

SUPPLEMENTAL MATERIAL FOR
Synthesis, biological evaluation, and molecular modeling studies of potent human neutrophil
elastase (HNE) inhibitors

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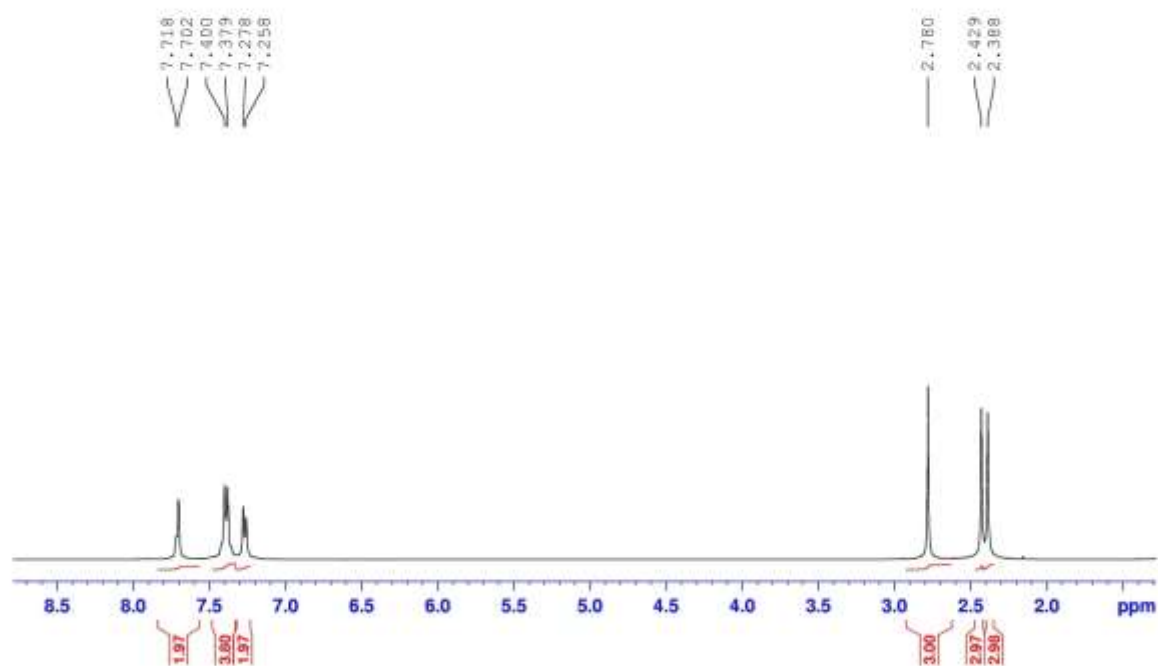
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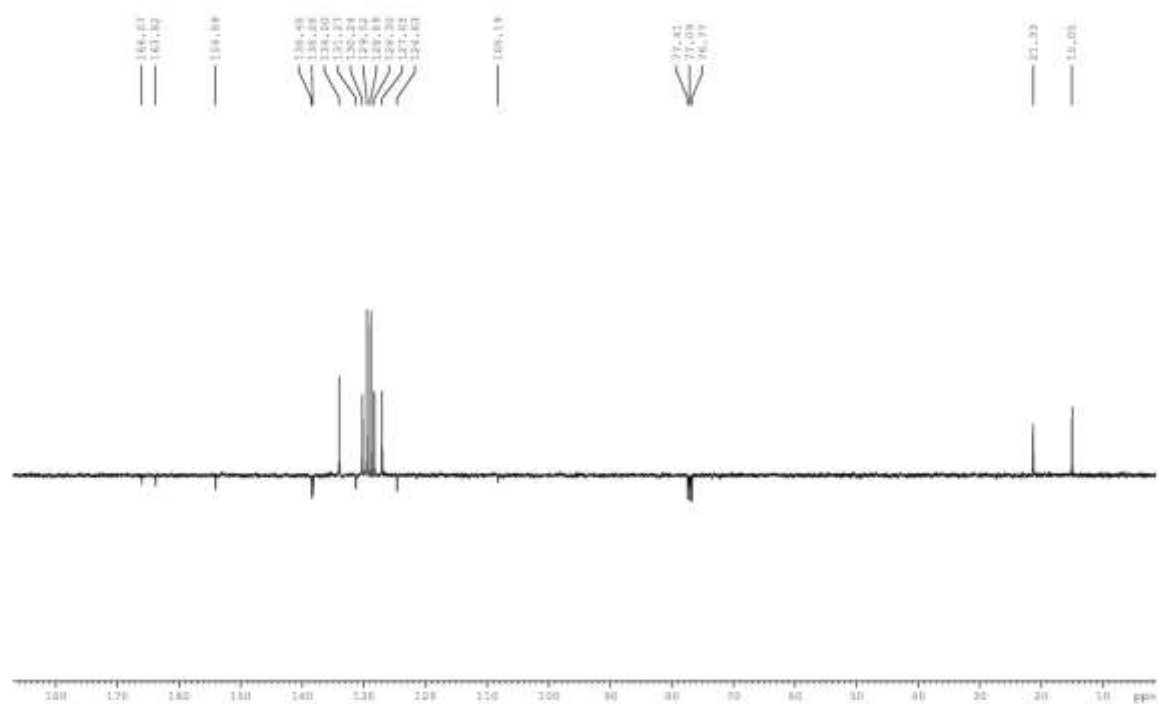
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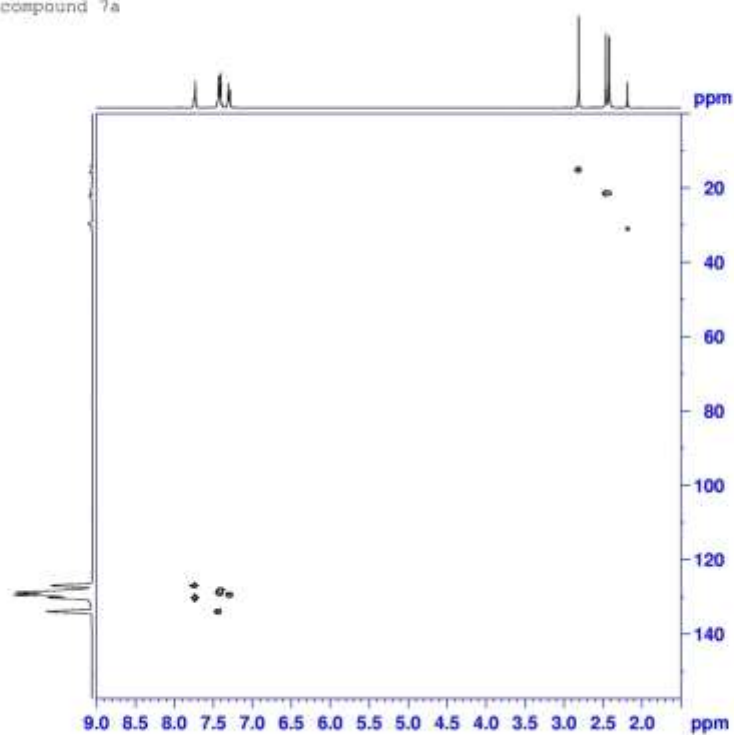
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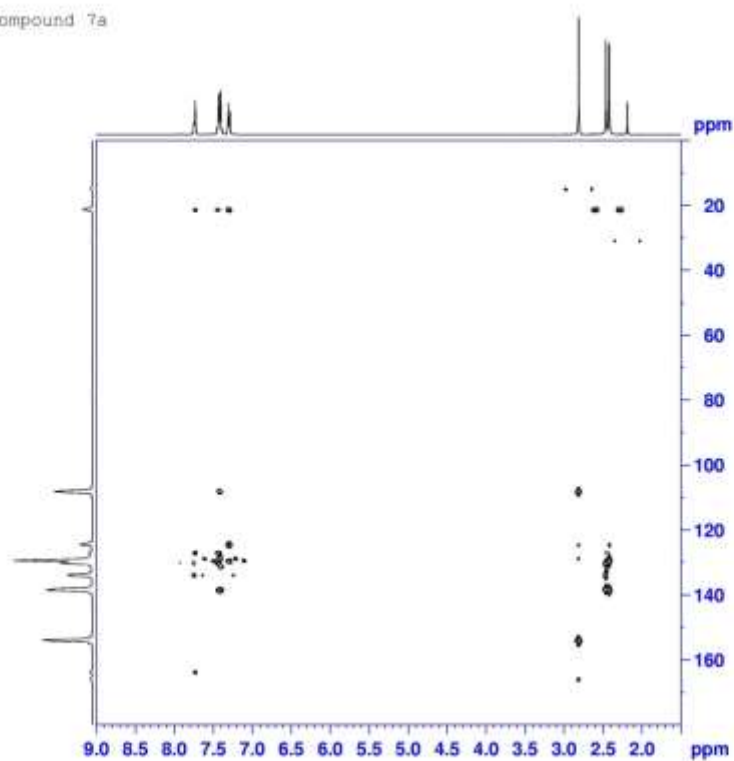
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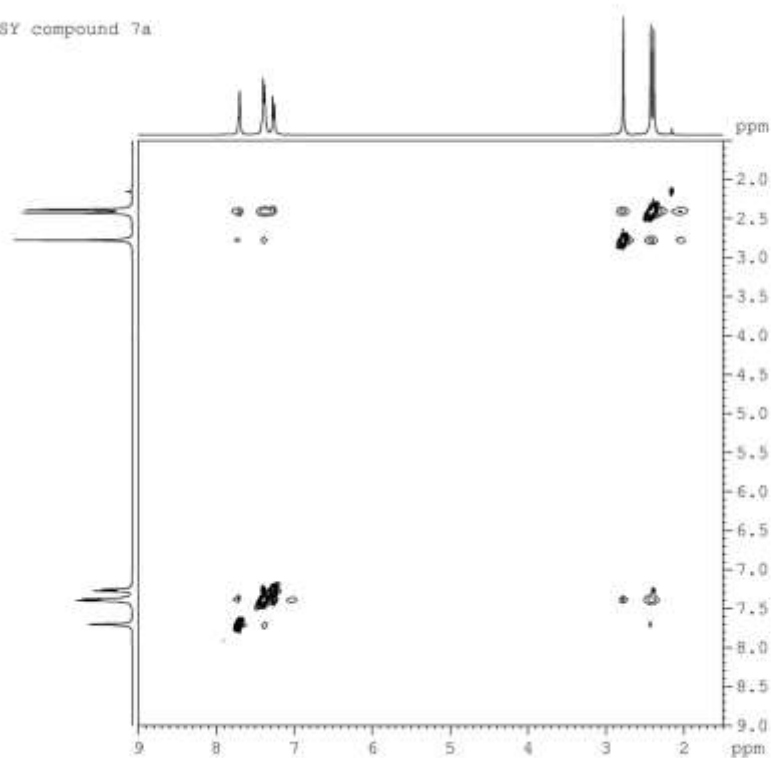
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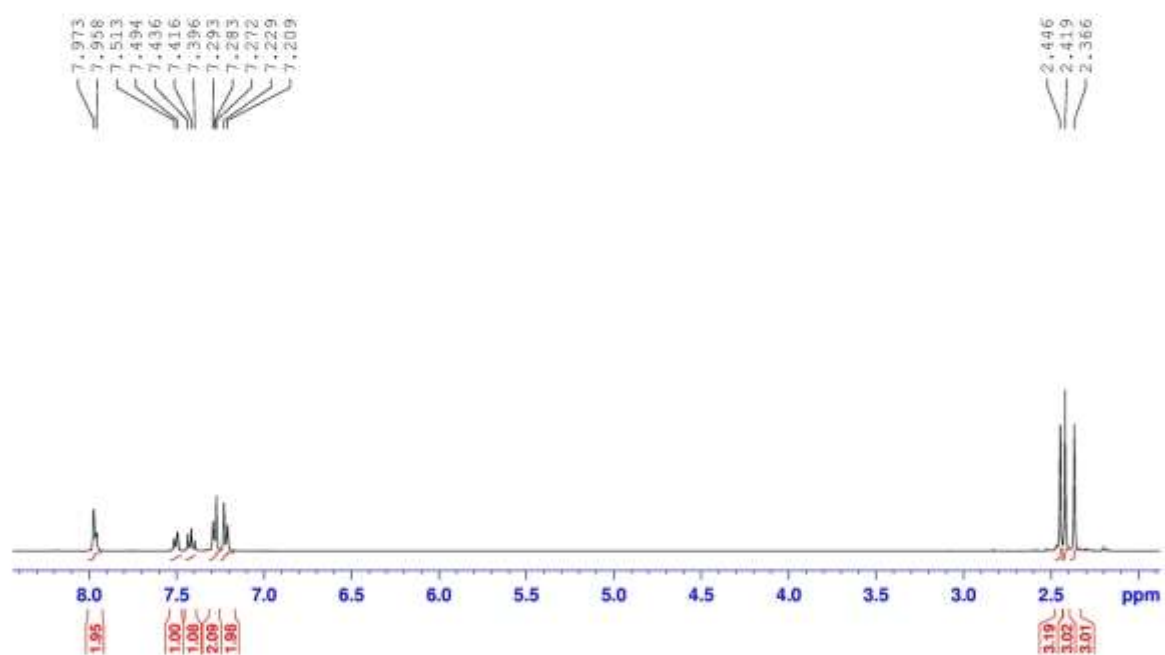
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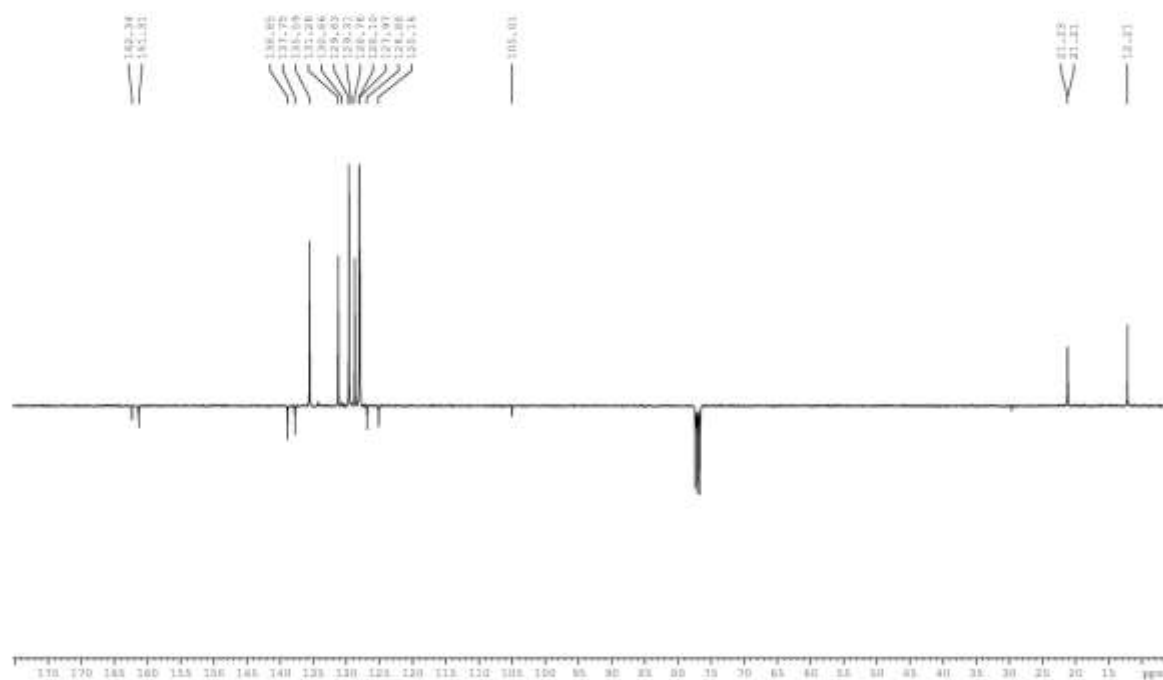
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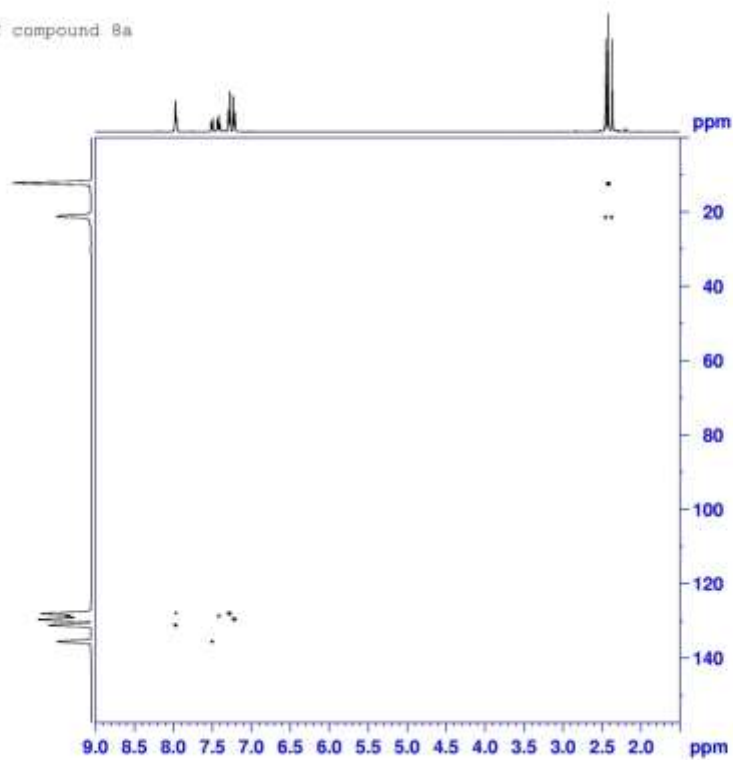
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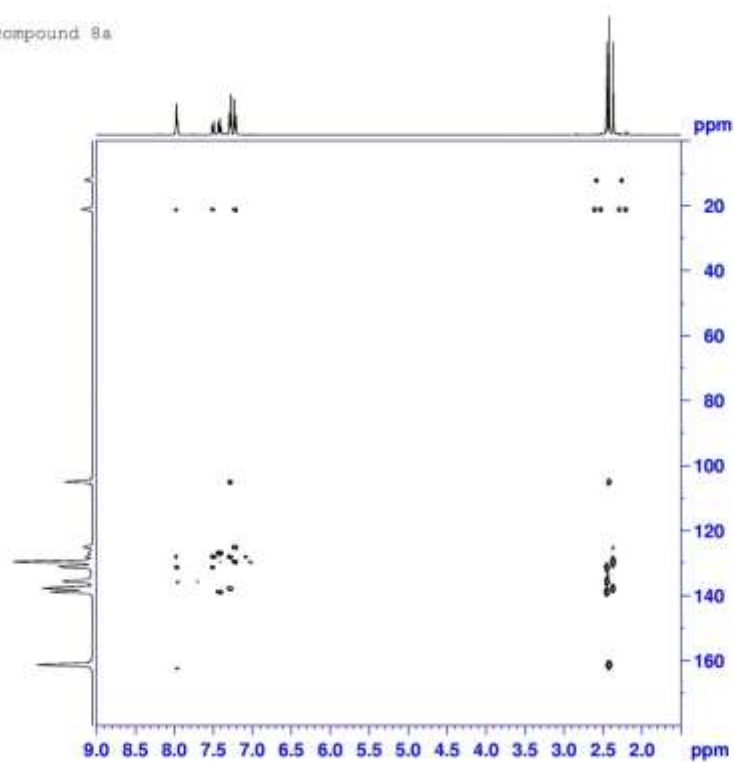
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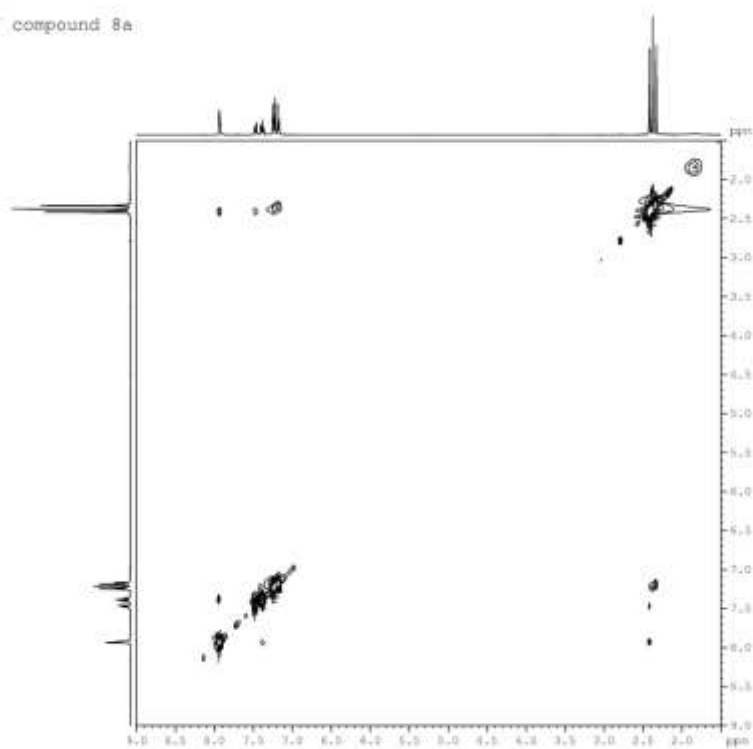
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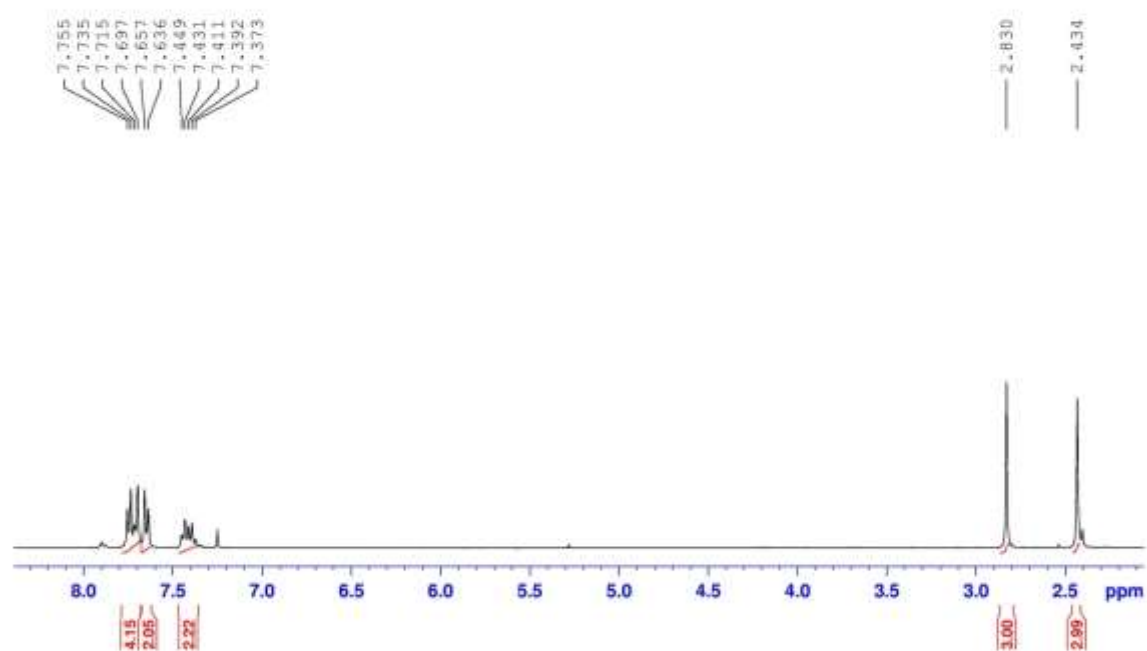
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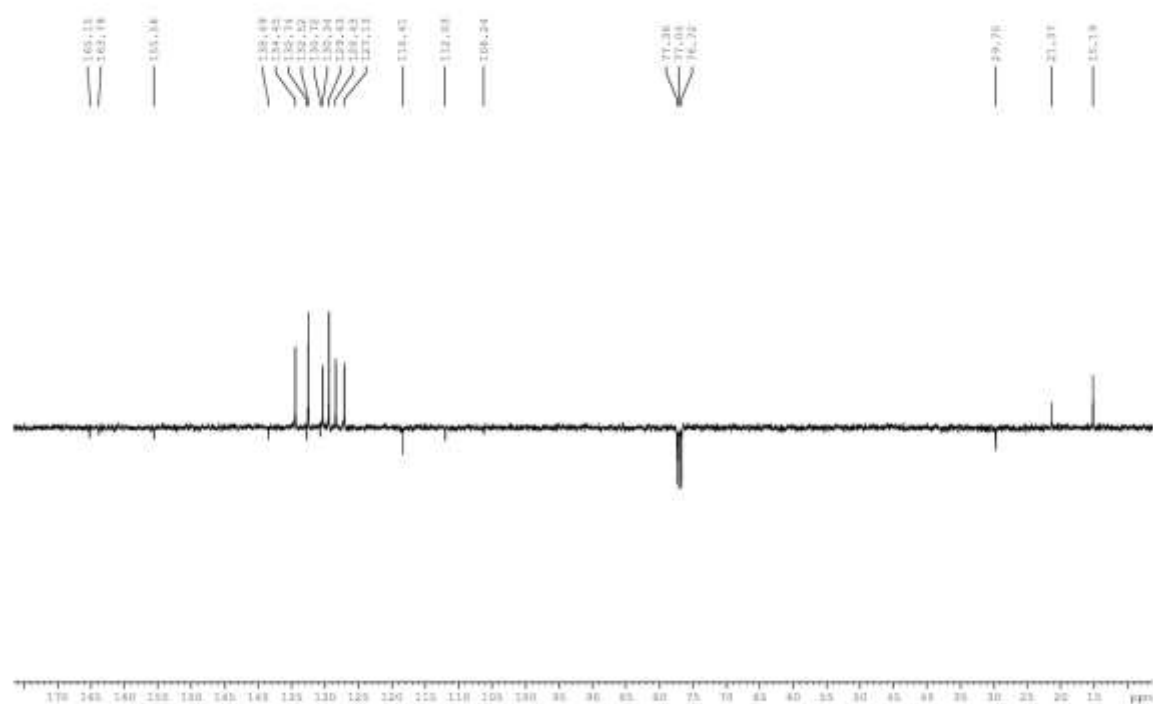
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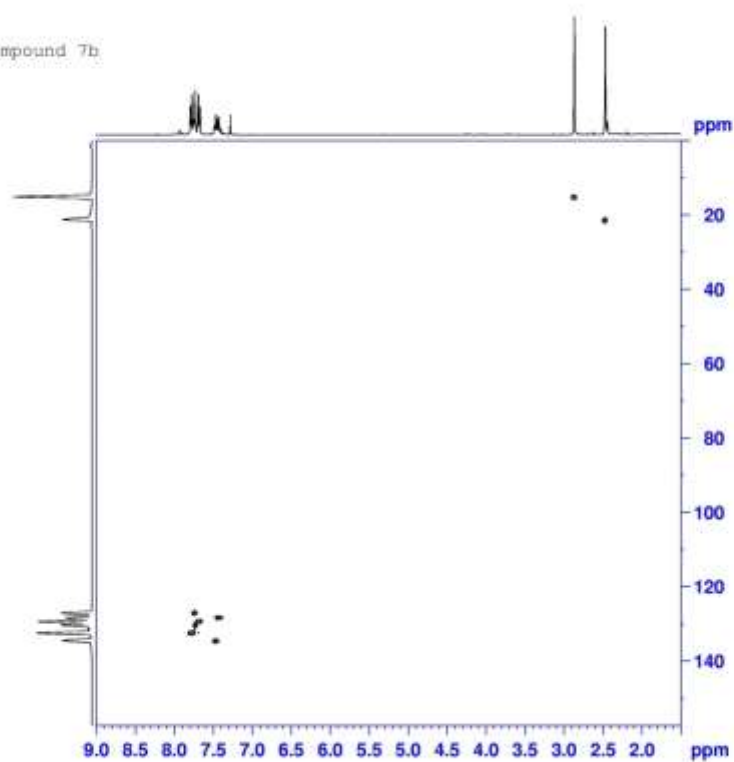
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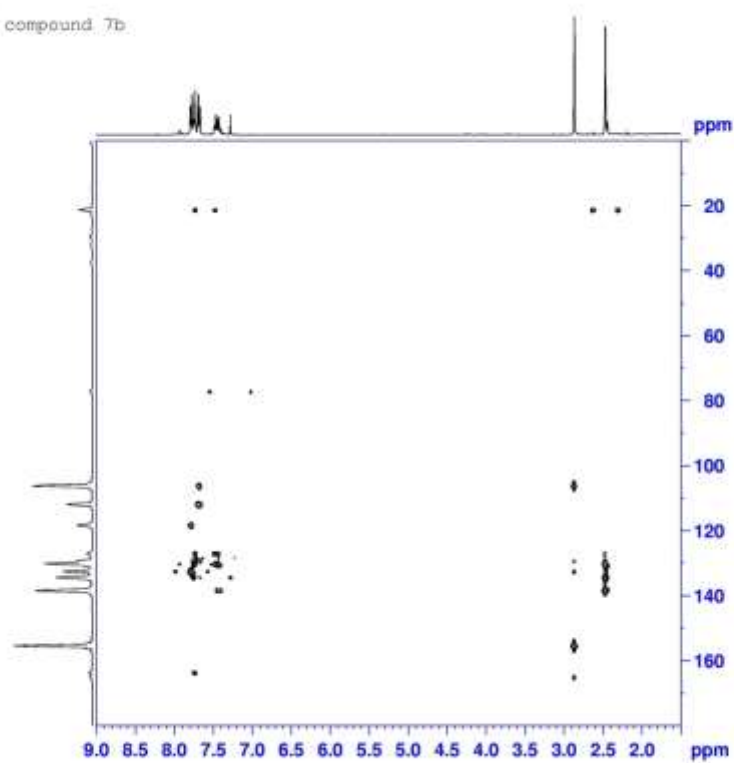
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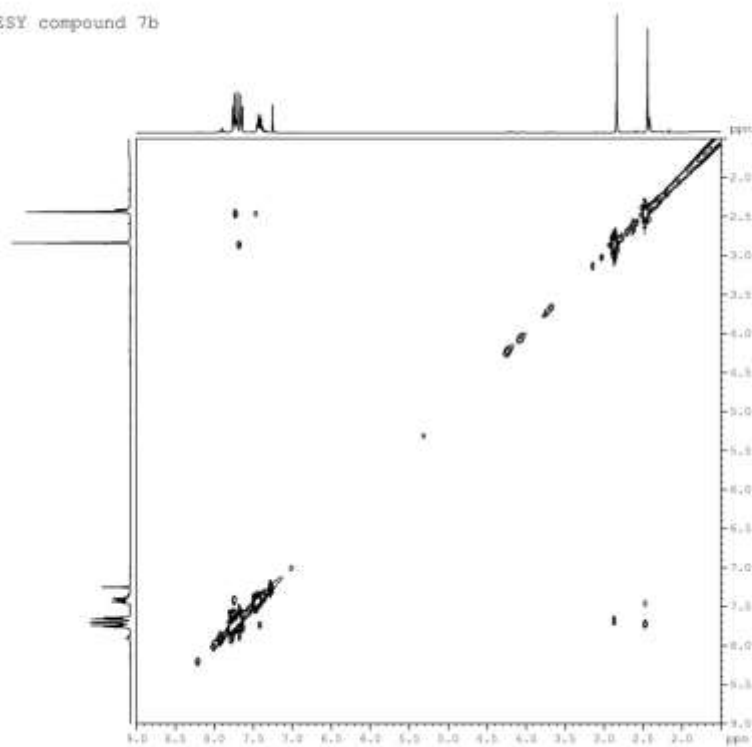
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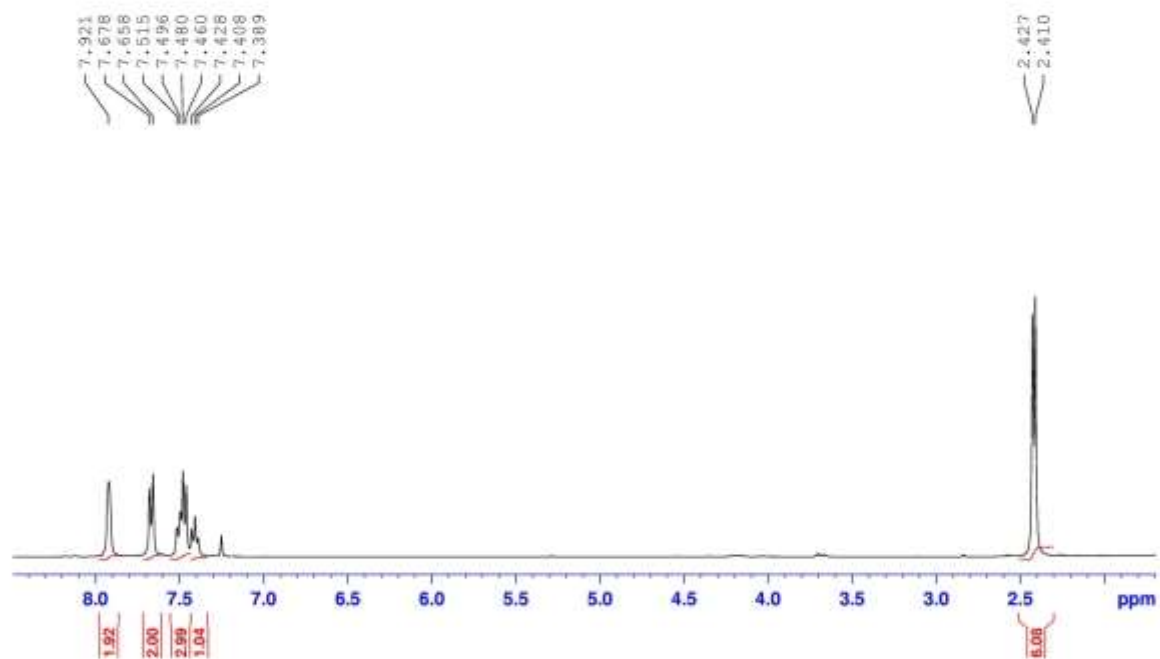
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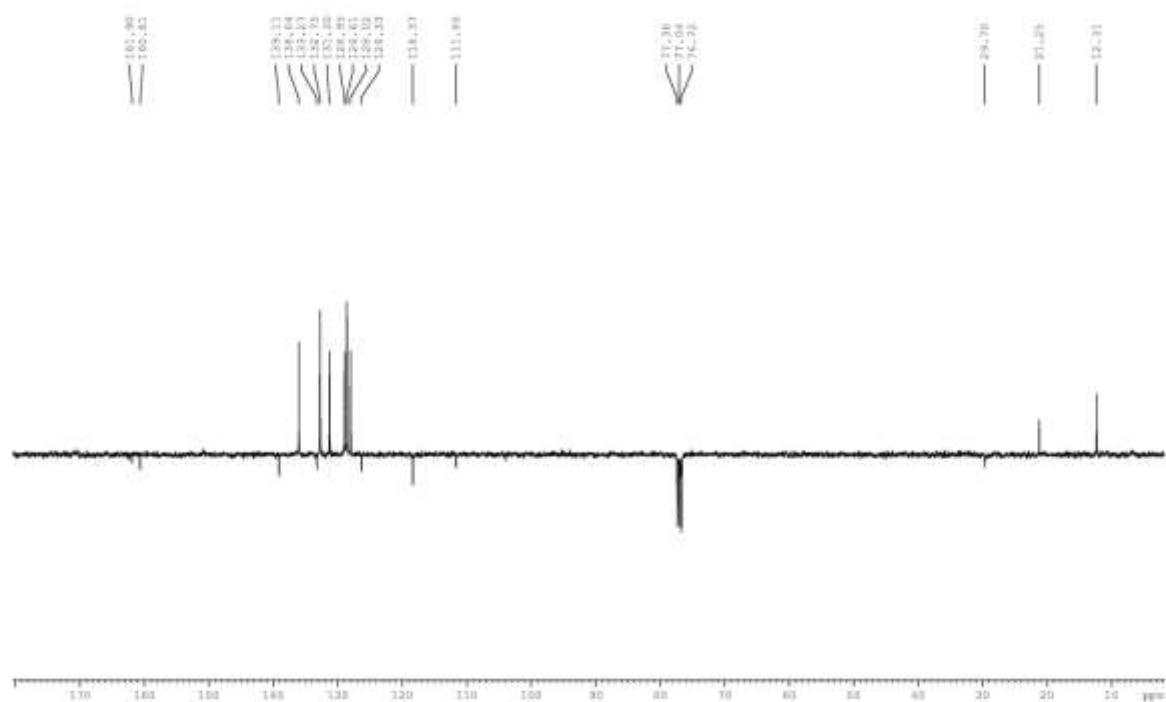
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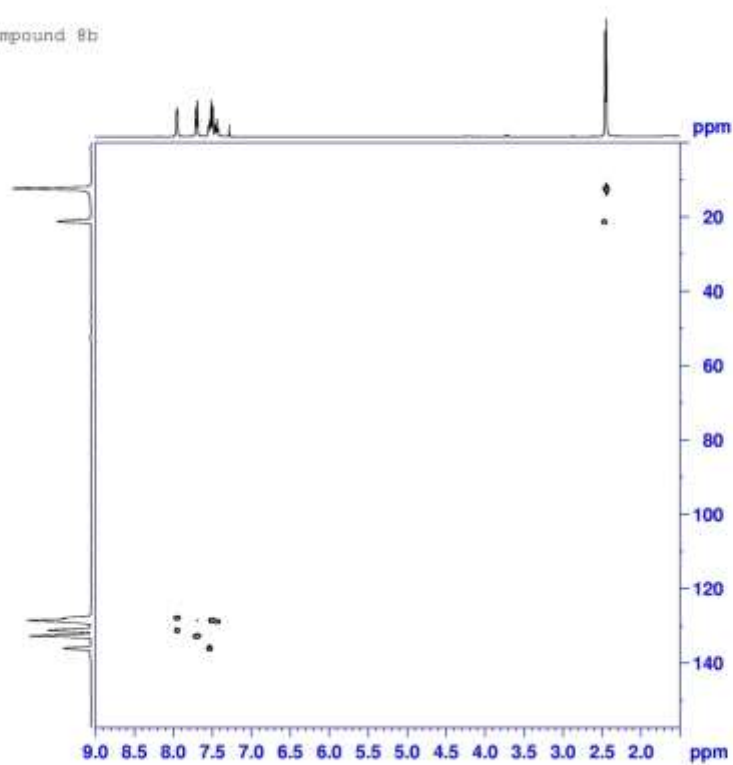
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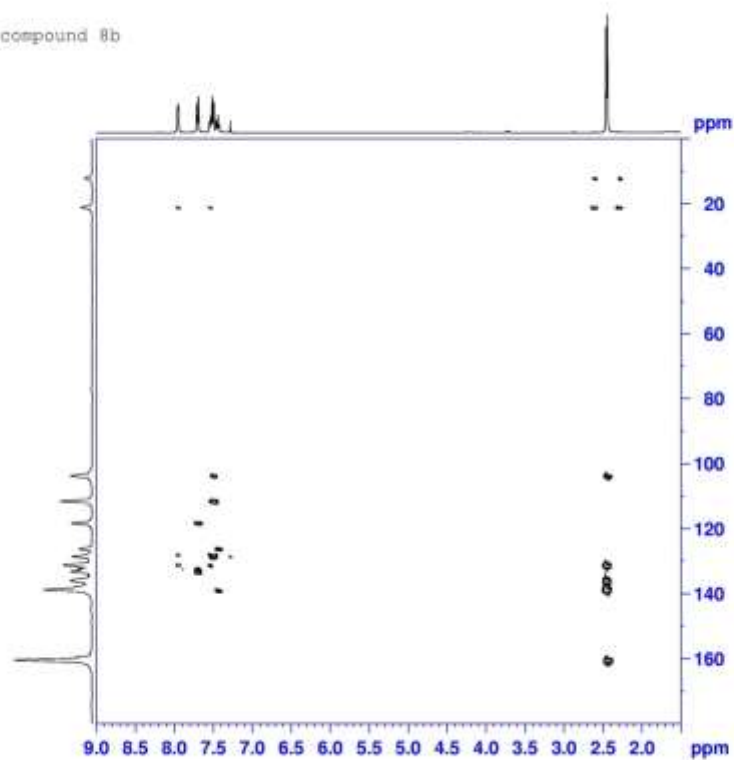
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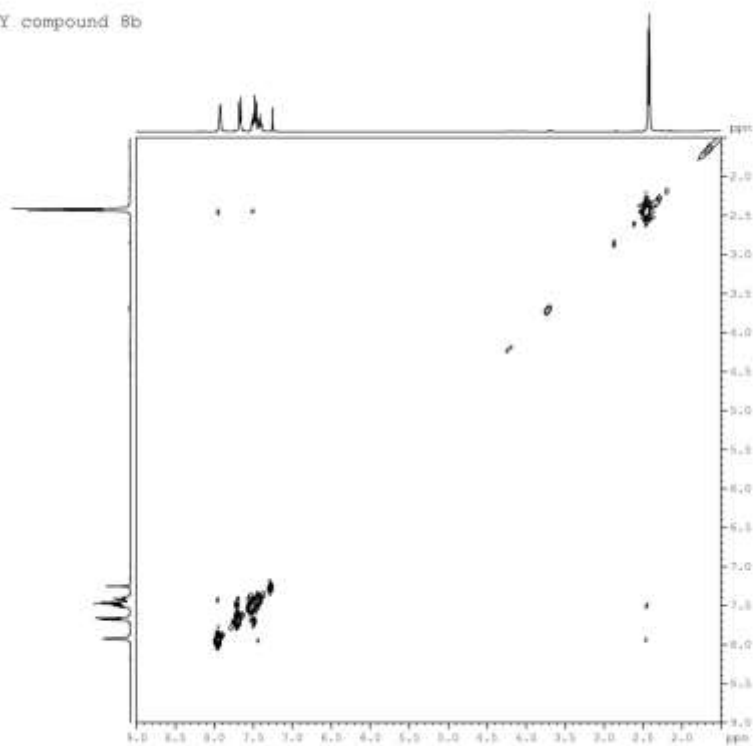
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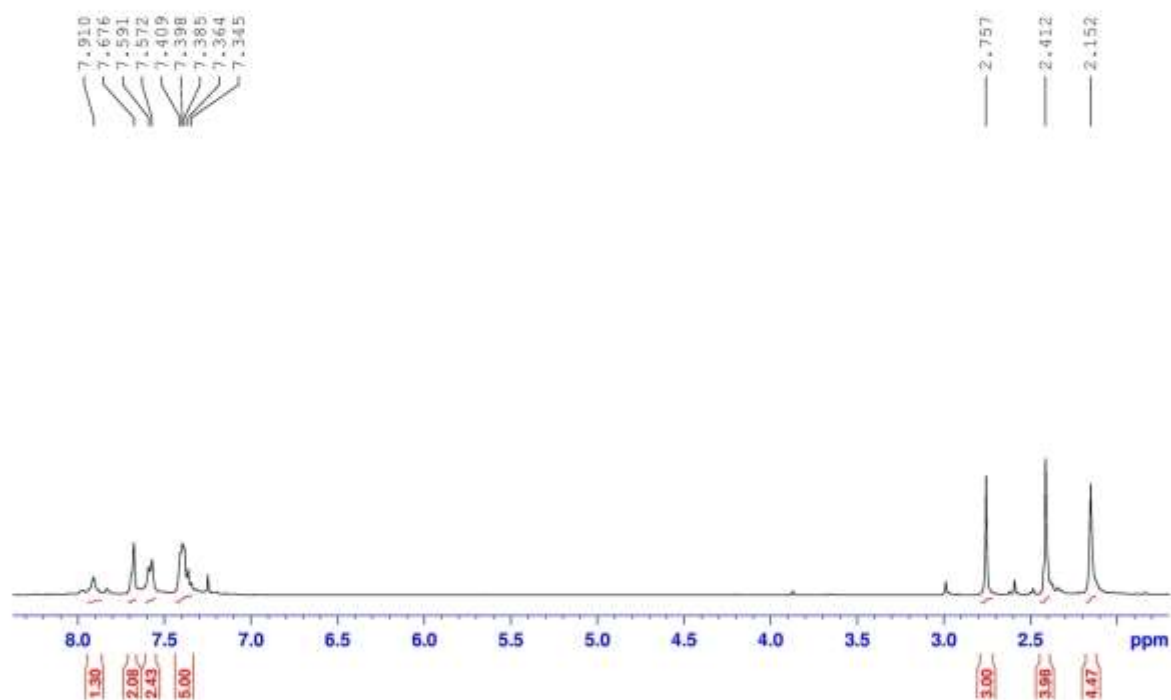
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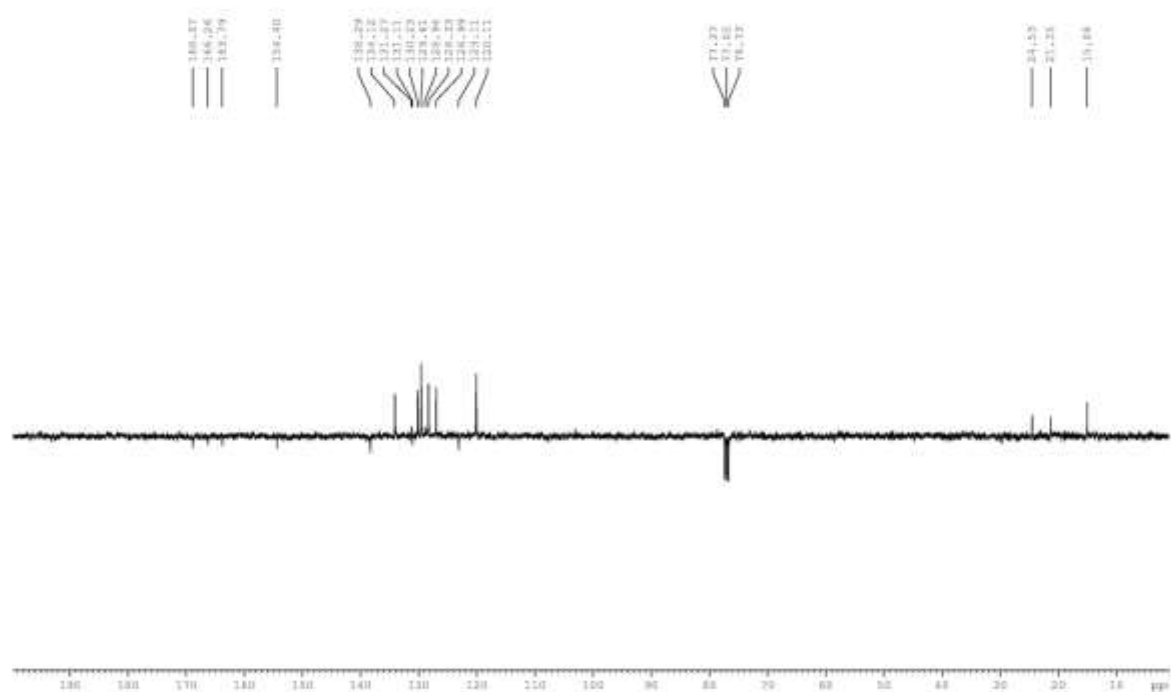
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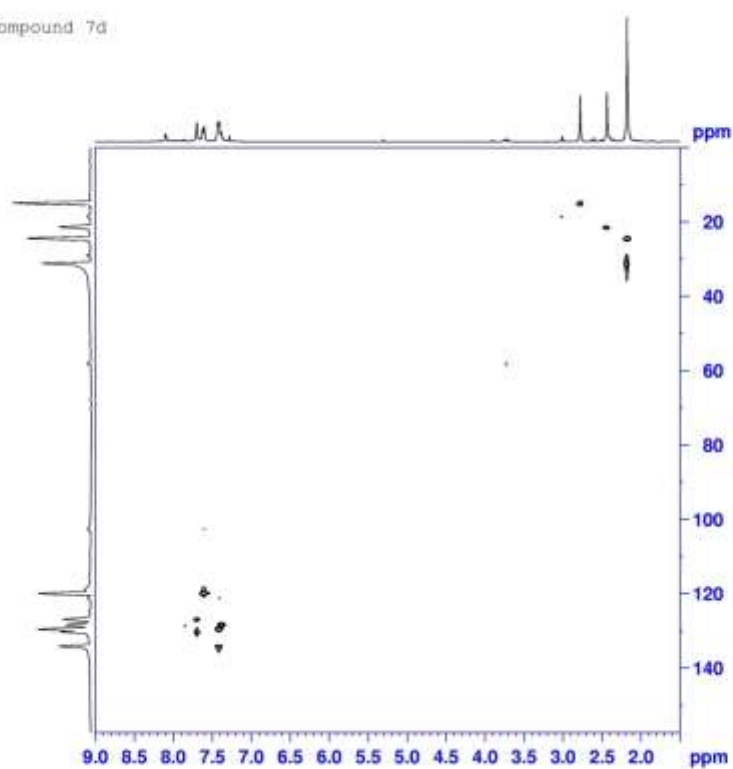
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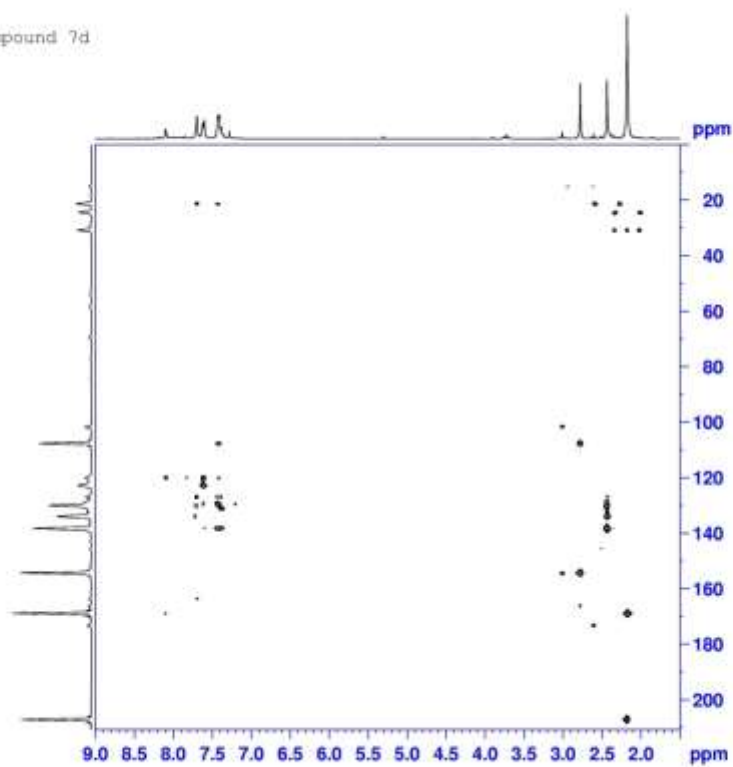
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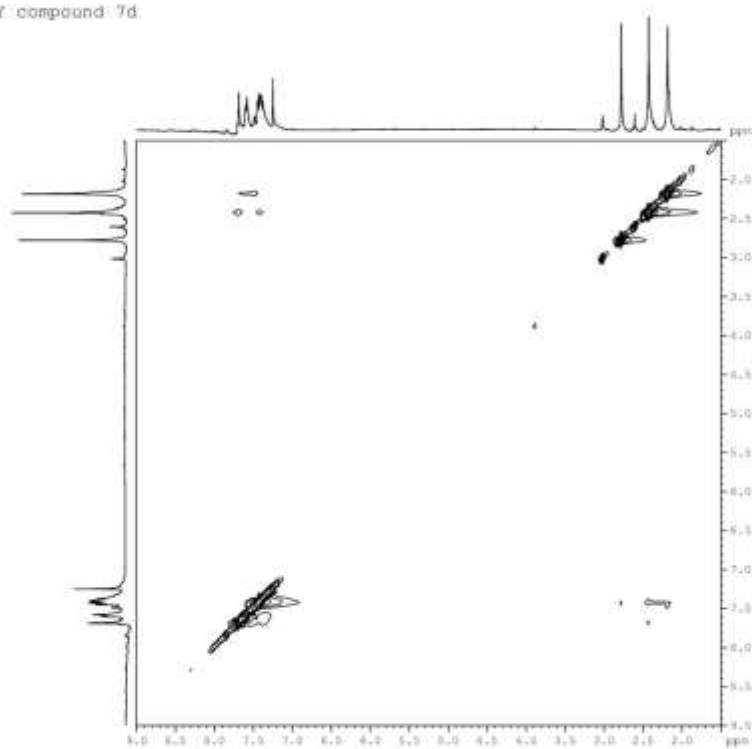
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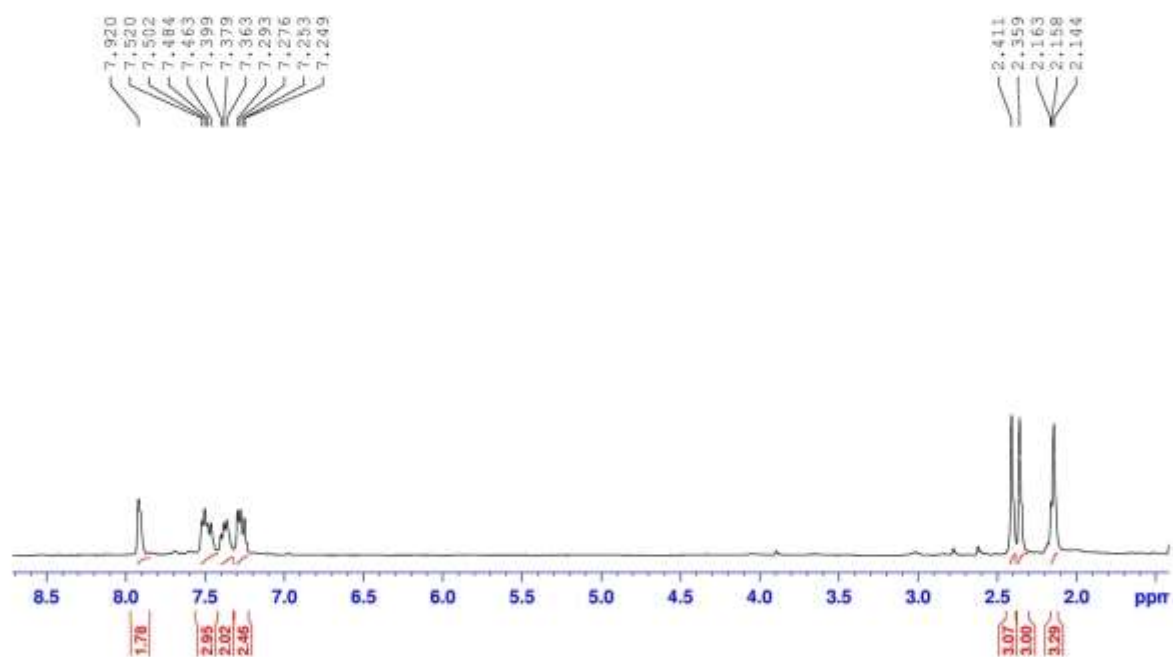
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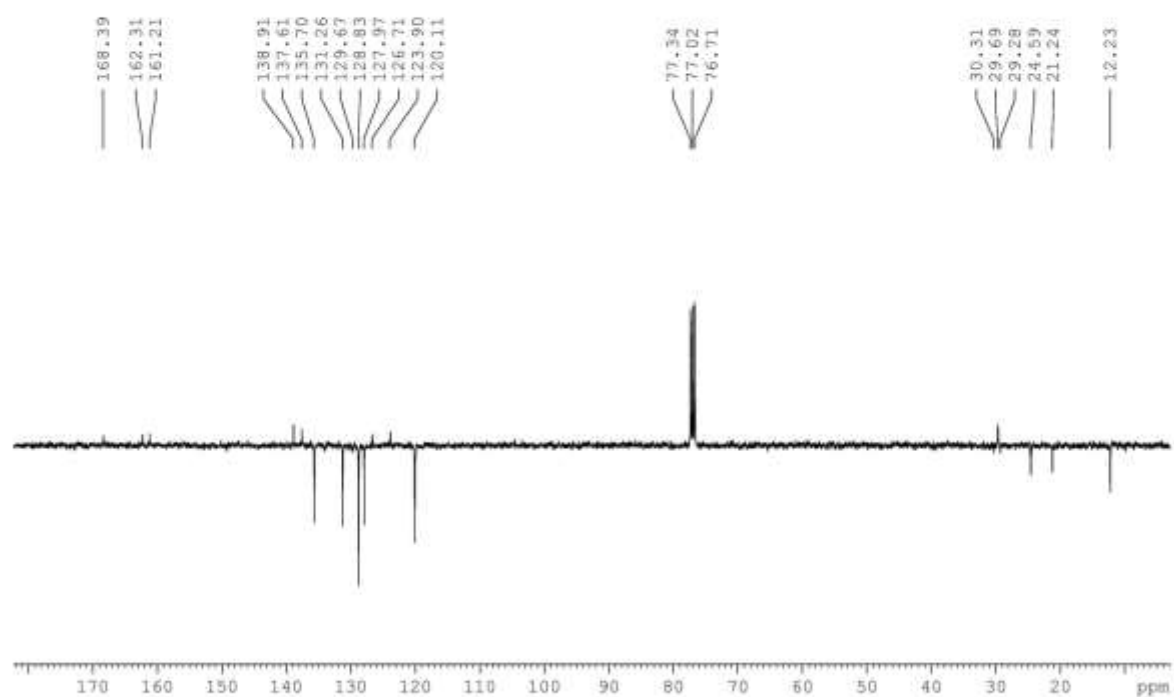
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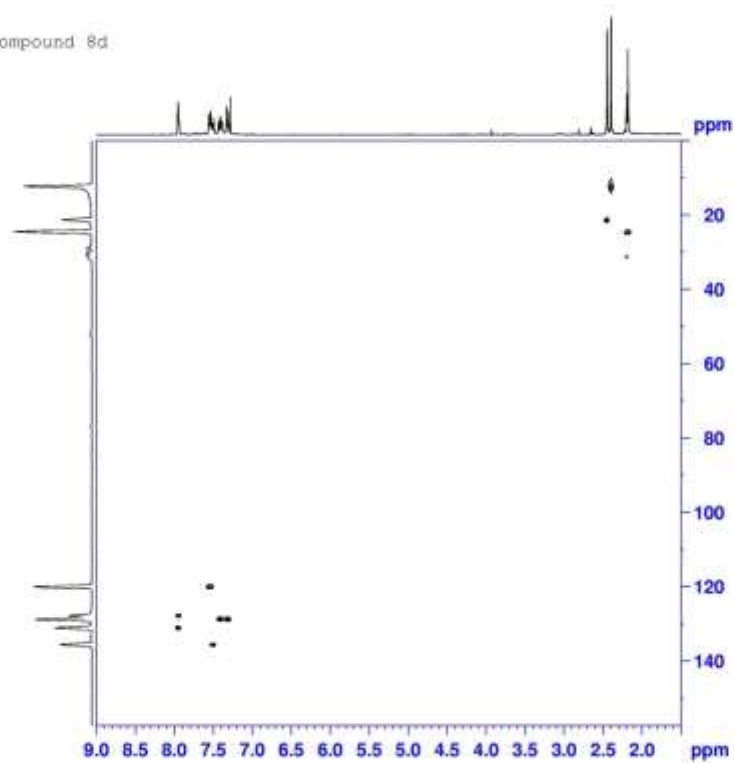
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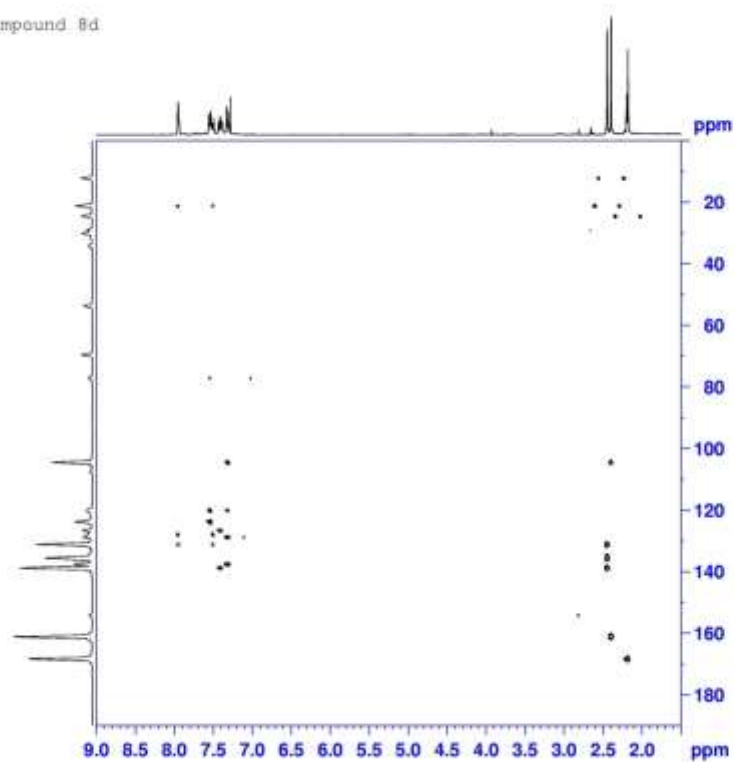
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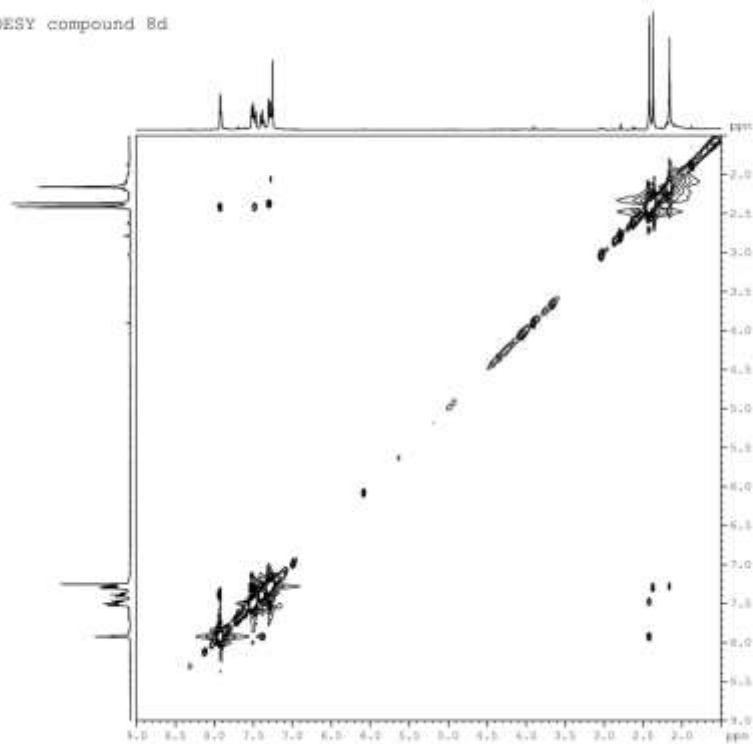
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HMBC compound 8d



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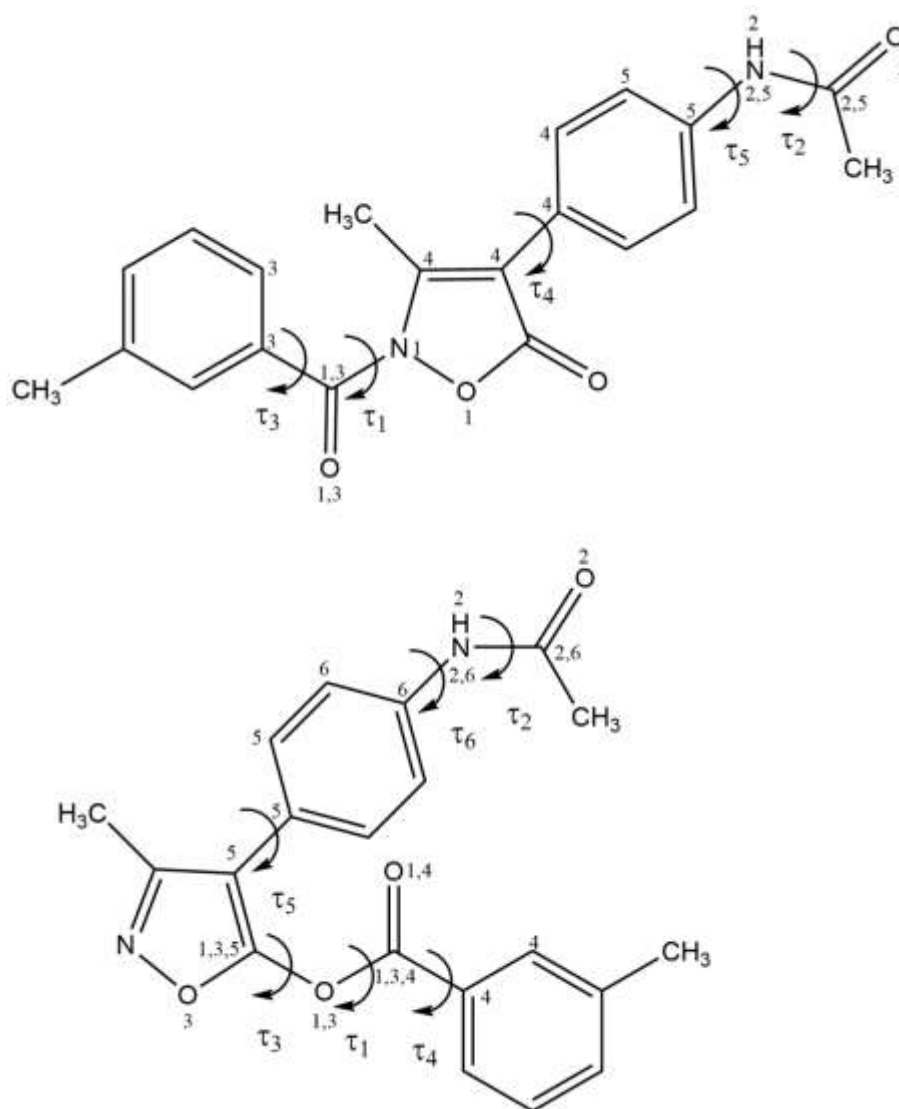


Fig. S1. Schematic drawing of **7d** and **8d** showing the torsion angles defining the molecular conformation monitored during MD simulations (numbers identify the set of atoms which define a given torsion).

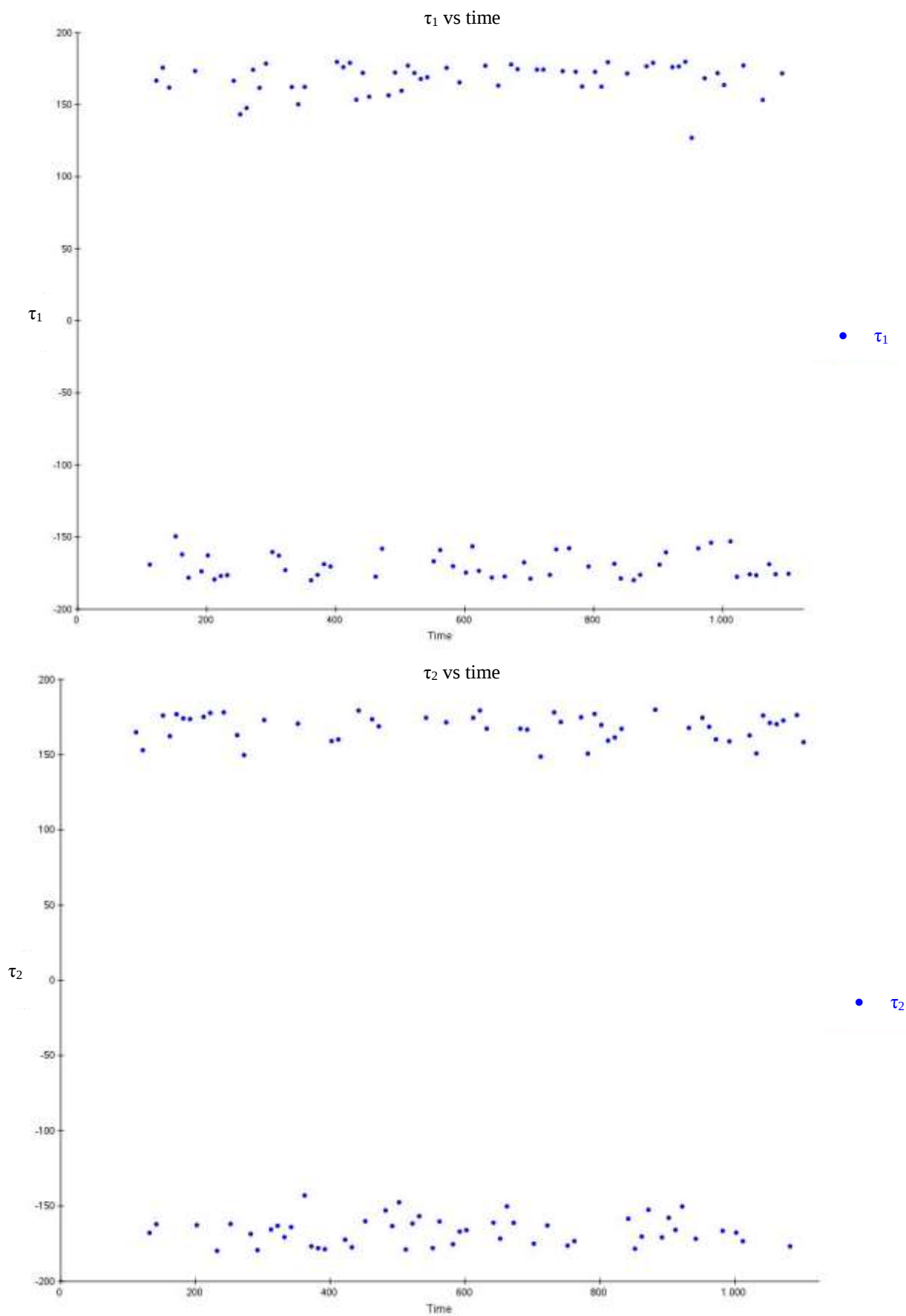


Fig. S2. Dihedral angle τ_1 - τ_2 distribution for **7d** during MD simulation (T=600K, $\epsilon=4r$).

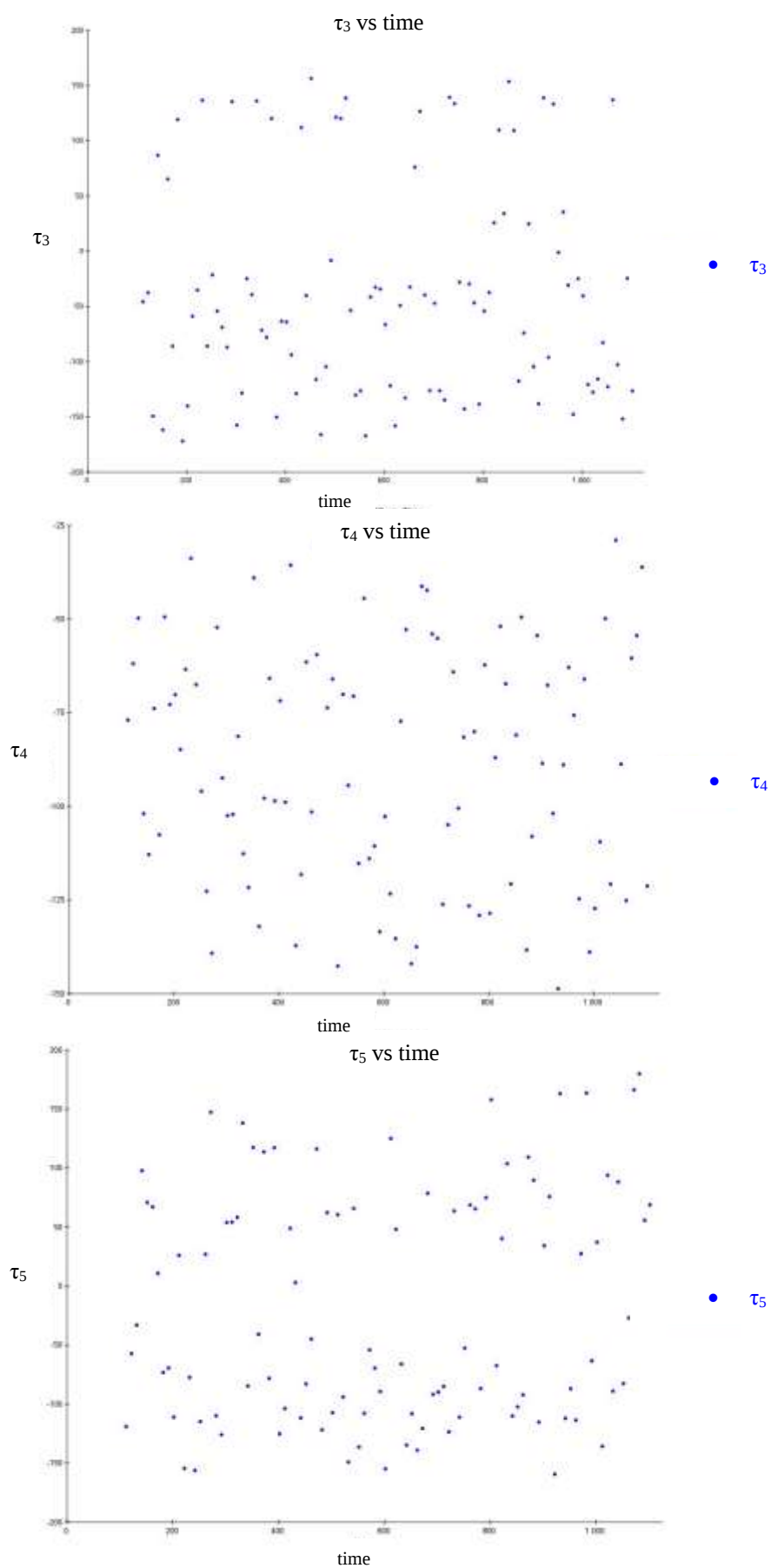


Fig. S3. Dihedral angle τ_3 - τ_5 distribution for **7d** during MD simulation (T=600K, $\epsilon=4r$).

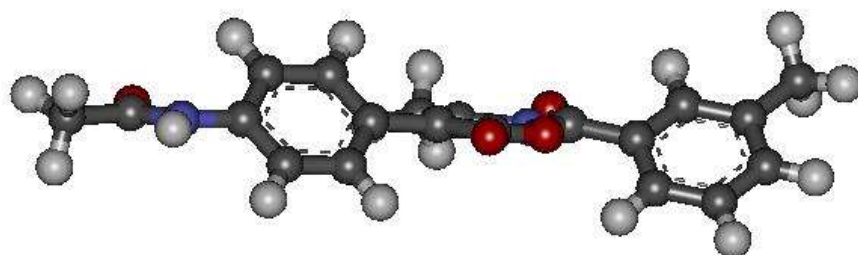


Fig. S4. View of the lowest energy conformer of **7d** as found from MD simulations (T=600K, $\epsilon=4\epsilon$).

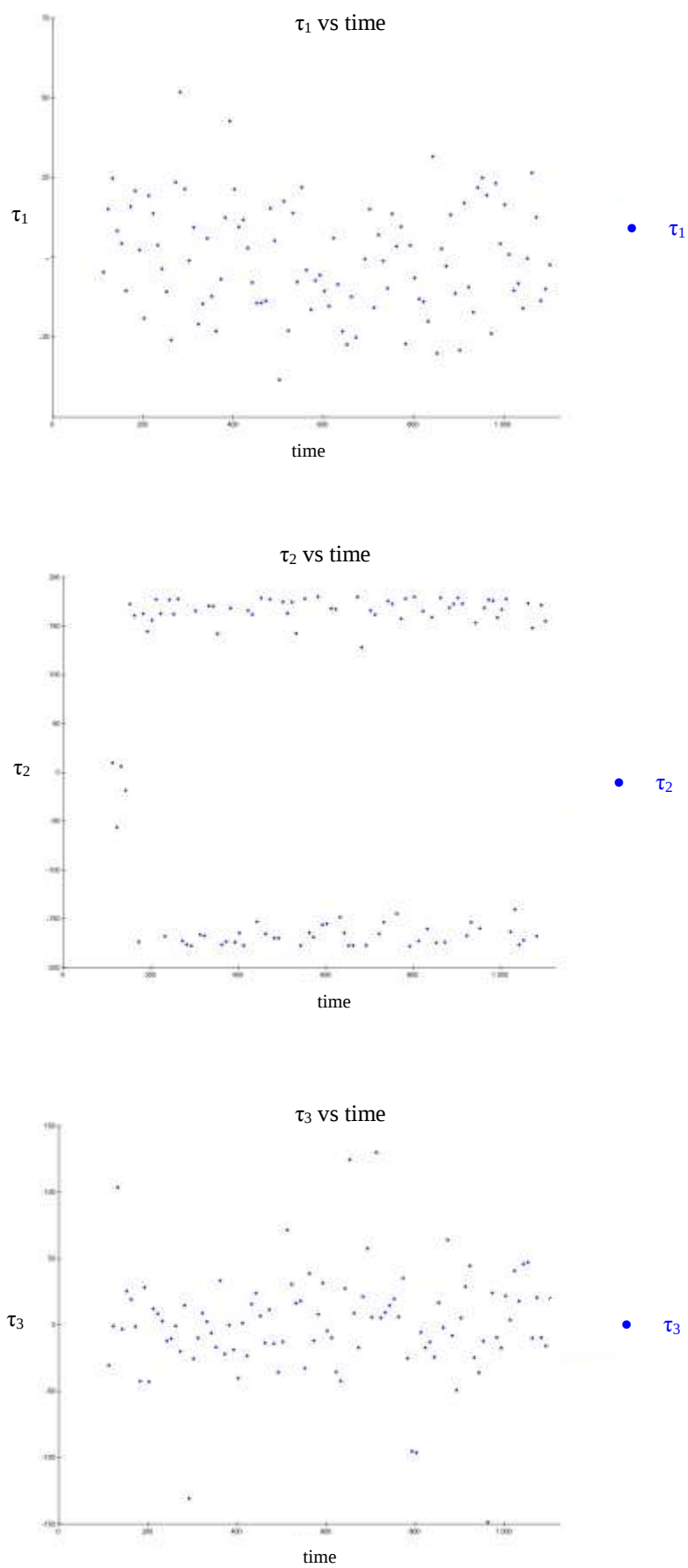


Fig. S5. Dihedral angle τ_1 - τ_3 distribution for **8d** during MD simulation (T=600K, $\epsilon=4r$).

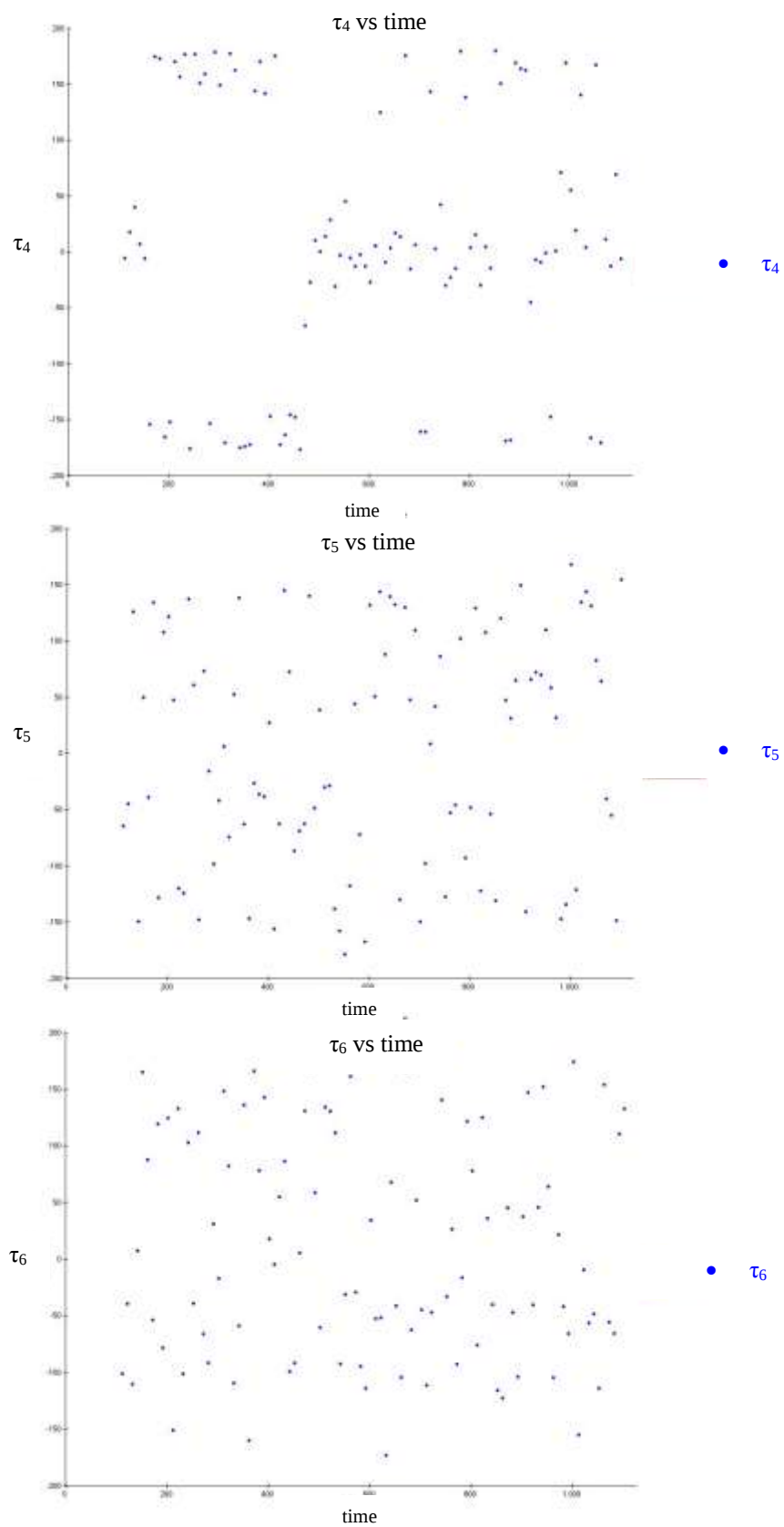


Fig. S6. Dihedral angle τ_4 - τ_6 distribution for **8d** during MD simulation (T=600K, $\epsilon=4r$).

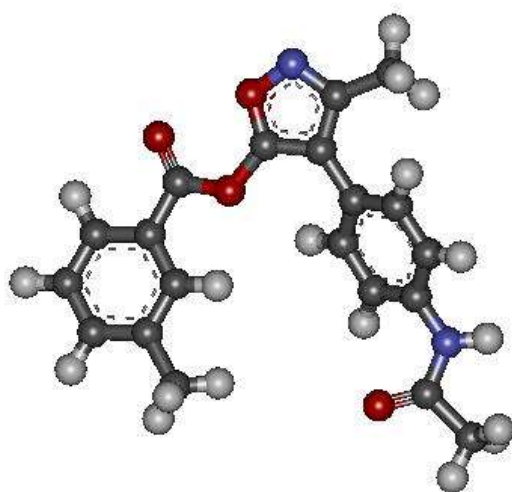


Fig. S7. View of the lowest energy conformer of **8d** as found from MD simulations (T=600K, $\epsilon=4$ r).



Fig. S8. View of the lowest energy conformer of **7d** as found from QC (B3LYP functional).

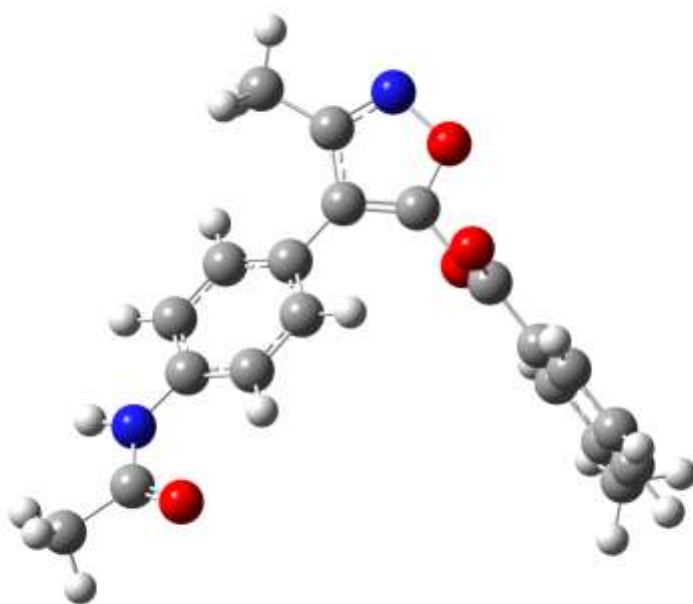


Fig. S9. View of the lowest energy conformer of **8d** as found from QC (B3LYP functional).

Table S1. Elemental analysis

Comp.	Formula (MW)	Anal. Calcd.				Anal. Found		
		C	H	N		C	H	N
2	C ₁₈ H ₁₇ NO ₂ (279.33)	77.40	6.13	5.01		77.94	6.15	5.03
3a	C ₁₆ H ₁₃ NO ₅ S (331.34)	58.00	3.95	4.23		58.23	3.96	4.25
3b	C ₂₁ H ₂₁ NO ₆ S (415.46)	60.71	5.09	3.37		60.95	5.11	3.38
3c	C ₂₁ H ₂₂ N ₂ O ₅ S (414.47)	60.85	5.35	6.76		61.09	5.37	6.78
4a	C ₁₈ H ₁₅ NO ₃ (293.32)	73.71	5.15	4.78		74.00	5.17	4.80
4b	C ₁₄ H ₁₃ NO ₃ (243.26)	69.12	5.39	5.76		69.39	5.41	5.78
4c	C ₁₈ H ₁₅ NO ₃ (293.32)	73.71	5.15	4.78		74.00	5.17	4.80
4d	C ₁₈ H ₁₅ NO ₃ (293.32)	73.71	5.15	4.78		74.00	5.17	4.80
4e	C ₁₈ H ₁₂ F ₃ NO ₃ (347.29)	62.25	3.48	4.03		62.50	3.49	4.05
4f	C ₁₈ H ₁₅ NO ₅ S (357.38)	60.49	4.23	3.92		60.73	4.25	3.93
4g	C ₁₈ H ₁₂ N ₂ O ₃ (304.30)	71.05	3.97	9.21		71.33	3.98	9.25
4h	C ₁₈ H ₁₂ N ₂ O ₃ (304.30)	71.05	3.97	9.21		71.33	3.98	9.25
4i	C ₂₂ H ₂₂ N ₂ O ₄ (378.42)	69.83	5.86	7.40		70.10	5.88	7.43
4l	C ₂₂ H ₂₁ NO ₅ (379.41)	69.64	5.58	3.69		69.91	5.60	3.70
4m	C ₂₃ H ₁₇ NO ₃ (355.39)	77.73	4.82	3.94		78.04	4.84	3.95
4n	C ₁₉ H ₁₅ NO ₃ (305.33)	74.74	4.95	4.59		75.03	4.97	4.61
4o	C ₁₈ H ₁₅ NO ₃ (293.32)	73.71	5.15	4.78		74.00	5.17	4.80
4p	C ₁₄ H ₁₃ NO ₃ (243.26)	69.12	5.39	5.76		69.39	5.41	5.78
4q	C ₁₇ H ₁₃ NO ₃ (279.29)	73.11	4.69	5.02		73.40	4.71	5.04
4r	C ₁₃ H ₁₁ NO ₃ (229.23)	68.11	4.84	6.11		68.38	4.86	6.13

4s	C ₁₇ H ₁₂ N ₂ O ₅ (324.29)	62.96	3.73	8.64		63.21	3.74	8.67
4t	C ₁₃ H ₁₀ N ₂ O ₅ (274.23)	56.94	3.68	10.22		57.16	3.69	10.26
4u	C ₁₉ H ₁₆ N ₂ O ₄ (336.34)	67.85	4.79	8.33		68.12	4.81	8.36
7a	C ₁₉ H ₁₇ NO ₃ (307.34)	74.25	5.58	4.56		74.55	5.60	4.58
7b	C ₁₉ H ₁₄ N ₂ O ₃ (318.33)	71.69	4.43	8.80		71.97	4.45	8.83
7c	C ₁₈ H ₁₄ N ₂ O ₅ (338.31)	63.90	4.17	8.28		64.15	4.19	8.31
7d	C ₂₀ H ₁₈ N ₂ O ₄ (350.37)	68.56	5.18	8.00		68.83	5.20	8.03
7e	C ₂₂ H ₂₀ N ₂ O ₄ (376.41)	70.20	5.36	7.44		70.48	5.38	7.47
8a	C ₁₉ H ₁₇ NO ₃ (307.34)	74.25	5.58	4.56		74.55	5.60	4.58
8b	C ₁₉ H ₁₄ N ₂ O ₃ (318.33)	71.69	4.43	8.80		71.97	4.45	8.83
8d	C ₂₀ H ₁₈ N ₂ O ₄ (350.37)	68.56	5.18	8.00		68.83	5.20	8.03
8e	C ₂₂ H ₂₀ N ₂ O ₄ (376.41)	70.20	5.36	7.44		70.48	5.38	7.47
10a	C ₁₀ H ₉ NO ₃ (191.18)	62.82	4.74	7.33		63.07	4.76	7.36
10b	C ₁₂ H ₁₃ NO ₃ (219.24)	65.74	5.98	6.39		66.00	6.00	6.41
10c	C ₁₅ H ₁₁ NO ₃ (253.25)	71.14	4.38	5.53		71.42	4.40	5.55