005 \text{ kcal.mol}^{-1}$$

polar

← Growth of face (110)

Comparison between the SEM image and morphology prediction

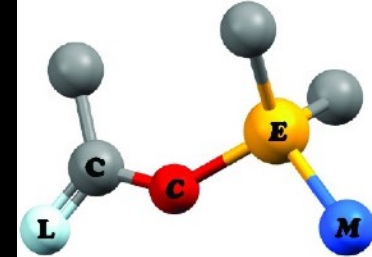
<i>hkl</i>	Multiplicity	d_{hkl} (Å)	$E_{att}(\text{Total})$ (kcal mol ⁻¹)	Total Facet Area (%)
(110)	4	9.6624	-120.7005	43.74
(200)	2	11.3564	-126.2876	24.32
(101)	4	5.9917	-180.9218	18.57
(011)	4	6.8973	-177.3885	13.37

Morphologically Important (MI) faces
% depends on d_{hkl} as well as on E_{att}

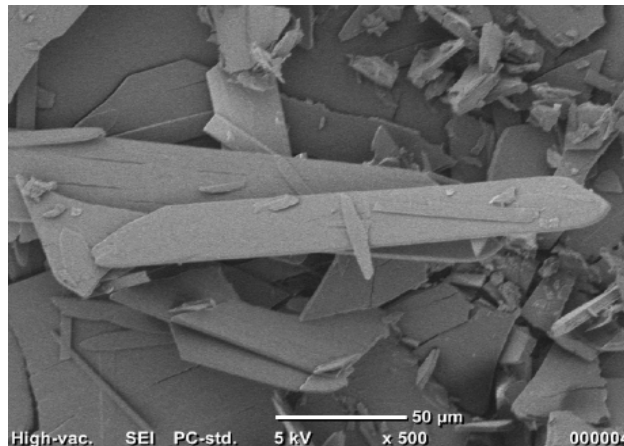
Strategy: use of less polar solvents, with water excess, during the crystallization of **LASSBio-1773**

Morphology prediction

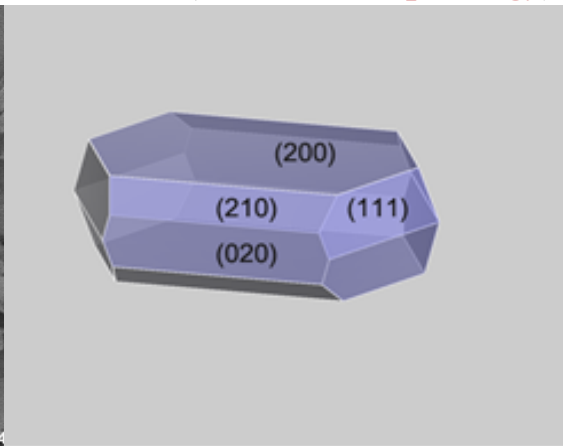
LASSBio-1774



SEM



Simulation (*Growth morphology*)



Potential effect of solvents in the growth mechanism of crystal faces



(200) $E_{att} = -87.4231 \text{ kcal.mol}^{-1}$

(210) $E_{att} = -154.3096 \text{ kcal.mol}^{-1}$

(020) $E_{att} = -178.4827 \text{ kcal.mol}^{-1}$

(111) $E_{att} = -235.5099 \text{ kcal.mol}^{-1}$

Comparison between the SEM image and morphology prediction

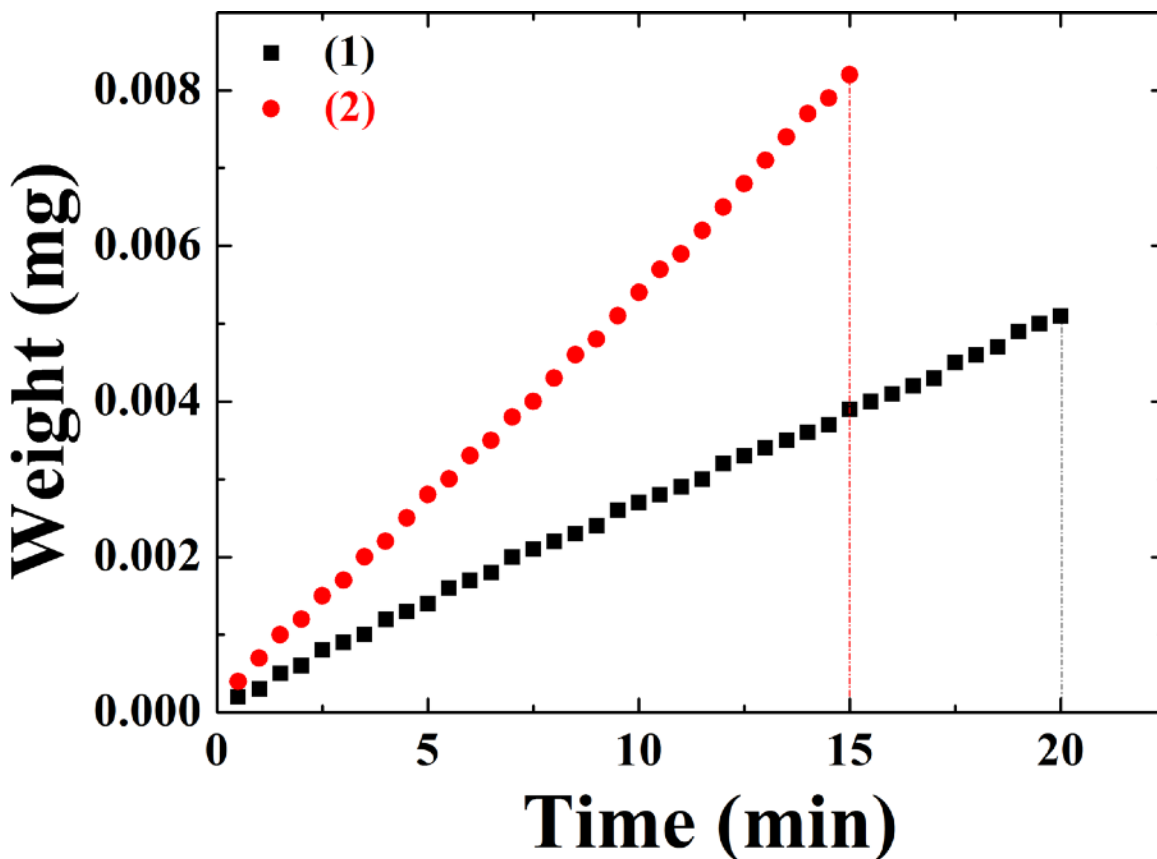
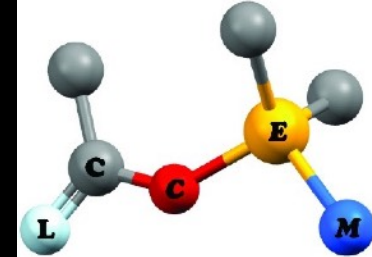
<i>hkl</i>	Multiplicity	<i>d</i> _{<i>hkl</i>} (Å)	<i>E</i> _{<i>att</i>} (Total) (kcal mol ⁻¹)	Total Facet Area (%)
(200)	2	14.4516	-87.4231	44.39
(210)	4	10.4192	-154.3096	17.12
(20)	2	7.5179	-178.4827	11.19
(111)	8	7.5441	-235.5099	27.30

polar



Favors the solubility in polar solvents

Intrinsic dissolution rate (IDR)



LASSBio-1773

IDR : 6.4(6) $\mu\text{g}\cdot\text{cm}^{-2}\cdot\text{min}^{-1}$

LASSBio-1774

IDR: 15.9(6) $\mu\text{g}\cdot\text{cm}^{-2}\cdot\text{min}^{-1}$

Poor solubility

values below 100 $\mu\text{g}\cdot\text{cm}^{-2}\cdot\text{min}^{-1}$

Comparison between the dissolution profiles

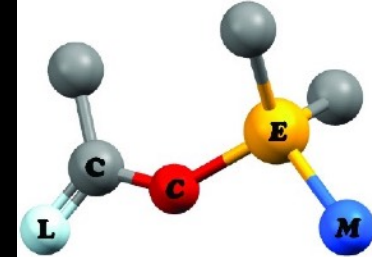
*Experimental conditions: dissolution medium - phosphate buffer (pH = 6.8) at 37°C



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LASSBio-1755

Synthesis procedure



Journal of Molecular Structure 1146 (2017) 735–743

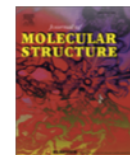


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Journal of Molecular Structure

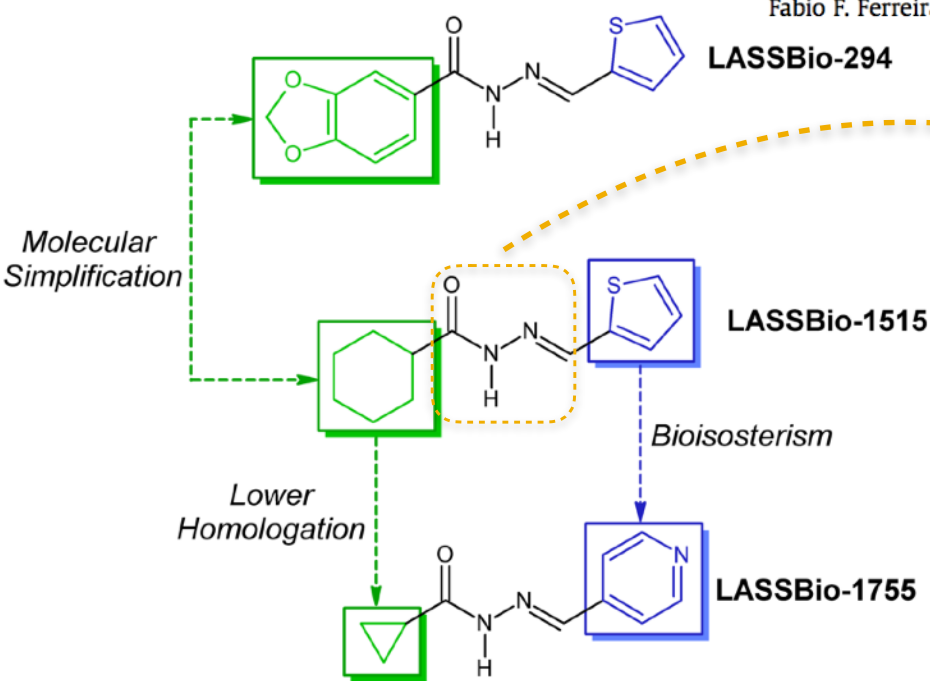
journal homepage: <http://www.elsevier.com/locate/molstruc>



A combined experimental and *in silico* characterization to highlight additional structural features and properties of a potentially new drug



Isadora T.S. Bastos ^a, Fanny N. Costa ^b, Tiago F. Silva ^c, Eliezer J. Barreiro ^{c, d},
Lidia M. Lima ^{c, d}, Delson Braz ^e, Giuseppe M. Lombardo ^f, Francesco Punzo ^f,
Fabio F. Ferreira ^{b, *}, Regina C. Barroso ^a



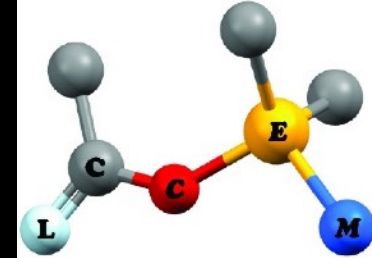
N-acylhydrazone privileged structure



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LASSBio-1755

Structure determination



1) Indexing procedure

✓ Crystal system

Triclinic

✓ unit cell parameters

$a = 4.8594(2) \text{ \AA}$;

$b = 9.3162(5) \text{ \AA}$;

$c = 11.3701(5) \text{ \AA}$;

$\alpha = 73.400(4)^\circ$;

$\beta = 78.088(3)^\circ$;

$\gamma = 82.640(4)^\circ$

✓ unit cell volume

$V = 479.70(4) \text{ \AA}^3$

✓ formula unit(s) per unit cell

$Z = 2$

$Z' = 1$

2) Space group determination

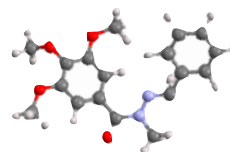
✓ space group

$P\bar{1}$



3) Structure determination

simulated annealing



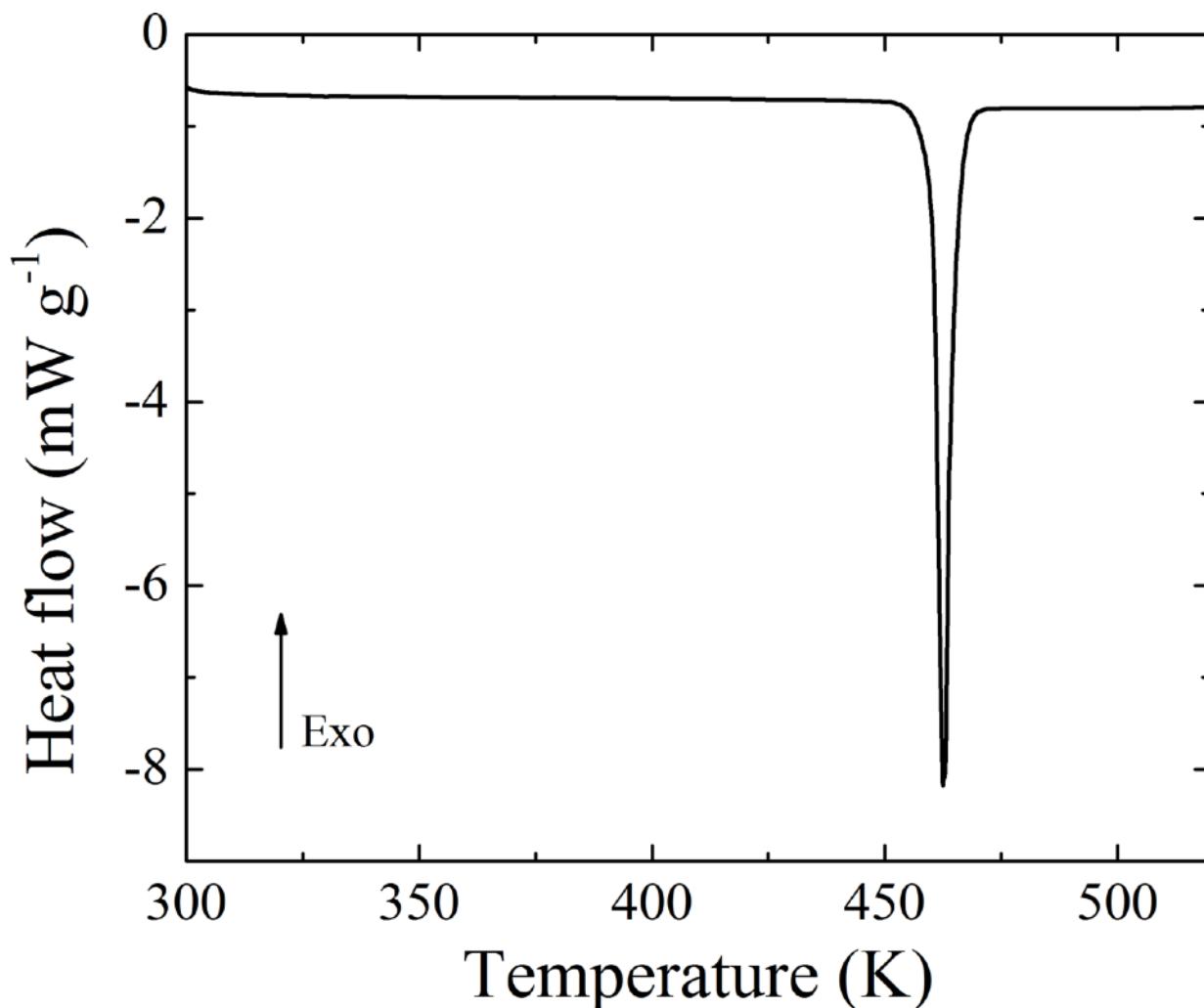
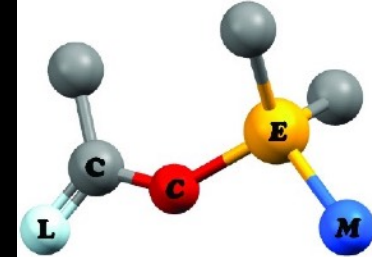
4) Rietveld refinement



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LASSBio-1755

DSC scan



well-defined melting behavior
with a sharp minimum

melting temperature
(460.81 K)

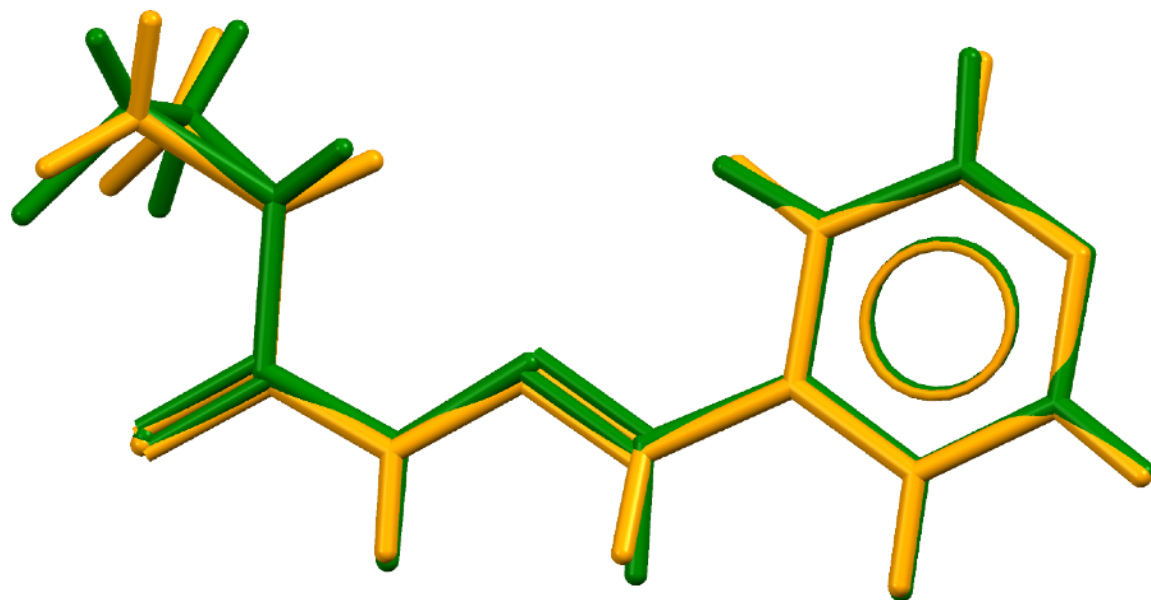
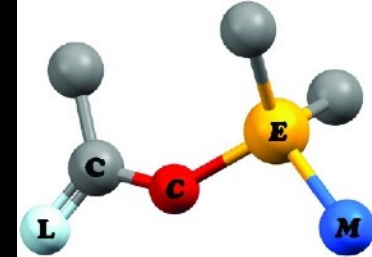
high purity and a
relative good stability

enthalpy of fusion
 $\Delta H_f = 30.43 \text{ kJ mol}^{-1}$

No tendency for
polymorph formation

X-ray powder diffraction

Overlay of structures



RMS = 0.0189
MD

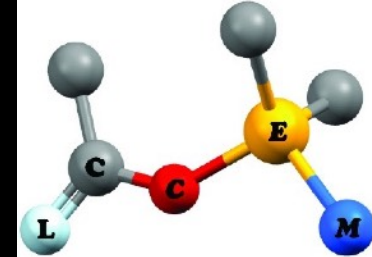
RMS = 0.0952
XRPD



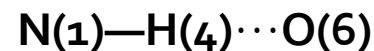
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Hydrogen interactions

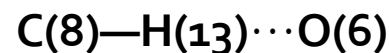


Intramolecular bond

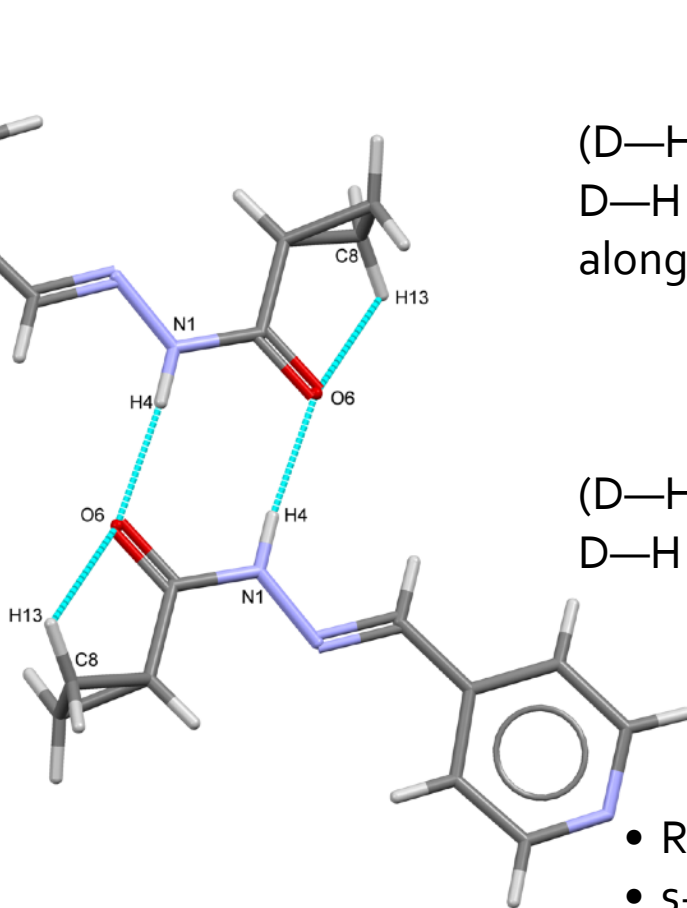


(D—H = 0.86(2) Å, H···A = 1.97(2) Å, D···A = 2.78(2) Å
 D—H···A = 156.4(19)°
 along x, 1 y, 1 z

weak intermolecular bond



(D—H = 0.96(3) Å, H···A = 2.42(2) Å, D···A = 2.83(3) Å
 D—H···A = 105.8(16)°



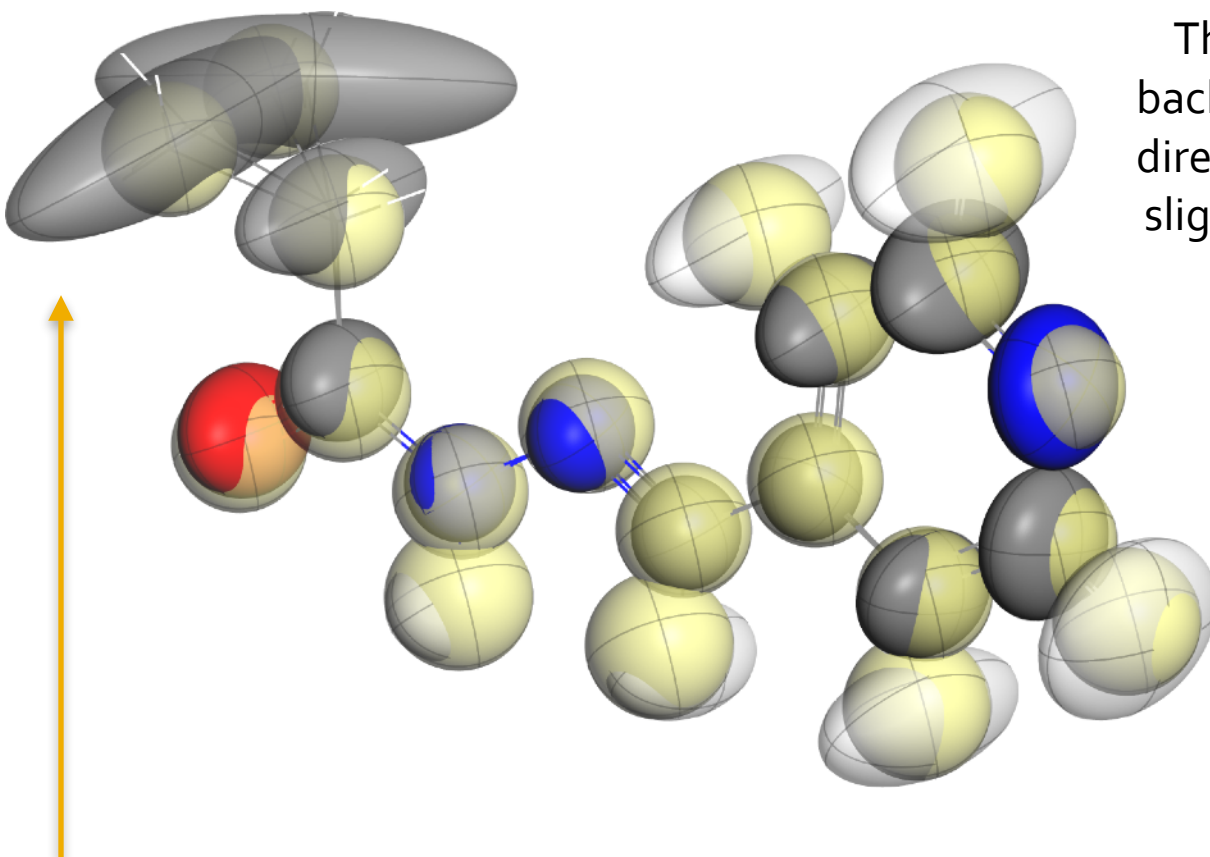
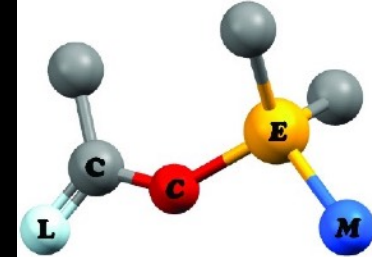
- Relative configuration *E* of the imine double bond
- *s-cis* conformation of the amide function of the *N*-acylhydrazone compound



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LASSBio-1755

ADPs calculation

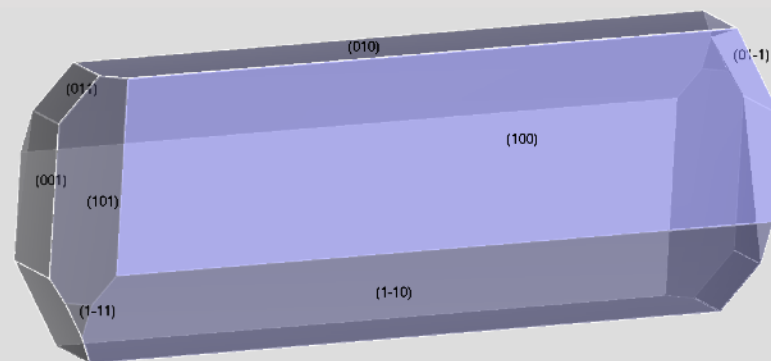
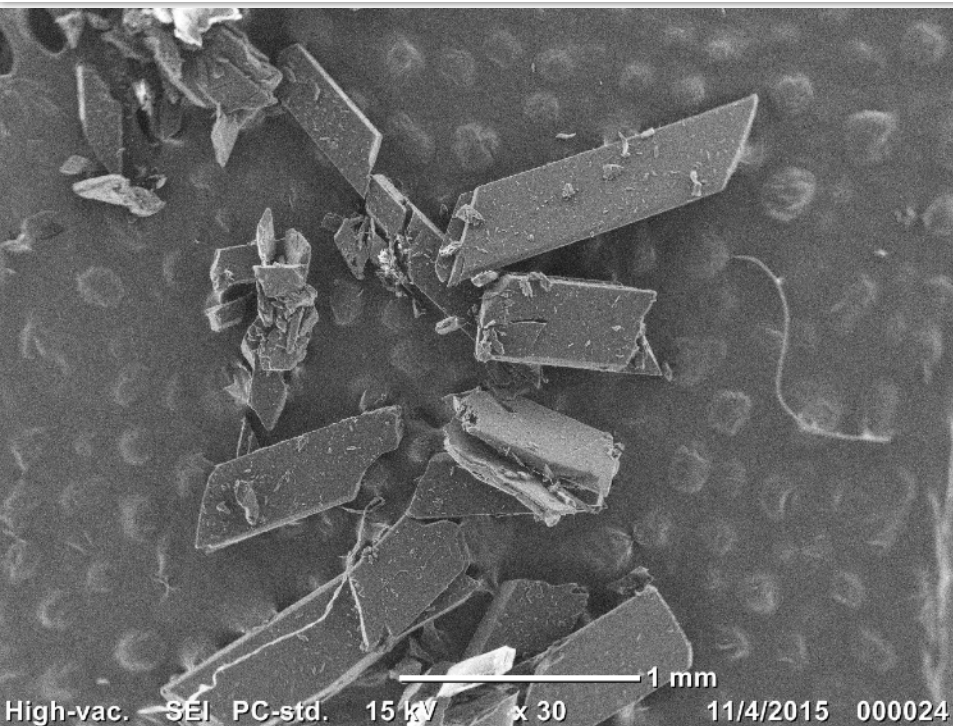
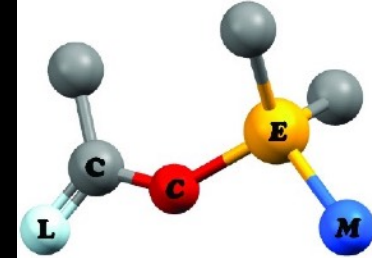


The adp motions in the molecule backbone points mostly in the same direction, while the two rings show a slightly different movement around their equilibrium

cycloalkylic moiety does not show a fairly good overall estimation of the app volume

LASSBio-1755

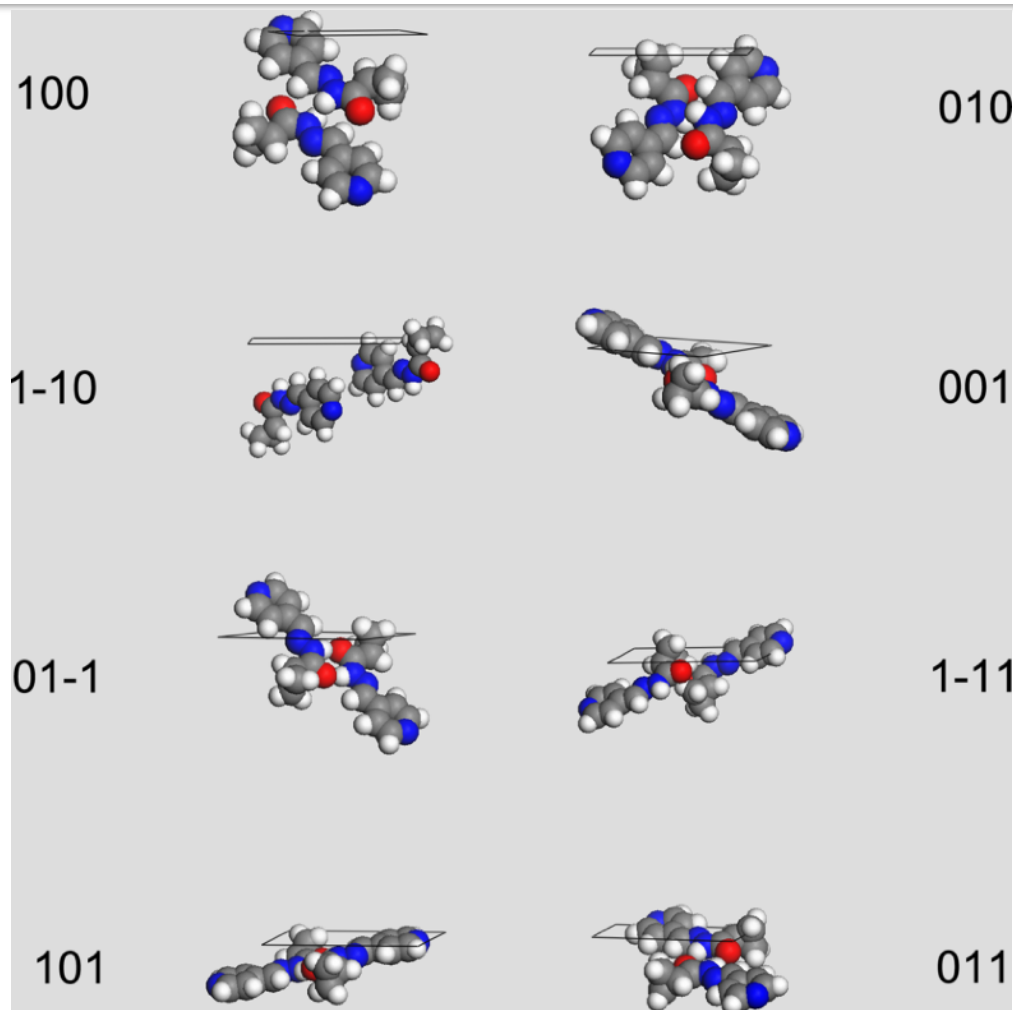
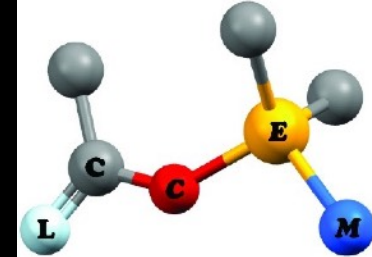
Morphology prediction



- Very good agreement between the experimentally inferred SEM images and the computationally derived crystal habit
- Growth rate of the crystal face is inversely proportional to the interplanar spacing d_{hkl}

LASSBio-1755

Morphologically important faces



Morphologically important (MI) faces have the largest d_{hkl} values

Use of the attachment energy (E_{att})

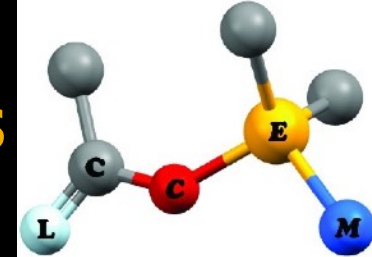
hkl	Multiplicity	d_{hkl} (Å)	E_{att} (Total) (kcal mol ⁻¹)	Total Facet Area (%)
(001)	2	10.6842	-16.2911	29.71
(010)	2	8.9036	-15.1595	29.24
(011)	2	7.9662	-16.7300	20.17
(101)	2	4.6534	-35.9330	7.21
(111)	2	4.4862	-37.6315	4.95
(112)	2	3.9747	-36.2963	0.22
(102)	2	3.9100	-32.0516	6.27
($\bar{1}11$)	2	3.8375	-33.8921	2.23



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A series of different compounds

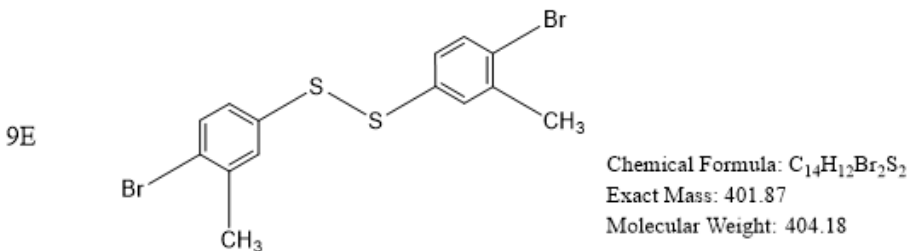
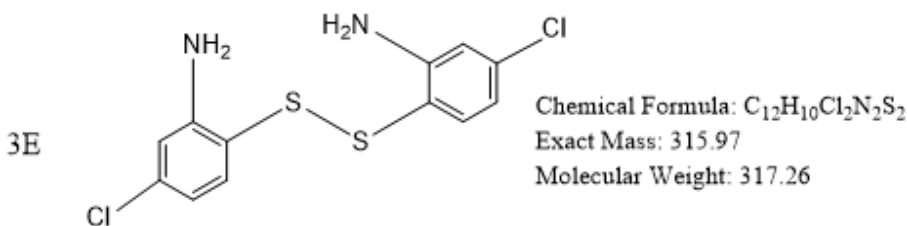
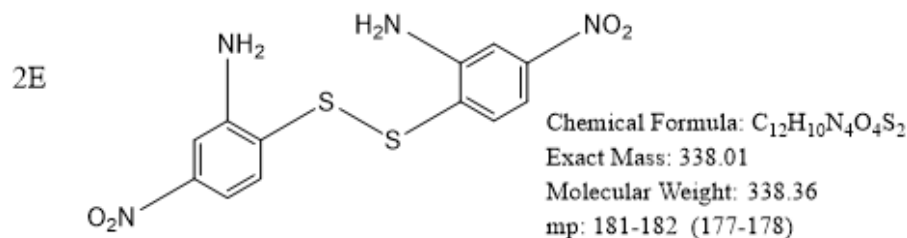
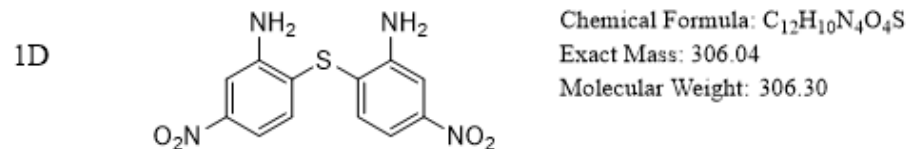
Morphologically important faces - on going work



Single S bond → double single S bond

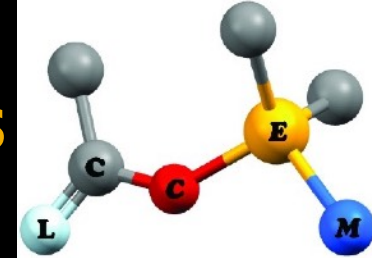
Change of functional groups from NO₂ to Cl

Complete change of functional groups

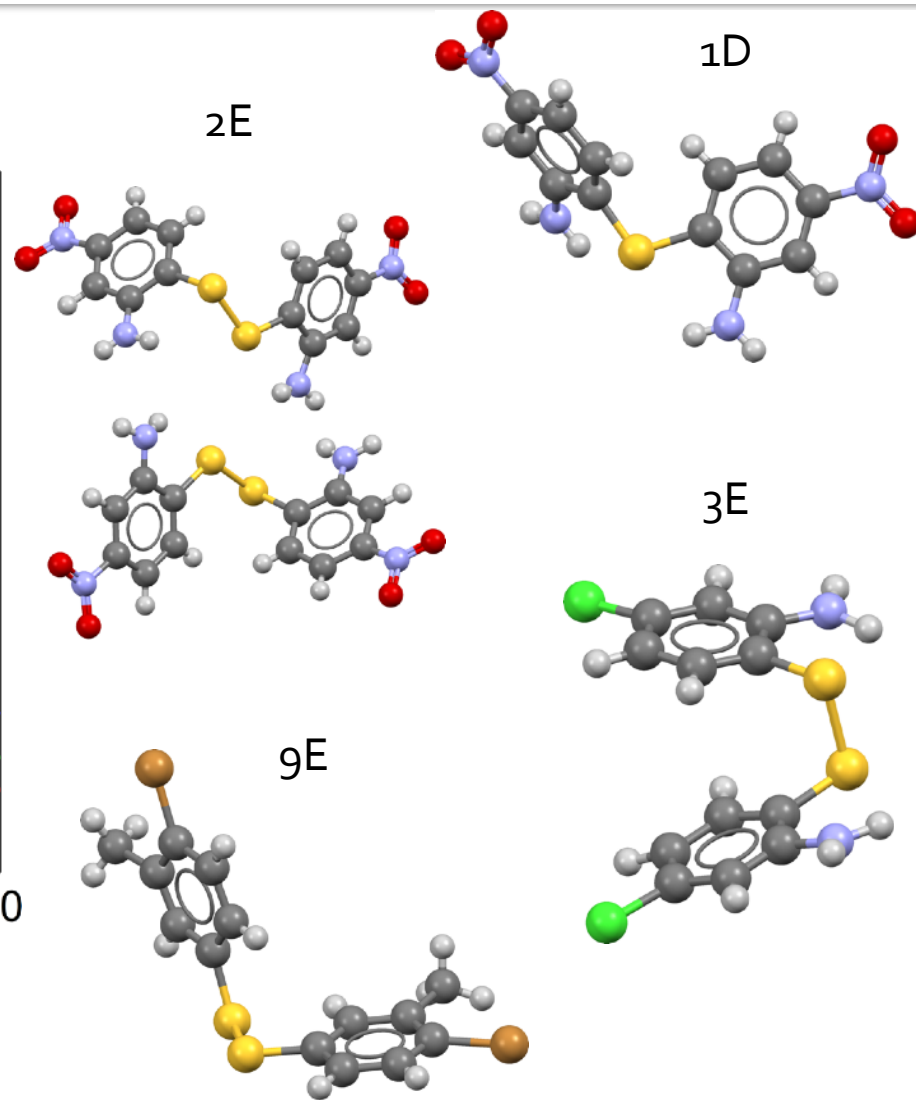
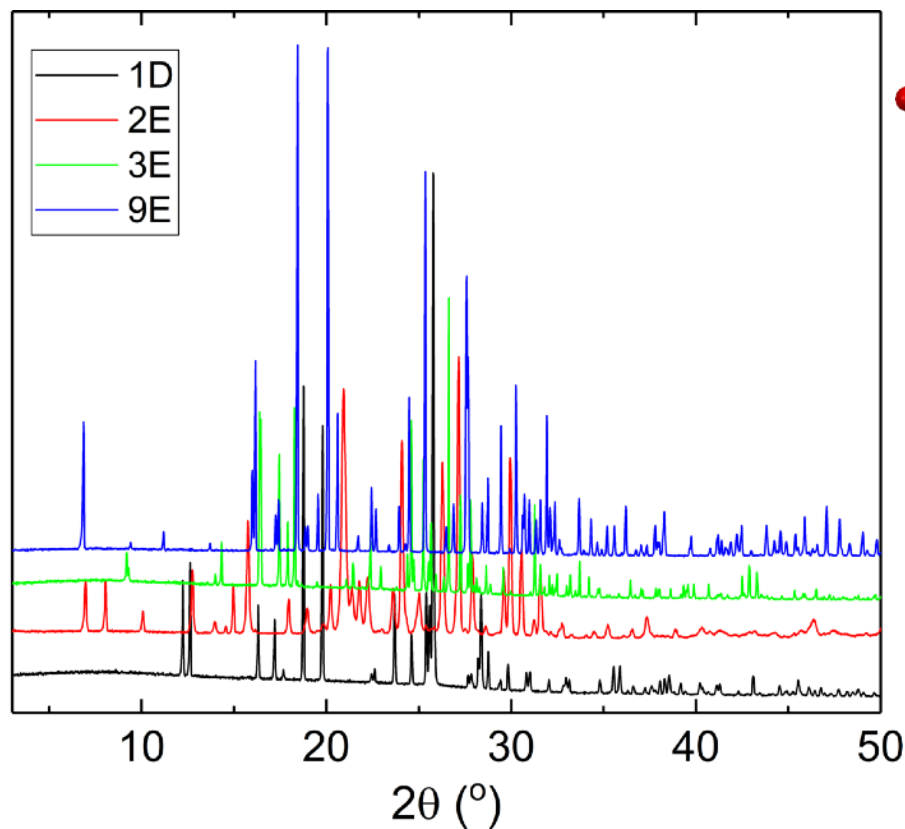


A series of different compounds

Morphologically important faces - on going work



Intensity (arb. units)

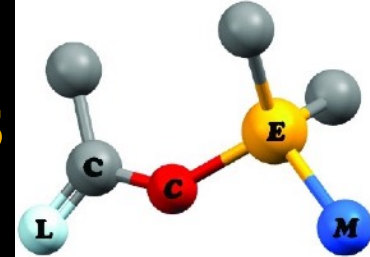




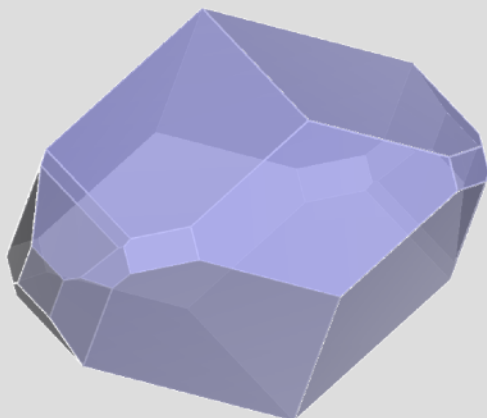
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A series of different compounds

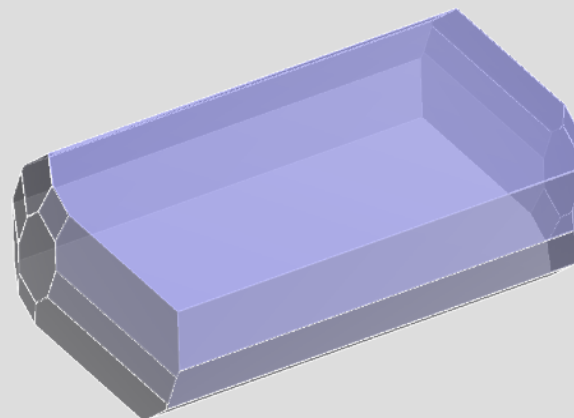
Morphologically important faces - on going work



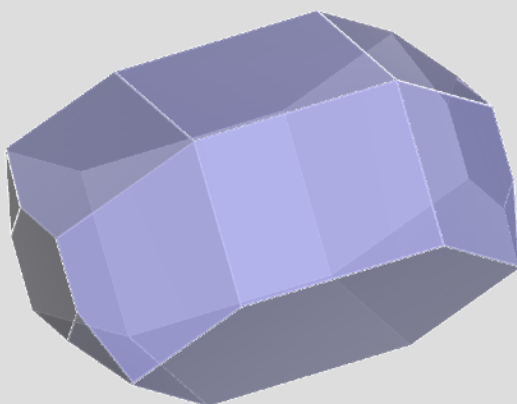
1D



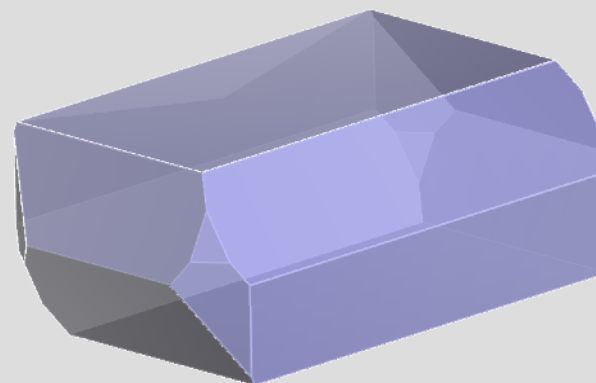
2E



3E

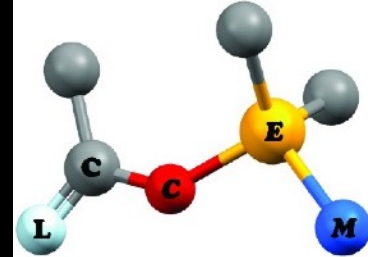


9E



The case of spironolactone

ongoing work



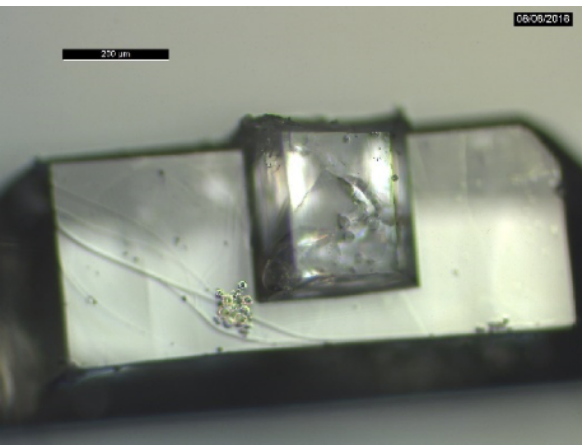
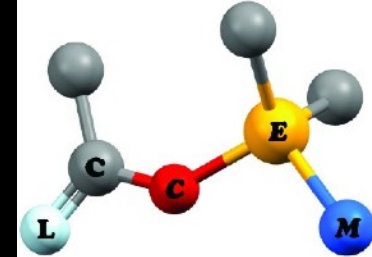
- A diuretic steroidal aldosterone agonist
- Poor water solubility and dissolution rate
- Two polymorphs and 5 solvates described in the literature
 - only 4 crystal structures reported
- Recrystallized from:
 - **acetone**: WUWROW (CSD Refcode)
 - **acetonitrile**: KIKWUW (CSD Refcode)
 - ethyl acetate (only space group and unit cell parameters reported so far)



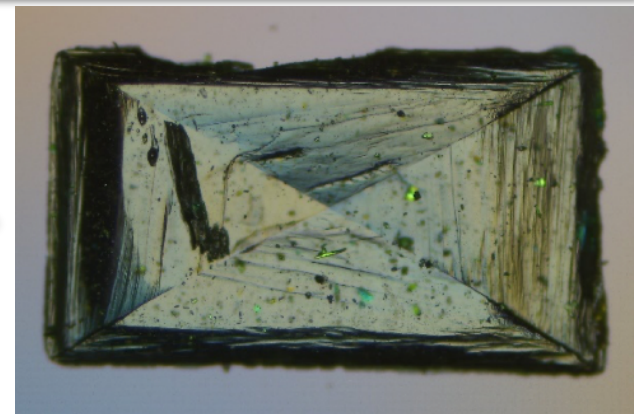
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The case of spironolactone

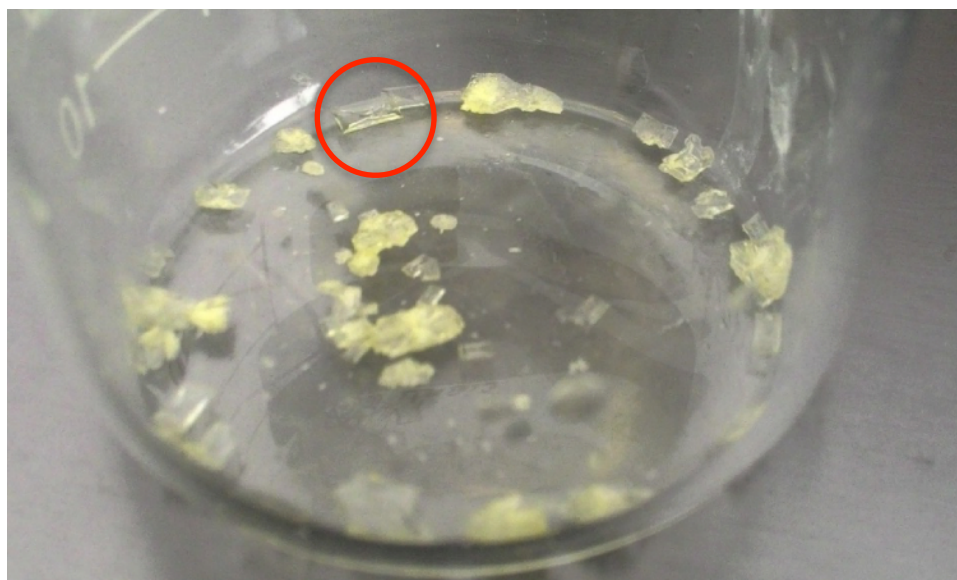
ongoing work



← Optical microscopy →



Spironolactone recrystallized in acetone



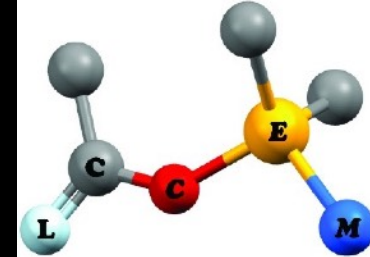
← Mobile phone camera



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The case of spironolactone

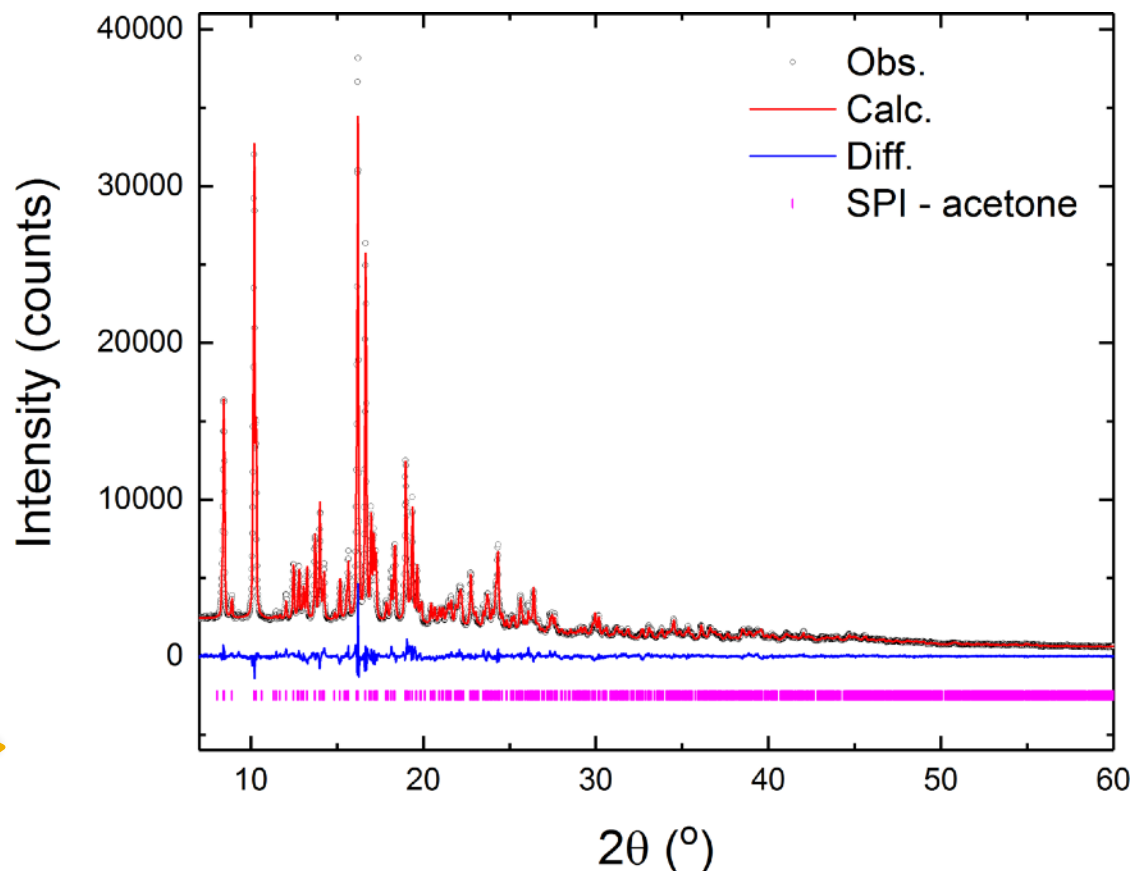
ongoing work



Agafonov [1] describes forms I and II → we have obtained the hydrated form, described by Takata [2]

- Structure contains disorder
- Fractional coordinates not refined

ATPRCL01	Spironolactone Form II
ATPRCL10	Spironolactone Form I
WUWROW	Spironolactone hydrate

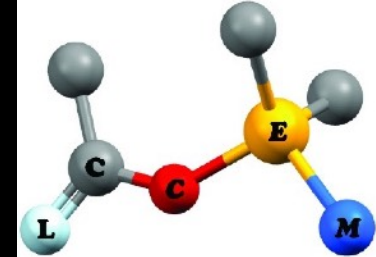


[1] V. Agafonov, B. Legendre, N. Rodier, Acta Crystallogr., Sect. C: Cryst. Struct. Commun. (1989), 45, 1661

[2] N. Takata, R. Takano, H. Uekusa, Y. Hayashi, K. Terada, Cryst. Growth Des. (2010), 10, 2116

The case of spironolactone

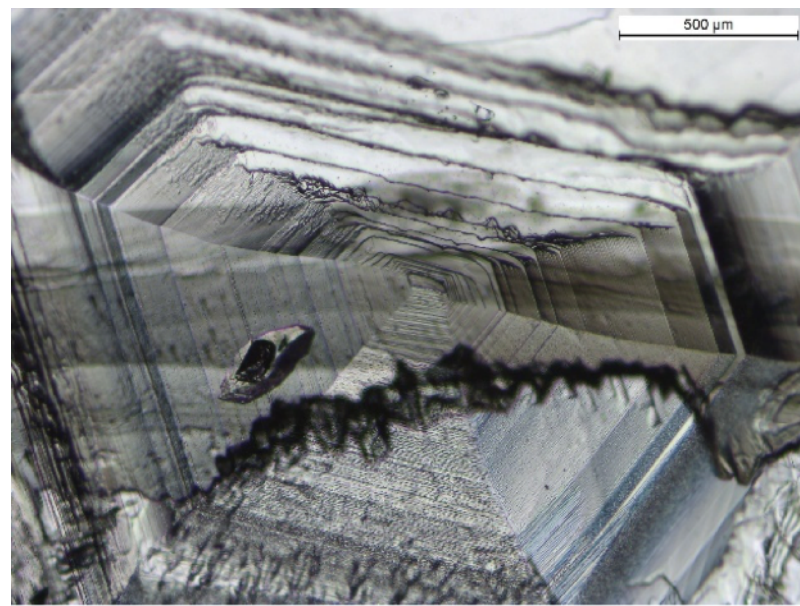
ongoing work



← Mobile phone camera

Spironolactone recrystallized in acetonitrile

Optical microscopy →

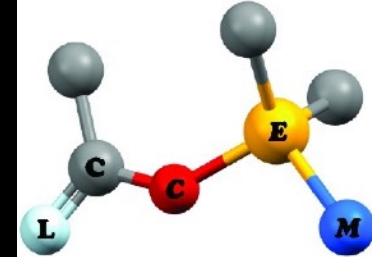




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The case of spironolactone

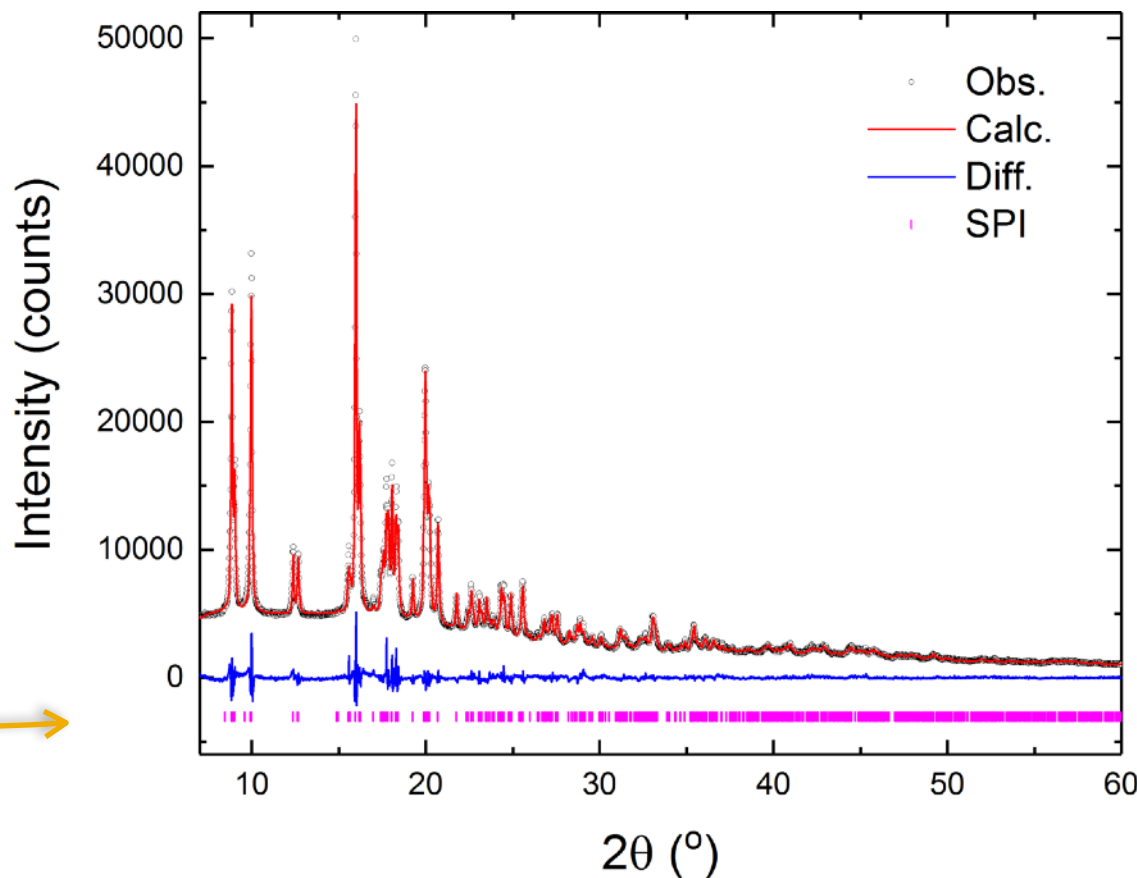
ongoing work



Form III described by Agafonov et al. [1]

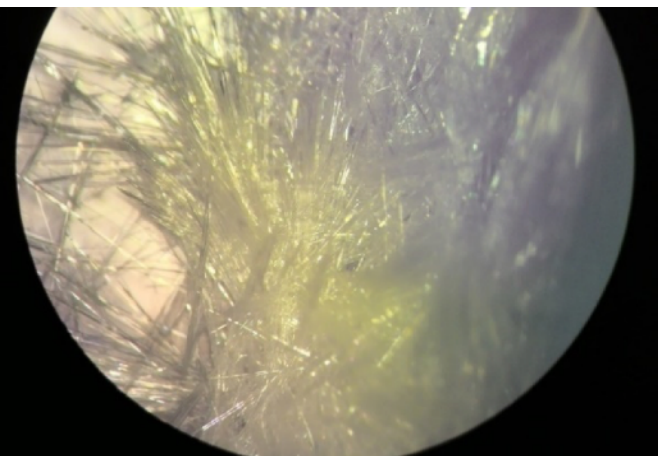
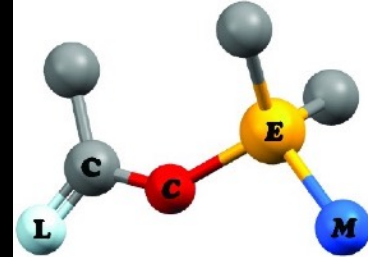
- Structure contains disorder
- Fractional coordinates not refined

KIKWUW: Spironolactone acetonitrile solvate
(Form III)

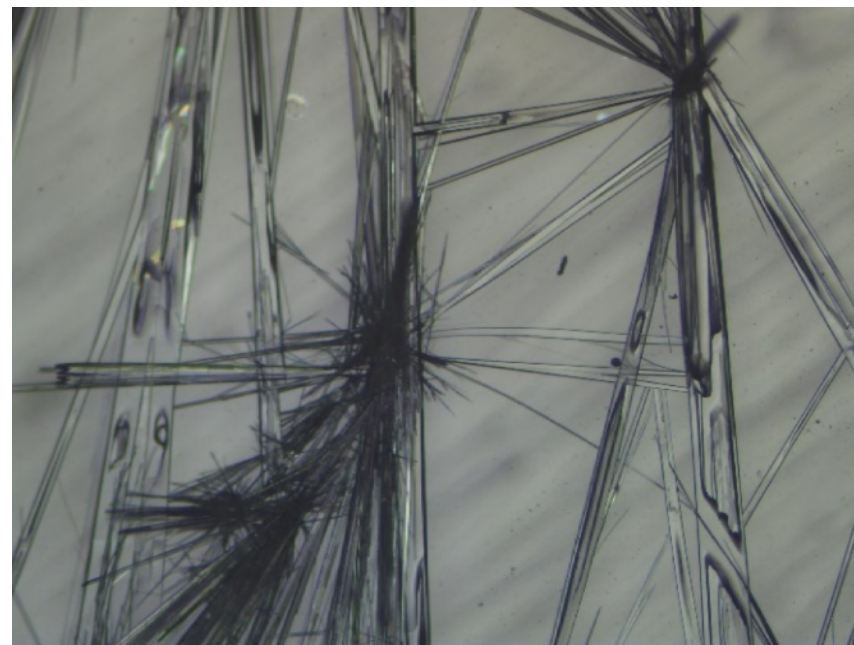
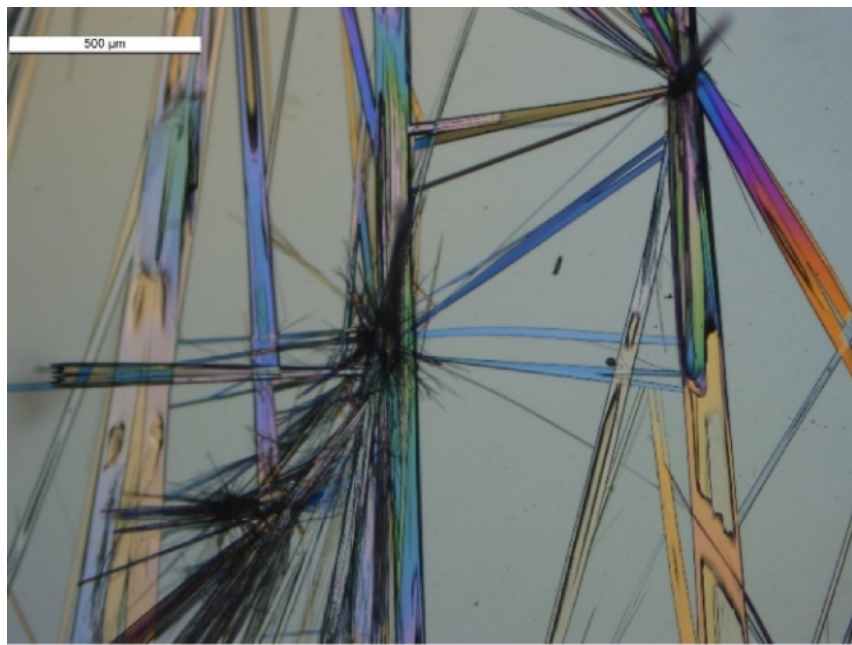


The case of spironolactone

ongoing work

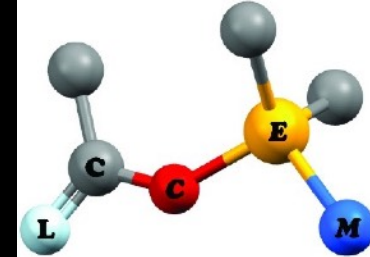


Spironolactone recrystallized
in ethyl acetate



The case of spironolactone

ongoing work



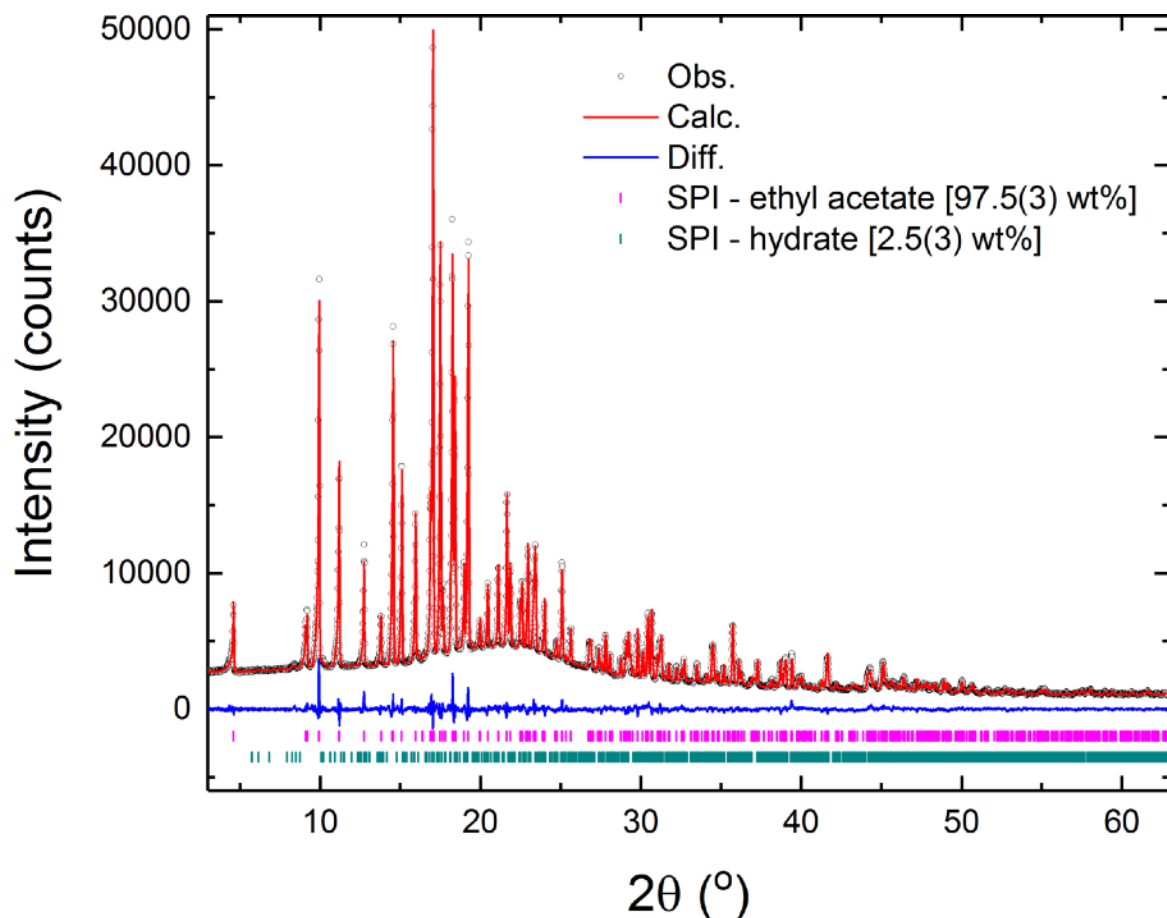
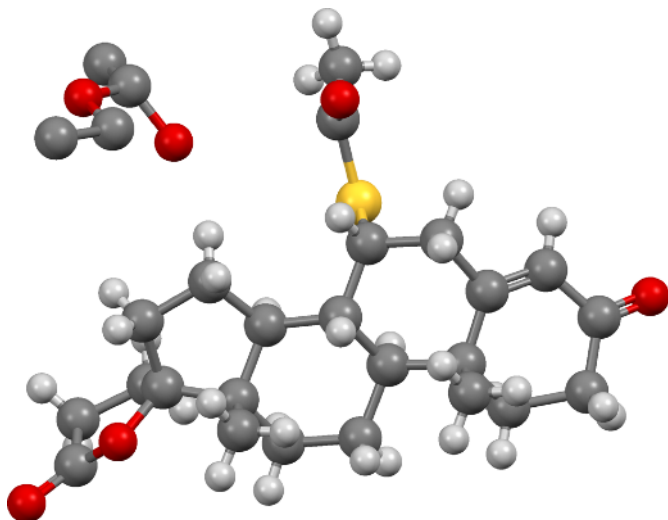
Hydrated form described by Takata [1]

WUWROW Spironolactone hydrate

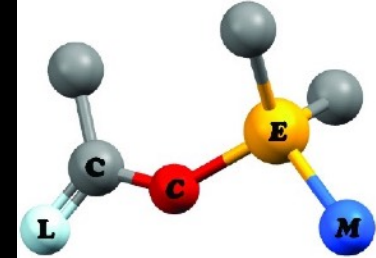
Spironolactone ethyl acetate



structure proposed in our group



Summary

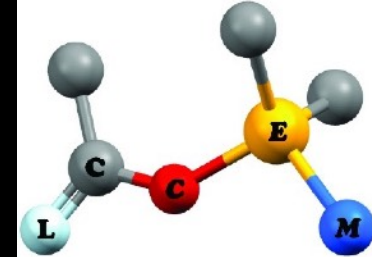


- LASSBio-1773 and LASSBio-1774: anhydrous and hydrated samples => tendency for a methylated compound to retain solvents within the crystal structure?!
- LASSBio-1755: ADPs calculation proved to be a useful approach using powder data
- XRPD is a fast tool to avoid the bottleneck provided by the need of harvesting good quality crystals
- Crystal Morphology Prediction (CMP) as a way to get valuable information regarding crystal habit
- Low cost POLYMORPH/SOLVATE SCREENING





Acknowledgements



Thank you!

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<http://nano.ufabc.edu.br>