

Antiproliferative, Pro-apoptotic Activity And Tubulin Dynamics Modulation of 1*H*-benzimidazol-2-yl Hydrazones in Human Breast Cancer Cell Line MDA-MB-231

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hydrazone **5d** in imino and amino tautomeric form

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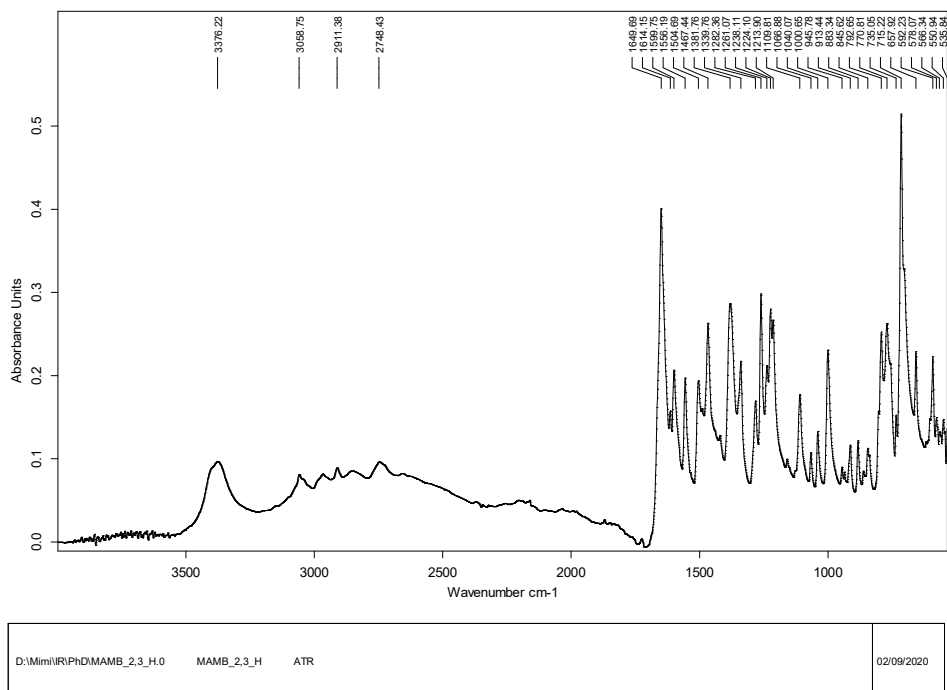


Fig. S1a. ATR-IR spectrum of compound **5c**

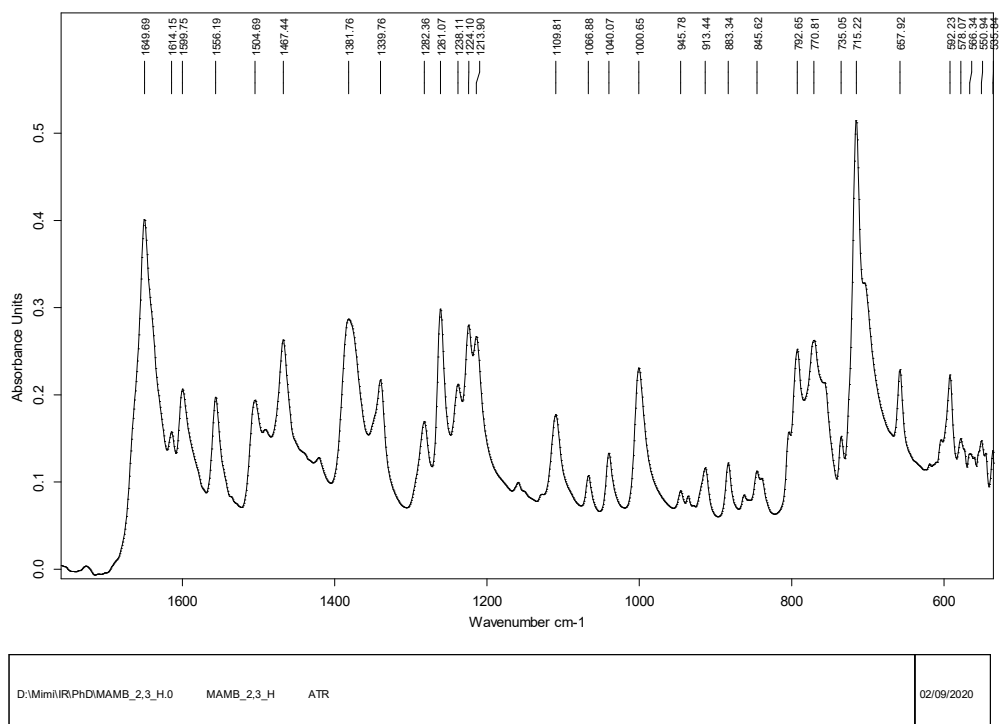


Fig. S1b. ATR-IR spectrum of compound **5c**

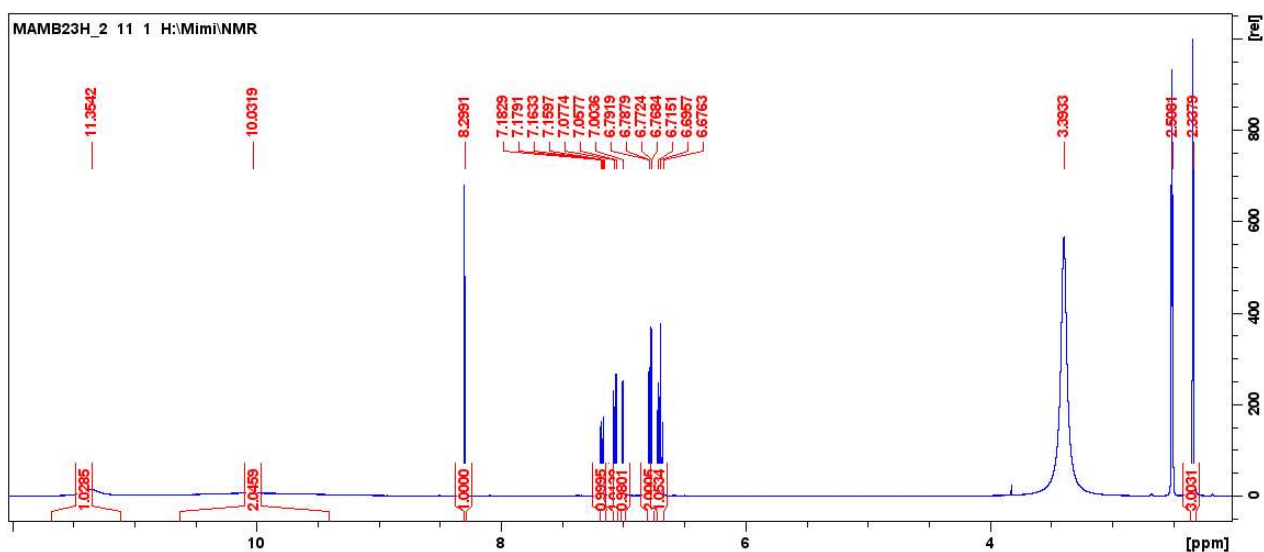


Fig. S2. ^1H NMR spectrum of compound **5c** in $\text{DMSO}-d_6$

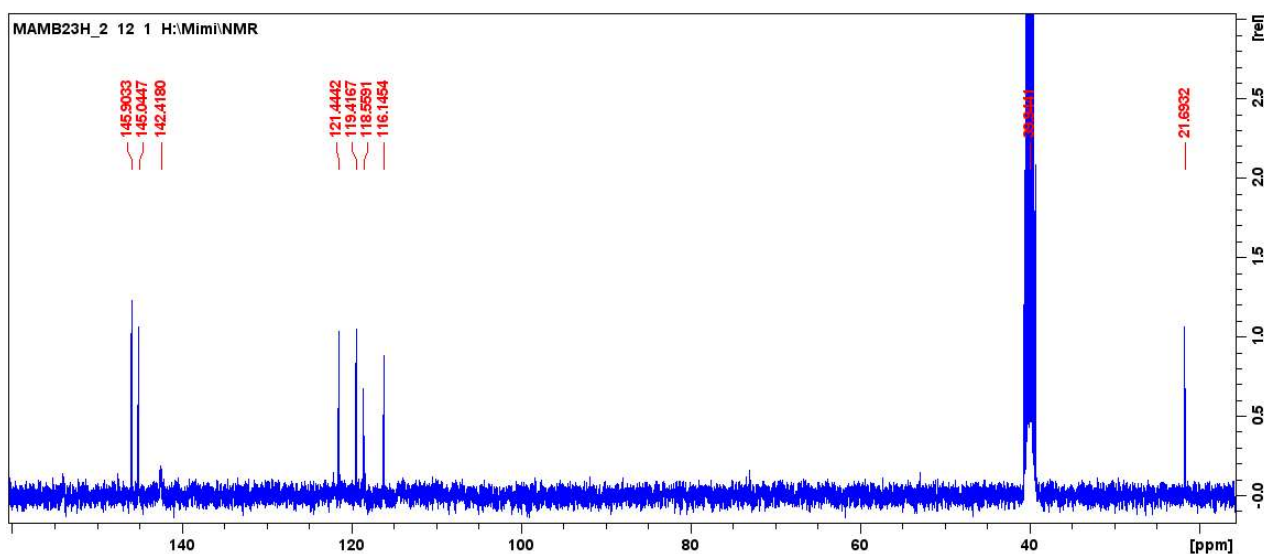


Fig. S3. ^{13}C NMR spectrum of compound **5c** in $\text{DMSO}-d_6$

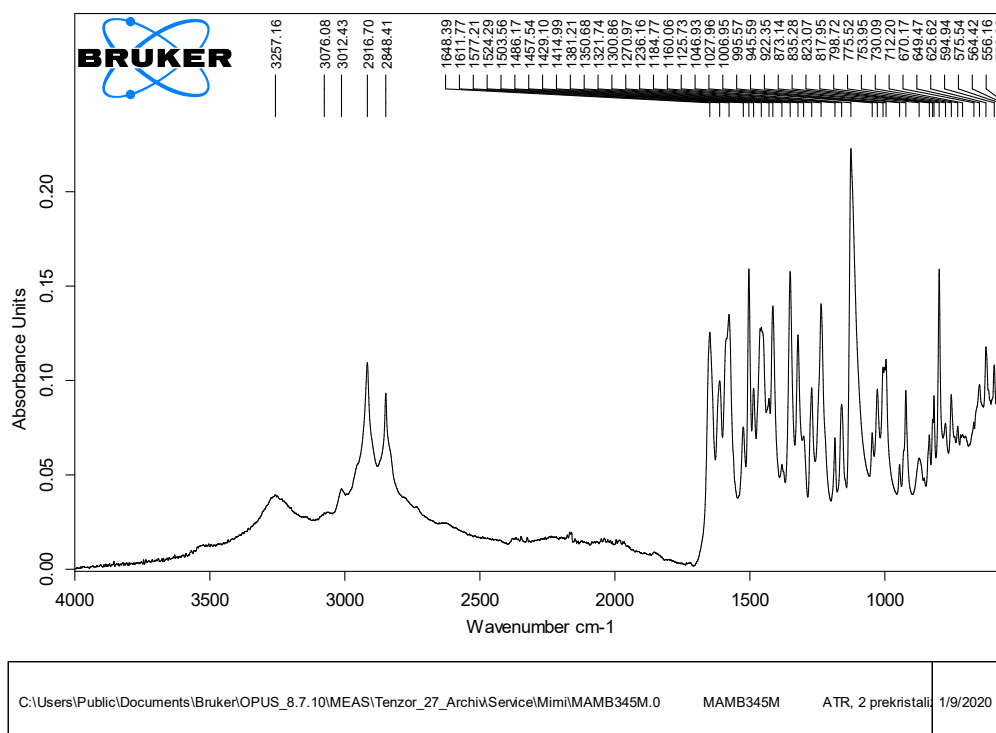


Fig. S4a. ATR-IR spectrum of compound **5d**

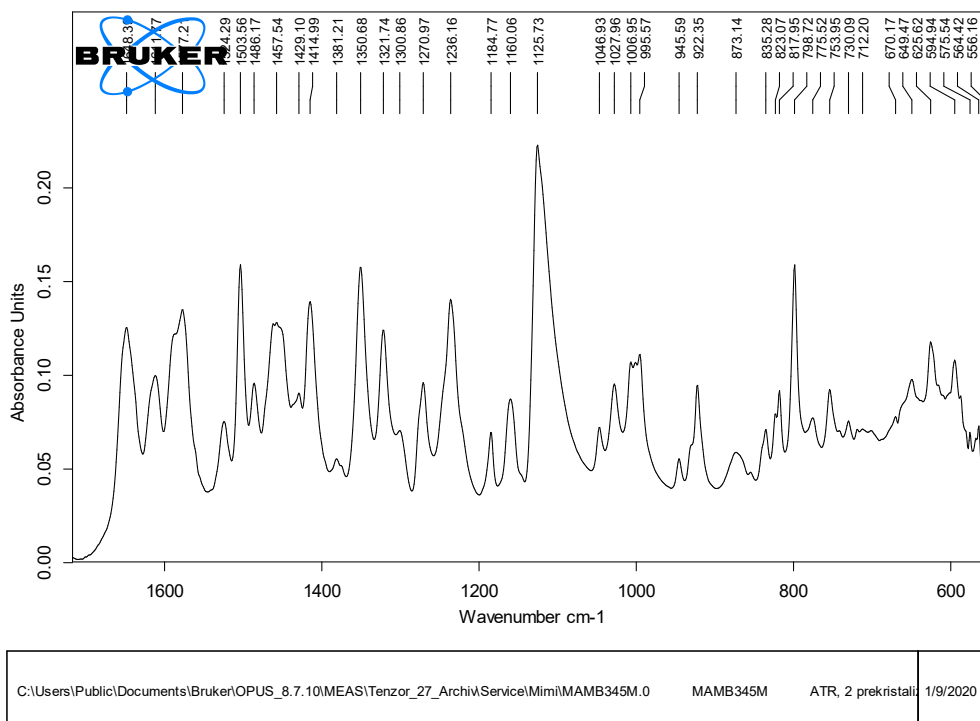


Fig. S4b. IR spectrum of compound **5d**

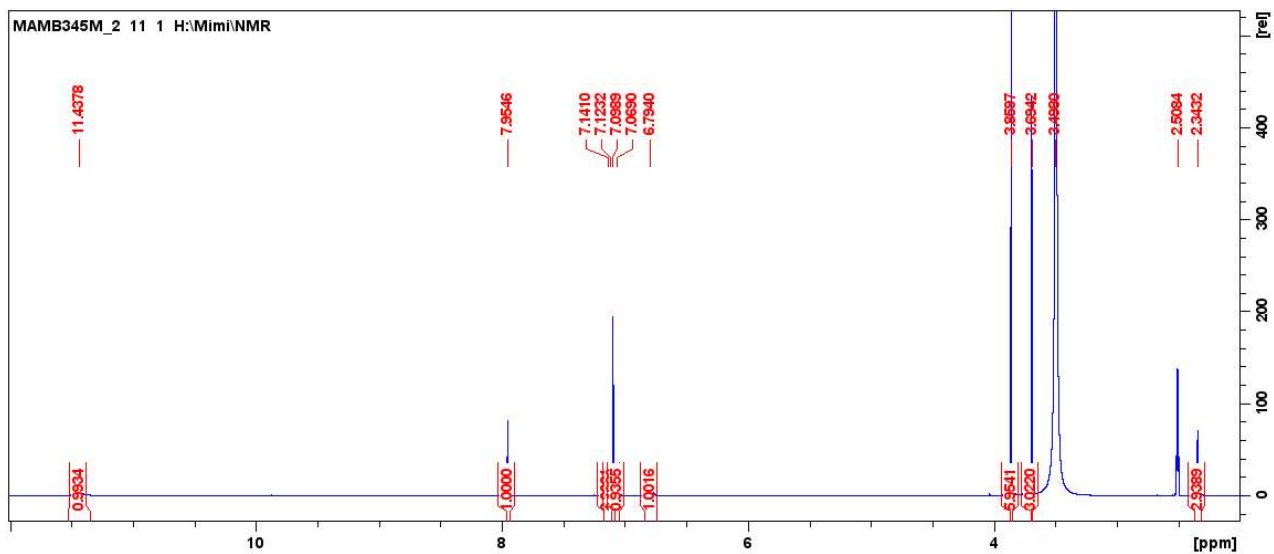


Fig. S5. ^1H NMR spectrum of compound **5d** in $\text{DMSO-}d_6$

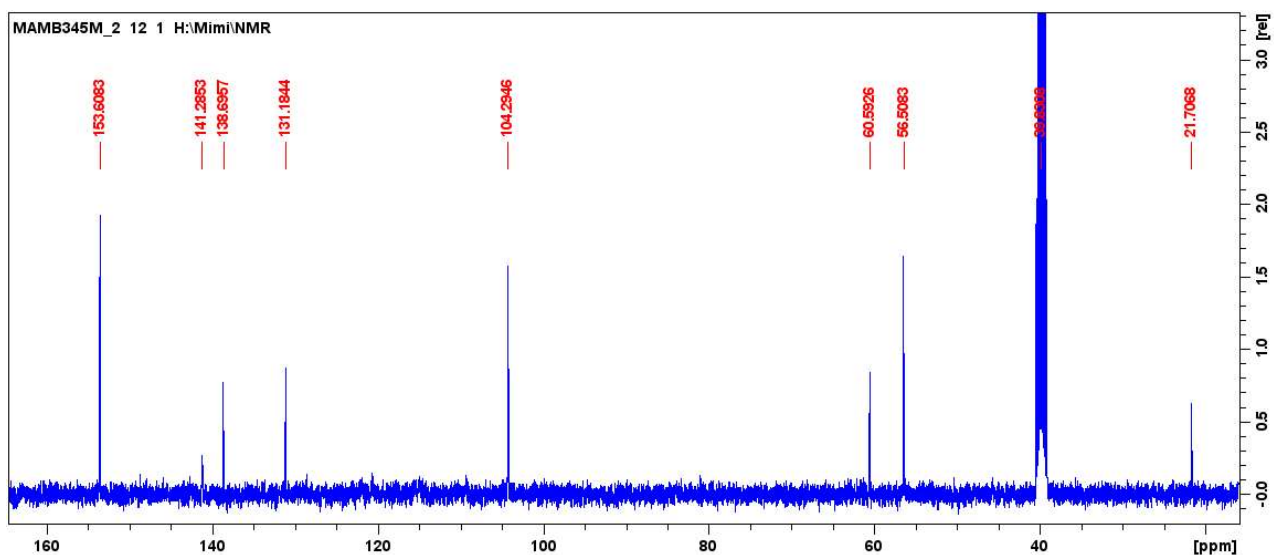
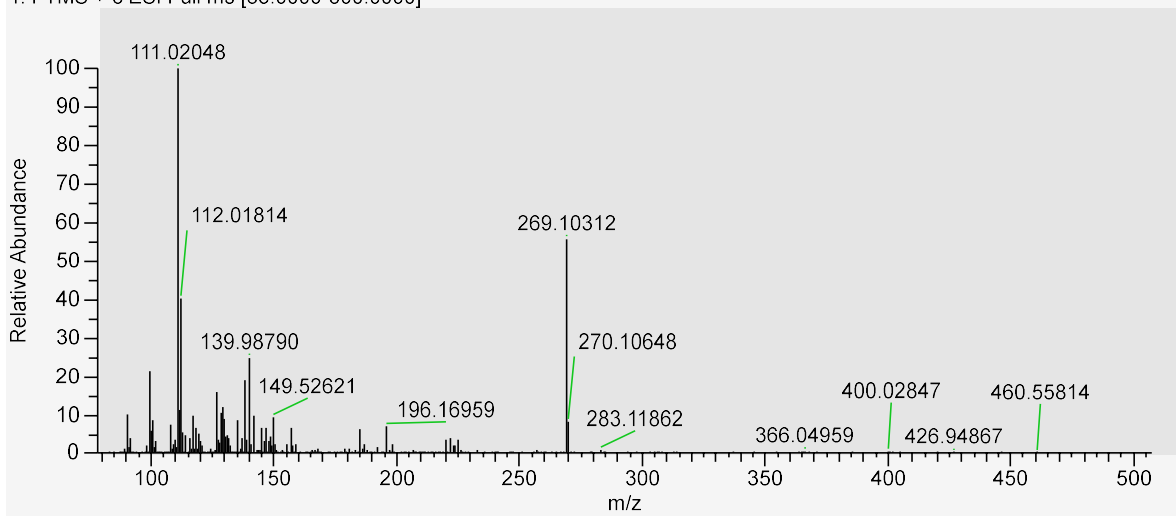


Fig. S6. ^{13}C NMR spectrum of compound **5d** in $\text{DMSO-}d_6$

MAB23H_20240507224148 #43 RT: 0.40 AV: 1 NL: 1.45E+008
T: FTMS + c ESI Full ms [83.0000-500.0000]



MAB23H_20240507224148 #42 RT: 0.39 AV: 1 NL: 2.95E+007
T: FTMS - c ESI Full ms [83.0000-500.0000]

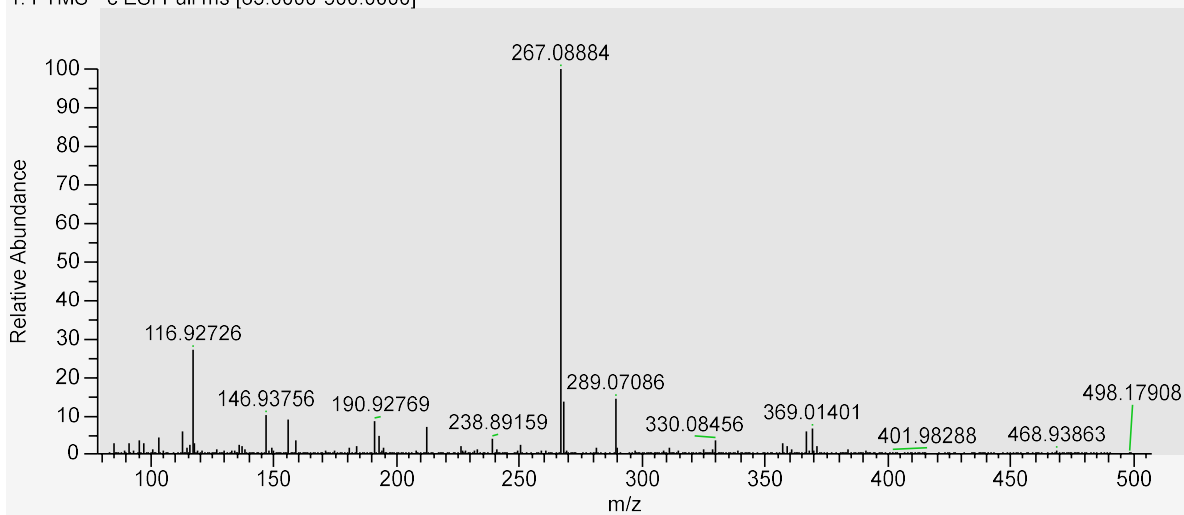
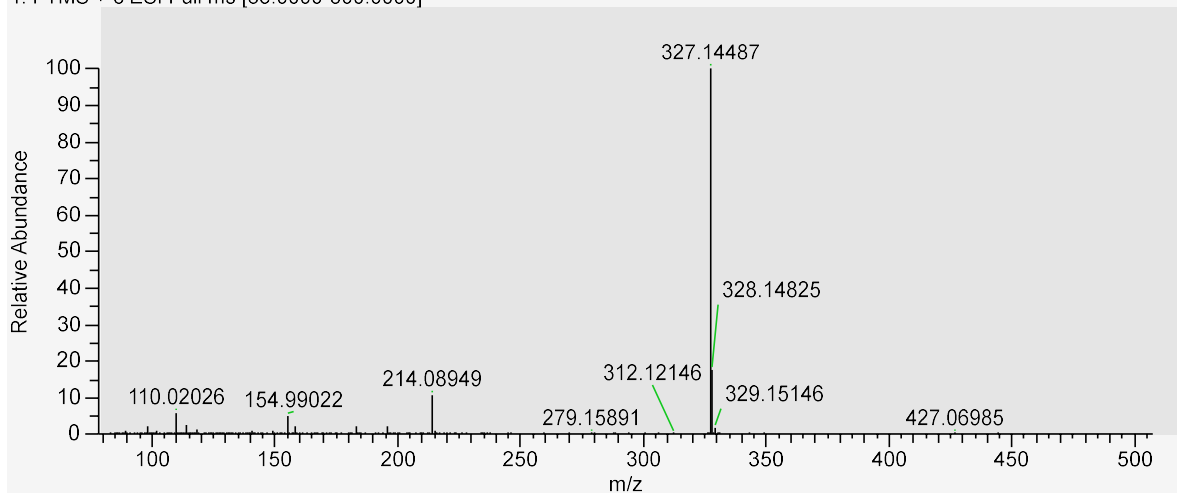


Figure S7. HRMS spectra of 5a

MAB345M #705 RT: 6.87 AV: 1 NL: 3.77E+008
T: FTMS + c ESI Full ms [83.0000-500.0000]



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T: FTMS - c ESI Full ms [83.0000-500.0000]

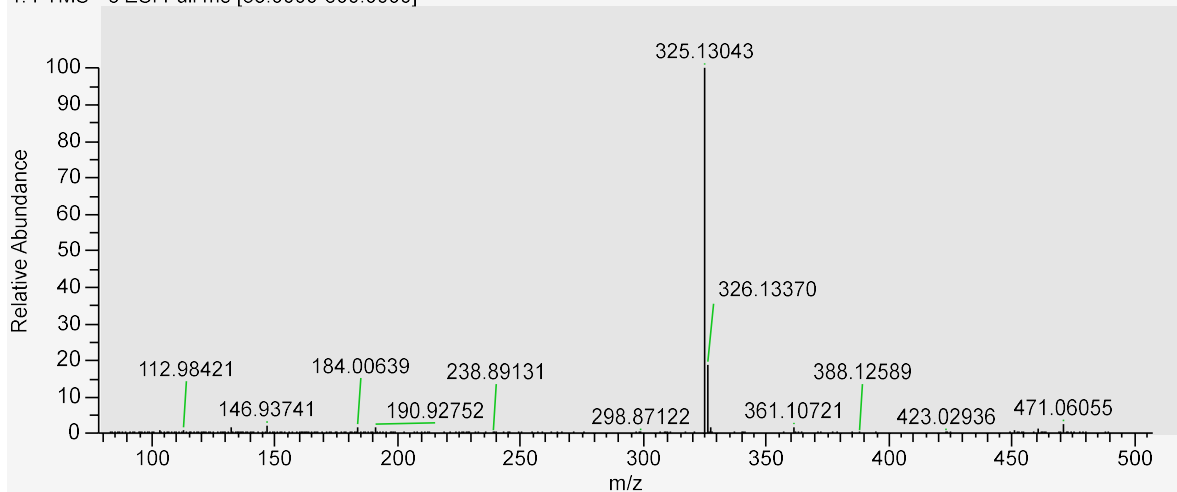
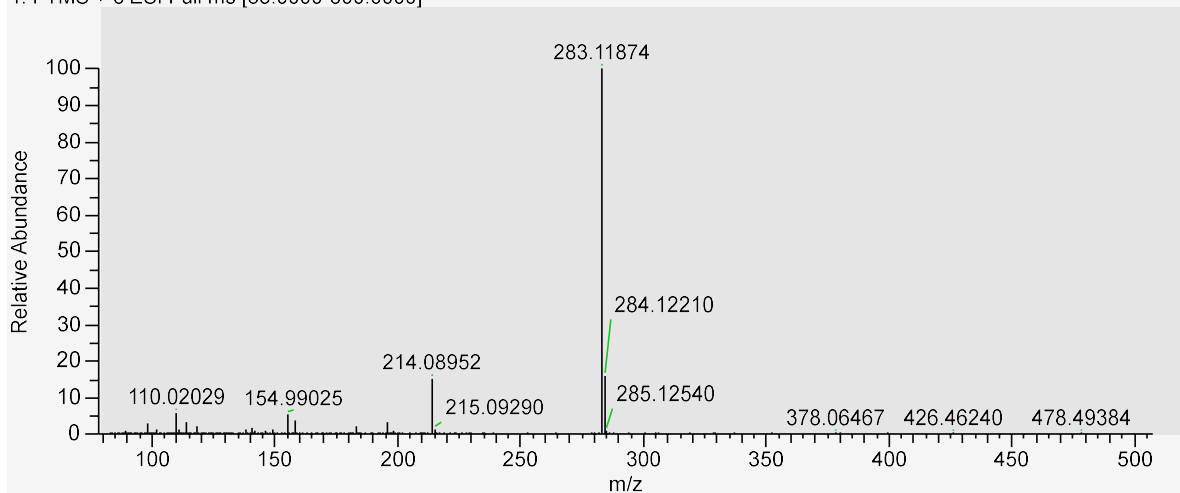


Figure S8. HRMS spectra of **5b**

MAMB23H #677 RT: 6.59 AV: 1 NL: 2.71E+008
T: FTMS + c ESI Full ms [83.0000-500.0000]



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T: FTMS - c ESI Full ms [83.0000-500.0000]

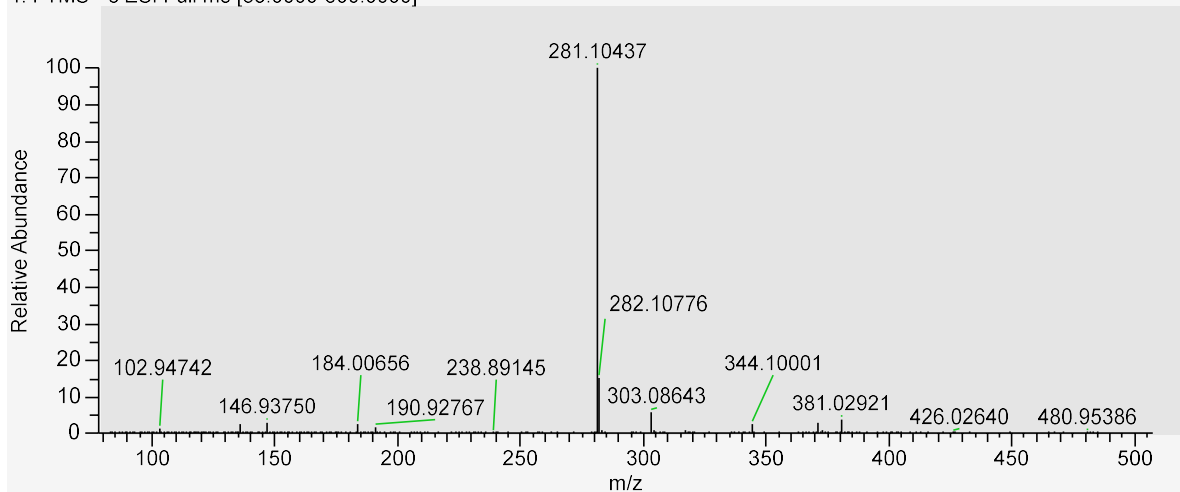
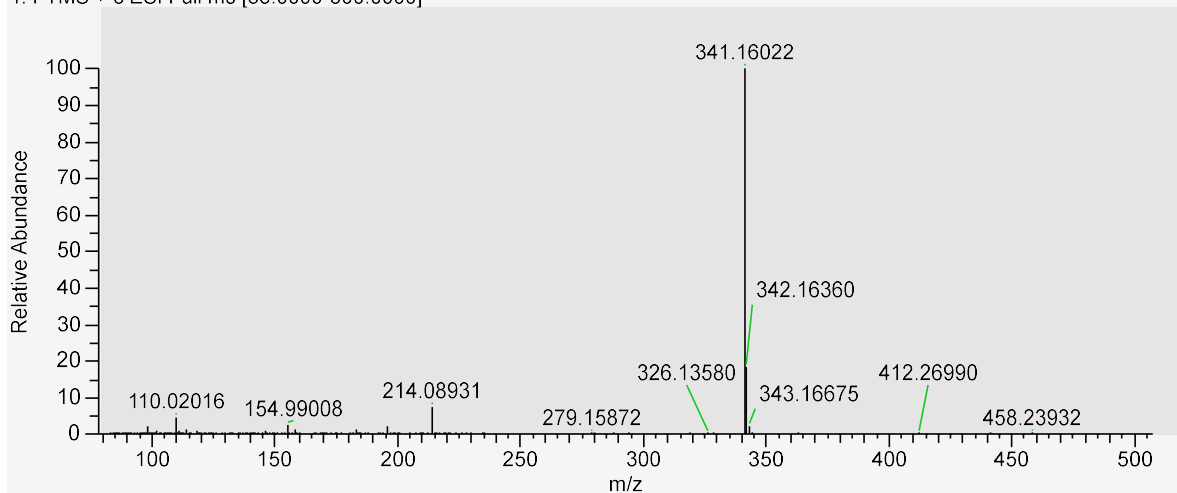


Figure S9. HRMS spectra of **5c**

MAMB345M #727 RT: 7.08 AV: 1 NL: 4.31E+008
T: FTMS + c ESI Full ms [83.0000-500.0000]



MAMB345M #728 RT: 7.09 AV: 1 NL: 1.09E+008
T: FTMS - c ESI Full ms [83.0000-500.0000]

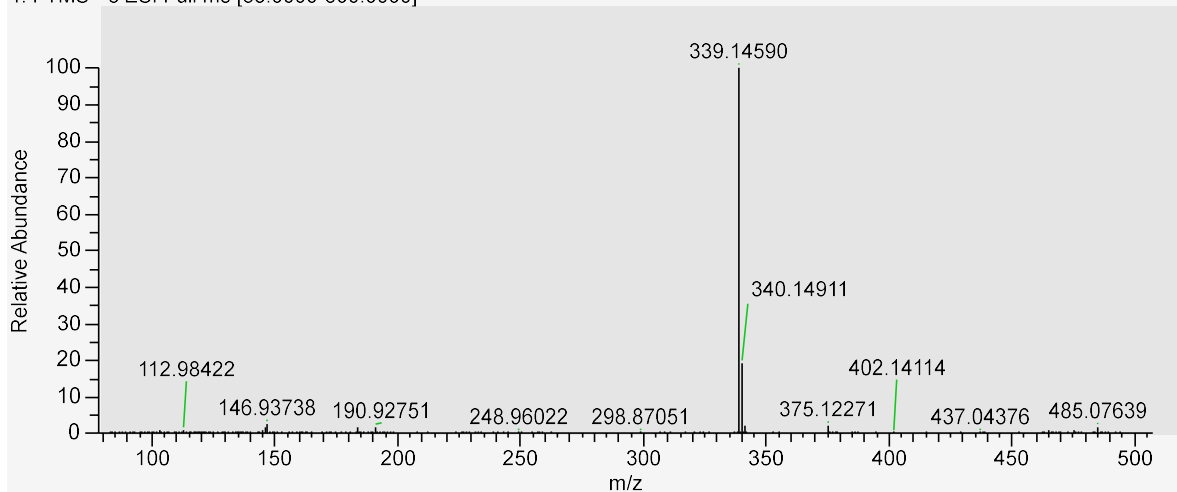


Figure S10. HRMS spectra of **5d**

Table S1. Optimized molecular structure, total Gibbs energy (in a.u.) and relative Gibbs energies (in kJ mol⁻¹) of the possible isomers of compound **5c** in imino tautomeric form, obtained at B3LYP/6-311++G(d,p) level of theory in gas phase and water.

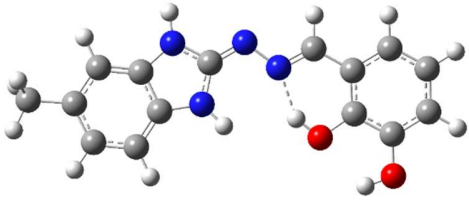
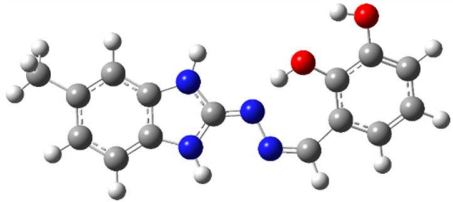
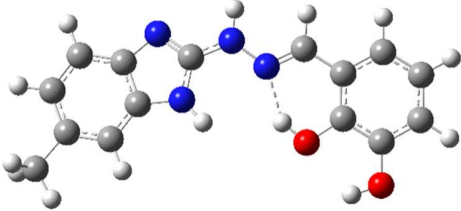
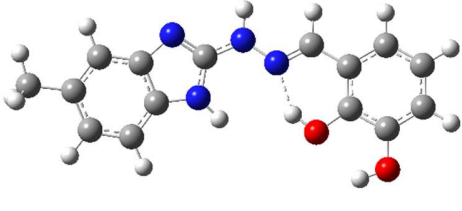
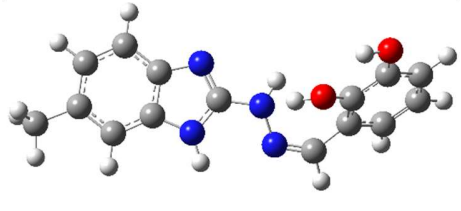
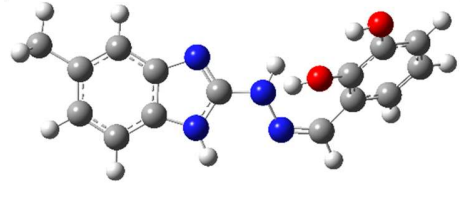
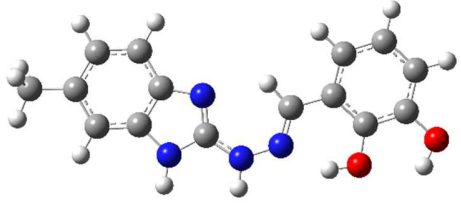
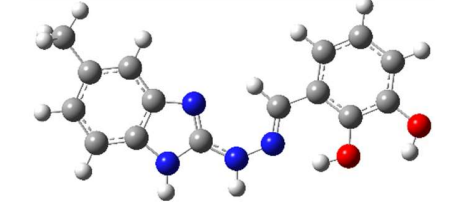
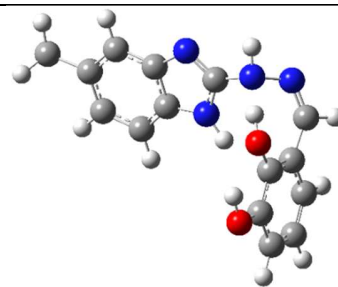
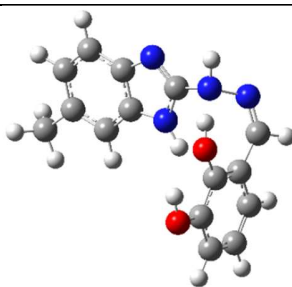
Medium	Imino tautomeric form	ΔG
		
	<i>E, s-trans</i>	
gas phase	G _{tot} : -949.516836	-
water	G _{tot} : -949.533431	-
		
	<i>Z, s-trans</i>	
gas phase	G _{tot} : -949.514769	5.42
water	G _{tot} : -949.530744	7.05

Table S2. Optimized molecular structure, total Gibbs energies (in a.u.) and relative Gibbs energies (in kJ.mol⁻¹) of the possible isomers of compound **5c** in amino tautomeric form, obtained at B3LYP/6-311++G(d,p) level of theory in gas phase and water.

Medium		6-methyl isomer	5-methyl isomer
			
gas phase	<i>E, s-trans</i>		
	G_{tot}	-949.513467	-949.512992
	ΔG	-	1.24
water	<i>E, s-trans</i>		
	G_{tot}	-949.531327 a.u.	-949.531095
	ΔG	-	0.60
			
gas phase	<i>Z, s-trans</i>		
	G_{tot}	-949.501294	-949.501873
	ΔG	1.52	-
water	<i>Z, s-trans</i>		
	G_{tot}	-949.516863	-949.51753
	ΔG	1.75	-
			
gas phase	<i>E, s-cis</i>		
	G_{tot}	-949.507653	-949.509719
	ΔG	5.42	-
water	<i>E, s-cis</i>		
	G_{tot}	-949.525408	-949.525125
	ΔG	-	0.74



		<i>Z, s-cis</i>	<i>Z, s-cis</i>
gas phase	G_{tot}	-949.493624	-949.493092
	ΔG	-	1.39
water	G_{tot}	-949.509446	-949.509564
	ΔG	0.30	-

Table S3. Summarized total Gibbs energies (in a.u.), relative Gibbs energies (in a.u. and kJ.mol⁻¹) and Boltzmann population (in %) for the tautomeric and isomeric forms of compound **5c**, obtained at B3LYP/6-311++G(d,p) level of theory in different media.

Tautomer/isomer	G_{tot}	ΔG (a.u.)	ΔG (kJ.mol⁻¹)	Boltzmann population
<i>Gas</i>				
imino (E,s-trans)	-949.516836	0	0	86.42
imino (Z,s-trans)	-949.514769	0.002067	5.43	9.67
6-isomer				
amino (E,s-trans)	-949.513467	0.003369	8.85	2.43
amino (Z,s-trans)	-949.501294	0.015542	40.81	
amino (E,s-cis)	-949.507653	0.009183	24.11	
amino (Z,s-cis)	-949.493624	0.019843	52.10	
5-isomer				
amino (E,s-trans)	-949.512992	0.003844	10.09	1.48
amino (Z,s-trans)	-949.501873	0.014963	39.29	
amino (E,s-cis)	-949.509719	0.007117	18.69	
amino (Z,s-cis)	-949.493092	0.023744	62.34	
<i>Water</i>				
imino (E,s-trans)	-949.533431	0	0	79.97
imino (Z,s-trans)	-949.530744	0.002687	7.05	4.65
6-isomer				
amino (E,s-trans)	-949.531327	0.002104	5.52	8.63
amino (Z,s-trans)	-949.516863	0.016568	43.50	
amino (E,s-cis)	-949.525408	0.008023	21.06	
amino (Z,s-cis)	-949.509446	0.023985	62.97	
5-isomer				
amino (E,s-trans)	-949.531095	0.002336	6.13	6.75
amino (Z,s-trans)	-949.51753	0.015901	41.75	
amino (E,s-cis)	-949.525125	0.008306	21.81	
amino (Z,s-cis)	-949.509564	0.023867	62.66	
<i>DMSO</i>				
imino (E,s-trans)	-949.532308	0	0	53.25
6-isomer				
amino (E,s-trans)	-949.531449	0.000859	2.26	21.40
5-isomer				
amino (E,s-trans)	-949.531607	0.000701	1.84	25.35

Table S4. Optimized molecular structure, total Gibbs energy (in a.u.) and relative Gibbs energies (in kJ mol⁻¹) of the possible isomers of compound **5d** in imino tautomeric form, obtained at B3LYP/6-311++G(d,p) level of theory in gas phase and water.

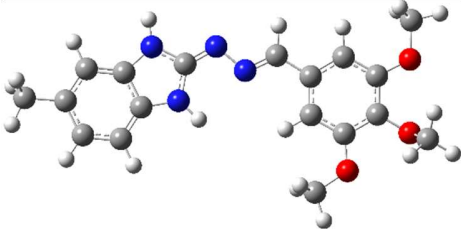
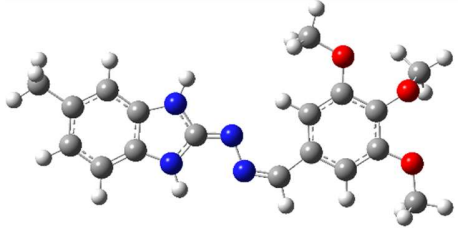
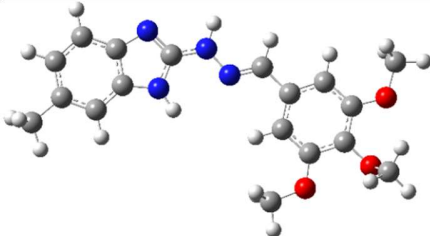
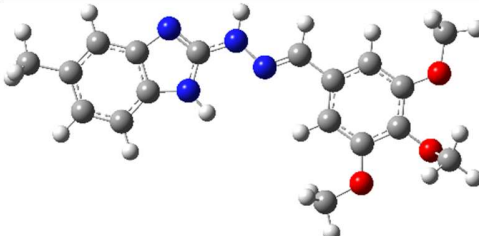
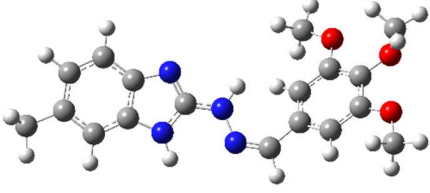
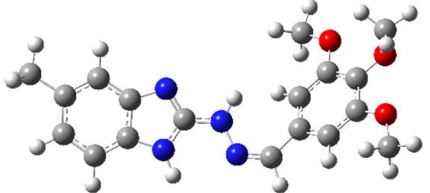
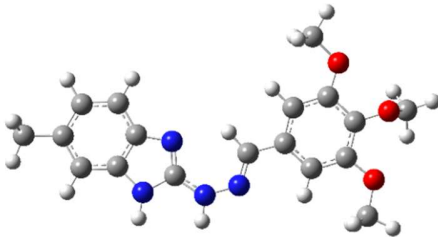
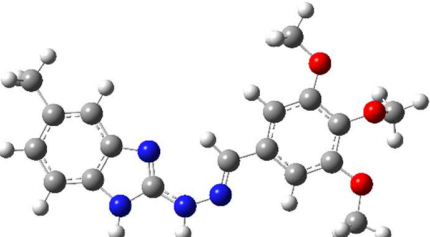
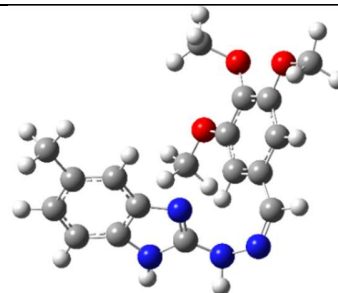
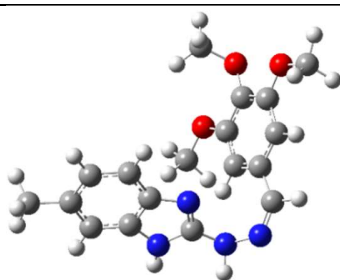
Medium	Imino tautomeric form	ΔG
		
	<i>E, s-trans</i>	
gas phase	G _{tot} : -1142,585534	-
water	G _{tot} : -1142,605514	-
		
	<i>Z, s-trans</i>	
gas phase	G _{tot} : -1142,582568	7.78
water	G _{tot} : -1142,604003	15.43

Table S5. Optimized molecular structure, total Gibbs energies (in a.u.) and relative Gibbs energies (in kJ.mol⁻¹) of the possible isomers of compound **5d** in amino tautomeric form, obtained at B3LYP/6-311++G(d,p) level of theory in gas phase and water.

Medium		6-methyl isomer	5-methyl isomer
			
gas phase	<i>E, s-trans</i>		
	G_{tot}	-1142.589796	-1142.589791
	ΔG	-	0.01
water	G_{tot}	-1142.610250	-1142.611019
	ΔG	-	2.01
			
gas phase	<i>Z, s-trans</i>		
	G_{tot}	-1142.583297	-1142.582924.
	ΔG	-	0.97
water	G_{tot}	-1142.602714	-1142.602627
	ΔG	-	0.22
			
gas phase	<i>E, s-cis</i>		
	G_{tot}	-1142.580100	-1142.580055
	ΔG	-	0.11
water	G_{tot}	-1142.601929	-1142.601816
	ΔG	-	0.29

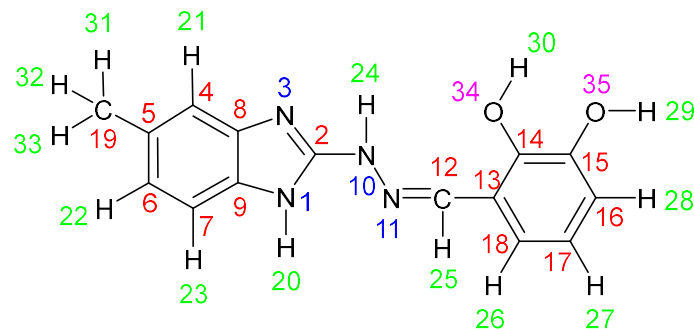


gas phase	G_{tot}	<i>Z, s-cis</i> -1142.564804	<i>Z, s-cis</i> -1142.565265
	ΔG	1.21	-
water	G_{tot}	-1142.587979	-1142.589262
	ΔG	3.36	-

Table S6. Summarized total Gibbs energies (in a.u.), relative Gibbs energies (in a.u. and kJ.mol⁻¹) and Boltzmann population (in %) for all tautomeric and isomeric forms of compound **5d**, obtained at B3LYP/6-311++G(d,p) level of theory in gas phase and water.

Tautomer/isomer	G _{tot}	ΔG (a.u.)	ΔG (kJ.mol ⁻¹)	Boltzmann population
<i>Gas</i>				
imino (E,s-trans)	-1142.585534	0.004262	11.19	
imino (Z,s-trans)	-1142.582568	0.007228	18.98	
6-isomer				
amino (E,s-trans)	-1142.589796	0	0	51.10
amino (Z,s-trans)	-1142.583297	0.006499	17.06	
amino (E,s-cis)	-1142.580100	0.009696	25.46	
amino (Z,s-cis)	-1142.564804	0.024992	65.62	
5-isomer				
amino (E,s-trans)	-1142.589791	5.10 ⁻⁶	0.01	49.90
amino (Z,s-trans)	-1142.582924	0.006872	18.04	
amino (E,s-cis)	-1142.580055	0.009741	25.57	
amino (Z,s-cis)	-1142.565265	0.024531	64.41	
<i>Water</i>				
imino (E,s-trans)	-1142.605514	0.005505	14.45	
imino (Z,s-trans)	-1142.604003	0.007016	18.42	
6-isomer				
amino (E,s-trans)	-1142.610250	0.000769	2.02	30.69
amino (Z,s-trans)	-1142.602714	0.008305	21.80	
amino (E,s-cis)	-1142.601929	0.00909	23.87	
amino (Z,s-cis)	-1142.587979	0.02304	60.49	
5-isomer				
amino (E,s-trans)	-1142.611019	0	0	69.31
amino (Z,s-trans)	-1142.602627	0.008392	22.03	
amino (E,s-cis)	-1142.601816	0.009203	24.16	
amino (Z,s-cis)	-1142.589262	0.021757	57.12	
<i>DMSO</i>				
imino (E,s-trans)	-1142.602682	0.007986	20.97	
6-isomer				
amino (E,s-trans)	-1142.610668	0	0	59.95
5-isomer				
amino (E,s-trans)	-1142.610288	0.00038	1.00	40.05

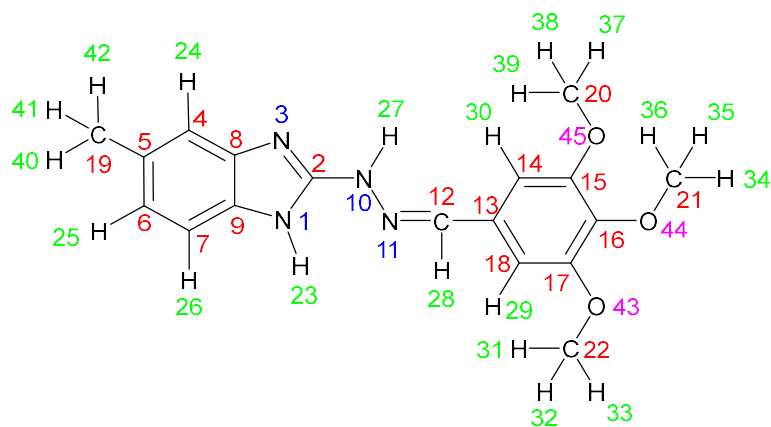
Table S7. Chemical shifts (in ppm) for the imino form, 5-methyl isomer sand 6-methyl isomer of the amino form of compound **5c** predicted by GIAO method using B3LYP functional and 6-311++G** basis set calculations in DMSO solvent and experimental data for **5c**.



B3LYP/6-311++G** DMSO							
C Atom №	$\delta_{\text{exp. in}}$ DMSO- d_6	Imino form		5-methyl isomer in amino form		6-methyl isomer in amino form	
		chem. shift	rel. to TMS	chem. shift	rel. to TMS	chem. shift	rel. to TMS
5	121.44	55.7767	128.8	43.6309	140.9	54.6946	129.9
4	118.55	70.5148	114.0	61.477	123.1	61.8959	122.7
6	142.41	42.6670	141.9	56.1349	128.4	44.8265	139.7
8	145.04	48.2187	136.3	33.4201	151.1	35.7915	148.8
7	116.14	69.0386	115.5	69.5873	114.9	69.0442	115.5
9	142.41	46.1206	138.4	46.1212	138.4	44.0197	140.5
2	145.90	25.3236	159.2	28.2886	156.2	28.6183	155.9
12	145.04	25.8879	158.7	34.7429	149.8	34.65	149.9
13	119.41	57.4188	127.1	59.3038	125.3	58.9294	125.6
18	121.44	57.0677	127.5	57.4073	127.1	56.7223	127.8
14	145.90	31.2122	153.4	32.7338	151.8	32.0896	152.5
17	121.44	58.167	126.4	56.7322	127.8	57.089	127.5

15	145.90	31.3152	153.2	30.4474	154.1	30.824	153.7
16	118.55	63.8011	120.8	62.0727	122.5	62.2033	122.3
19	31.69	160.9968	23.6	160.8786	23.7	160.8177	23.7
H Atom							
N ₂							
21	6.99	24.7542	7.2	24.4974	7.4	24.4566	7.5
23	7.17-7.16	24.7444	7.2	24.4042	7.5	24.5319	7.4
20	11.36	24.1828	7.8	23.694	8.3	23.6377	8.3
25	8.28	24.9762	6.9	23.9583	8.0	23.8575	8.1
24	-	23.32	8.6	23.9835	7.9	23.9284	8.0
22	7.06-7.04	24.8098	7.1	24.8379	7.1	24.571	7.4
27	6.69-6.67	24.9239	7.0	21.6753	10.3	24.8774	7.1
30	9.34	20.0768	11.9	24.8807	7.1	21.6214	10.3
26	6.77-6.75	24.9005	7.0	24.6416	7.3	24.9113	7.1
28	6.77-6.75	24.8502	7.1	26.3282	5.6	24.7245	7.2
29	9.34	26.3549	5.6	29.3535	2.6	26.2931	5.7
32	2.32	29.3897	2.6	29.7879	2.1	29.2003	2.7
31	2.32	29.396	2.6	29.3012	2.6	29.2892	2.7
33	2.32	29.8073	2.1	24.6493	7.3	29.8911	2.1

Table S8. Chemical shifts (in ppm) for the 5-methyl isomer and 6-methyl isomer of the amino form of compound **5d** predicted by GIAO method using B3LYP functional and 6-311++G** basis set calculations in DMSO solvent and experimental data for **5d**.



C Atom No	$\delta_{\text{Exp.}}$ in DMSO- d_6	6-methyl isomer in amino form		5-methyl isomer in amino form	
		chem. shift	rel. to TMS	chem. shift	rel. to TMS
5	131.18	55.0135	129.5	43.9386	140.6
4	131.18	61.8759	122.7	61.7245	122.8
6	138.69	45.0612	139.5	57.0824	127.5
8	153.60	35.3683	149.2	32.6431	151.9
7	104.29	69.3288	115.2	69.9332	114.6
9	138.69	44.0731	140.5	46.0336	138.5
2	153.60	26.9442	157.6	26.7588	157.8
12	141.28	37.8856	146.7	37.6092	146.9
13	138.69	45.8021	138.8	46.2601	138.3
18	104.29	82.4458	102.1	82.2531	102.3
14	104.29	74.7445	109.8	73.3204	111.2
17	153.60	20.4700	164.1	20.3526	164.2

15	153.60	21.3767	163.2	20.9407	163.6
16	141.28	37.7587	146.8	36.8681	147.7
20	56.50	126.9329	57.6	126.3548	58.2
22	56.50	126.2986	58.3	125.771	58.8
21	60.59	121.2851	63.3	121.3134	63.2
19	21.70	160.8036	23.8	160.6998	23.9
H Atom №					
24	7.17-7.05	24.5066	7.4	24.5080	7.4
26	7.17-7.05	24.5864	7.4	24.4192	7.5
23	11.38	23.2296	8.7	23.2109	8.7
28	7.95	24.2383	7.7	24.1798	7.8
27	-	23.8798	8.1	23.9441	8.0
29	7.17-7.05	24.5861	7.4	24.7363	7.2
30	6.82-6.76	25.4045	6.5	24.5438	7.4
39	3.89-3.70	28.1858	3.8	25.3911	6.5
38	3.89-3.70	28.1533	3.8	28.1066	3.8
37	3.89-3.70	27.7777	4.2	28.126	3.8
32	3.89-3.70	28.0180	3.9	27.7053	4.2
31	3.89-3.70	28.0352	3.9	27.9863	3.9
33	3.89-3.70	27.6460	4.3	27.9857	3.9
25	7.17-7.05	24.6049	7.3	27.6808	4.3
36	3.89-3.70	28.3043	3.6	28.0531	3.9
34	3.89-3.70	28.0007	3.9	28.4023	3.5
35	3.89-3.70	27.8828	4.1	28.0407	3.9
40	2.35	29.2756	2.7	29.2027	2.7

41	2.35	29.7639	2.2	29.2743	2.6
42	2.35	29.4015	2.5	29.9443	2.0

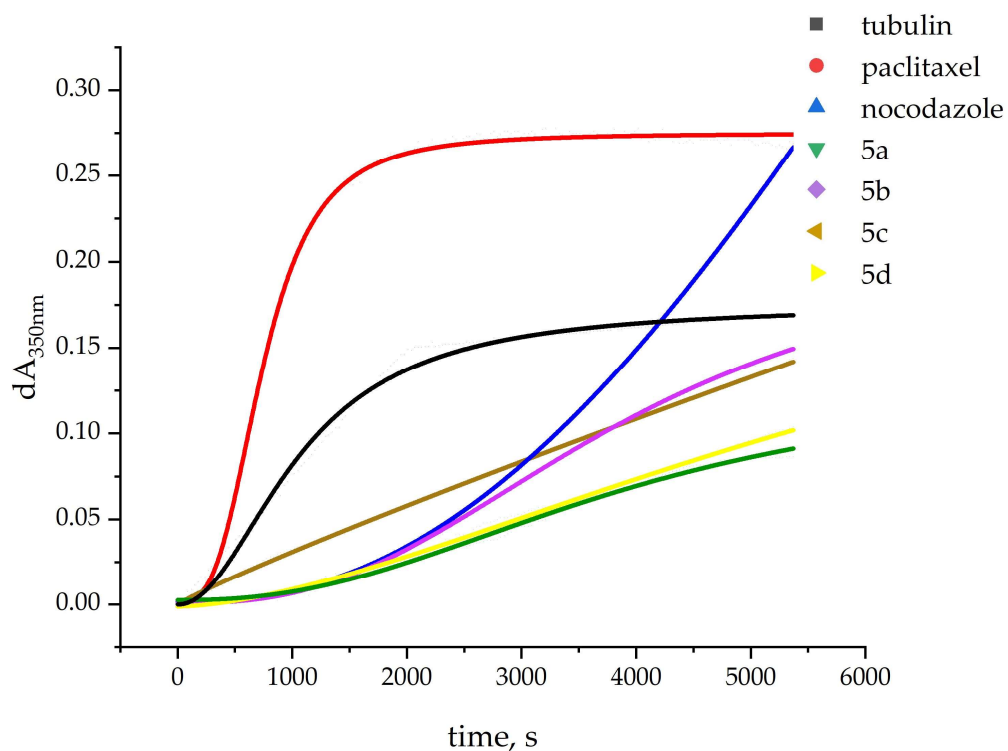


Figure S11. Tubulin (final 54 μM) in presence of PB-GTP buffer, 10 μM paclitaxel, 10 μM nocodazole and 10 μM of compounds **5a-d**. Reaction was conducted in a thermostated spec-trophotometric chamber at 37° C for 90 min. Turbidity was measured at 350 nm every 30 sec. Data for compounds **5a** and **5b** from [37].