

SUPPORTING INFORMATION

High water soluble agrichemicals by engineered organic salts for reducing adverse environmental impacts

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Table S1. Crystal data and refinement parameters of AMT and AMT salts

Compound	AMT	AMT FUM	AMT MAL-I	AMT MAL-II	(AMTMAL) ·2H ₂ O
Empirical formula	C ₁₈ H ₃₄ N ₁₀ S ₂	C ₁₃ H ₂₁ N ₅ O ₄ S	C ₃₉ H ₆₃ N ₁₅ O ₁₂ S ₃	C ₁₃ H ₂₁ N ₅ O ₄ S	C ₁₃ H ₂₅ N ₅ O ₆ S
Formula weight	454.67	342.95	343.41	343.41	379.44
Temperature/K	120	120.0	120.0	298	298
Crystal system	Monoclinic	Monoclinic	Monoclinic	Monoclinic	Triclinic
Space group	P2 ₁ /c	P2 ₁ /c	P2 ₁ /c	P2 ₁ /c	P $\bar{1}$
a/Å	12.018(4)	18.132(3)	13.495(3)	8.5300(3)	8.8250(4)
b/Å	22.054(6)	11.105(2)	16.262(3)	9.8310(3)	9.7640(5)
c/Å	9.300(3)	16.920(3)	23.781(5)	22.6570(8)	12.0800(6)
α/°	90	90	90	90	107.075(4)
β/°	101.617(5)	95.693(3)	99.226(4)	111.5740(10)	100.749(3)
γ/°	90	90	90	90	96.427(3)
Volume/Å ³	2414.5(12)	3390.1(10)	5151.4(17)	1766.87(10)	962.04(8)
Z, Z'	4, 2	4, 2	4, 3	4, 1	2, 1
ρ _{calc} /cm ³	1.251	1.344	1.328	1.291	1.310
μ/mm ⁻¹	0.247	0.218	0.215	0.209	0.206
F(000)	976.0	1452.0	2184.0	728.0	404.0
Crystal size/mm ³	0.253 x 0.076 x 0.021	0.277 x 0.12 x 0.121	0.218 x 0.096 x 0.04	0.36 x 0.30 x 0.19	0.170 x 0.160 x 0.10
2θ range for data collection/°	5.8 to 52.744	4.516 to 52.044	4.98 to 51.358	7.13 to 50.786	5.912 to 52.41
Index ranges	-15 ≤ h ≤ 15 -27 ≤ k ≤ 27 -11 ≤ l ≤ 11	-22 ≤ h ≤ 22 -13 ≤ k ≤ 13 -20 ≤ l ≤ 20	-16 ≤ h ≤ 16, -19 ≤ k ≤ 19, -28 ≤ l ≤ 28	-10 ≤ h ≤ 10 -11 ≤ k ≤ 11 -27 ≤ l ≤ 27	-10 ≤ h ≤ 10 -12 ≤ k ≤ 11 -15 ≤ l ≤ 14

Reflections collected	50224	67624	101144	6249	6996
Independent reflections	4917 [$R_{\text{int}} = 0.1074$, $R_{\text{sigma}} = 0.0515$]	6669 [$R_{\text{int}} = 0.0414$, $R_{\text{sigma}} = 0.0267$]	9760 [$R_{\text{int}} = 0.0436$, $R_{\text{sigma}} = 0.0210$]	3227 [$R_{\text{int}} = 0.0182$, $R_{\text{sigma}} = 0.0279$]	3806 [$R_{\text{int}} = 0.0776$, $R_{\text{sigma}} = 0.0876$]
Data/restraints/parameters	4917/17/292	6669/128/472	9760/26/651	3227/0/213	3806/0/249
Goodness-of-fit on F^2	1.030	1.089	1.028	1.011	1.027
Final R indexes [$I \geq 2\sigma(I)$]	$R_1 = 0.0562$, $wR_2 = 0.1369$	$R_1 = 0.0955$, $wR_2 = 0.2342$	$R_1 = 0.0545$, $wR_2 = 0.1436$	$R_1 = 0.0590$, $wR_2 = 0.1761$	$R_1 = 0.0651$, $wR_2 = 0.1754$
Final R indexes [all data]	$R_1 = 0.0944$, $wR_2 = 0.1595$	$R_1 = 0.1009$, $wR_2 = 0.2379$	$R_1 = 0.0671$, $wR_2 = 0.1531$	$R_1 = 0.0789$, $wR_2 = 0.1934$	$R_1 = 0.1077$, $wR_2 = 0.2094$
Largest diff. peak/hole / $e \text{ \AA}^{-3}$	0.64/-0.52	1.51/-0.66	1.38/-1.15	0.75/-0.28	0.24/-0.35

Table S2. The pK_a differences between AMT and the conformer (carboxylic acids). Following this approach, the salt formation is expected for both cases, even to HFUM that ΔpK_a is less than 2-3 pK_a units^[1].

Compound	pK_{a1}	$\Delta pK_a = pK_a(\text{AMT}) - pK_{a1}(\text{acid})$
Ametryn (AMT)	4.10	-
Fumaric acid (HFUM)	3.03	1.07
Maleic acid (HMAL)	1.83	2.27

Table S3. Geometric parameters of the H-bonds in the AMT and AMT salts.

Interaction	D···A(Å)	H···A(Å)	D–H···A(°)	Symmetry Code
AMT				
N2B–H2B···N3A	3.077(3)	2.249(2)	161.38(17)	x+1,+y,+z
N4A–H4A···N1B	3.033(3)	2.182(2)	170.44(18)	x-1,+y,+z
N4B–H4B···N1A	3.007(4)	2.155(3)	170.79(17)	x,y,z
N2A–H2A···N3B	3.126(4)	2.269(3)	174.92(19)	x,y,z
C9A–H9AB···N2B	3.487(5)	2.730(3)	136.28(25)	x-1,+y,+z
C9A–H9AA···N4A	3.613(5)	2.673(3)	166.40(26)	x,-y+1/2,+z-1/2
AMT FUM				
N5–H5···O1A	2.687(4)	1.810(3)	173.66(22)	x,y,z
N4A–H4A···O2A	2.771(4)	1.903(3)	168.36(22)	x,y,z
N5B–H5B···O1B	2.783(11)	1.910(10)	171.04(36)	x,y,z
N2B–H2B···O2B	2.522(18)	1.654(15)	167.84(69)	x,y,z
O4A–H4AE···O1A	2.534(4)	1.696(3)	176.20(22)	-x+1,+y-1/2,-z+1/2+1
N2A–H2A···O3A	2.833(5)	2.025(4)	152.22(25)	-x+1,+y+1/2,-z+1/2+1
N4B–H4B···O3B	2.842(5)	2.059(4)	147.80(26)	-x+2,+y-1/2,-z+1/2
O4B–H4BE···O1B	2.748(11)	1.914(15)	171.90(55)	-x+2,+y+1/2,-z+1/2
C8B–H8BC···O3B	3.419(7)	2.677(4)	132.71(33)	-x+2,+y-1/2,-z+1/2
C11A–H11A···O4A	3.125(5)	2.496(3)	123.75(24)	-x+1,+y+1/2,-z+1/2+1
C4B–H4BD···S1A	3.856(21)	2.920(1)	140.28(50)	x,-y+1/2+1,+z+1/2
AMT MAL-I				
N3C–H3C···O2C	2.660(3)	1.810(2)	161.66(15)	x,y,z
N1B–H1B···O2B	2.687(3)	1.830(2)	163.94(15)	x,y,z
N4C–H1C···O1C	2.928(3)	2.053(2)	172.72(15)	x,y,z
N2B–H2B···O1B	2.841(3)	1.963(2)	174.92(16)	x,y,z
N4A–H4A···O1A	2.875(3)	1.999(2)	173.54(16)	-x,-y+1/2,-z+1/2
N3A–H3A···O2A	2.666(3)	1.797(2)	168.69(16)	-x,-y+1/2,-z+1/2
N2C–H2C···O4C	2.830(3)	1.952(2)	175.29(16)	x-1,+y,+z
N4B–H4B···O4B	2.921(3)	2.050(2)	170.08(16)	x+1,+y,+z
N2A–H2A···O4A	2.886(3)	2.015(2)	169.82(16)	-x+1,+y+1/2,-z+1/2

C7A–H7AB···O4B	3.346(4)	2.546(2)	138.83(23)	x,y,z
C5A–H5AE···O2B	3.447(9)	2.629(2)	141.11(49)	x,y,z
C9C–H9CB···O2B	3.341(3)	2.516(2)	141.75(17)	-x+1,-y+1,-z+1
C9C–H9CC···O2C	3.441(3)	2.497(2)	161.73(16)	-x+1,-y+1,-z+1
C9A–H9AA···O2B	3.413(3)	2.441(2)	171.24(16)	-x+1,+y-1/2,-z+1/2
C9A–H9AC···O2C	3.346(3)	2.538(2)	139.70(17)	-x+1,+y-1/2,-z+1/2
C9B–H9BA···O2A	3.453(3)	2.525(2)	158.04(17)	x+1,+y+1,+z
C7B–H7BB···O3A	3.553(12)	2.603(2)	161.42(50)	x+1,-y-1/2,+z+1/2
C7B–H7BC···O3C	3.312(9)	2.682(2)	122.42(50)	-x+2,-y+1,-z+1
C9A–H9AC···S1C	3.736(3)	2.941(1)	138.91(16)	-x+1,+y-1/2,-z+1/2
AMT MAL-II				
N1–H1···O2	2.696(3)	1.842(2)	171.87(15)	x,y,z
N2–H2···O1	2.855(4)	1.999(3)	173.49(17)	x,y,z
N4–H4···O3	3.008(3)	2.196(3)	159.29(17)	x,-y+1/2+1,+z+1/2
C9–H9A···O4	3.424(4)	2.488(2)	164.90(20)	-x+2,-y+2-1,-z+1
C9–H9B···O1	3.381(4)	2.589(2)	139.99(21)	-x+2,-y+1,-z+1
C11–H11···O3	3.490(6)	2.759(4)	136.22(22)	-x+2,-y+1,-z+1
(AMTMAL)-2H₂O				
N3–H3···O2	2.701(2)	1.846(2)	172.27(13)	x,y,z
N4–H4···O1	2.870(3)	2.011(2)	176.67(16)	x,y,z
N2–H2···OW1	3.011(4)	2.187(3)	160.47(18)	-x,-y+1,-z
OW1–HW1B···OW2B	2.687(19)	1.965(19)	142.20(57)	x,y,z
OW2B–HW2A···O1	2.756(26)	2.003(2)	147.16(80)	-x+1,-y+1,-z+1
C12–H12···O3	3.417(4)	2.489(2)	175.39(20)	-x+2,-y+2,-z+2
C8–H8B···O3	3.594(3)	2.709(3)	153.42(13)	x-1,+y-1,+z
C9–H9B···O4	3.372(5)	2.552(3)	143.44(25)	-x+1,-y+2,-z+1

COMPLEMENTARY FIGURES AND TABLES

FIG. S1. ASYMMETRIC UNIT OF (A) AMT ($P2_1/c$, $Z' = 2$), (B) AMT-FUM ($P2_1/c$, $Z' = 2$), (C) AMT-MAL-I ($P2_1/c$, $Z' = 3$), (D) AMT-MAL-II ($P2_1/c$, $Z' = 1$) AND (E) (AMT-MAL)·H₂O ($P\bar{1}$, $Z' = 1$) WITH LABELLING THE SYMMETRY-INDEPENDENT MOLECULES. THE ELLIPSOID PLOT ARE DRAWN AT 50% PROBABILITY LEVEL. THE LOWEST OCCUPANCY PARTS OF DISORDERED FRAGMENTS ARE SHOWN WITH 50% TRANSPARENCY FOR CLARITY.

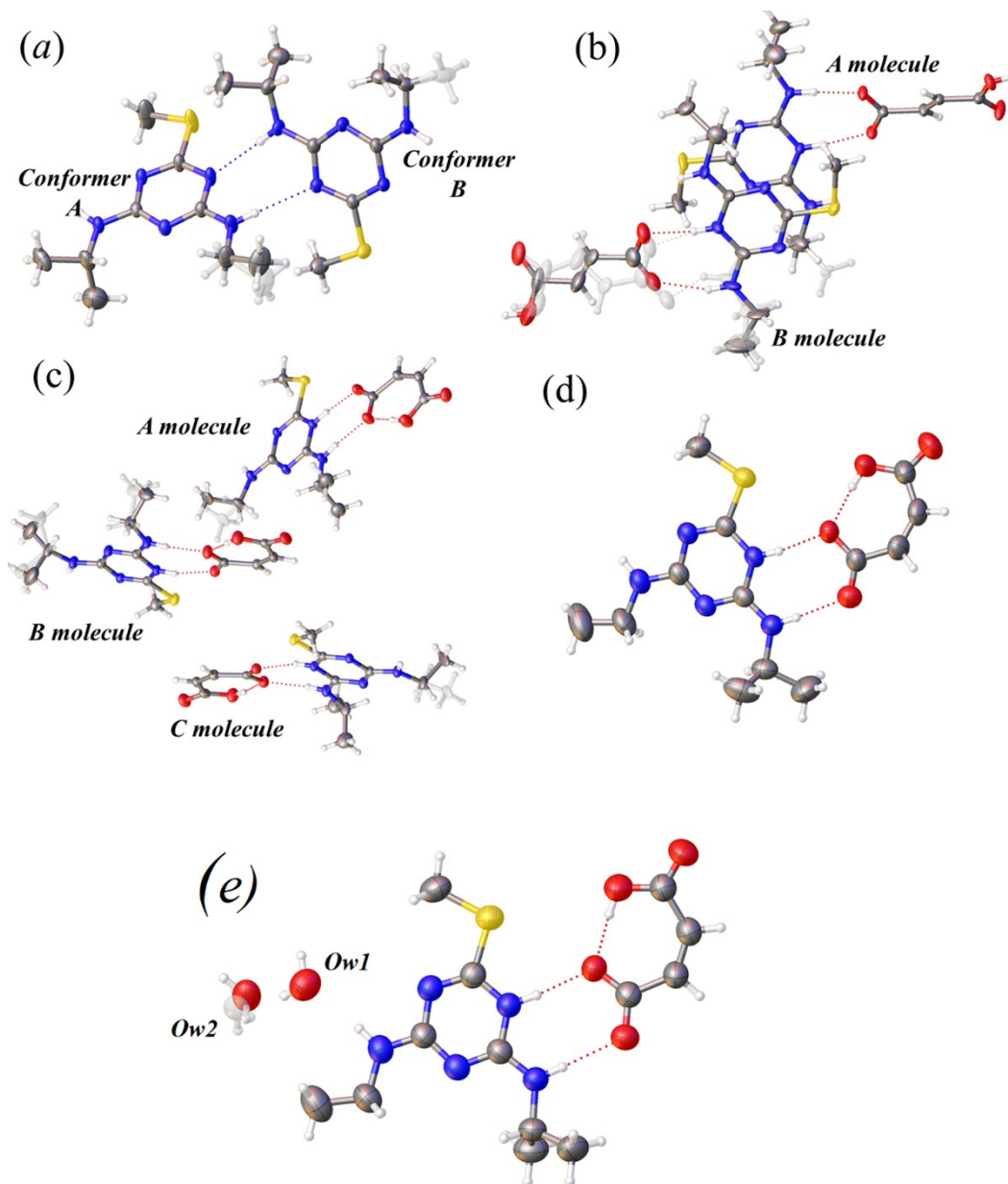


FIG. S2: ALONG THE [100] DIRECTION, ADJACENT AMT CHAINS STACKS PARALLEL TO EACH OTHER VIA $\text{CH}\cdots\pi$ (LIGHT GREEN DASH LINE) AND $\text{S}\cdots\pi$ (ORANGE DASH LINE) INTERACTIONS RESULTING IN DOMAINS OF LAYERED STRUCTURE.

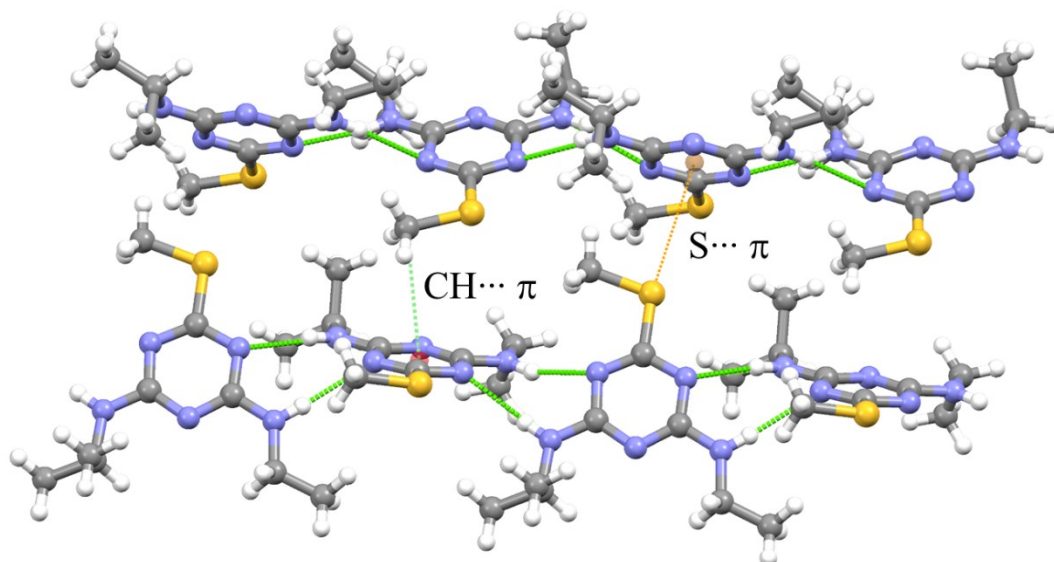


Fig. S3. In the unit cell of AMT ($Z' = 2$), the A and B conformers are alternately arranged along the [100] direction, such that, the independent molecules are offset each other by 55.34° . In this figure, the red plane defines to that formed by the main plane of conformer A while the blue one is define by conformer B. The molecular overlay of highest occupancy parts of A and B conformers from AMT superimposed through s-triazine groups.

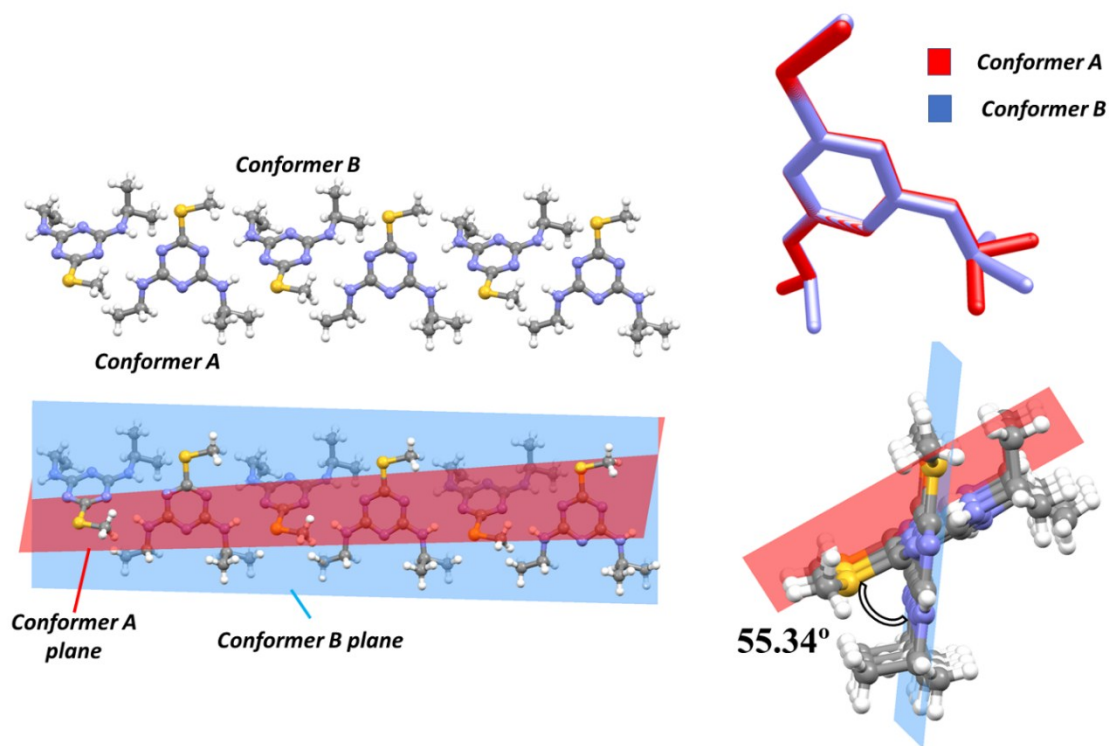


FIG. S4: FOURIER DIFFERENCE MAP FOR THE (A) MOLECULE A AND B FROM AMT-FUM SALT. IT SEEN THE PROTONATION OF AMT MOLECULE ON THE N5-ATOM FOR BOTH CASES. (B) MOLECULAR CONFORMATIONS OF AMT CATIONS (A AND B MOLECULES) IN AMT-FUM SALT. HIGHEST OCCUPANCY PARTS CONSIDERED IN THE PRESENTATION.

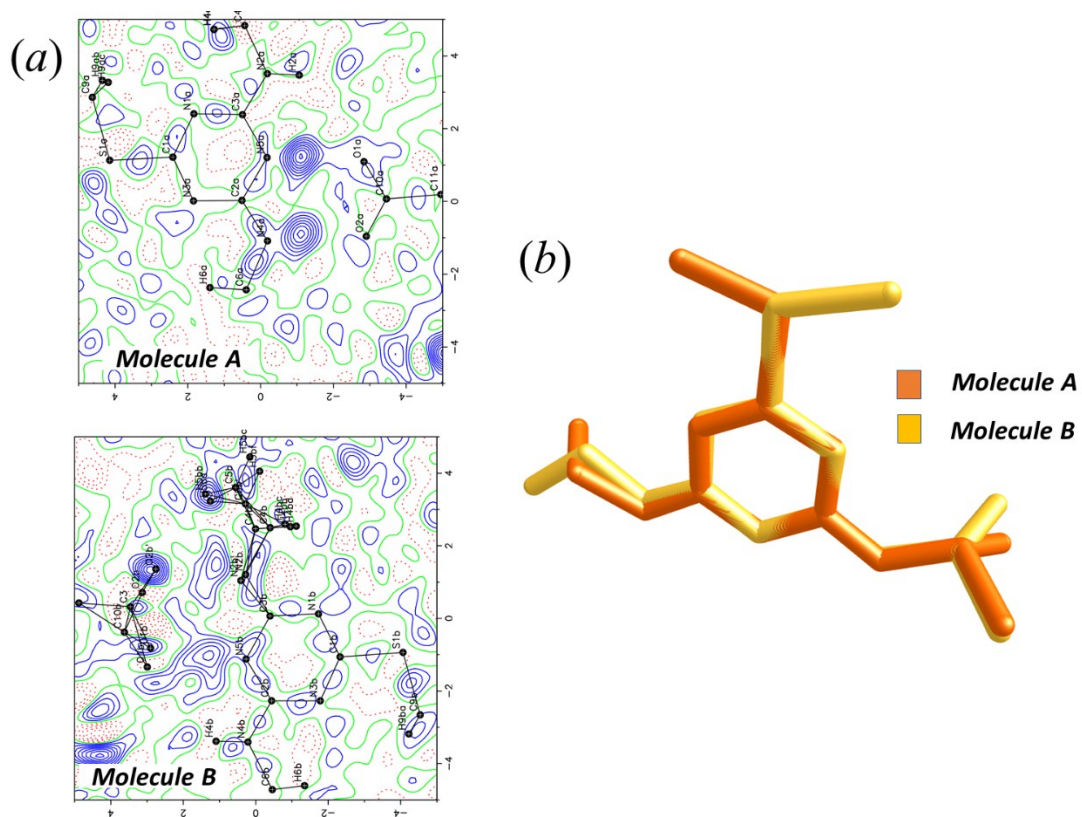


Fig. S5 - (a) Fourier difference maps of both A and B molecules of AMT-MAL-I showing the presence of an H-atom attached to N5 atom ones for both cases. (b) Overlay of the AMT⁺ conformers of AMT-MAL-I using the s-triazine groups as the common reference.

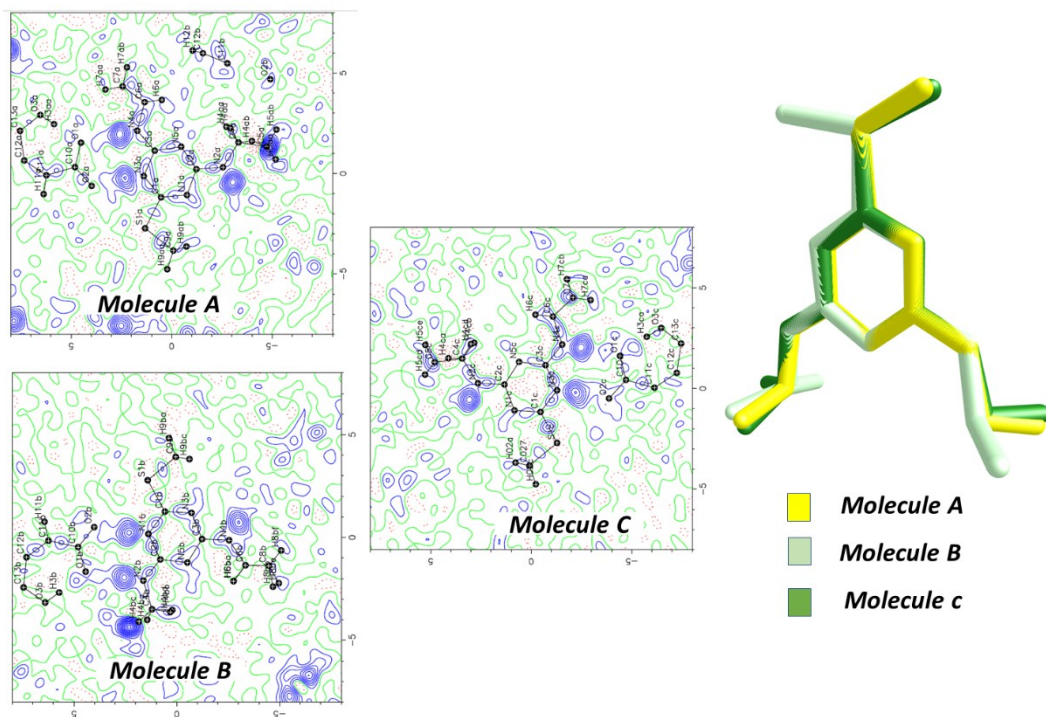


Fig. S6 - Beside the conformational and ionization multiplicity of AMT^+ , the polymorphism of ametryn maleate is also associated to diversity of cation-anion assemblies. (a) The ionic pair in AMT-MAL-I interact that way carboxylic of acid and the sulfur-methyl groups of AMT are cis-oriented. Opposite situation are found in AMT-MAL-II. (c) $\pi \cdots \pi$ interaction between AMT^+ molecules in AMT-MAL-I.

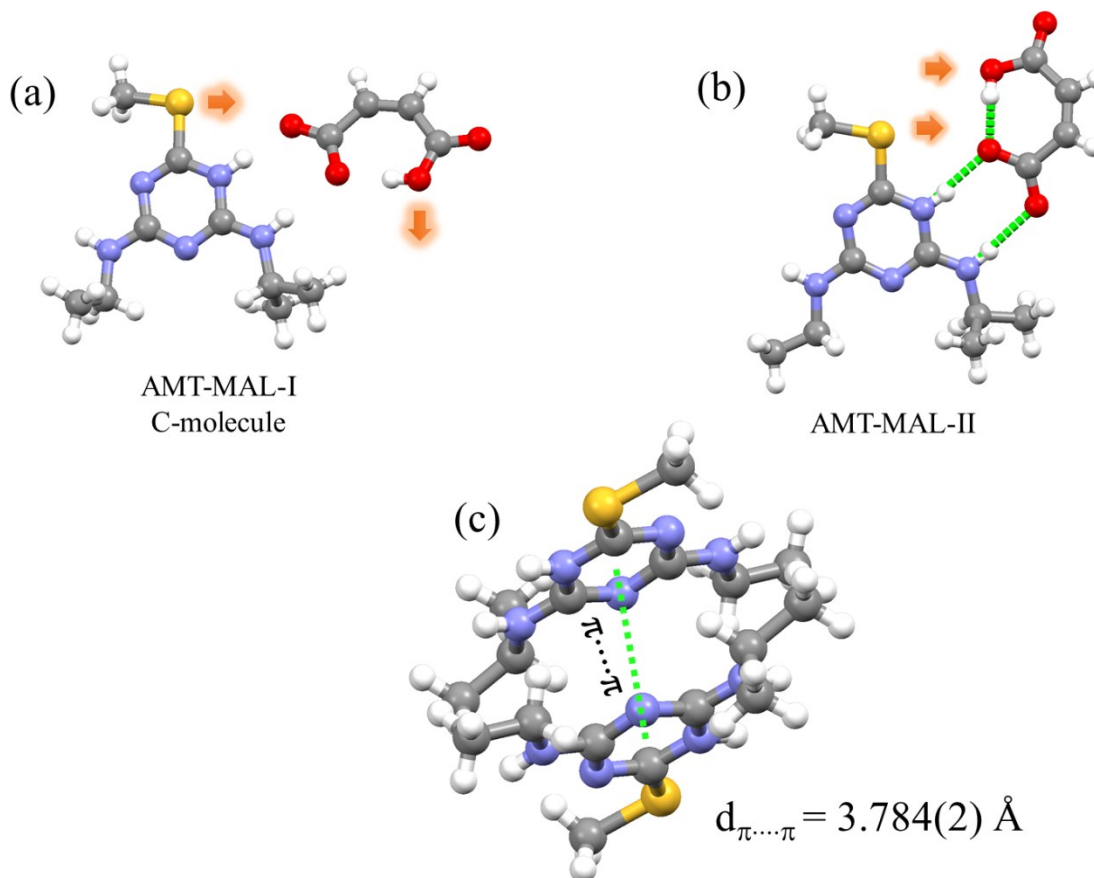


Fig. S7 - (a) The packing of (AMT-MAL) \cdot 2H $_2$ O illustrating the water channels (blue circles - viewed along [100] direction). (b) The (H $_2$ O) $_2$ clusters featuring in the crystal structure (water channel core). (c) π \cdots π Interactions between AMT $^+$ \cdots H $_2$ O assemblies.

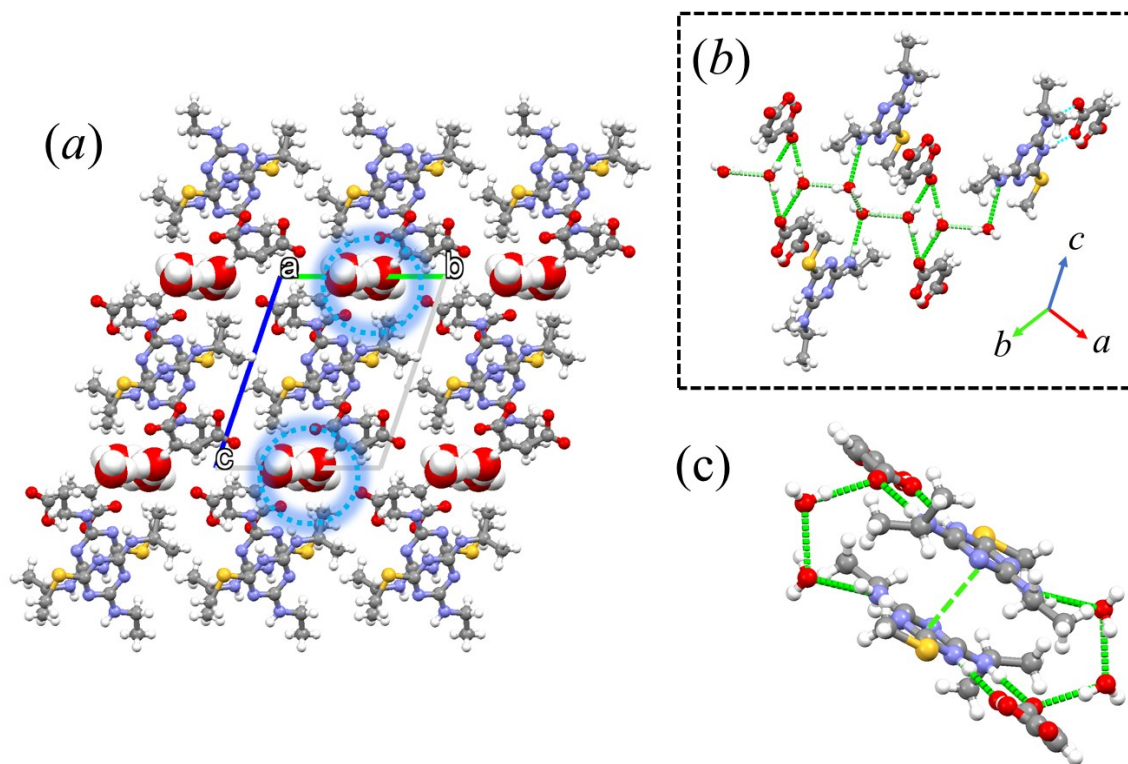


Figure S8 - Diffractograms of AMT and its salts. For all compounds, the experimental and calculated PXRD patterns show an excellent agreement which indicates the purity (unique phase) of the samples.

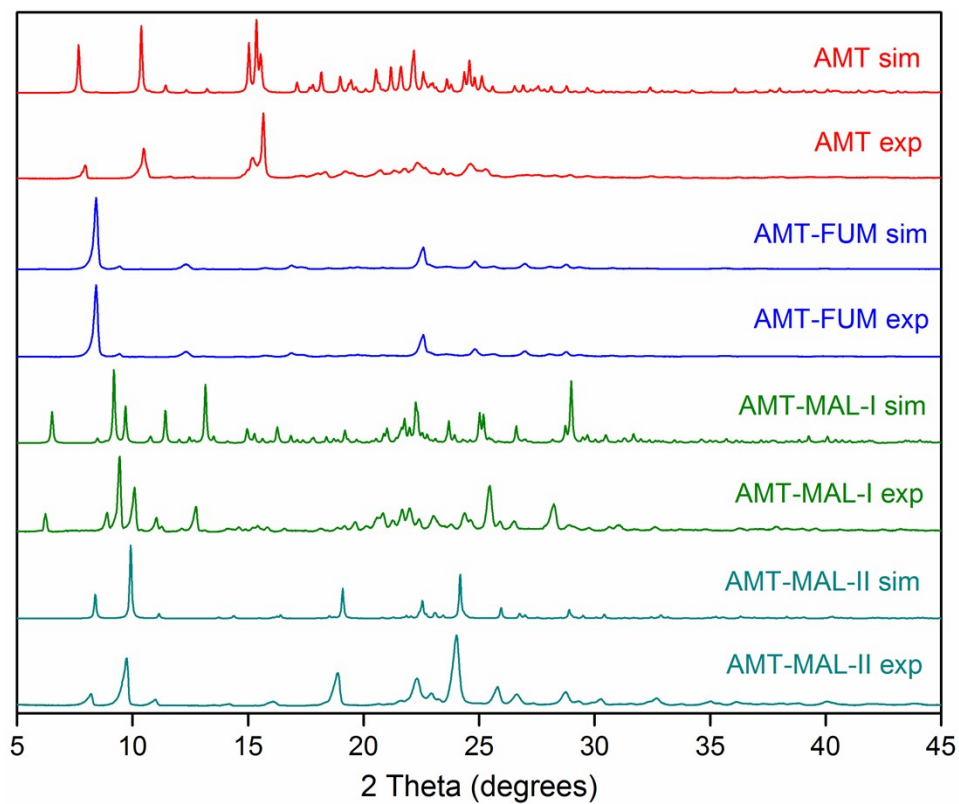


Figure S10 - Aqueous solubility (blue bars) and melting point (line) of AMT and its salts.

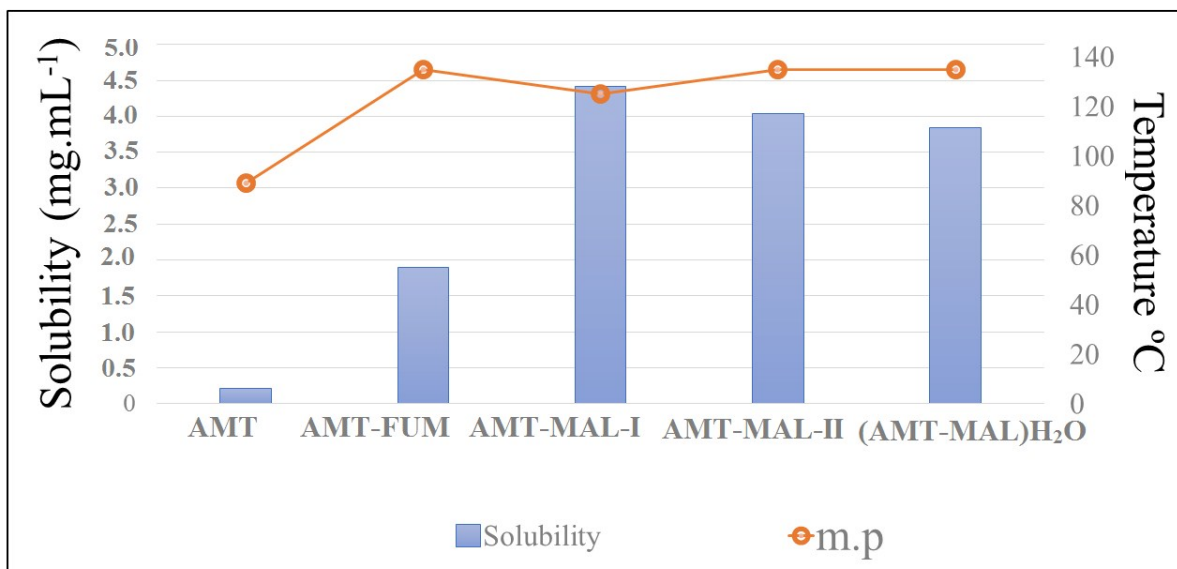


Table S3 - Energy differences between the AMT conformer from AMT and AMT-FUM

Ametryn	
	$E + E_{ZPVE}$, kcal. mol ⁻¹
A	-643115,0062
B	-643118,0213
Ametryn fumarate	
A*	-643424,1252
B*	-643426,9791
Ametryn Fumarate Deprotonated	
A†	-643108,0278
B†	-643108,8987
$[E(A^\dagger) + E(A^\dagger)_{ZPVE}] - [E(A) + E(A)_{ZPVE}]$	-6,9784275
$[E(B^\dagger) + E(B^\dagger)_{ZPVE}] - [E(B) + E(B)_{ZPVE}]$	-9,122595

Figure S11. Herbicide 4.98 to 51.358 application system



Similarly, applications of AMT salts were tested. These application solutions prepared dissolving an amounts of AMT-FUM and AMT-MAL-I salts equivalent to the to 1.0 kg ha⁻¹ dose in water in order to simulate the field application of 400 L ha⁻¹. After stirring, 150 rpm during 5 min, an aliquot of the solution was collected to determine the soluble concentration of AMT in solution before spraying. The GIW of these systems were compared with the control treatment, the 0 dose of AMT. All experiments were carried out in triplicate.

After 14 days of herbicide application, the plants were cut and dried in a forced-air oven at 70 °C for 72h to determine the dry matter content (DM). The herbicidal efficacy of AMT (at different doses), AMT-FUM and AMT-MAL-I were estimated through of the growth inhibition of *Brachiaria decumbens* after herbicide application. The Growth Inhibition index (ν) was calculated using the equation:

$$\nu = 100 - [(DM_x - DM_i)/(DM_{control} - DM_i)] \times 100 \quad \text{Equation 1}$$

where DM_x is the dry matter content of each sample, DM_i is the dry matter average content before AMT application, and $DM_{control}$ is the dry matter content of control treatment (0 dose of AMT).

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