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CRYSTAL ENGINEERING
MATERIALS

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Supporting information for article:

**Intermolecular Interaction Energies and Molecular Conformations
in *N*-Substituted 4-Aryl-2-methylimidazoles with Promising *in vitro*
Antifungal Activity**

**Nerith-Rocio Elejalde, Estefanía Butassi, Susana Zacchino, Mario A. Macías
and Jaime Portilla**

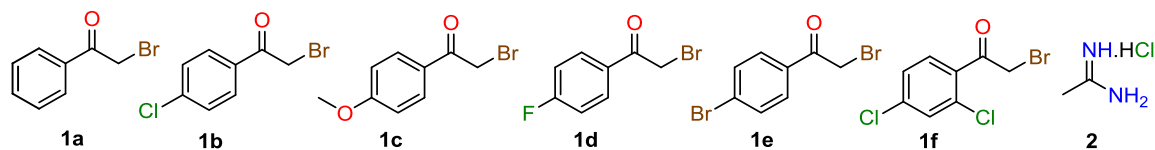
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1. Overview of reactants, substrates and products numbering
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S1. Overview of reactants, substrates and products numbering

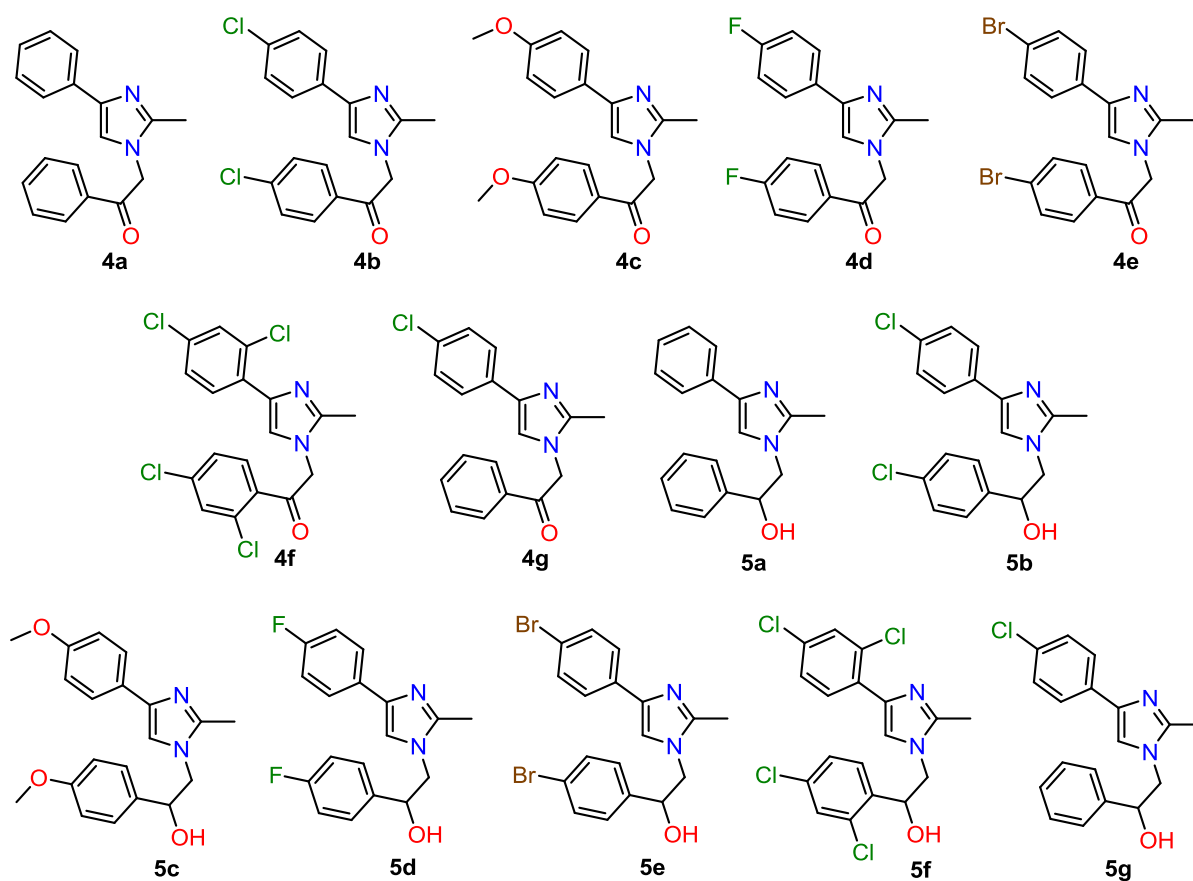
Reactants and substrates

α -Bromoacetophenones **1a–f** and acetamidine hydrochloride **2**



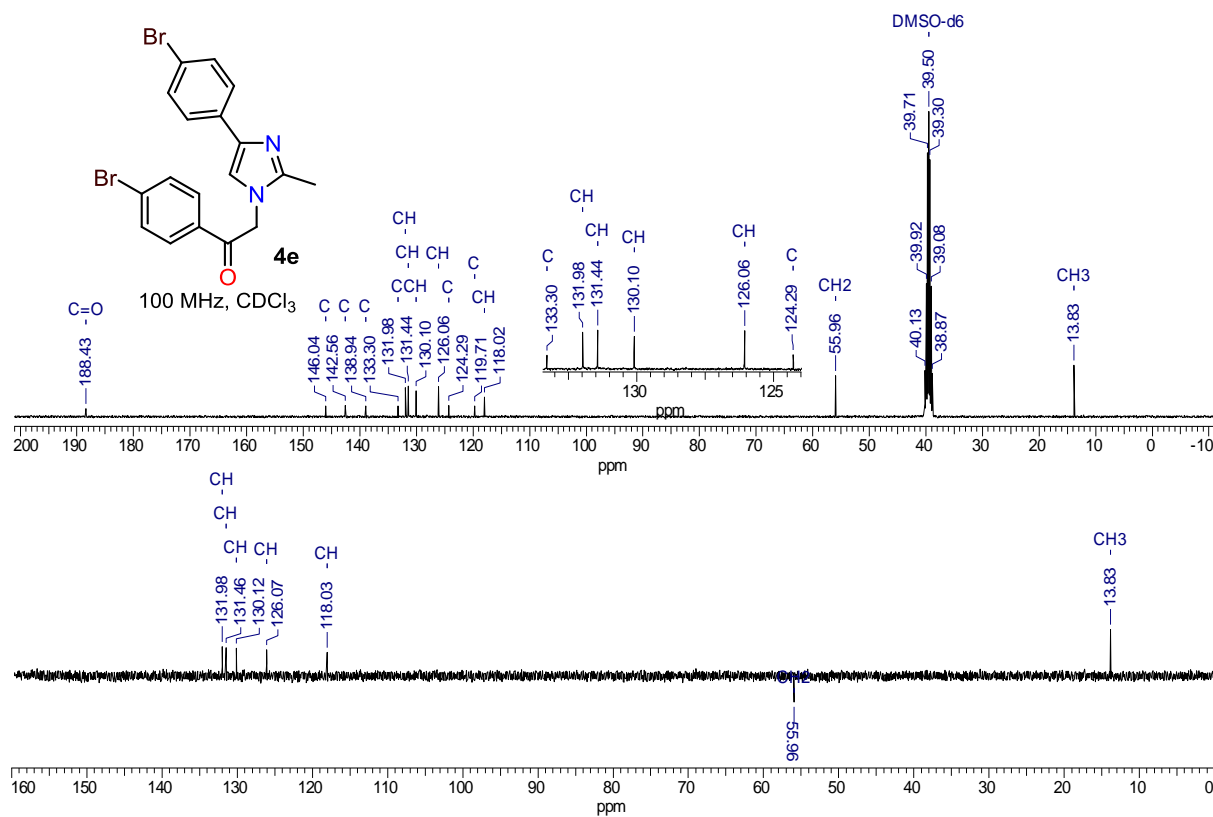
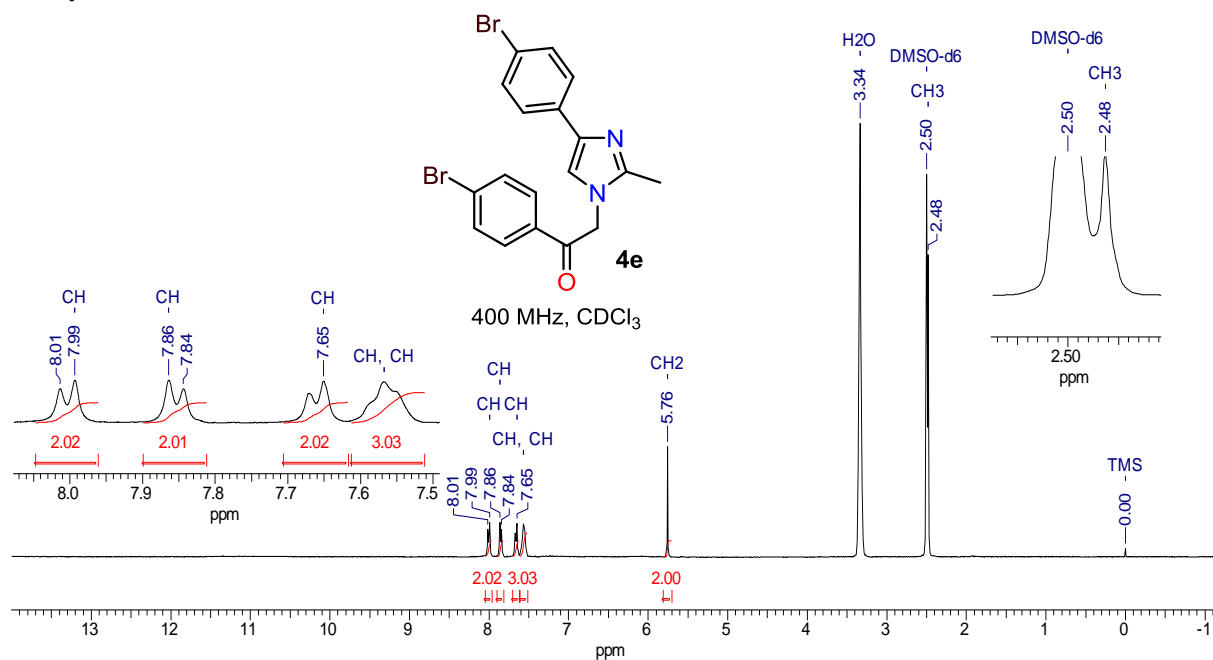
Products

N*-Phenacyl-2-methylimidazoles **4a–g** and *N*-(2-hydroxyethyl)-2-methylimidazoles **5a–g*

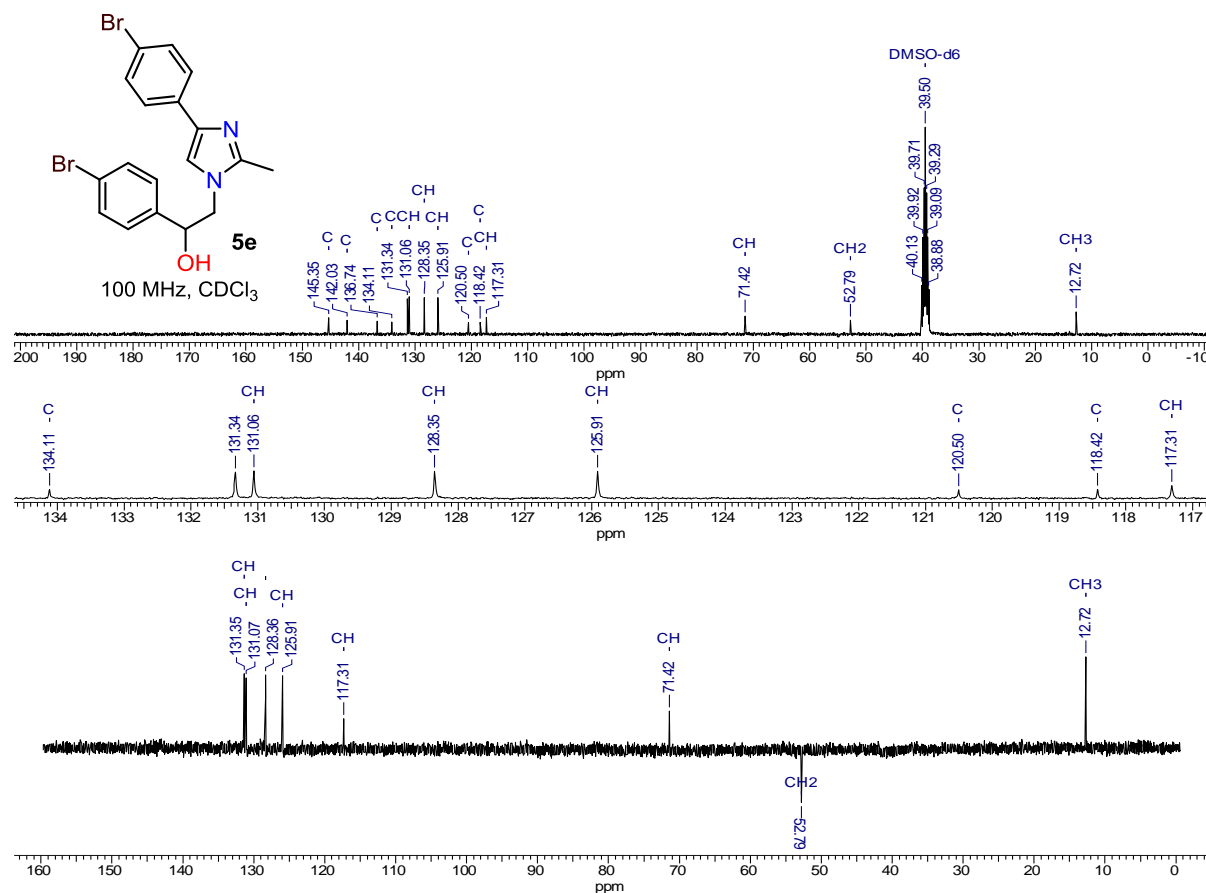
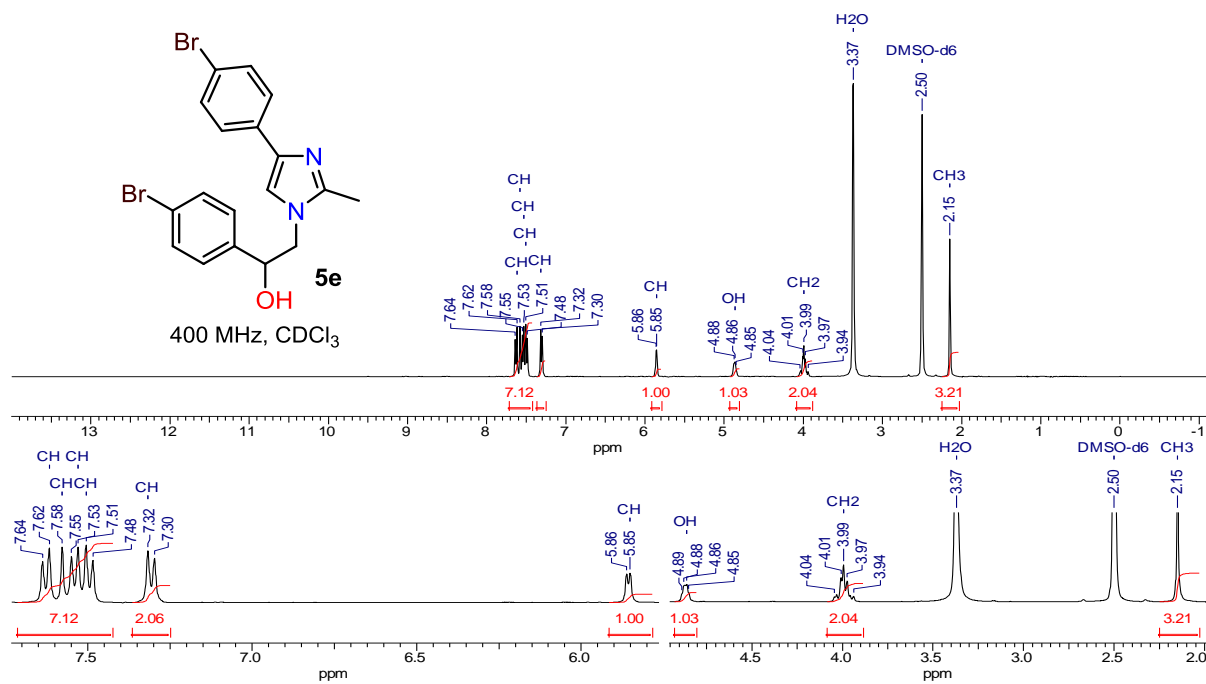


S2. Copies of NMR spectra of novel compounds 4e, 4f, 5e, and 5f

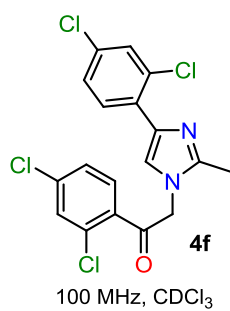
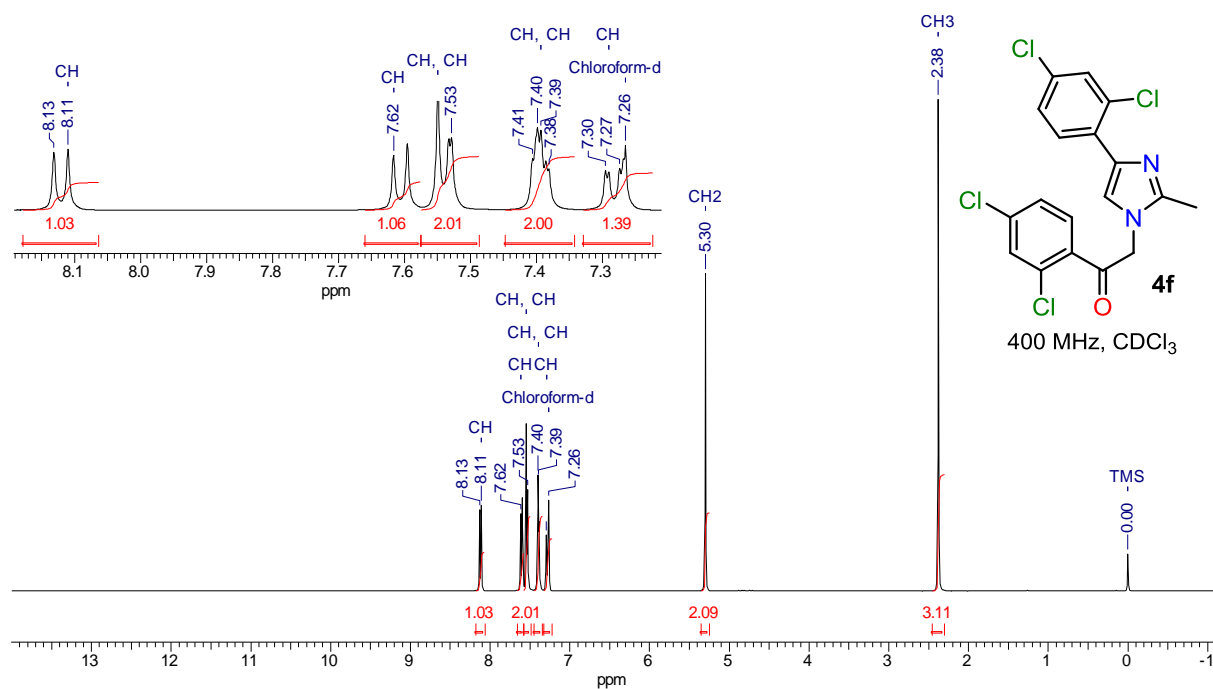
¹H, ¹³C{¹H} NMR and DEPT-135 of 4-(2,4-dichlorophenyl)-1-(2-(2,4-dichlorophenyl)-2-ethanone)-2-methyl-1*H*-imidazole (**4e**)

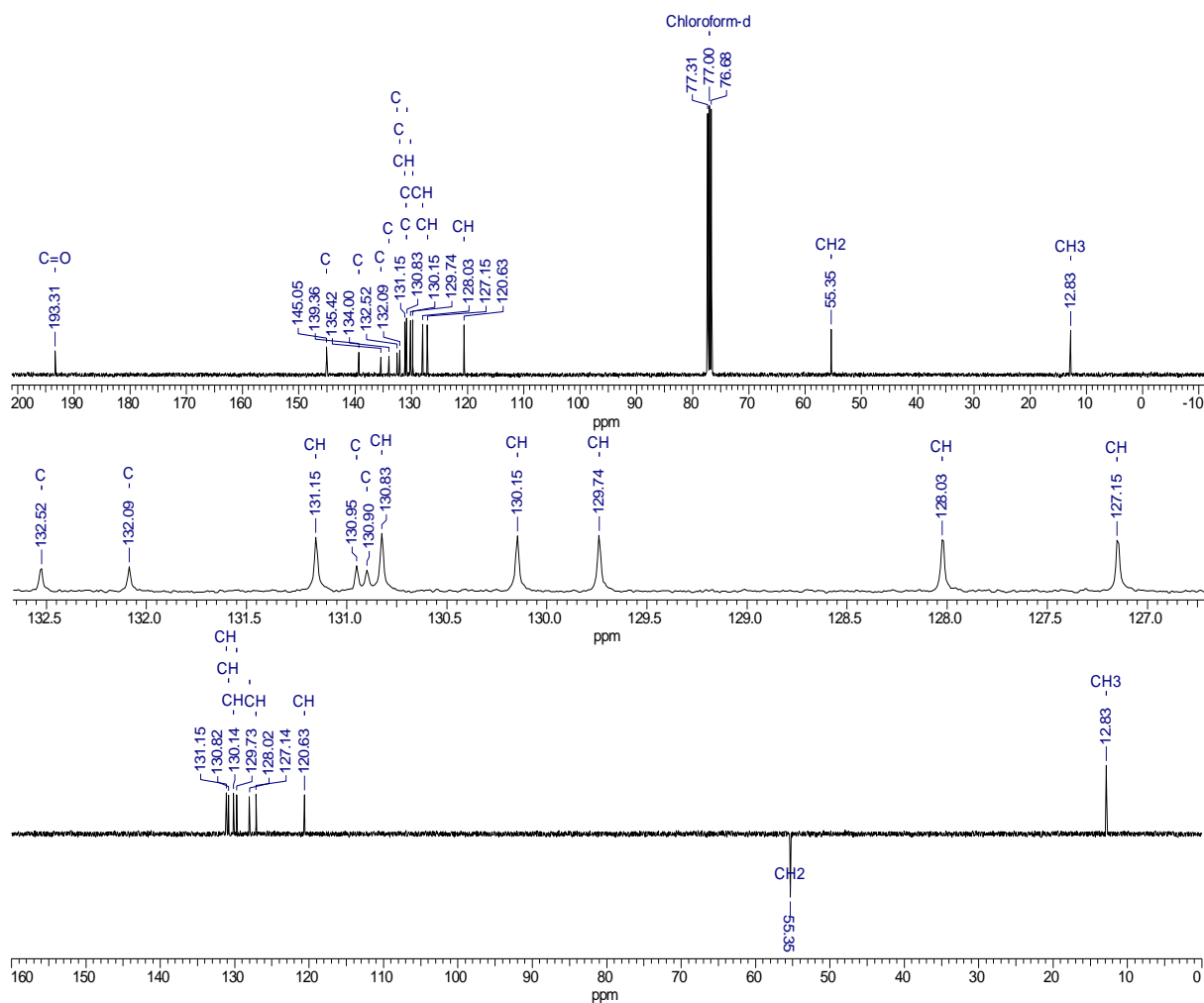


^1H , $^{13}\text{C}\{^1\text{H}\}$ NMR and DEPT-135 of 1-(4-bromophenyl)-2-(4-(4-bromophenyl)-2-methyl-1H-imidazol-1-yl)ethan-1-ol (**5e**)

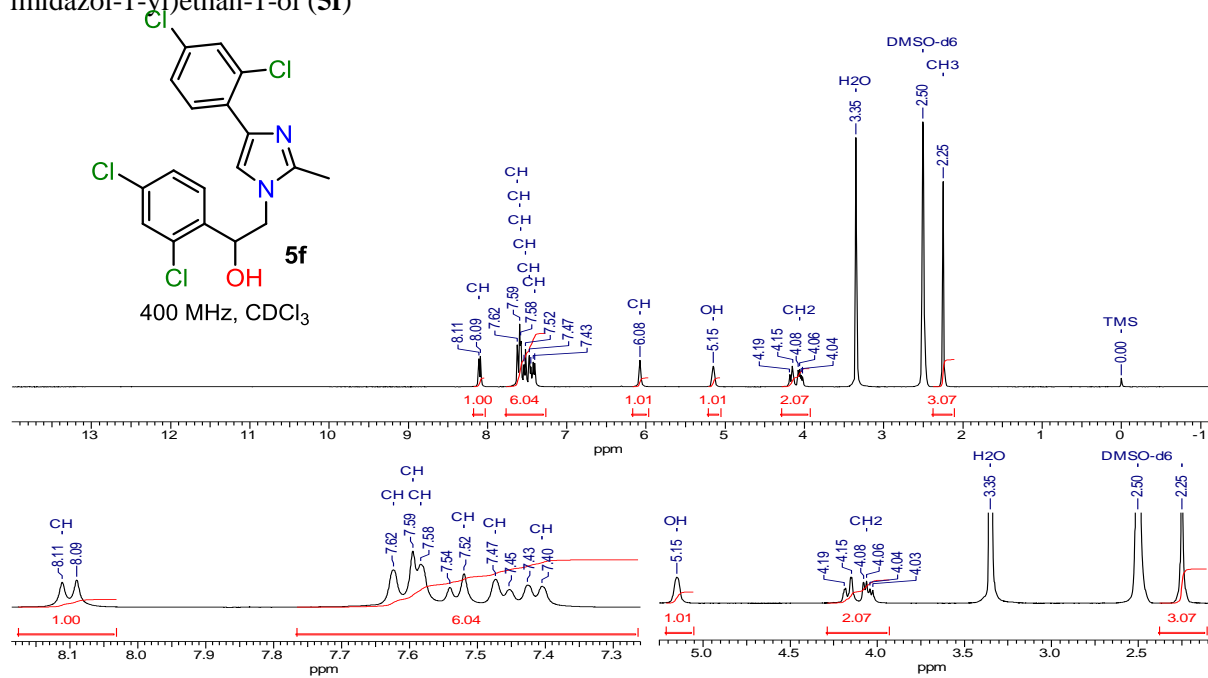


^1H , $^{13}\text{C}\{^1\text{H}\}$ NMR and DEPT-135 of 4-(2,4-dichlorophenyl)-1-(2-(2,4-dichlorophenyl)-2-ethanone)-2-methyl-1*H*-imidazole (**4f**)





^1H , $^{13}\text{C}\{^1\text{H}\}$ NMR and DEPT-135 of 1-(2,4-dichlorophenyl)-2-(4-(2,4-dichlorophenyl)-2-methyl-1*H*-imidazol-1-yl)ethan-1-ol (**5f**)



S3. HRMS analysis data of novel compounds 4e, 4f, 5e, and 5f

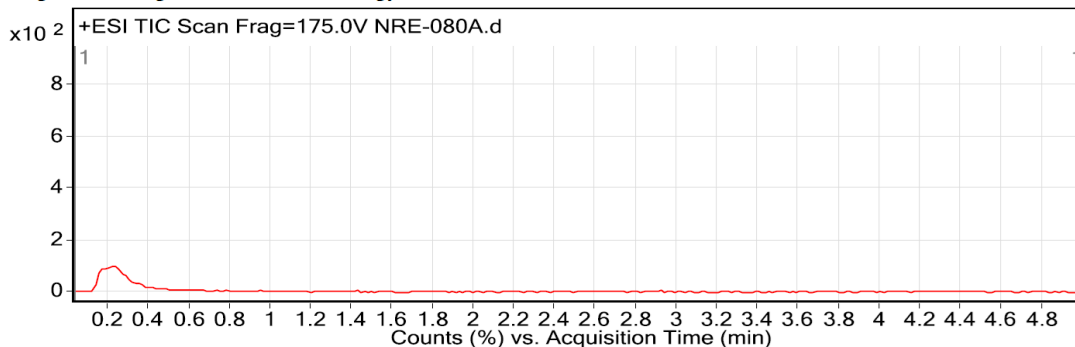
Qualitative Analysis Report

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IRM Calibration Status	Success	DA Method	Metodo-analisis-signaltonoise.m
Comment			

Sample Group Info.

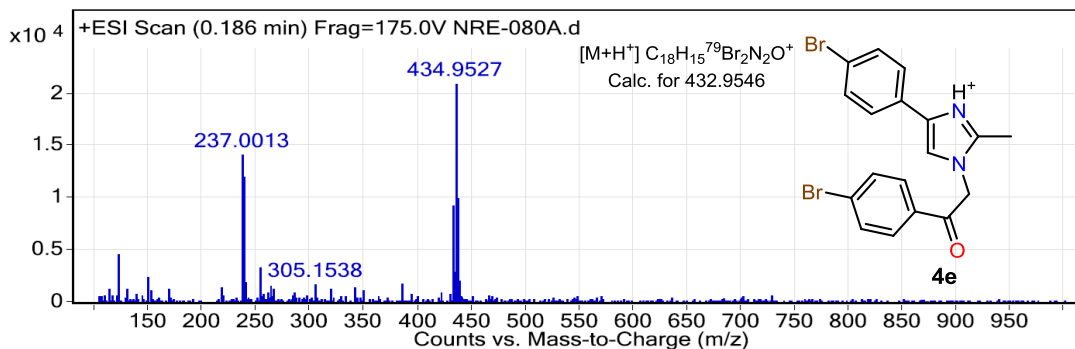
User Chromatograms

Fragmentor Voltage 175 Collision Energy 0 Ionization Mode ESI



User Spectra

Fragmentor Voltage 175 Collision Energy 0 Ionization Mode ESI



Peak List

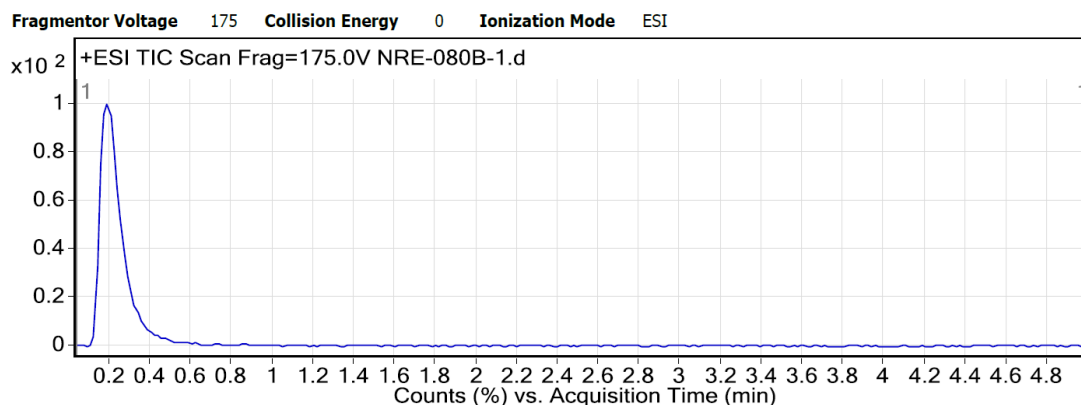
m/z	z	Abund
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237.0013	1	14186.1
237.0855		3029.3
238.9999	1	12102.9
253.0632		3368
432.9549	1	9334.9
433.9619	1	2957.5
434.9527	1	21008.3
435.9558	1	3716.5
436.9526	1	10015.2

Qualitative Analysis Report

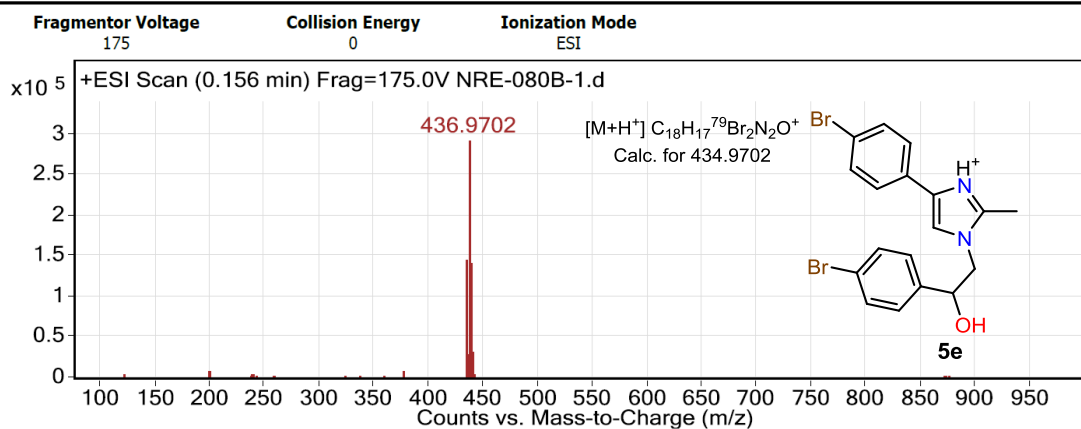
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Comment			

Sample Group **Info.**

User Chromatograms



User Spectra



Peak List

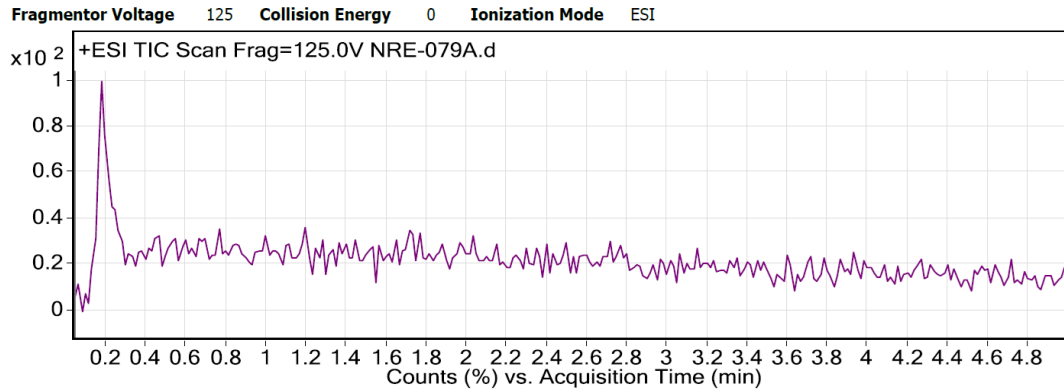
m/z	z	Abund
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435.9732	1	29615.1
436.9702	1	293207.7
437.0262		23875.3
437.0782		16226.3
437.1635		16520.1
437.9738	1	59808
438.968	1	142594.7
439.9696	1	32863.7

Qualitative Analysis Report

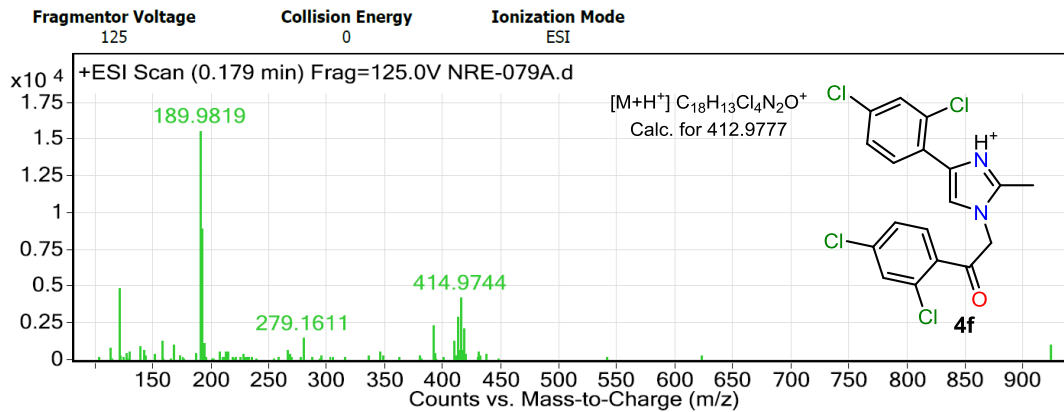
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Comment			

Sample Group **Info.**

User Chromatograms



User Spectra



Peak List

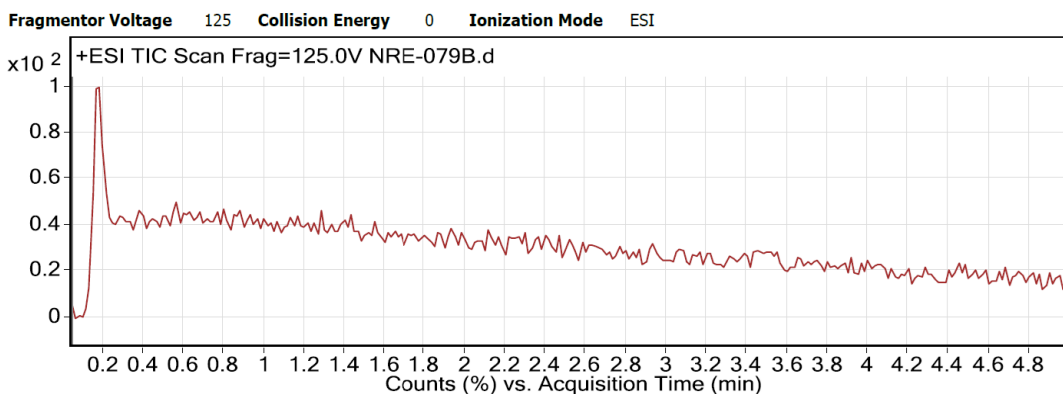
<i>m/z</i>	<i>z</i>	Abund
121.0859		4921.1
189.9819	1	15650.8
190.0214		1728.5
190.9866	1	1523.3
191.9796	1	9018.4
279.1611		1544.2
391.2846		2473.6
412.9789		3032
414.9744		4302.8
416.9715		2270.3

Qualitative Analysis Report

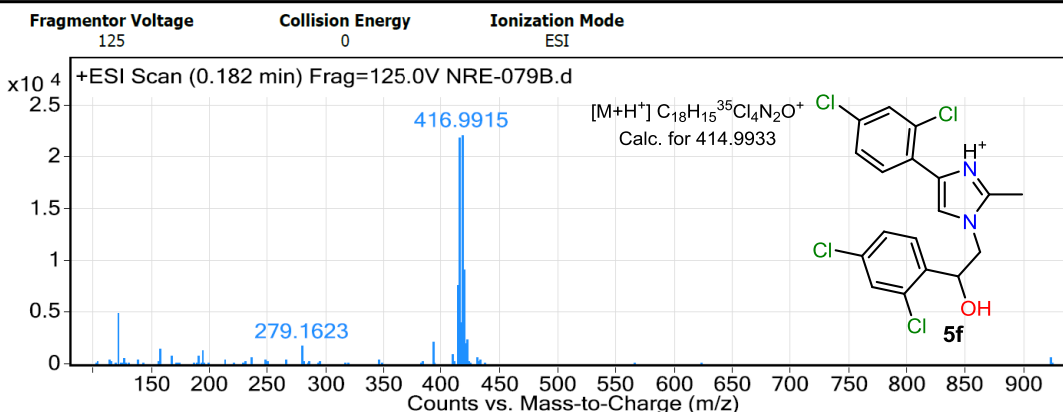
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Comment			

Sample Group **Info.**

User Chromatograms

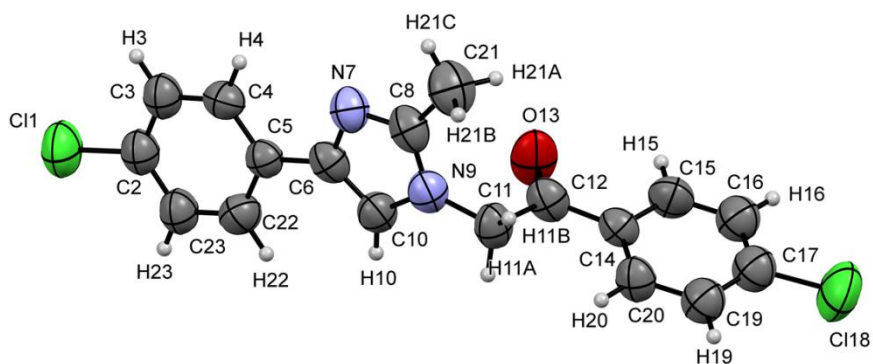


User Spectra

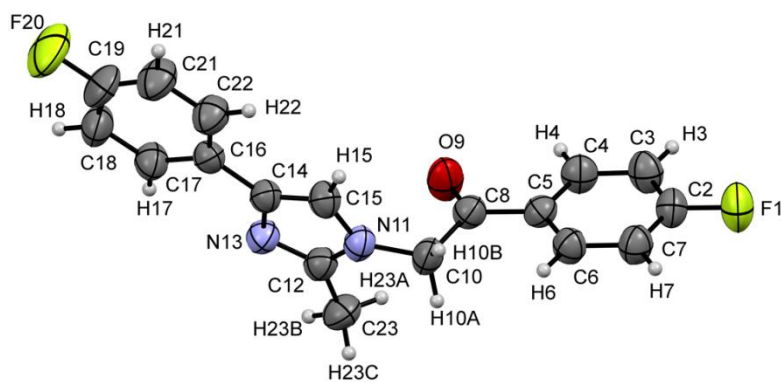


Peak List

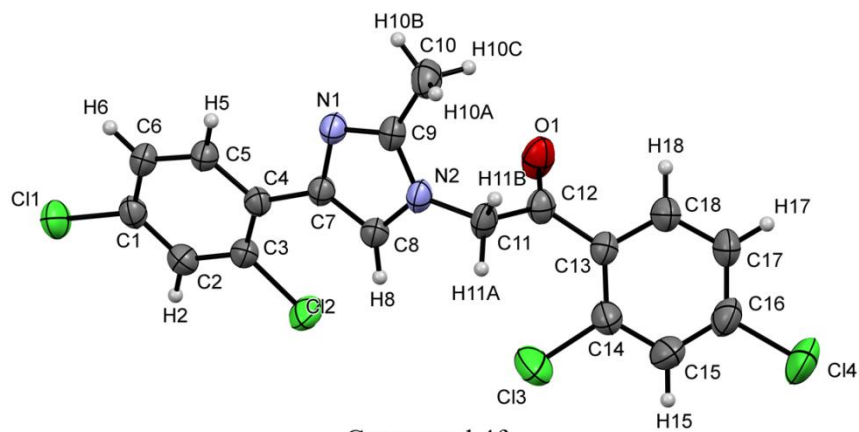
m/z	z	Abund
121.0865		5023.8
391.2915		2287.6
412.9802	1	7694.8
414.9919	1	21886.9
415.993	1	4057
416.9915	1	22239.7
417.0477		4107.8
417.9935	1	4059.2
418.99	1	9201.4
420.9894	1	2553.3

S4. ORTEP plots of compounds 4b, 4d, and 4f^a

Compound 4b



Compound 4d

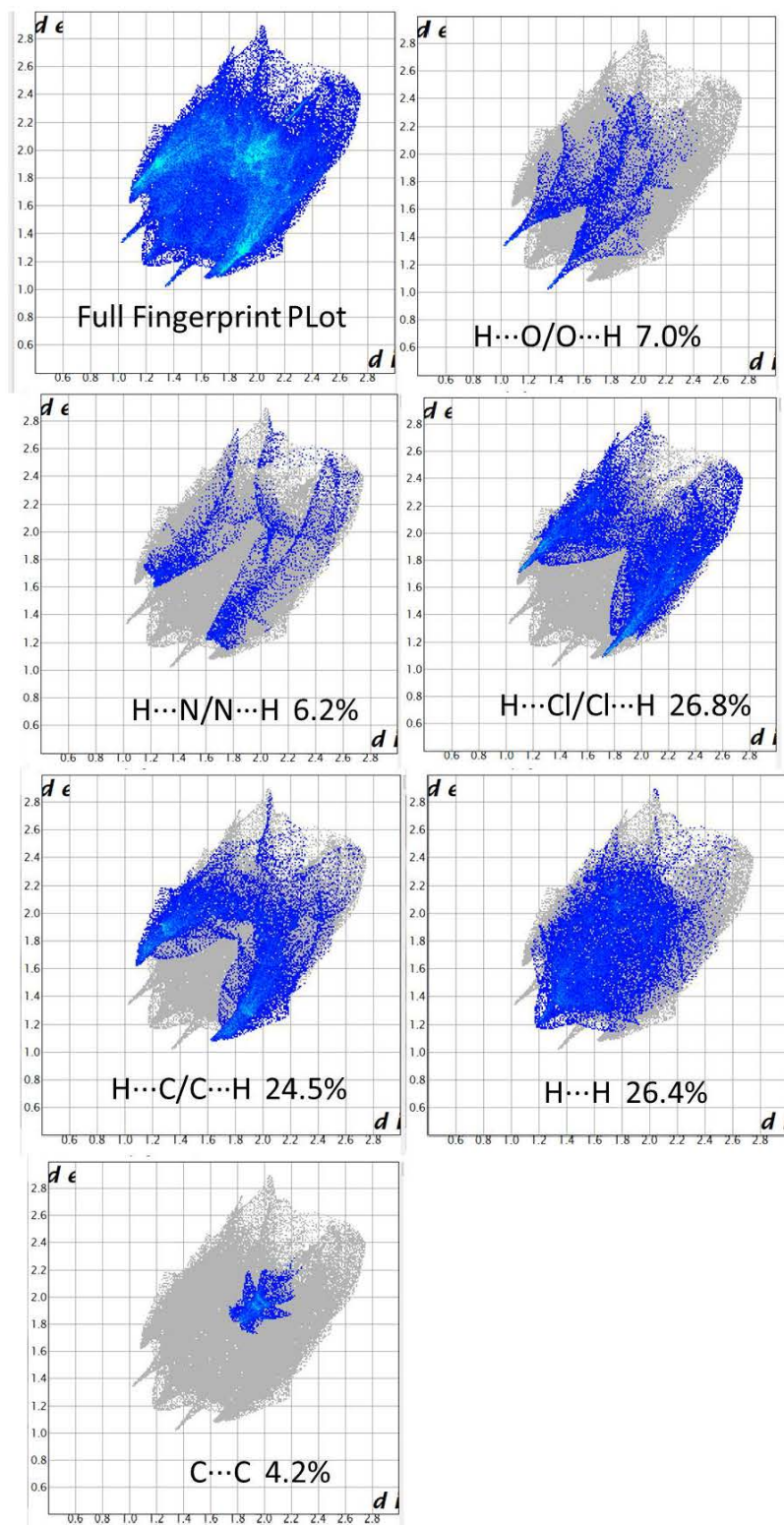


Compound 4f

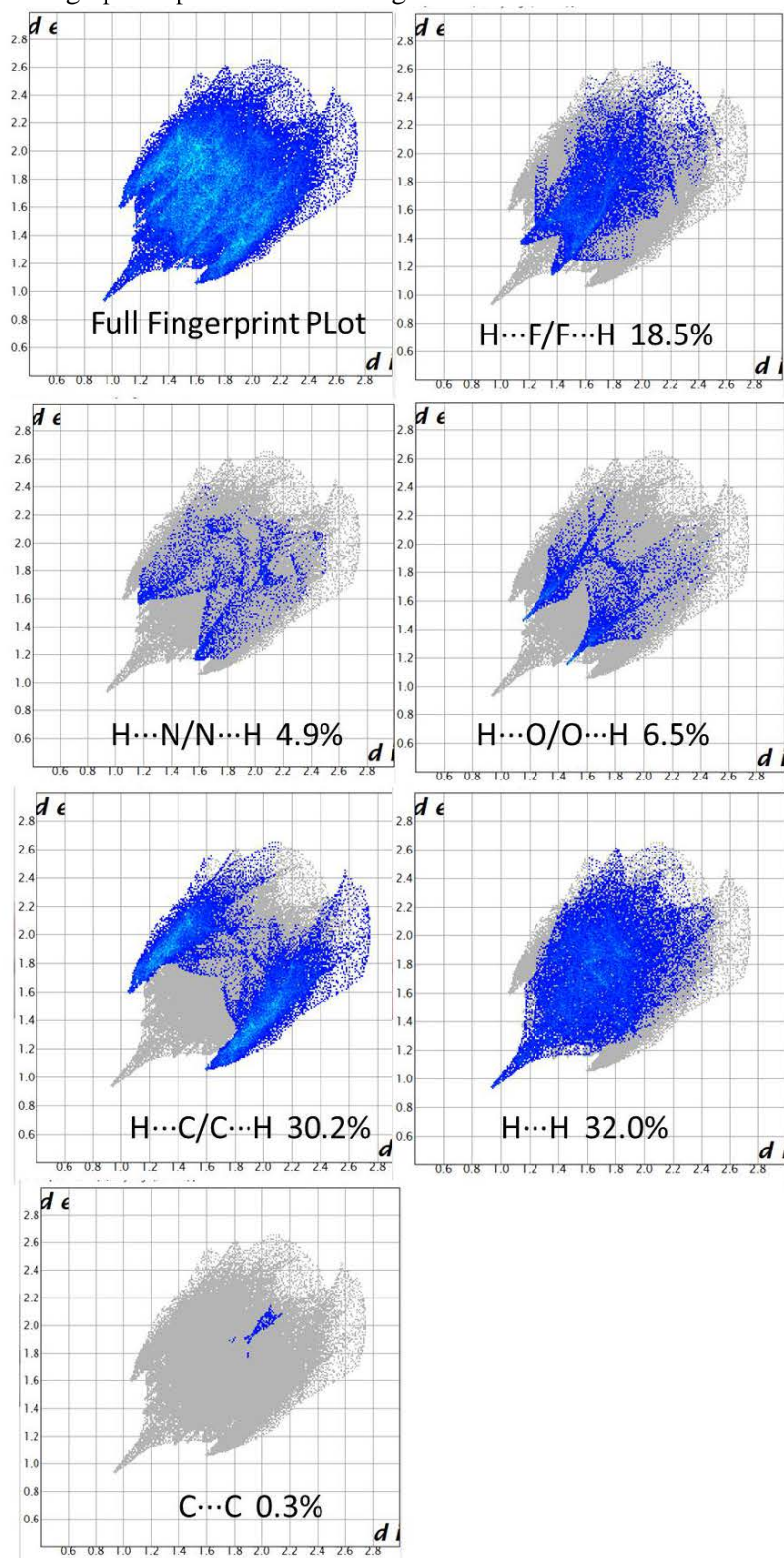
^a Showing the numbering scheme and displacement ellipsoids drawn at the 50% probability level

S5. Two-dimensional fingerprints plots of 4b, 4d, and 4f^a

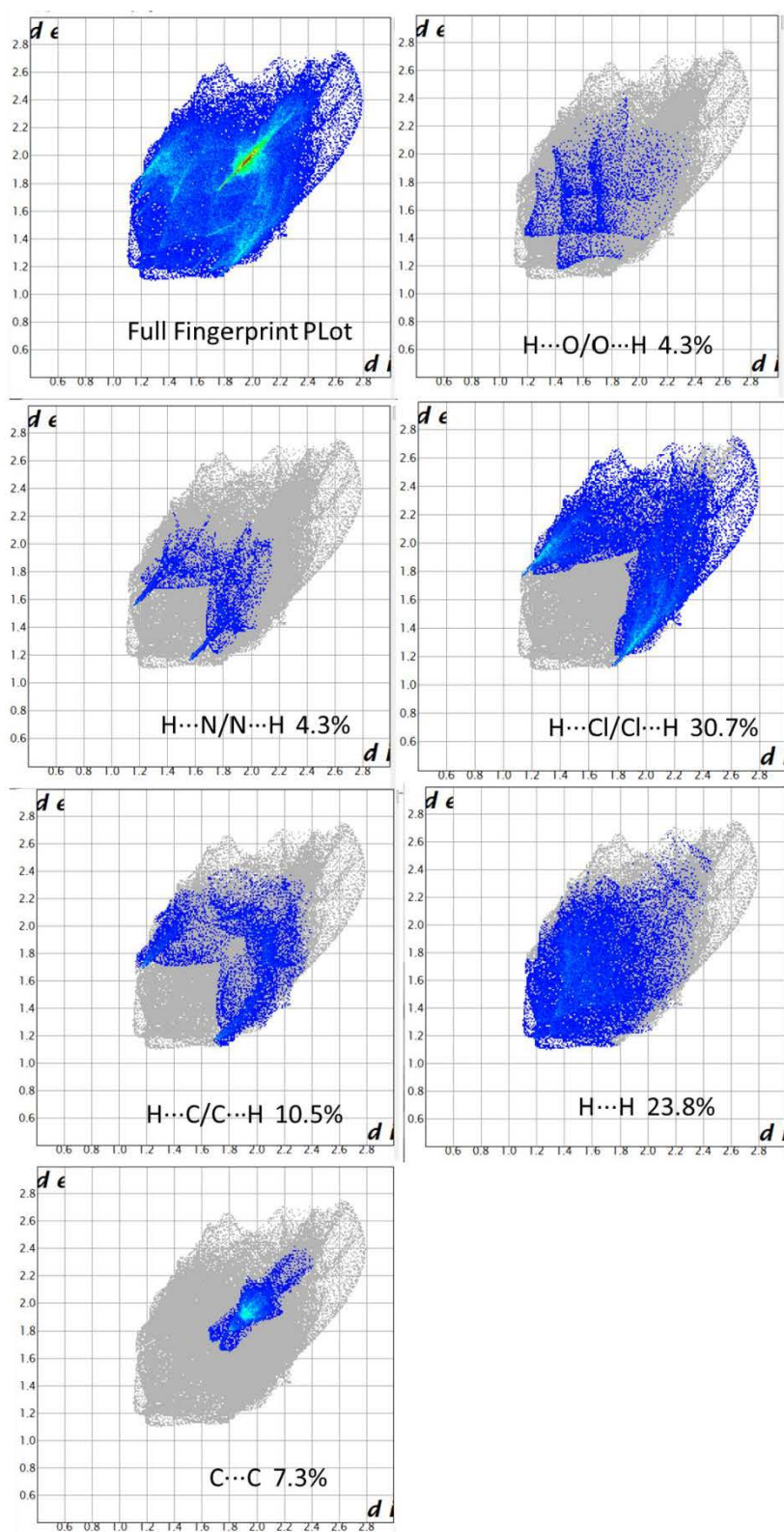
^a Two-dimensional fingerprints plots of **4b** showing contributions from different contacts



^a Two-dimensional fingerprints plots of **4d** showing contributions from different contacts



^a Two-dimensional fingerprints plots of **4f** showing contributions from different contacts



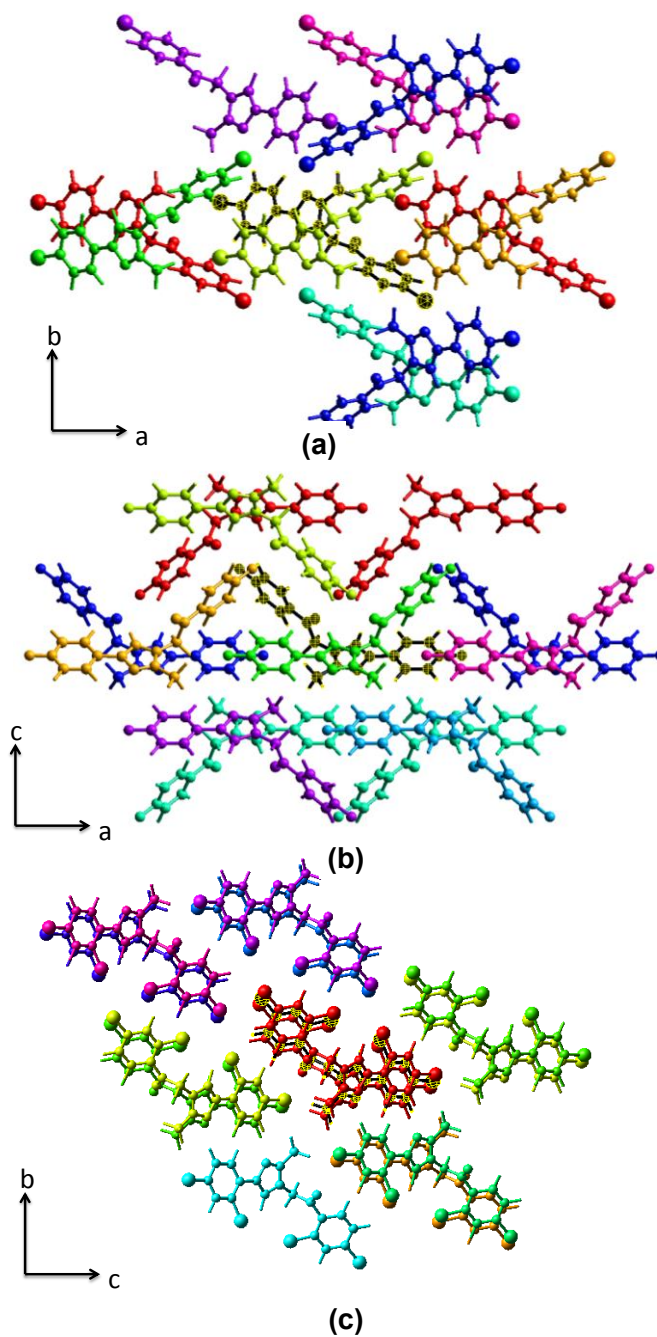
S6. CE-B3LYP interaction energies (kJ mol⁻¹) for 4b, 4d and 4f^a

4b							
<i>z</i>	<i>N</i>	<i>R</i>	<i>E_{ele}</i>	<i>E_{pol}</i>	<i>E_{dis}</i>	<i>E_{rep}</i>	<i>E_{tot}</i>
2	12.48	-5.9	-0.9	-19.8	13.6	-15.8	
2	15.45	-3.9	-0.5	-8.1	6.5	-7.6	
2	4.86	-22.3	-8.2	-60.8	36.1	-60.4	
2	10.95	-4.2	-1.9	-21.2	13.4	-16.0	
1	9.51	-4.5	-2.3	-40.6	22.3	-28.1	
2	9.66	-3.0	-0.6	-10.7	4.8	-10.0	
2	12.54	-0.1	-0.6	-7.0	4.2	-4.0	
1	11.01	-9.5	-2.9	-23.3	15.9	-22.7	
1	10.75	-1.6	-0.1	-2.0	0.1	-3.5	
4d							
<i>z</i>	<i>N</i>	<i>R</i>	<i>E_{ele}</i>	<i>E_{pol}</i>	<i>E_{dis}</i>	<i>E_{rep}</i>	<i>E_{tot}</i>
2	8.63	-7.0	-1.8	-23.3	10.7	-22.4	
2	12.46	-2.4	-0.9	-12.4	9.4	-8.2	
2	9.58	2.5	-1.7	-14.6	5.2	-8.1	
2	4.79	-22.8	-7.0	-50.6	28.3	-55.9	
2	9.36	-3.9	-1.9	-20.0	11.2	-16.0	
1	10.85	-10.6	-3.4	-24.2	27.7	-17.7	
2	11.98	-7.8	-1.5	-19.0	11.4	-18.9	
1	10.31	-2.0	-0.2	-5.1	1.3	-6.0	
2	13.34	0.9	-0.5	-7.8	1.5	-5.3	
4f							
<i>z</i>	<i>N</i>	<i>R</i>	<i>E_{ele}</i>	<i>E_{pol}</i>	<i>E_{dis}</i>	<i>E_{rep}</i>	<i>E_{tot}</i>
2	4.49	-17.3	-6.4	-97.0	55.0	-73.5	
1	11.96	-18.2	-3.3	-28.1	29.4	-27.9	
2	12.63	-7.6	-1.2	-20.5	14.6	-17.7	
2	10.97	-3.5	-1.9	-18.1	11.0	-14.0	
1	10.49	-4.2	-1.4	-27.3	11.6	-22.1	
1	11.53	-2.1	-0.3	-5.3	1.6	-6.1	
1	9.55	-5.7	-0.5	-20.4	16.0	-14.3	
1	16.65	-2.9	-0.4	-8.7	8.1	-6.0	
1	11.75	-1.2	-0.9	-13.4	10.3	-7.3	
1	18.52	0.0	-0.0	-3.6	3.6	-0.9	

^a *N* is the number of molecules with an *R* molecular centroid-to-centroid distance (Å) with an assigned *C* color-coded related to Figure 8. Electron density was calculated using B3LYP/6-31G(d,p) model energies. Note: (*) scale factors used to determine *E_{tot}*: *E_{ele}* = 1.057; *E_{pol}* = 0.740; *E_{dis}* = 0.871;

$$E_{rep} = 0.618.$$

^aColored representation of the centroid-to-centroid interactions observed in Table 6 for (a) **4b**, (b) **4d** and (c) **4f**. Molecules in black and marked in the center of each figure correspond to the asymmetric unit and reference for the interactions.



S7. Hydrogen-bond geometry (Å, °) for 4b, 4d, and 4f

Table S1 Table S1. Selected hydrogen-bonds for 4b, 4d, and 4f

Hydrogen-bond geometry (Å, °) for (4b)				
<i>D</i> — <i>H</i> ... <i>A</i>	<i>D</i> — <i>H</i>	<i>H</i> ... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> — <i>H</i> ... <i>A</i>
C4—H4...N7	0.93	2.57	2.903(5)	101.0
C11—H11B...O13 ⁱ	0.97	2.50	3.462(5)	170.1
C23—H23...Cg ₁ ⁱⁱ	0.93	3.00	3.931(5)	174.5
	Cg...Cg	Alpha*	Gamma*	mpd*
Cg ₁ ...Cg ₁ ⁱⁱⁱ	3.863(3)	0.0(2)	18.7	-3.6583(19)

Symmetry code: (i) $x, -y+3/2, z+1/2$; (ii) $-1+x, y, -1+z$; (iii) $2-x, 1-y, 2-z$
 *Alpha: dihedral angle between planes I and J (°)
 *mpd: perpendicular distance of Cg(I) on ring J (Å)
 *Gamma: angle Cg(I)→Cg(J) vector and normal to plane J (°)
 Cg₁ is the centroid of C14/C20 ring

Hydrogen-bond geometry (Å, °) for (4d)				
<i>D</i> — <i>H</i> ... <i>A</i>	<i>D</i> — <i>H</i>	<i>H</i> ... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> — <i>H</i> ... <i>A</i>
C17—H17...N13	0.93	2.57	2.903(5)	101.0
C3—H3...F1 ⁱ	0.93	2.59	3.212(5)	124.8
C6—H6...Cg ₁ ⁱⁱ	0.93	2.94	3.717(3)	66.0
C23—H23C...Cg ₁ ⁱⁱⁱ	0.96	2.94	3.713(4)	66.0

Symmetry code: (i) $1/2 - x, -1/2 + y, z$; (ii) $3/2 - x, -1/2 + y, z$; (iii) $-1/2 + x, 1/2 - y, 1 - z$
 Cg₁ is the centroid of C16/C22 ring

Hydrogen-bond geometry (Å, °) for (4f)				
<i>D</i> — <i>H</i> ... <i>A</i>	<i>D</i> — <i>H</i>	<i>H</i> ... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> — <i>H</i> ... <i>A</i>
C5—H5...N1	0.93	2.47	2.825(3)	103.0
C8—H8...C12	0.93	2.59	3.110(3)	116.0
C11—H11A...C13	0.97	2.57	3.268(3)	128.8
C6—H6...N1 ⁱⁱ	0.93	2.87	3.746(3)	157.6
	Cg...Cg	Alpha*	Gamma*	mpd*
Cg ₁ ...Cg ₁ ⁱ	4.4932(17)	0.00(15)	42.6	3.3084(11)
Cg ₂ ...Cg ₂ ⁱ	4.4931(17)	0.00(12)	36.5	3.6103(11)
Cg ₃ ...Cg ₃ ⁱ	4.493(2)	0.03(16)	32.5	3.7900(14)

Symmetry code: (i) $-1+x, y, z$; (ii) $-1 - x, -y, 1 - z$
 *Alpha: dihedral angle between planes I and J (°)
 *mpd: perpendicular distance of Cg(I) on ring J (Å)
 *Gamma: angle Cg(I)→Cg(J) vector and normal to plane J (°)
 Cg₁, Cg₂ and Cg₃ are the centroids of N1/C9, C1/C6 and C13/C18 rings, respectively

S8. Percentages of inhibition of 4e, 4f, and 5e-g

Table S2 Table S2. Percentages of inhibition of 4e, 4f, and 5e-g against *C. albicans* (*Ca*) and *C. neoformans* (*Cn*)^a

	Structure	Fungi	250	125	62.5	31.2	15.6	7.8	3.9	IC ₅₀
4e		<i>Ca</i>	12.8 ± 0.4	4.9 ± 1.5	4.2 ± 0.8	3.7 ± 0.1	2.4 ± 0.4	1.7 ± 0.5	2.0 ± 0.6	>250
		<i>Cn</i>	54.6 ± 1.0	27.2 ± 0.2	26.3 ± 0.4	18.4 ± 0.1	10.7 ± 0.2	6.6 ± 1.5	4.9 ± 1.4	250
4f		<i>Ca</i>	36.7 ± 0.2	26.9 ± 1.1	15.6 ± 0.1	6.4 ± 0.6	4.2 ± 0.7	4.1 ± 0.6	0 ± 0	>250
		<i>Cn</i>	65.3 ± 1.8	43.6 ± 1.5	25.9 ± 1.8	12.3 ± 1.2	9.8 ± 0.9	5.9 ± 0.5	2.8 ± 1.8	250
5e		<i>Ca</i>	31.5 ± 1.1	19.1 ± 1.1	11.1 ± 0.3	5.7 ± 0.2	6.3 ± 0.4	5.2 ± 0.9	0 ± 0	>250
		<i>Cn</i>	42.6 ± 1.4	34.2 ± 0.7	33.1 ± 0.4	29.6 ± 0.6	26.5 ± 0.2	23.3 ± 1.9	0 ± 0	>250
5f		<i>Ca</i>	55.1 ± 0.3	33.7 ± 0.1	21.1 ± 1.4	11.6 ± 0.5	10.4 ± 1.5	3.5 ± 0.3	0 ± 0	250
		<i>Cn</i>	100	100	100	100	83.3 ± 0.9	52.9 ± 0.4	40.4 ± 0.6	7.8
5g		<i>Ca</i>	49.9 ± 1.9	31.2 ± 0.7	19.1 ± 1.0	11.9 ± 1.5	8.9 ± 1.4	5.2 ± 0.2	0 ± 0	250
		<i>Cn</i>	38.6 ± 0.6	28.8 ± 0.5	21.4 ± 0.1	20.0 ± 1.2	15.9 ± 0.1	14.4 ± 0.5	11.0 ± 0.6	>250
AmpB		<i>Ca</i>	100	100	100	100	100	100	100	1.5
		<i>Cn</i>	100	100	100	100	100	100	100	1.0

^a *Ca* ATCC 10231 and *Cn* ATCC 32264. Dilutions are at the range 250–3.9 µg/mL. IC₅₀ value represents the concentration of each compound that inhibits 50 % of fungal growth. **AmpB**: Amphotericin B.

S9. HPLC analysis data of compounds 4e, 4f, and 5e-g

SHIMADZU
LabSolutions

Analysis Report

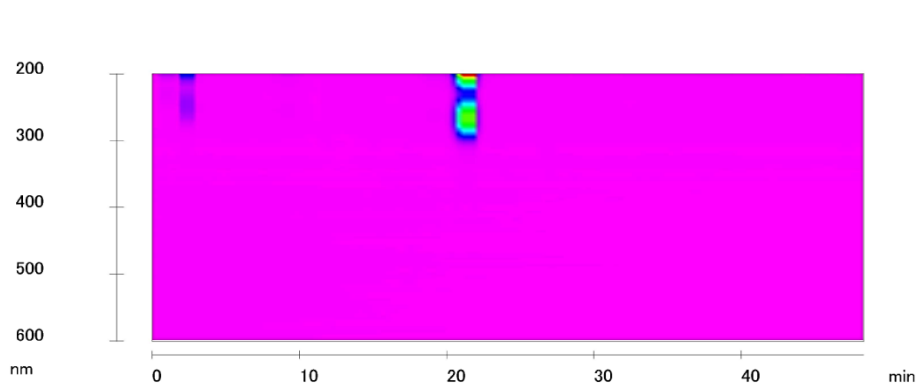
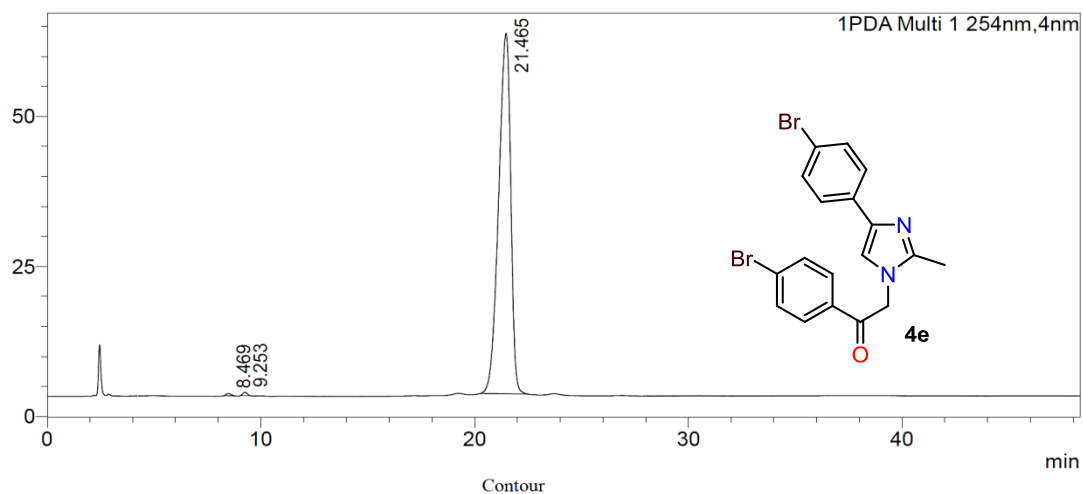
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 Method Filename : a.lcm
 Batch Filename :
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 Injection Volume : 20 uL
 Date Acquired : 12/04/2019 4:27:08 p. m.
 Date Processed : 12/04/2019 5:15:32 p. m.

Sample Type : Unknown
 Acquired by : System Administrator
 Processed by : System Administrator

<Chromatogram>

mAU



<Peak Table>

PDA Ch1 254nm

Peak#	Ret. Time	Area	Height	Name	Area%
1	8.469	6346	418		0.240
2	9.253	10185	660		0.386
3	21.465	2624093	66785		99.374
Total		2640624	67863		100.000

SHIMADZU
LabSolutions

Analysis Report

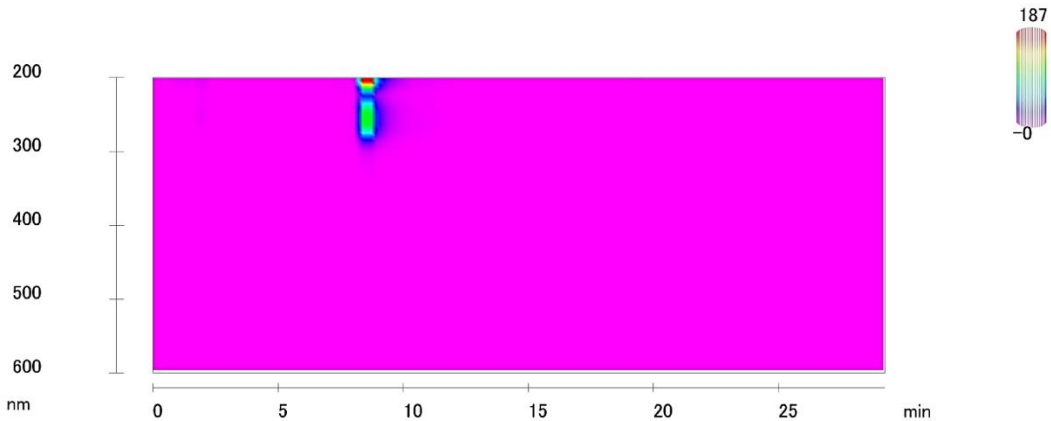
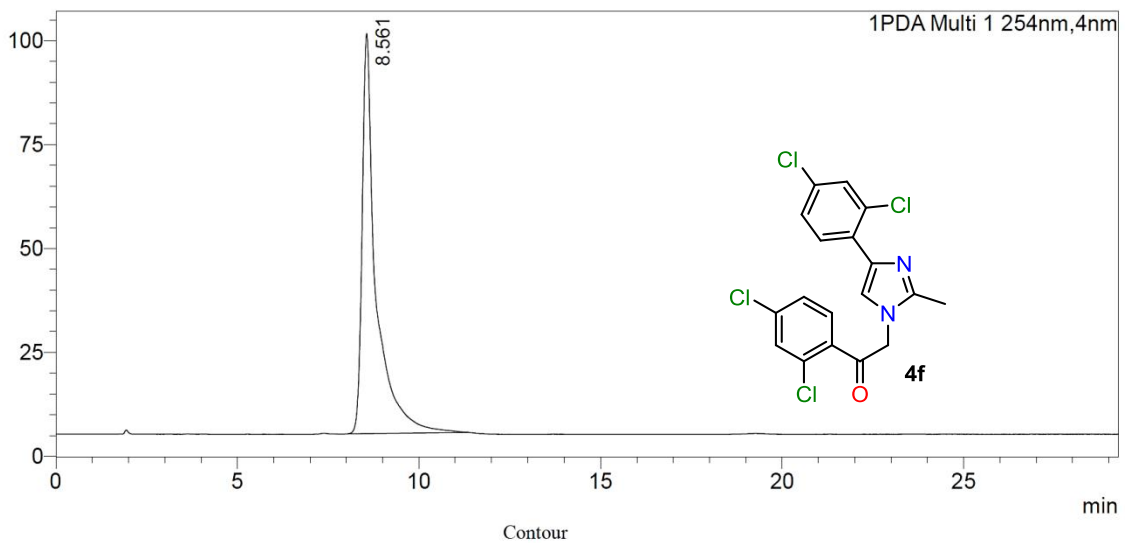
<Sample Information>

Sample Name : NRE-079A1
 Sample ID : NRE-079A1
 Data Filename : NRE-079A1.lcd
 Method Filename : a.lcm
 Batch Filename :
 Vial # : 1-1
 Injection Volume : 20 uL
 Date Acquired : 12/04/2019 3:06:59 p. m.
 Date Processed : 12/04/2019 3:36:17 p. m.

Sample Type : Unknown
 Acquired by : System Administrator
 Processed by : System Administrator

<Chromatogram>

mAU



<Peak Table>

PDA Ch1 254nm

Peak#	Ret. Time	Area	Height	Name	Area%
1	8.561	2840002	106984		100.000
Total		2840002	106984		100.000

SHIMADZU
LabSolutions

Analysis Report

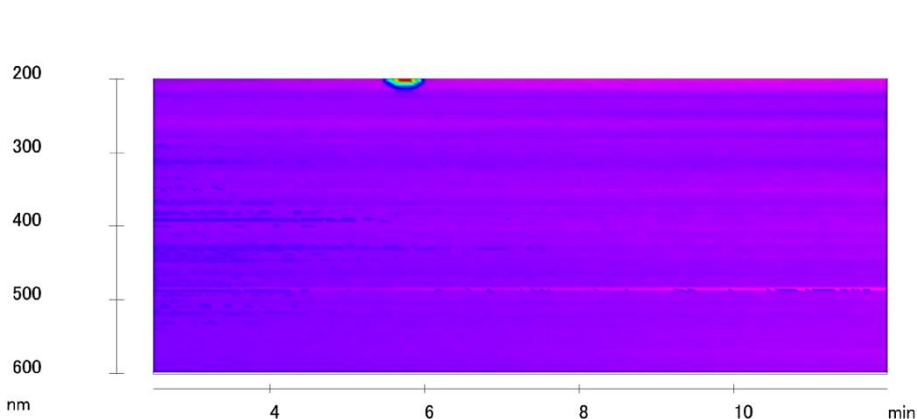
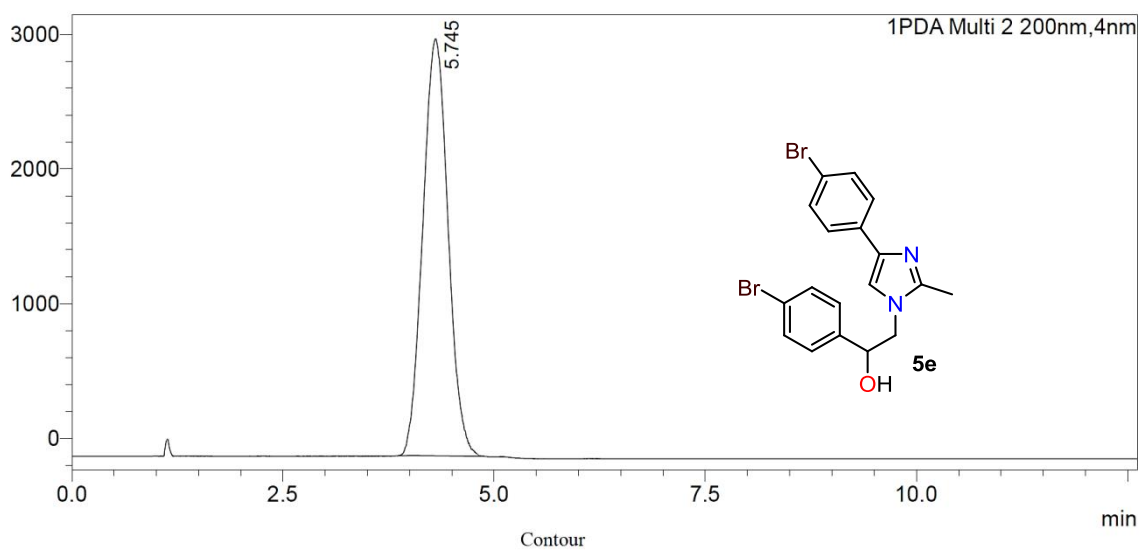
<Sample Information>

Sample Name : NRE-080B
 Sample ID : NRE-080B
 Data Filename : NRE-080B R2.lcd
 Method Filename : a.lcm
 Batch Filename :
 Vial # : 1-1
 Injection Volume : 20 uL
 Date Acquired : 10/05/2019 4:22:03 p. m.
 Date Processed : 13/05/2019 12:51:39 p. m.

Sample Type : Unknown
 Acquired by : System Administrator
 Processed by : System Administrator

<Chromatogram>

uAU



<Peak Table>

PDA Ch2 200nm

Peak#	Ret. Time	Area	Height	Name	Area%
1	5.745	53798	3440		100.000
Total		53798	3440		100.000



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Analysis Report

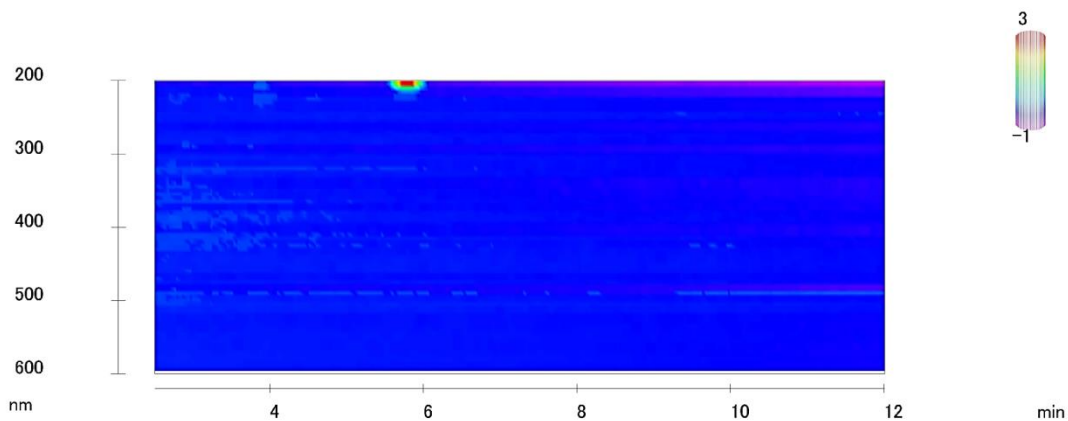
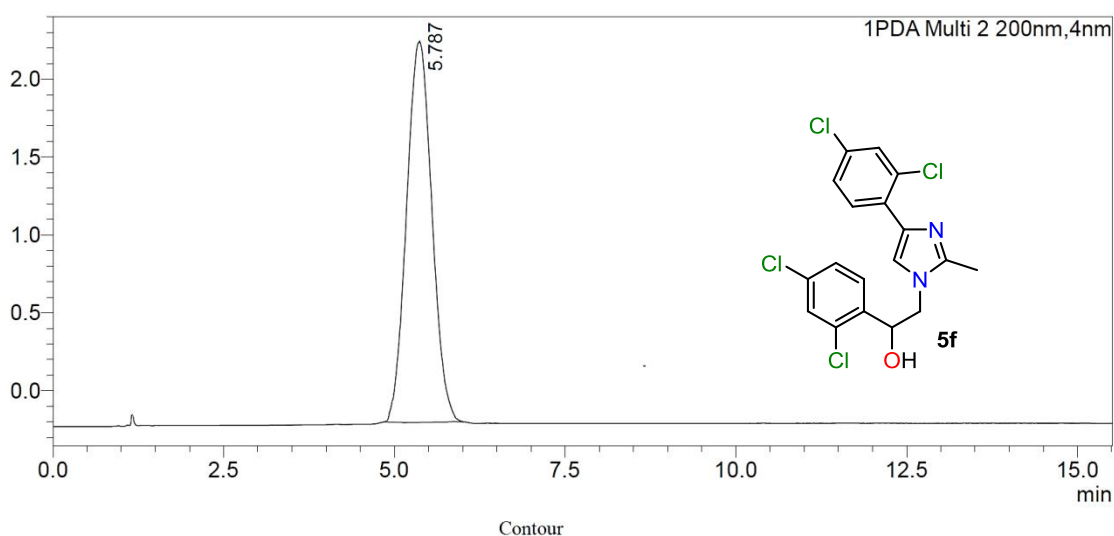
<Sample Information>

Sample Name : NRE-079B-R2
 Sample ID : NRE-079B-R2
 Data Filename : NRE-079B-R2.lcd
 Method Filename : a.lcm
 Batch Filename :
 Vial # : 1-1
 Injection Volume : 20 uL
 Date Acquired : 10/05/2019 3:42:19 p. m.
 Date Processed : 13/05/2019 12:53:44 p. m.

Sample Type : Unknown
 Acquired by : System Administrator
 Processed by : System Administrator

<Chromatogram>

mAU



<Peak Table>

PDA Ch2 200nm

Peak#	Ret. Time	Area	Height	Name	Area%
1	5.787	41947	2689		100.000
Total		41947	2689		100.000

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LabSolutions

Analysis Report

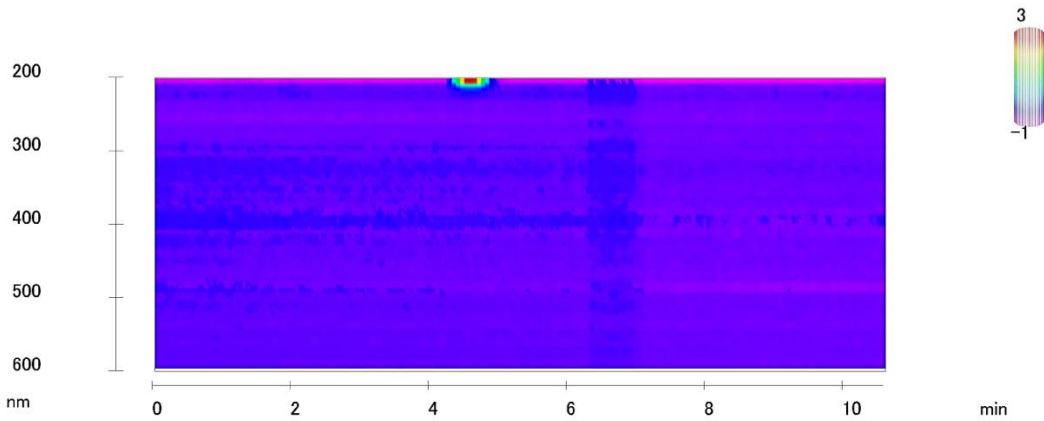
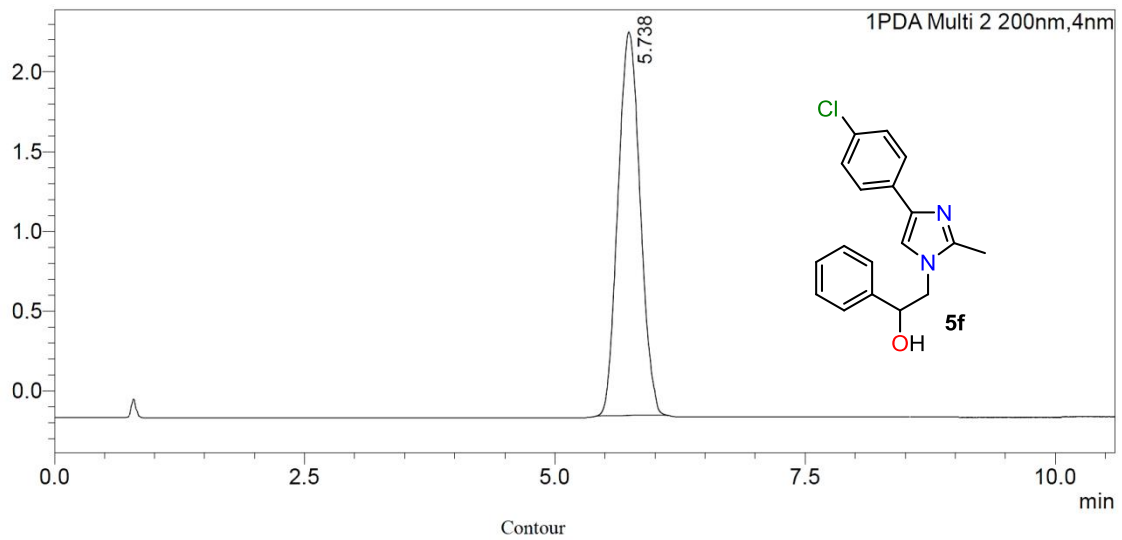
<Sample Information>

Sample Name : NRE-035E
 Sample ID : NRE-035E
 Data Filename : NRE-035E R2.lcd
 Method Filename : a.lcm
 Batch Filename :
 Vial # : 1-1
 Injection Volume : 20 uL
 Date Acquired : 10/05/2019 4:38:00 p. m.
 Date Processed : 10/05/2019 4:48:39 p. m.

Sample Type : Unknown
 Acquired by : System Administrator
 Processed by : System Administrator

<Chromatogram>

mAU



<Peak Table>

PDA Ch2 200nm

Peak#	Ret. Time	Area	Height	Name	Area%
1	5.738	41059	2669		100.000
Total		41059	2669		100.000