

A fragment-based approach for the development of G-quadruplex ligands: role of the amidoxime moiety

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Supplementary Material

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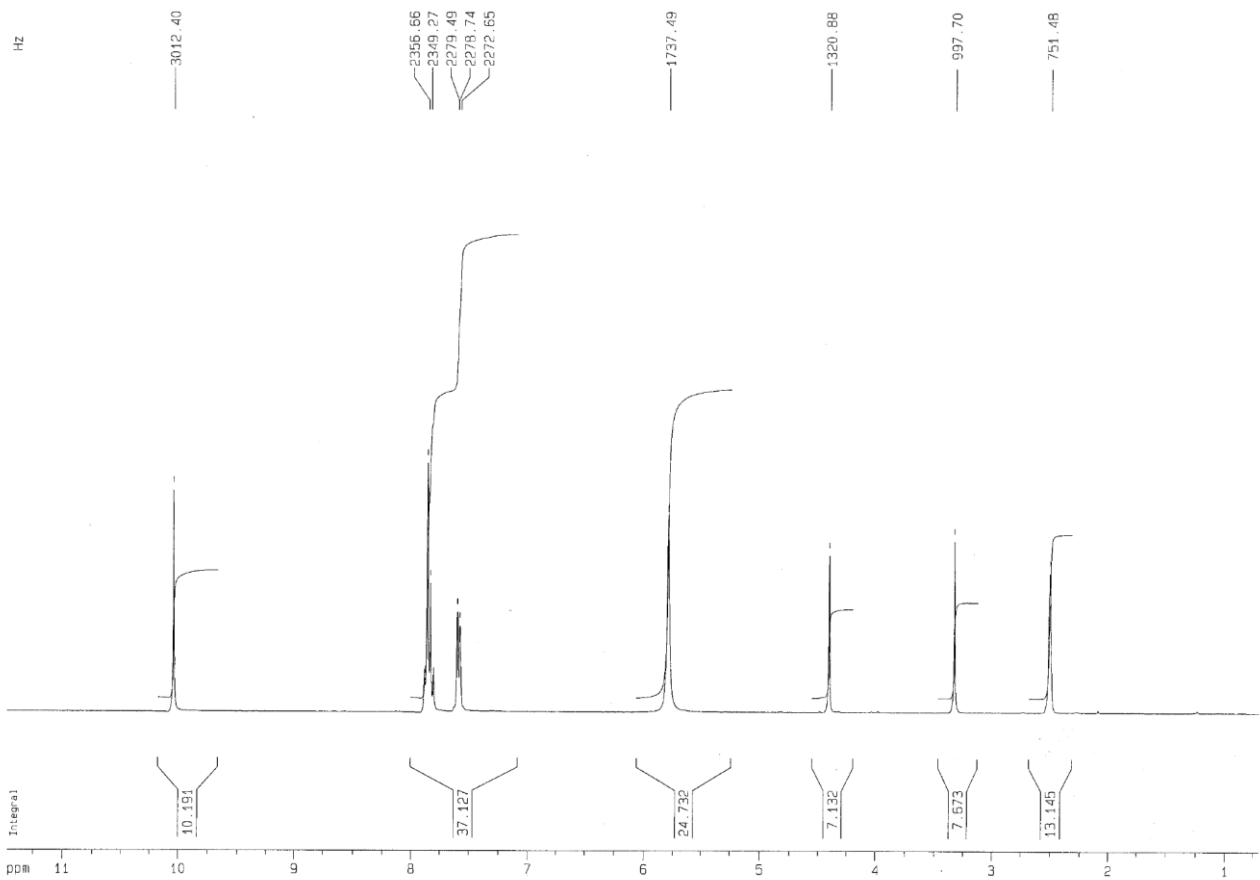
Table S1. Oligonucleotides used in this study

Application	Name	Sequence (5'→ 3')
FRET melting	LTR-III	FAM ¹ -GGGAGGCGTGGCCTGGGCGGGACTGGGG-TAMRA ²
	hTel	FAM ¹ -AGGGTTAGGGTTAGGGTTAGGG-TAMRA ²
	dsDNA	FAM ¹ -CTATAGCGCGCTATAG-TAMRA ²
FRET competition	LTR-III labeled	FAM ¹ -GGGAGGCGTGGCCTGGGCGGGACTGGGG-TAMRA ²
	LTR-III unlabeled	GGGAGGCGTGGCCTGGGCGGGACTGGGG
	hTel unlabeled	AGGGTTAGGGTTAGGGTTAGGG
CD	LTR-III	GGGAGGCGTGGCCTGGGCGGGACTGGGG
	hTel	AGGGTTAGGGTTAGGGTTAGGG
<i>Taq</i> polymerase stop assay	Taq primer	GGCAAAAAGCAGCTGCTTATATGCAG
	non-G4 cnt	TTGTCGTTAAAGTCTGACTGCGAGCTCTCAGATCCTGCATAT
		AAGCAGCTGCTTTTTGCC
	LTR-III	TTTTTGGGAGGCGTGGCCTGGGCGGGACTGGGGTTTTTCTGC
		ATATAAGCAGCTGCTTTTTGCC
hTel	TTTTTGGGTTAGGGTTAGGGTTAGGGTTTTTCTGCATATAAGC GCTTTTTGCC	

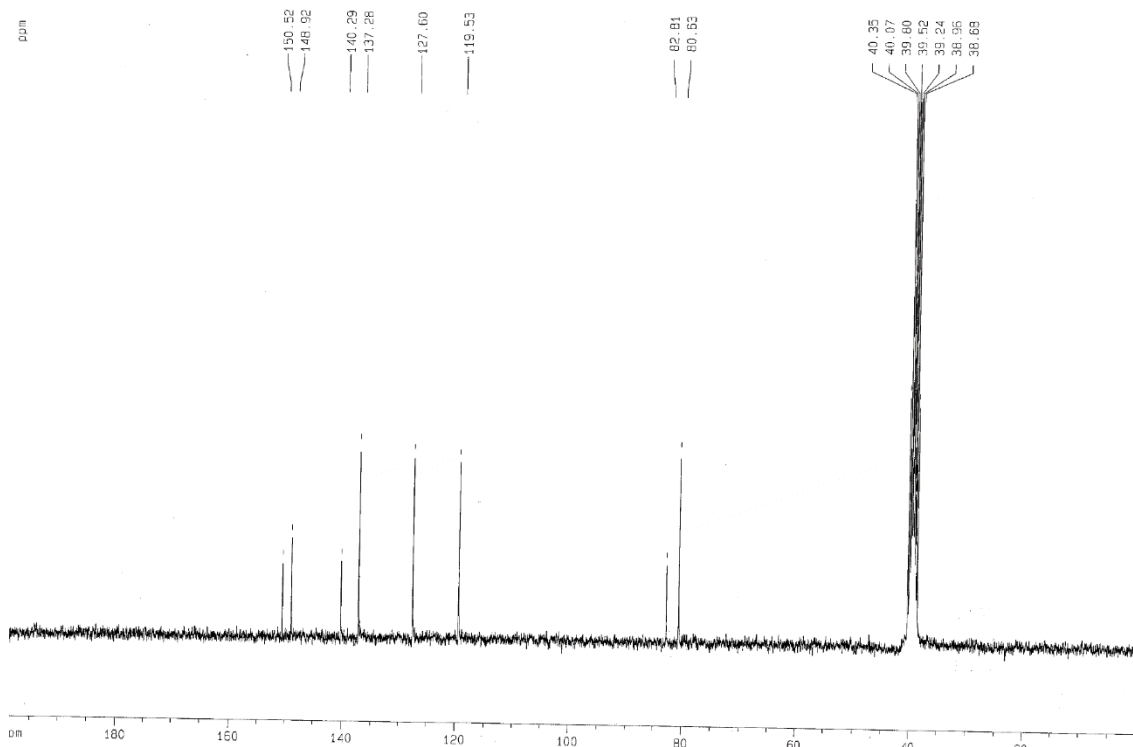
¹ 6-carboxyfluorescein. ² 6-carboxy-tetramethylrhodamine.

NMR characterization:

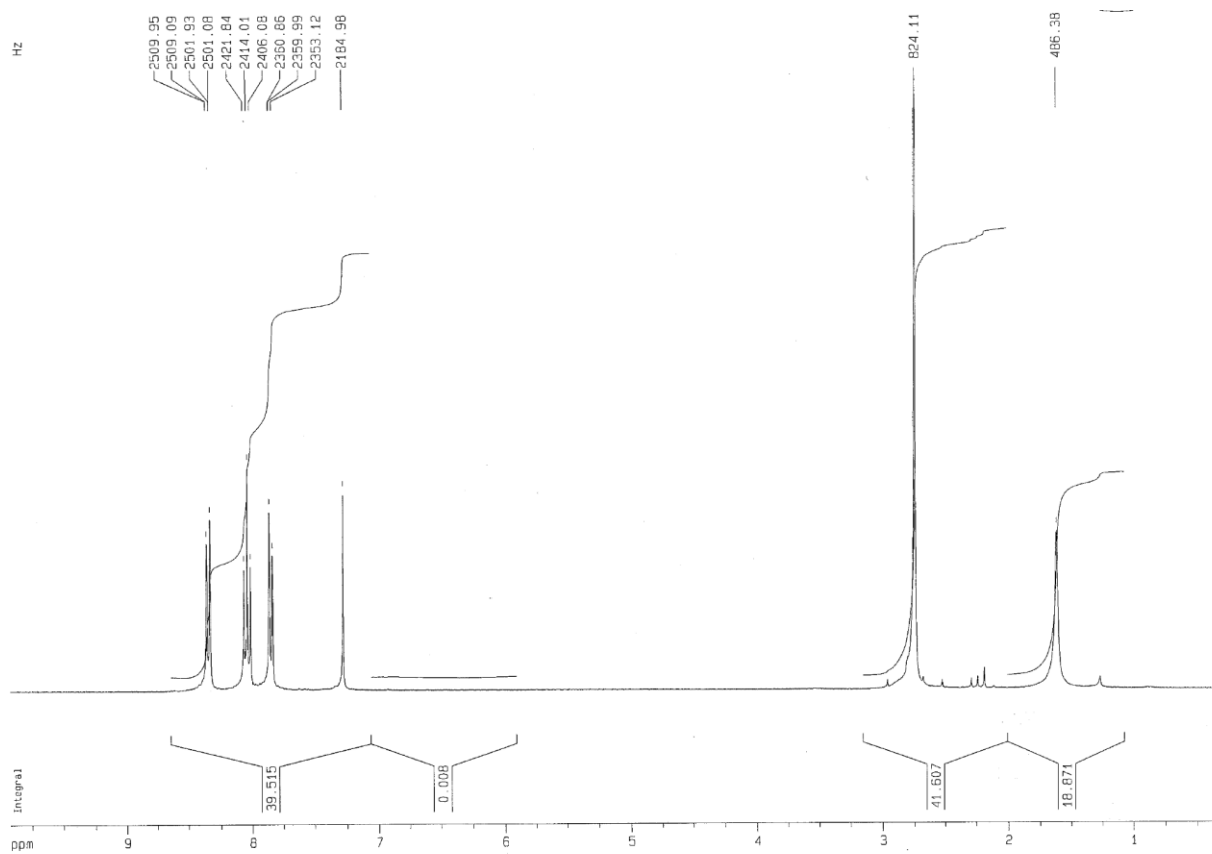
¹H-NMR – DMSO d6: Fragment 5



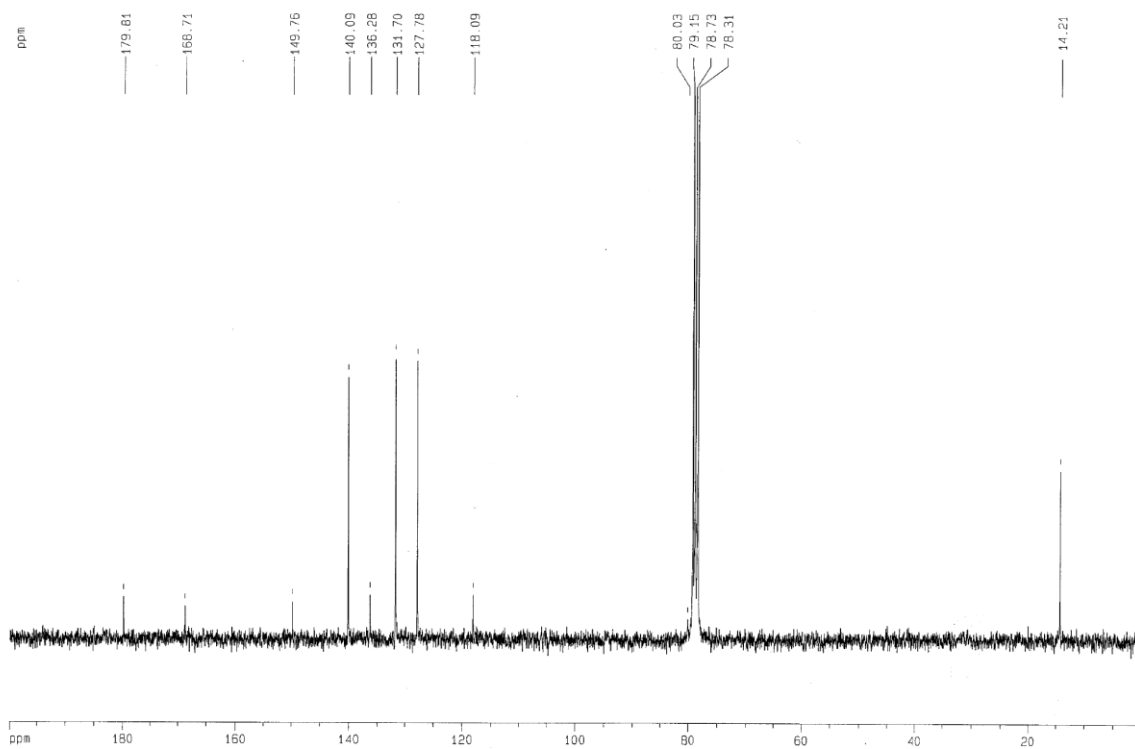
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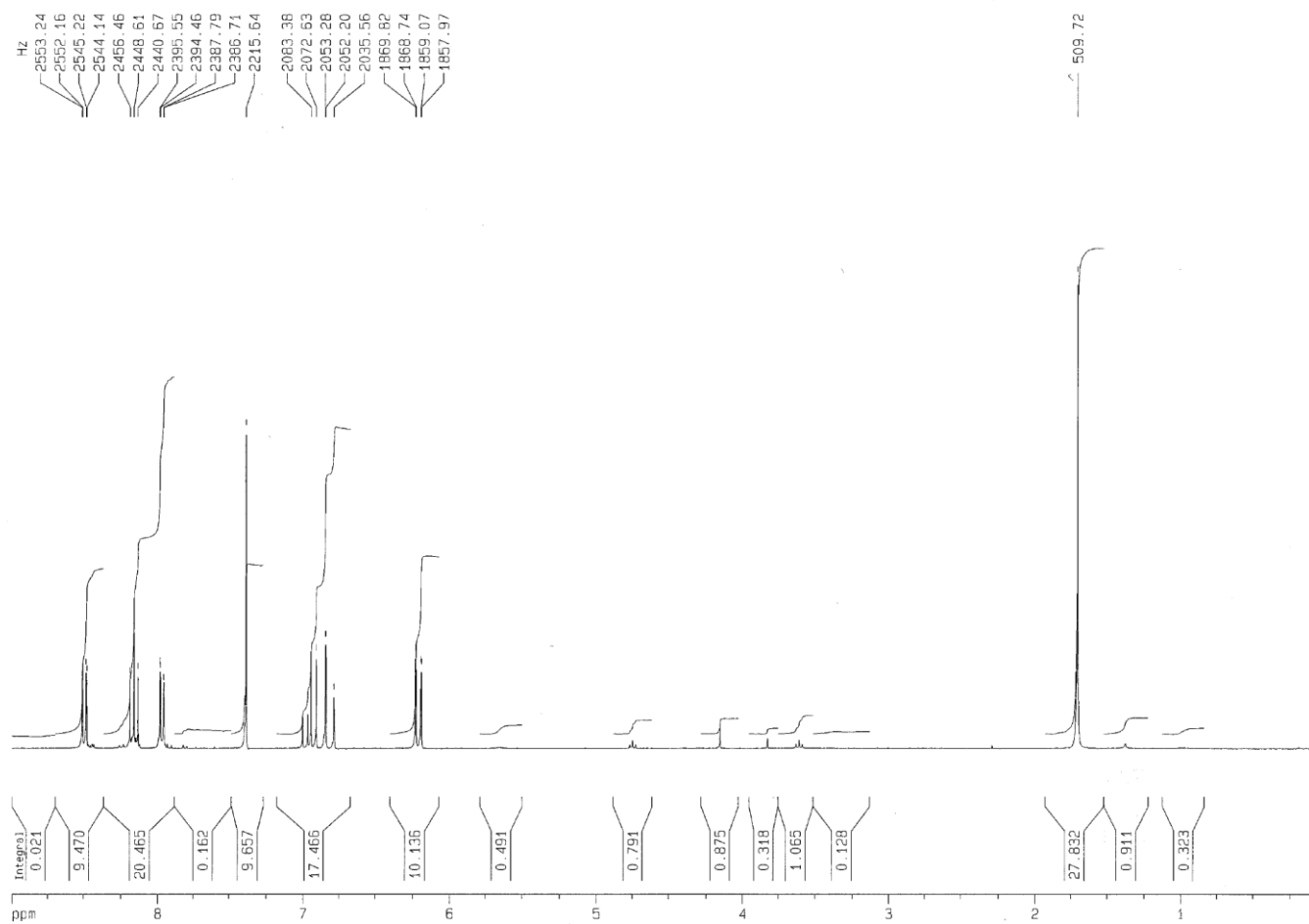
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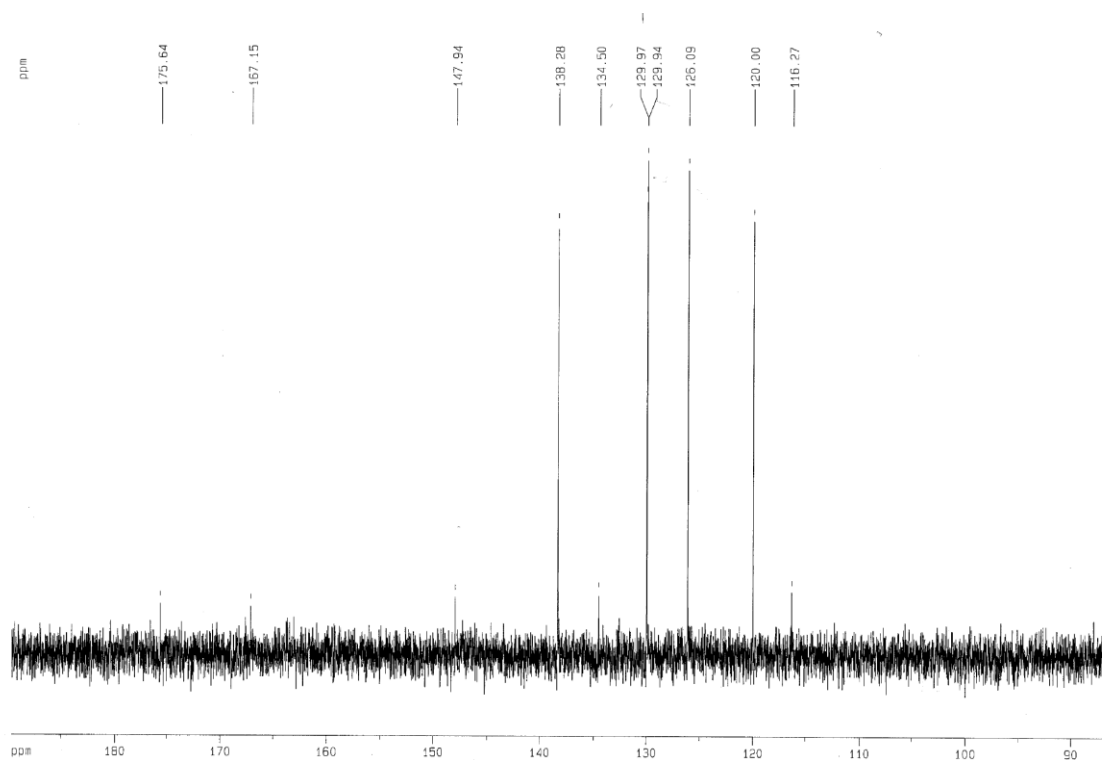
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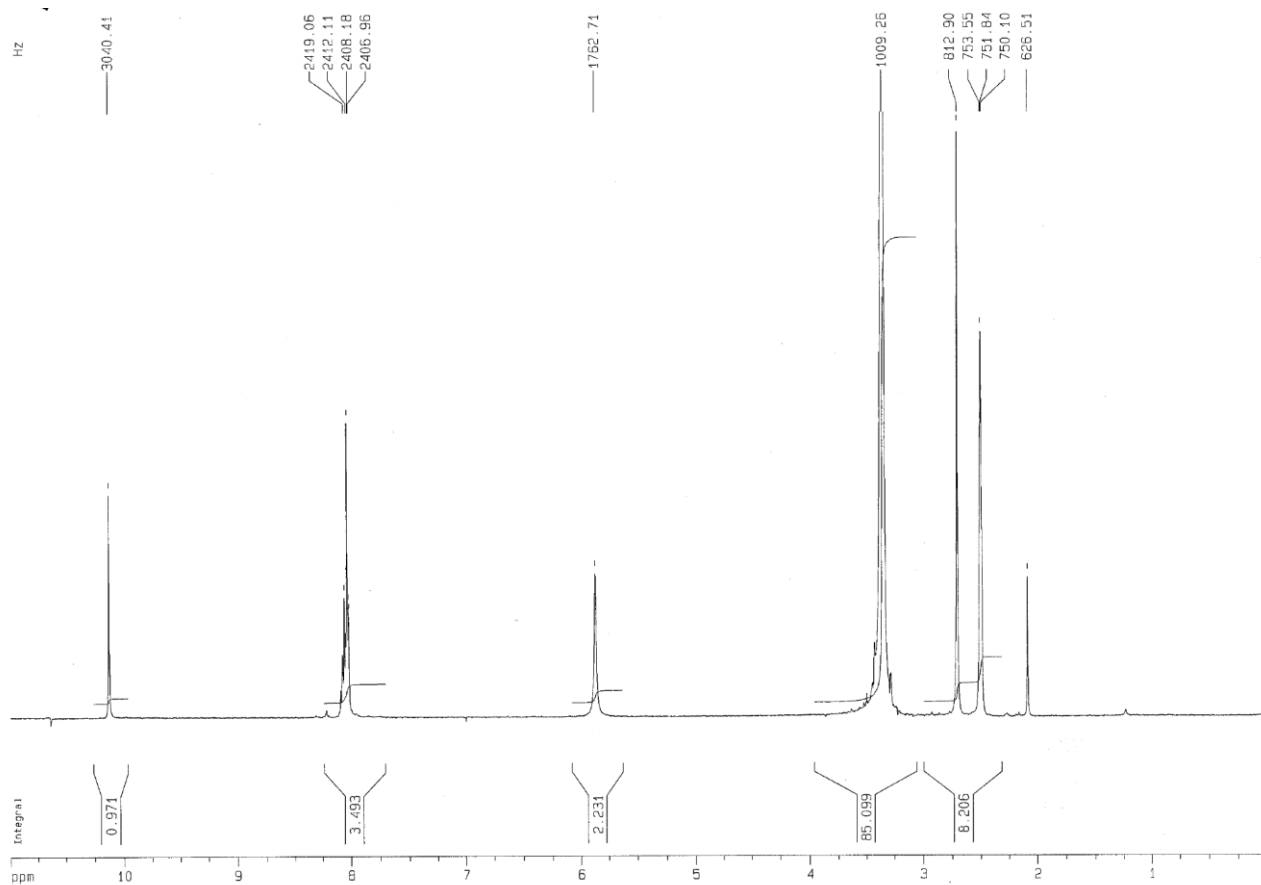
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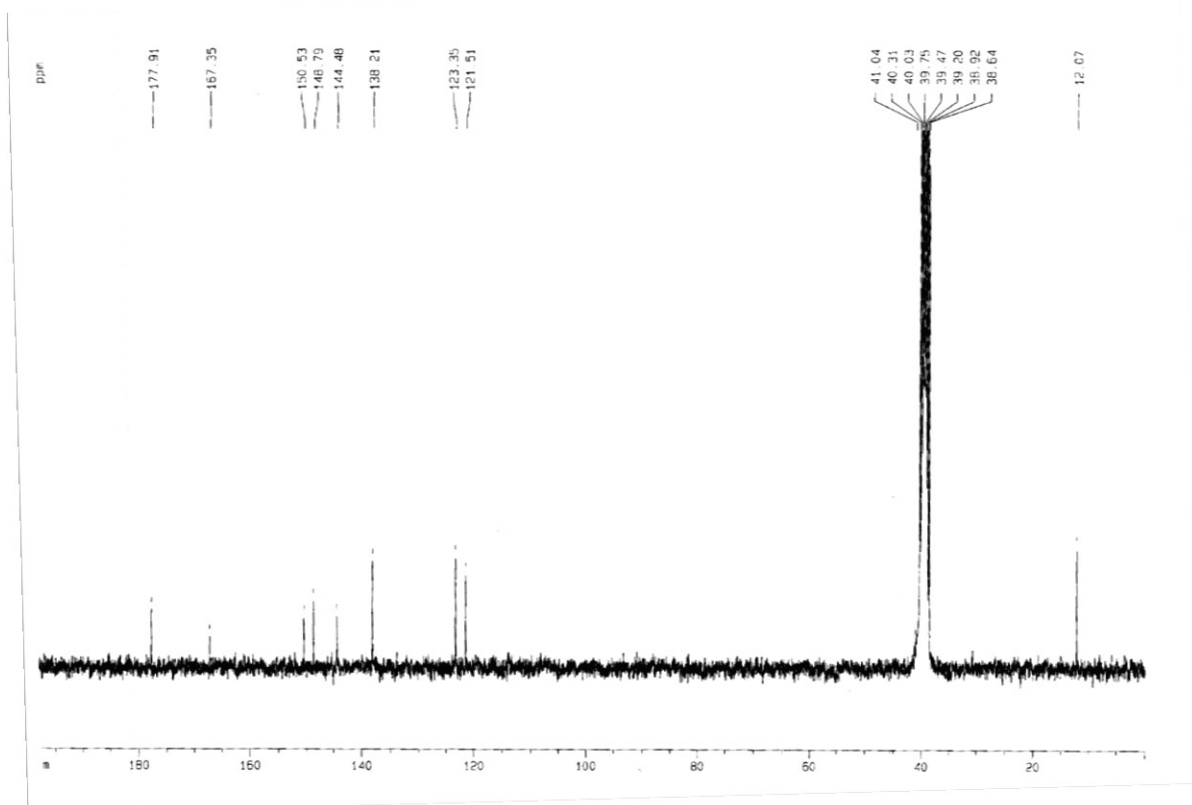
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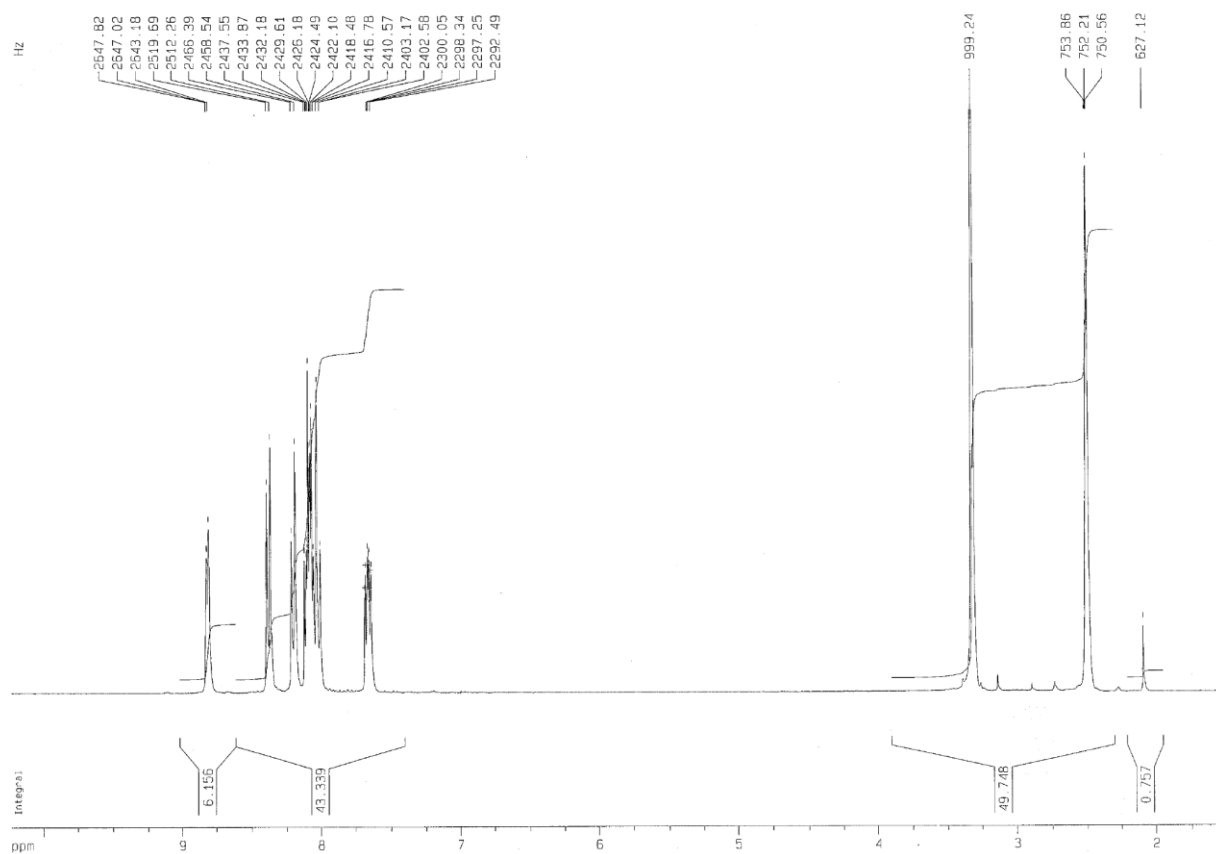
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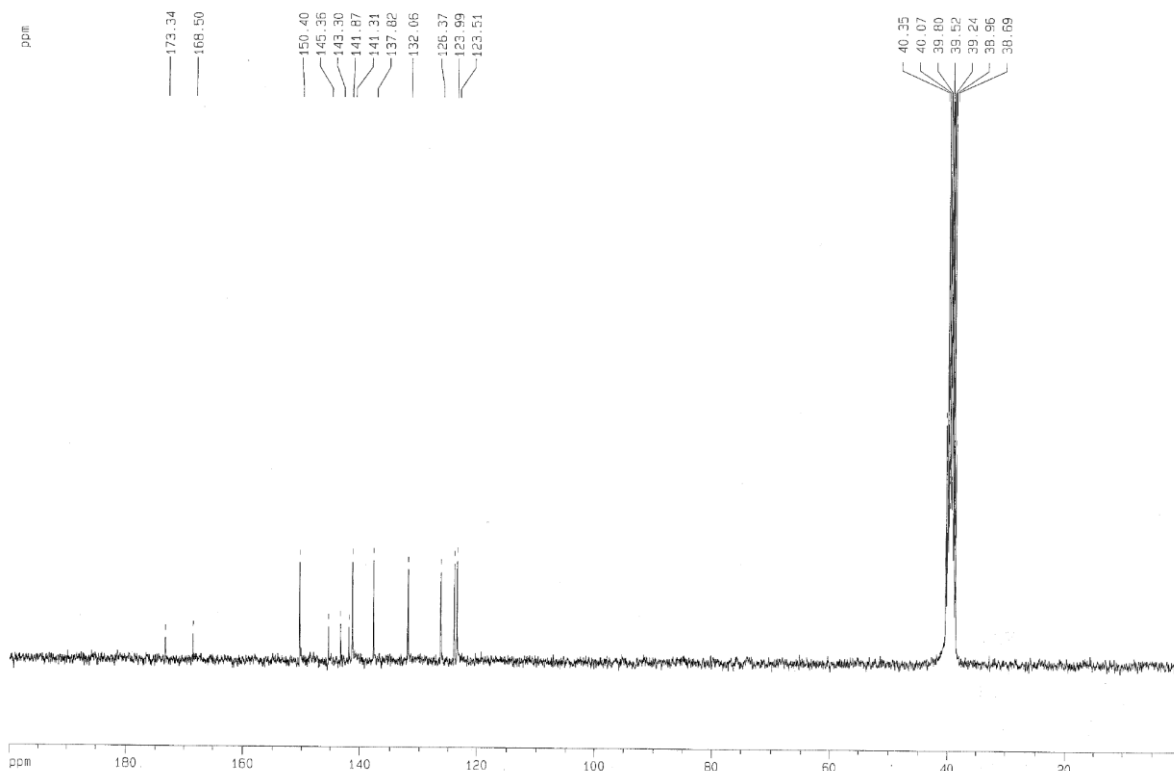
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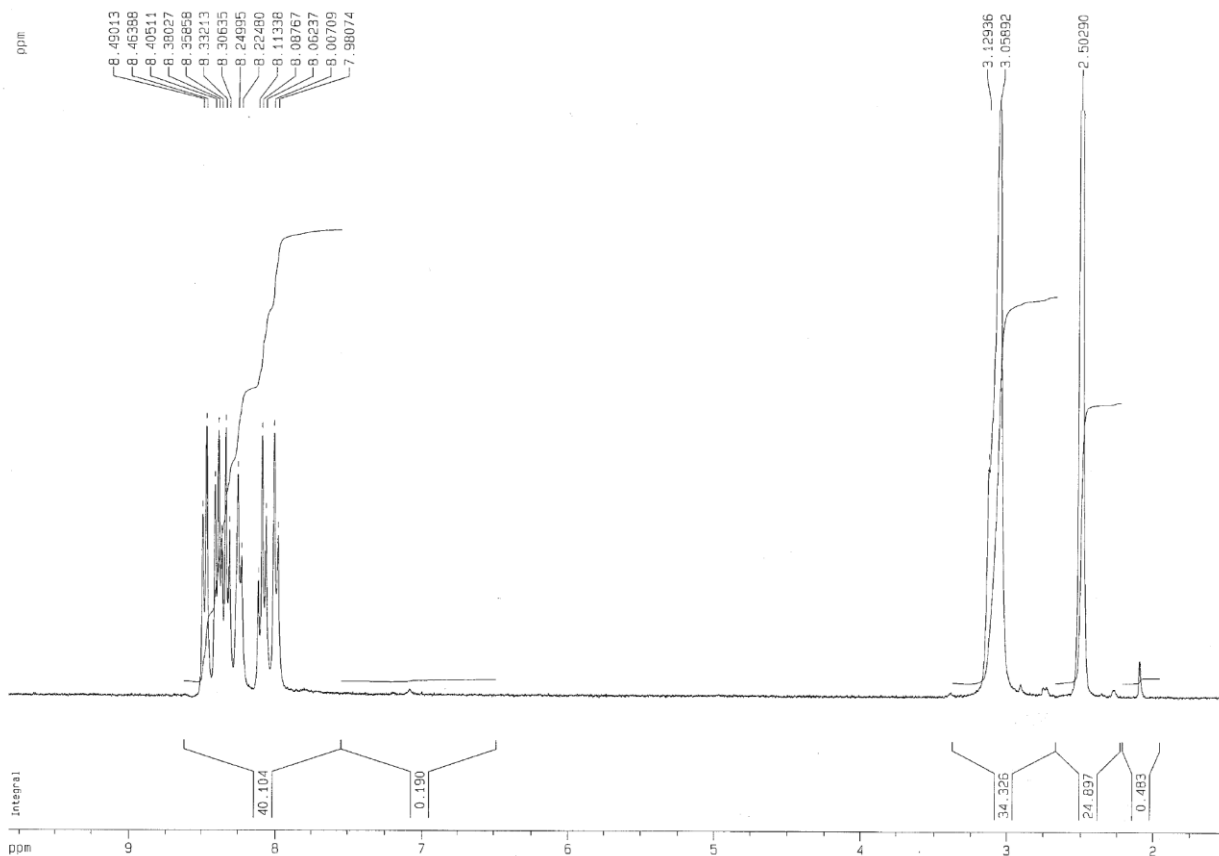
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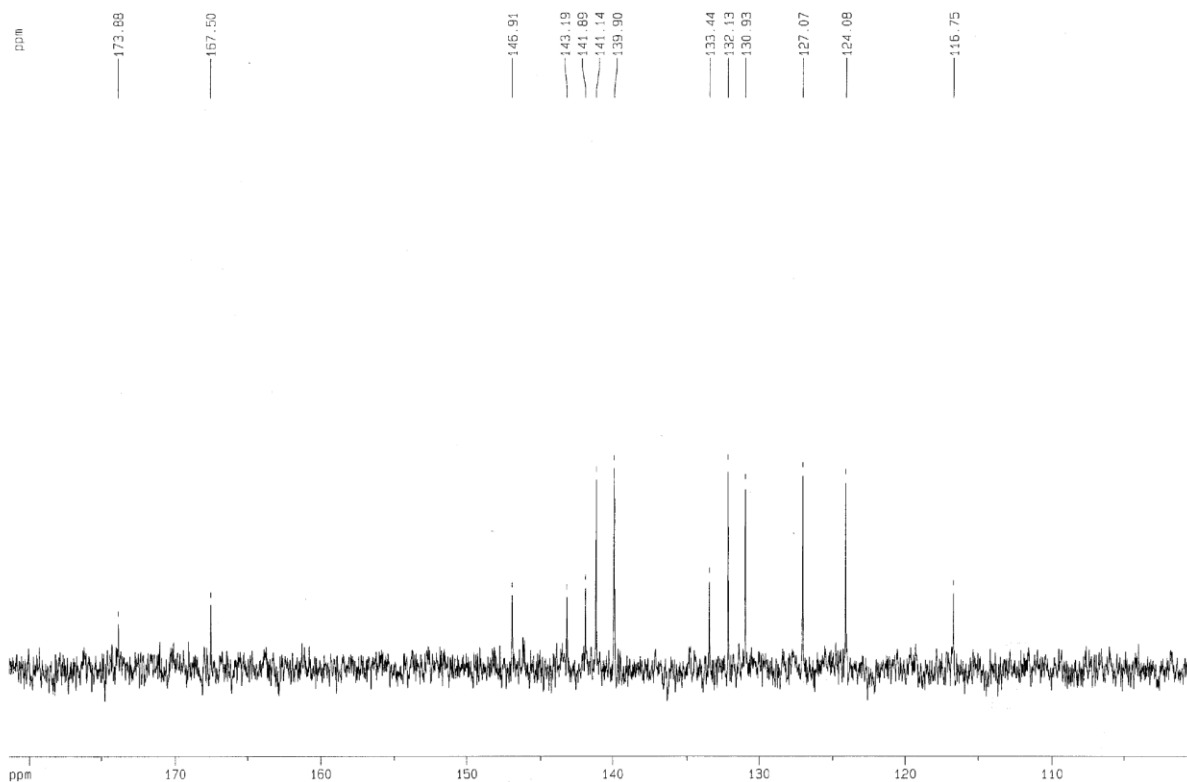
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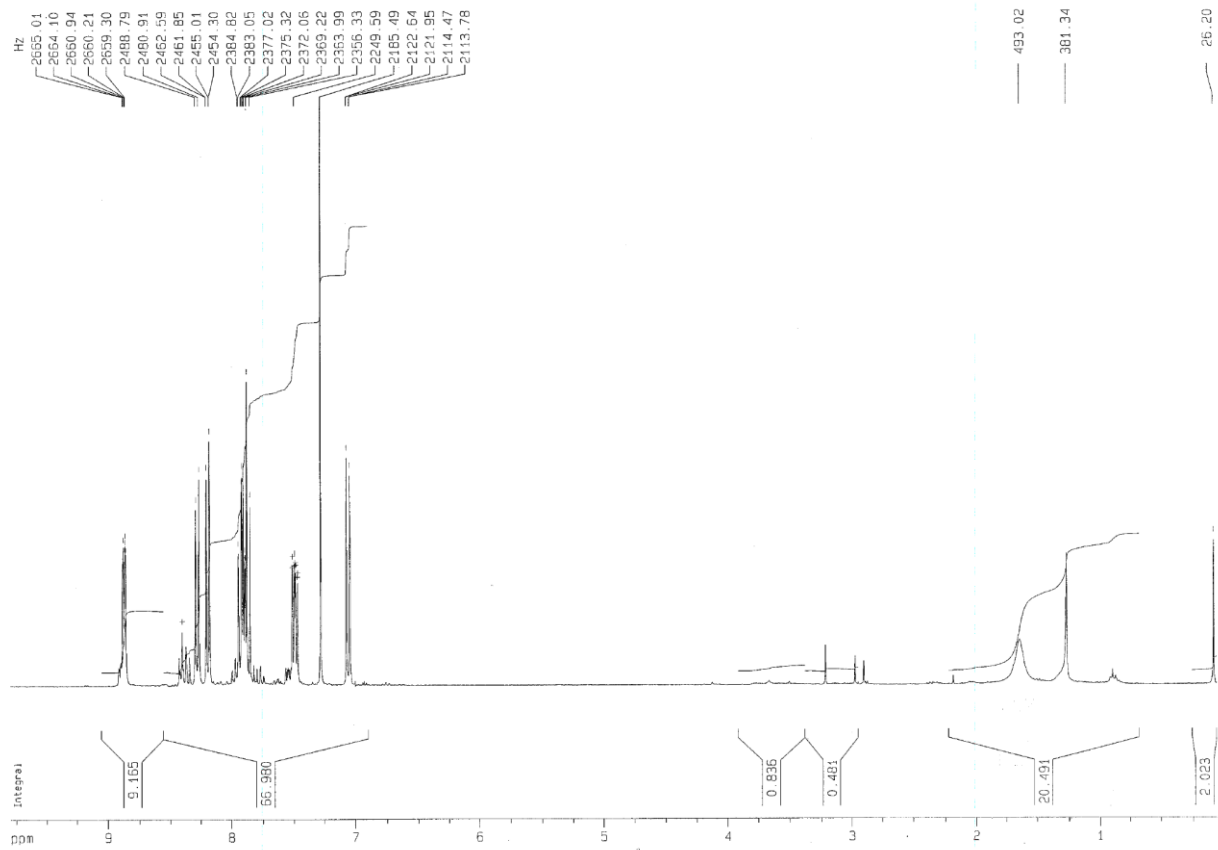
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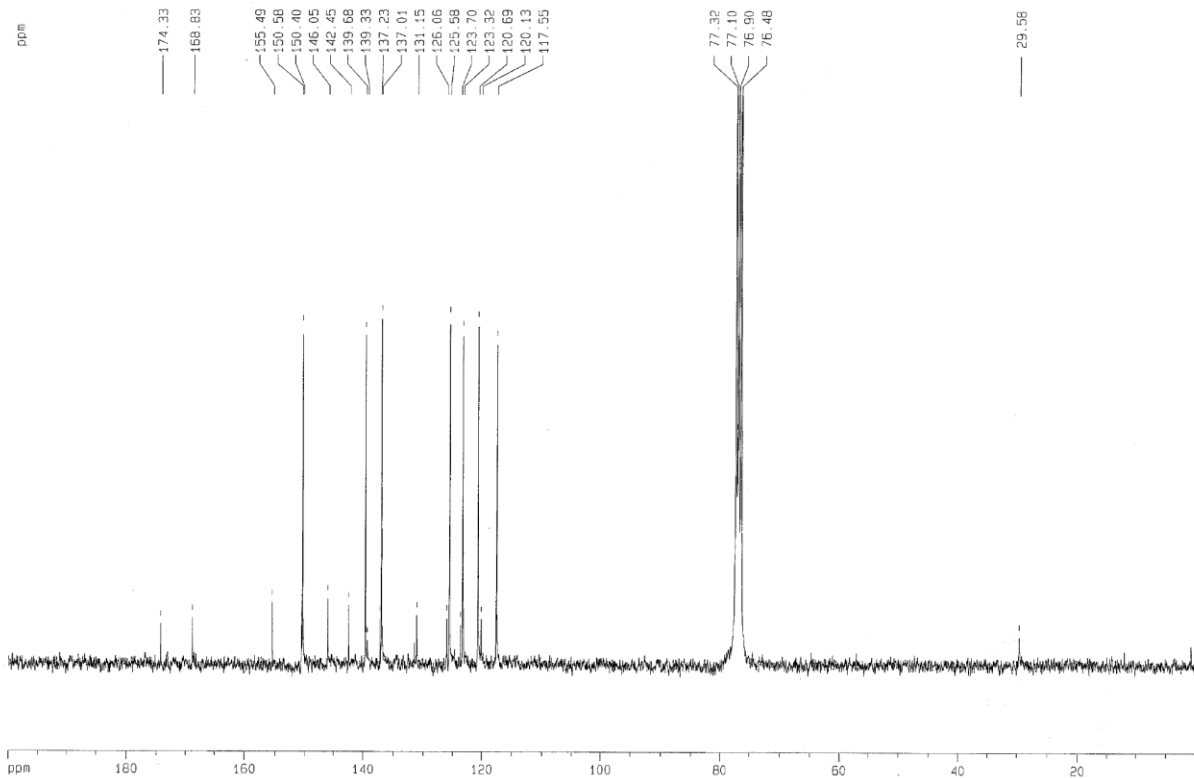
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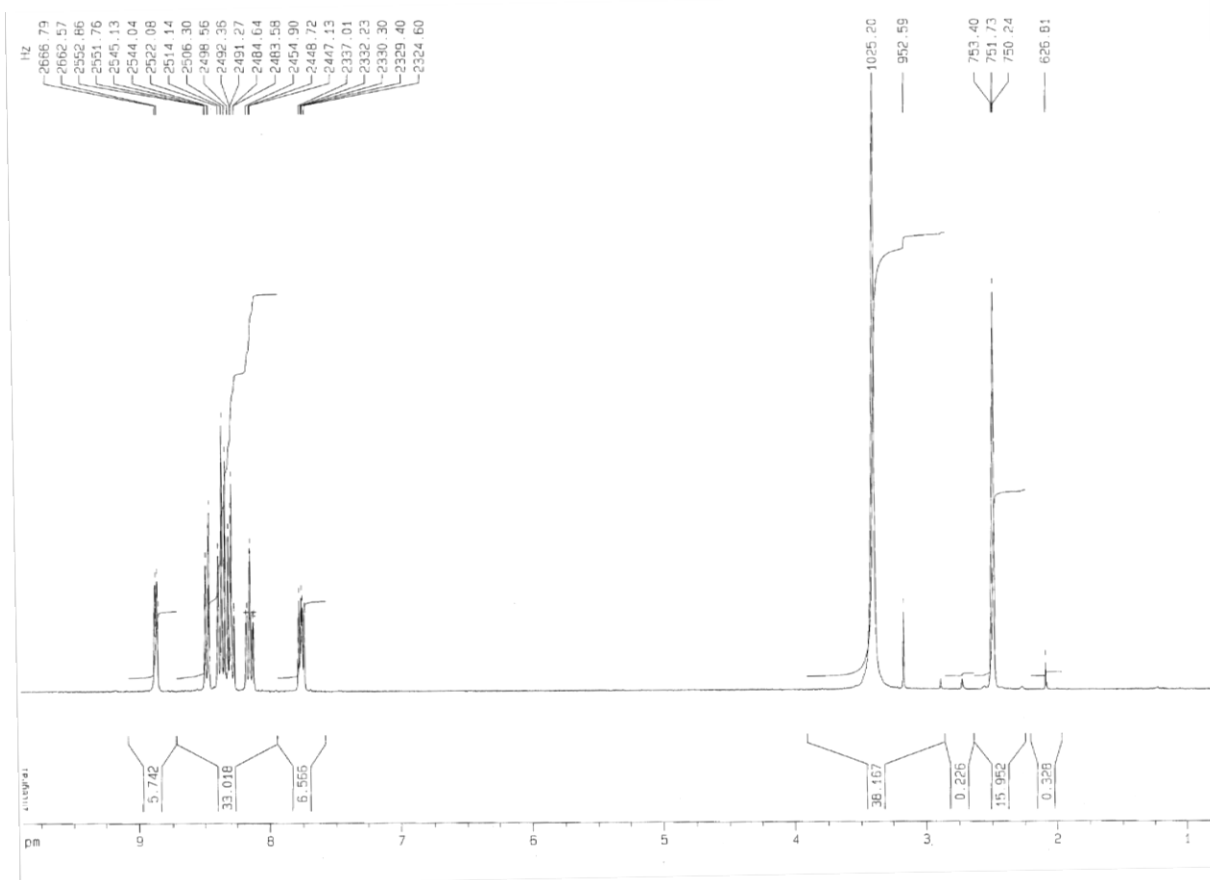
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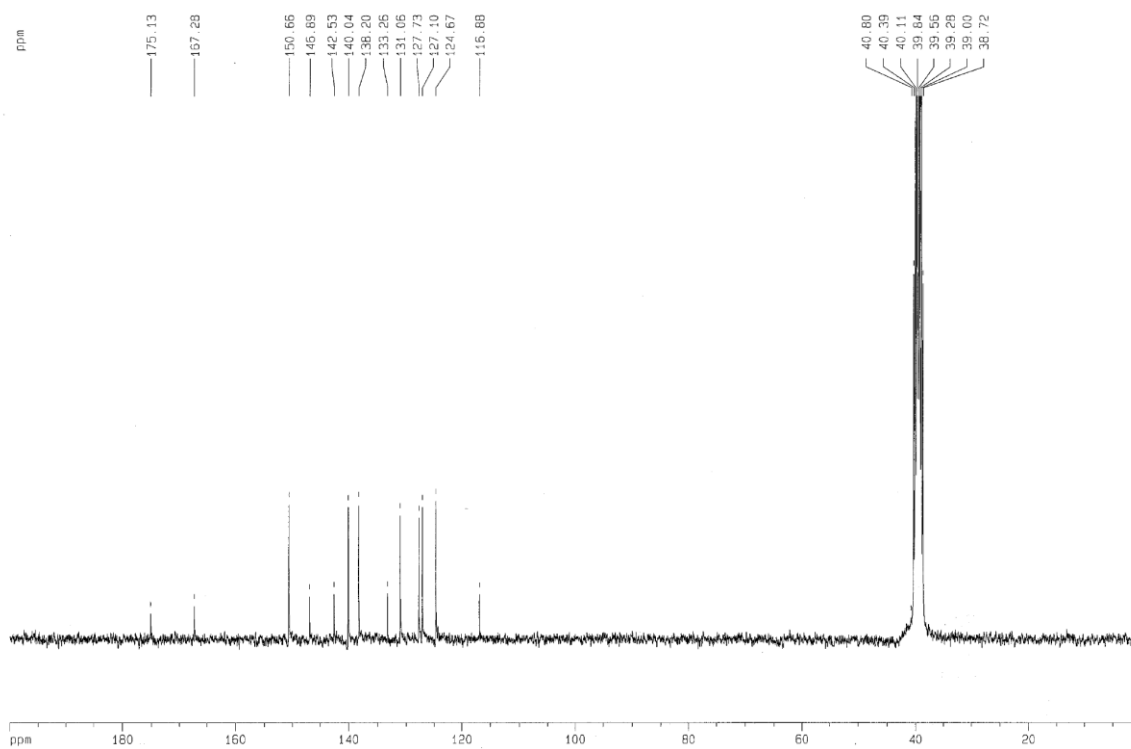
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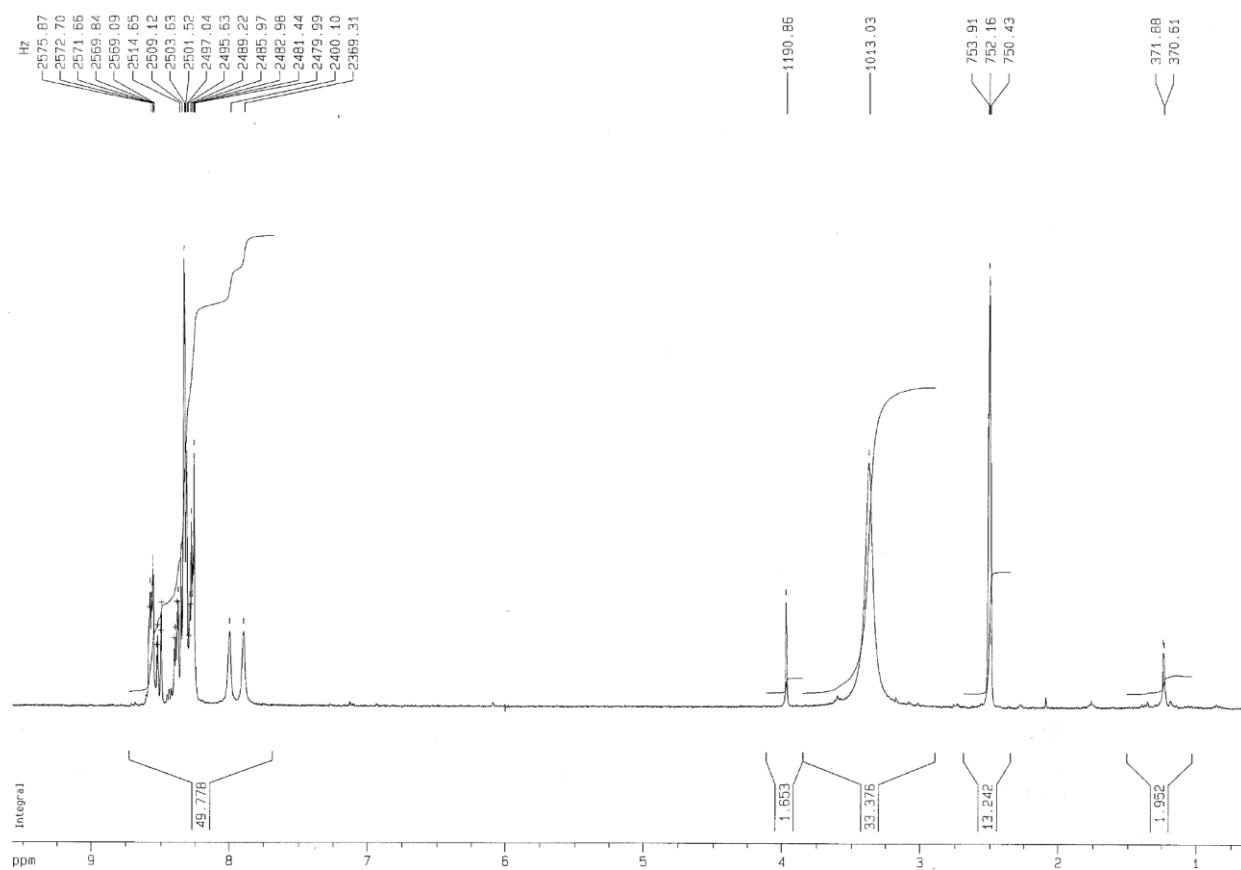
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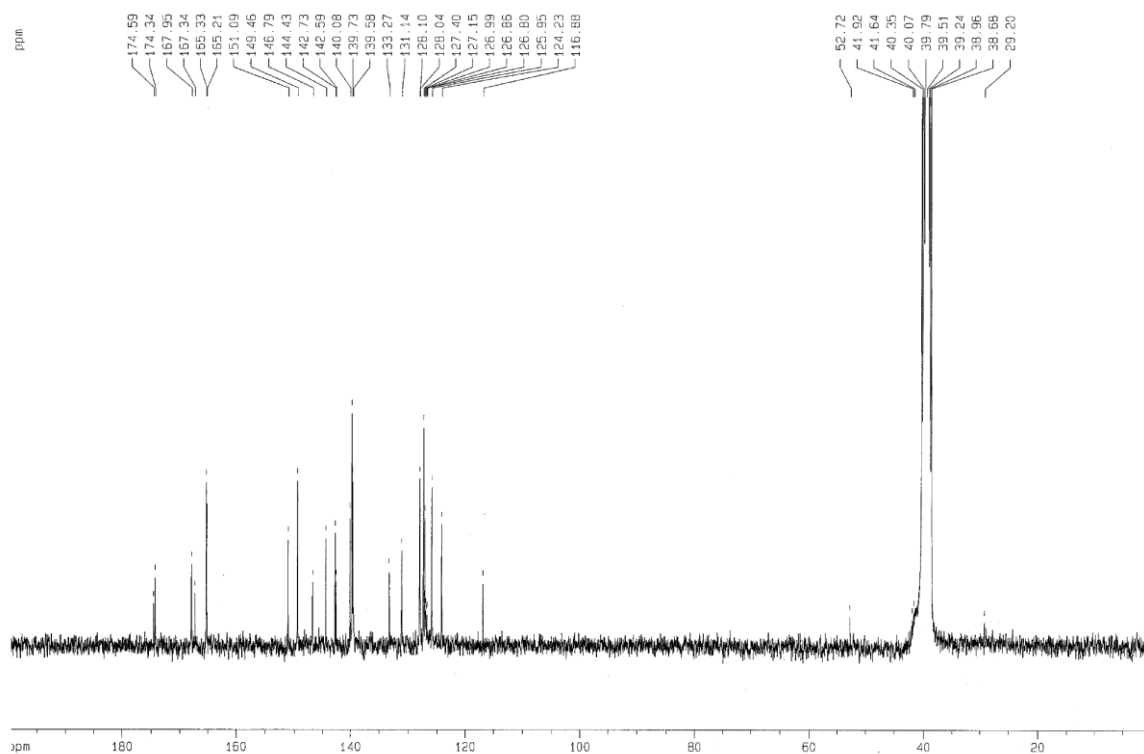
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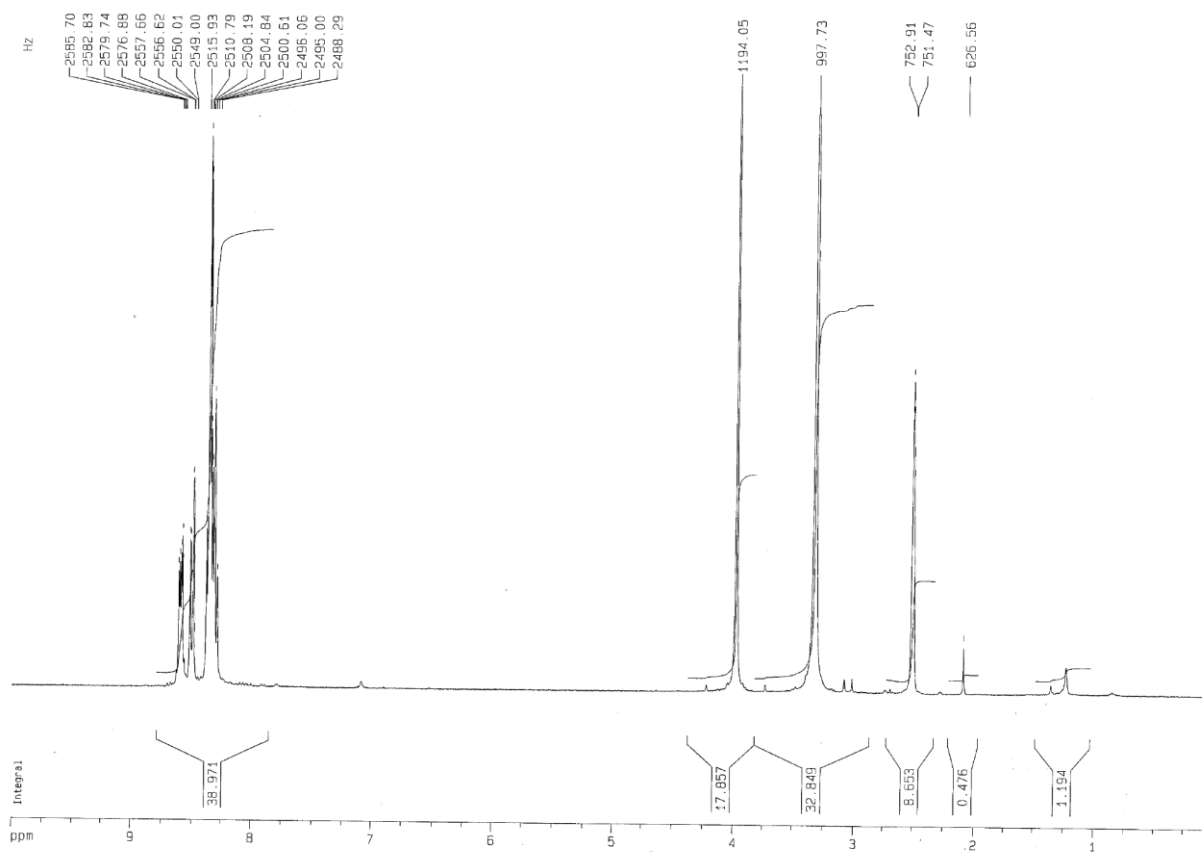
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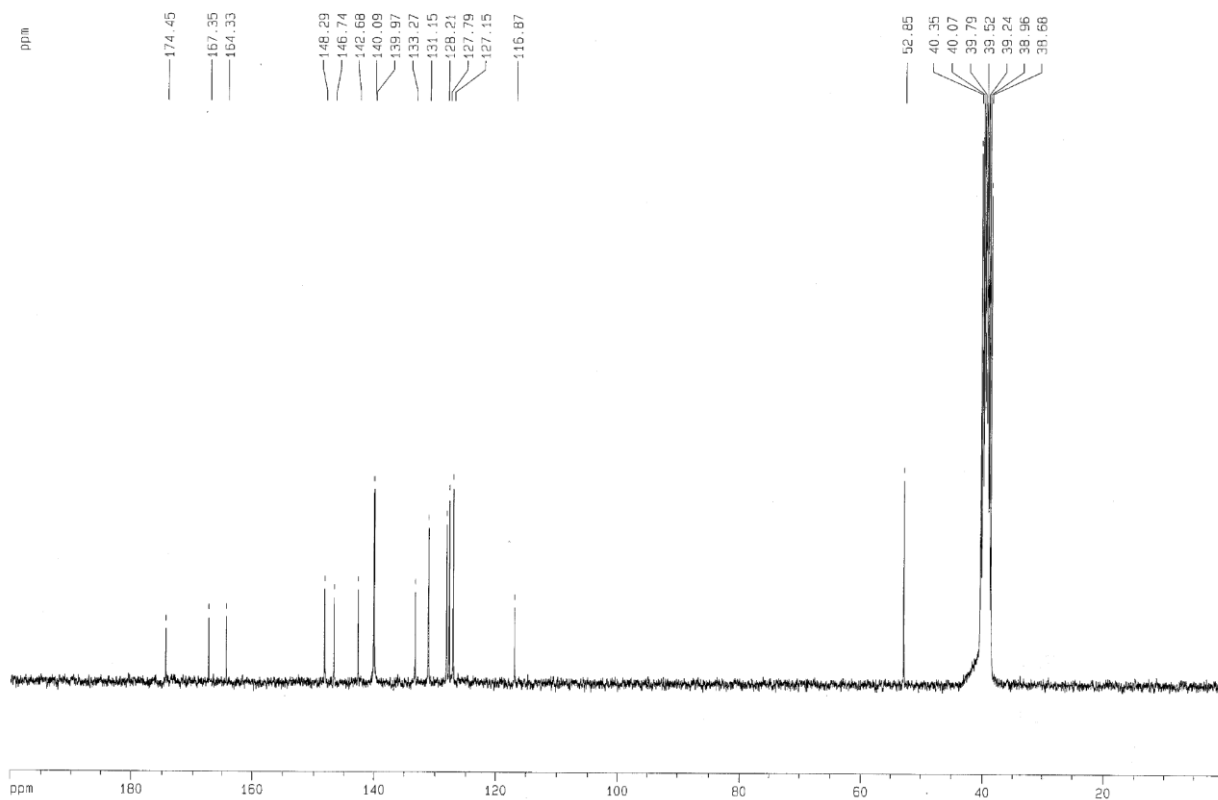
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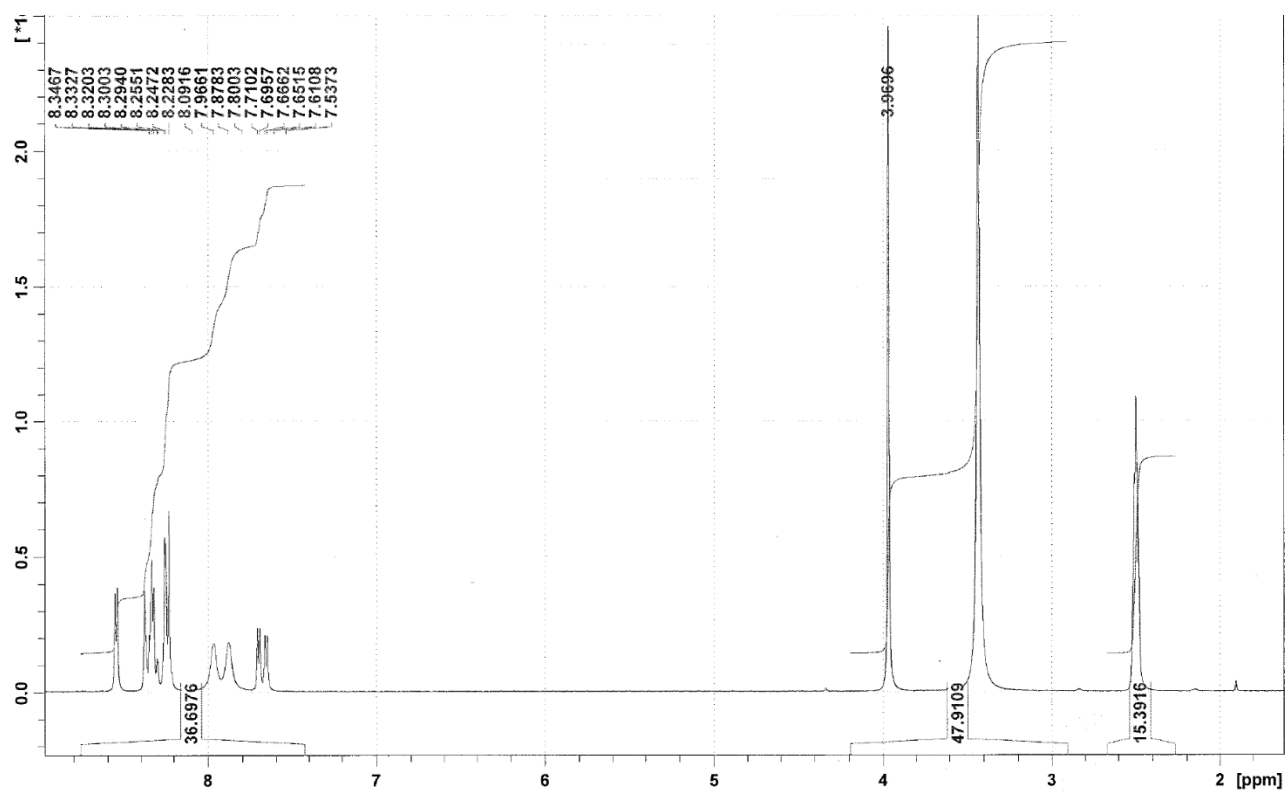
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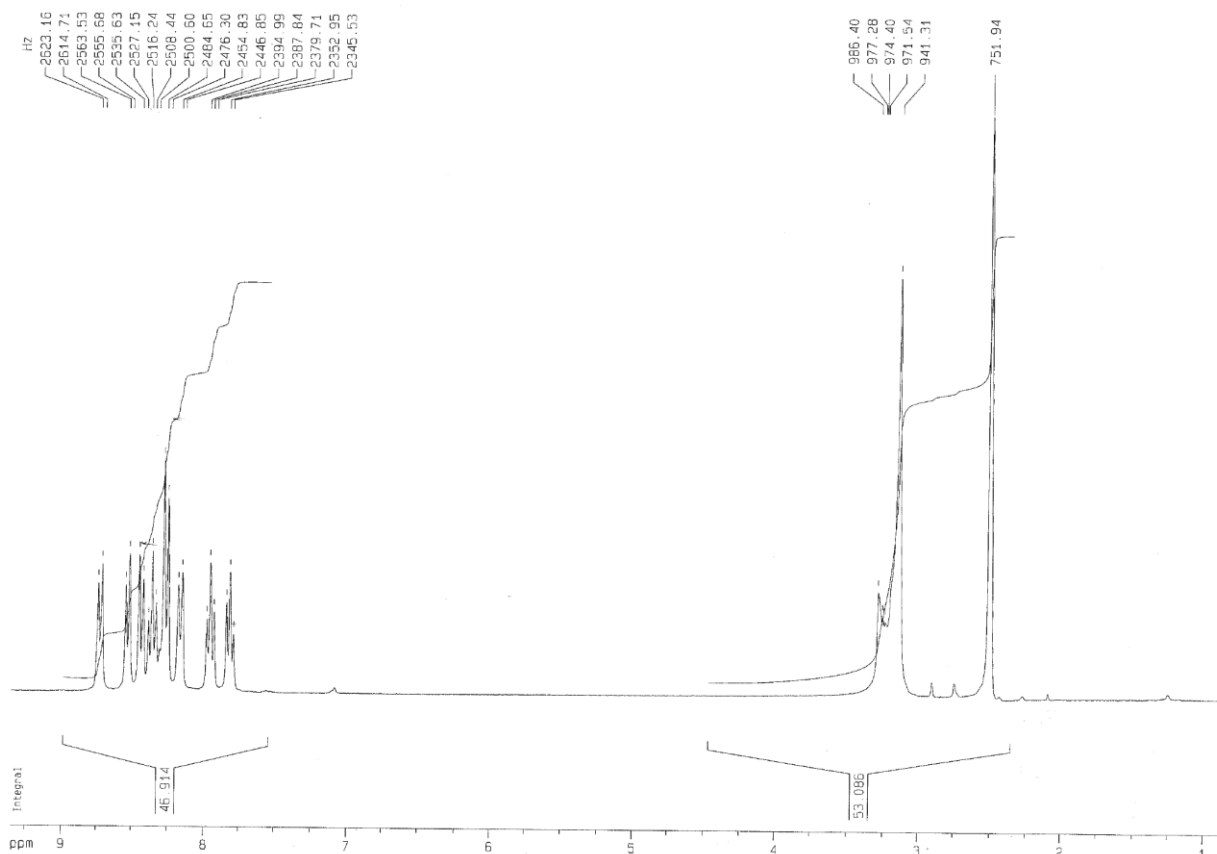
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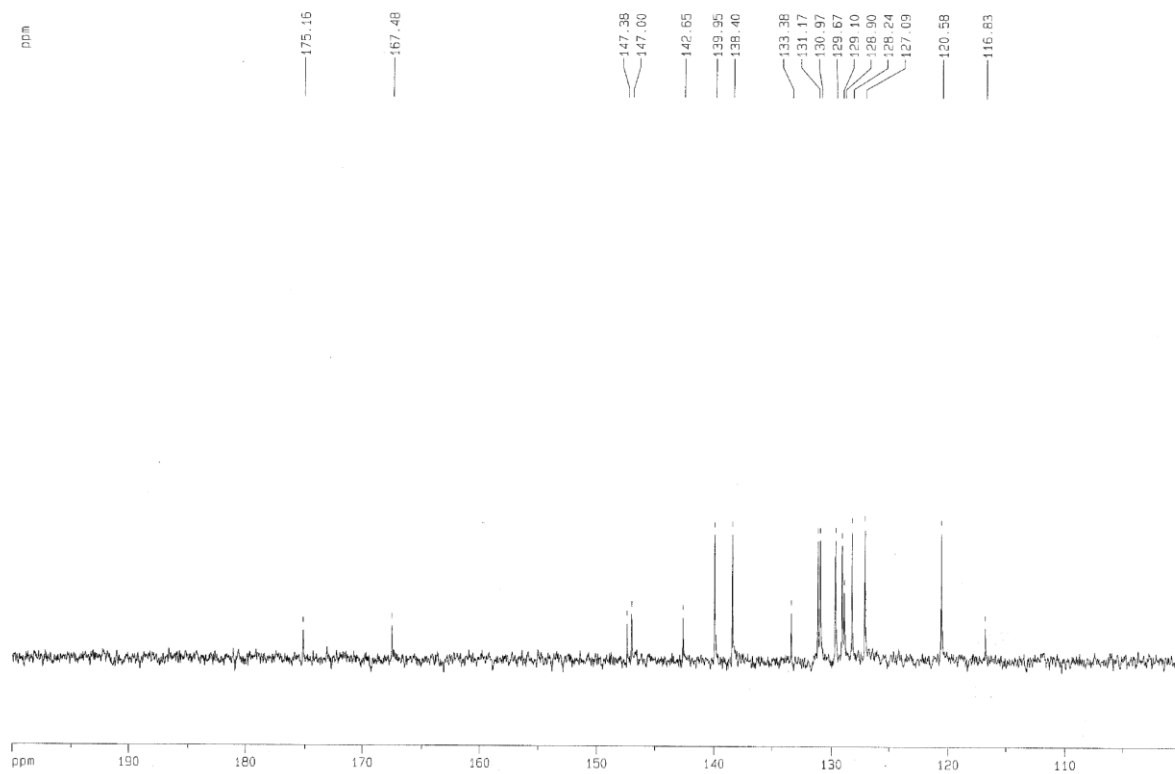
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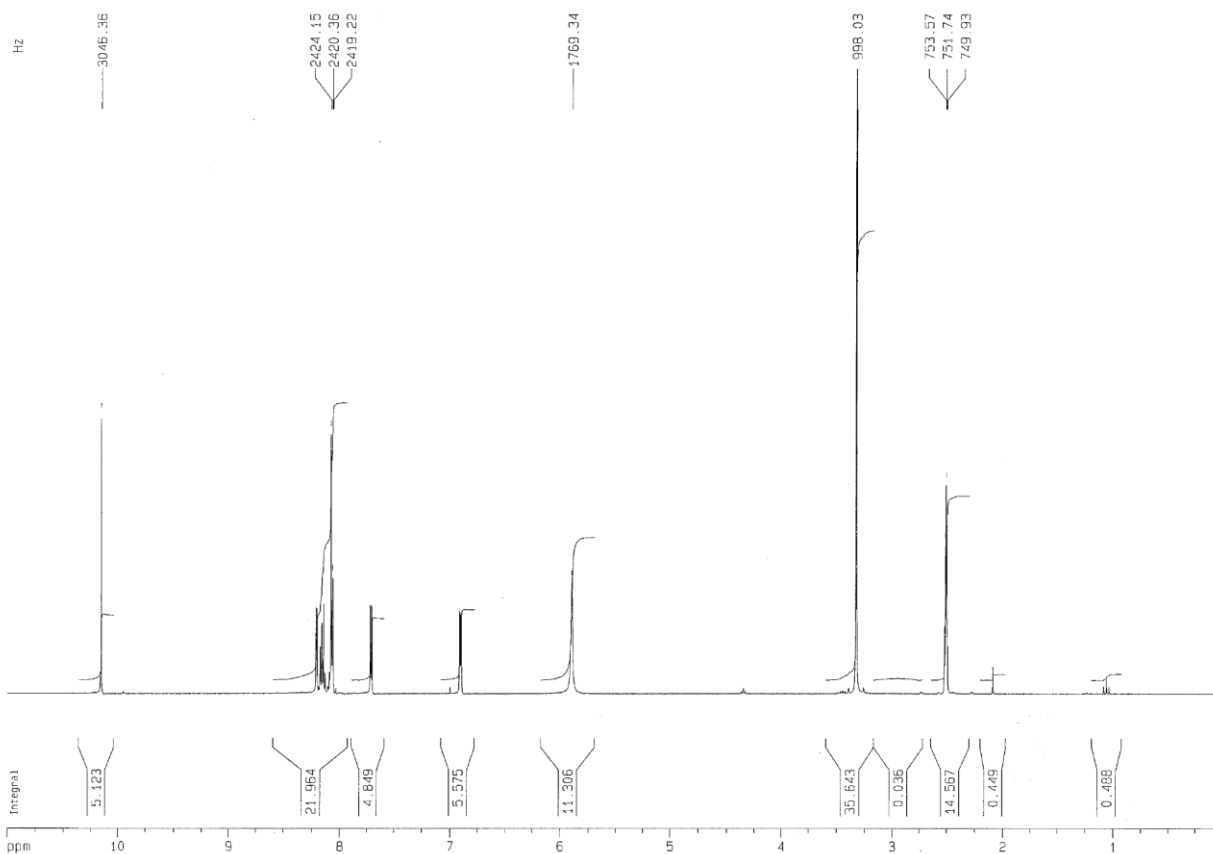
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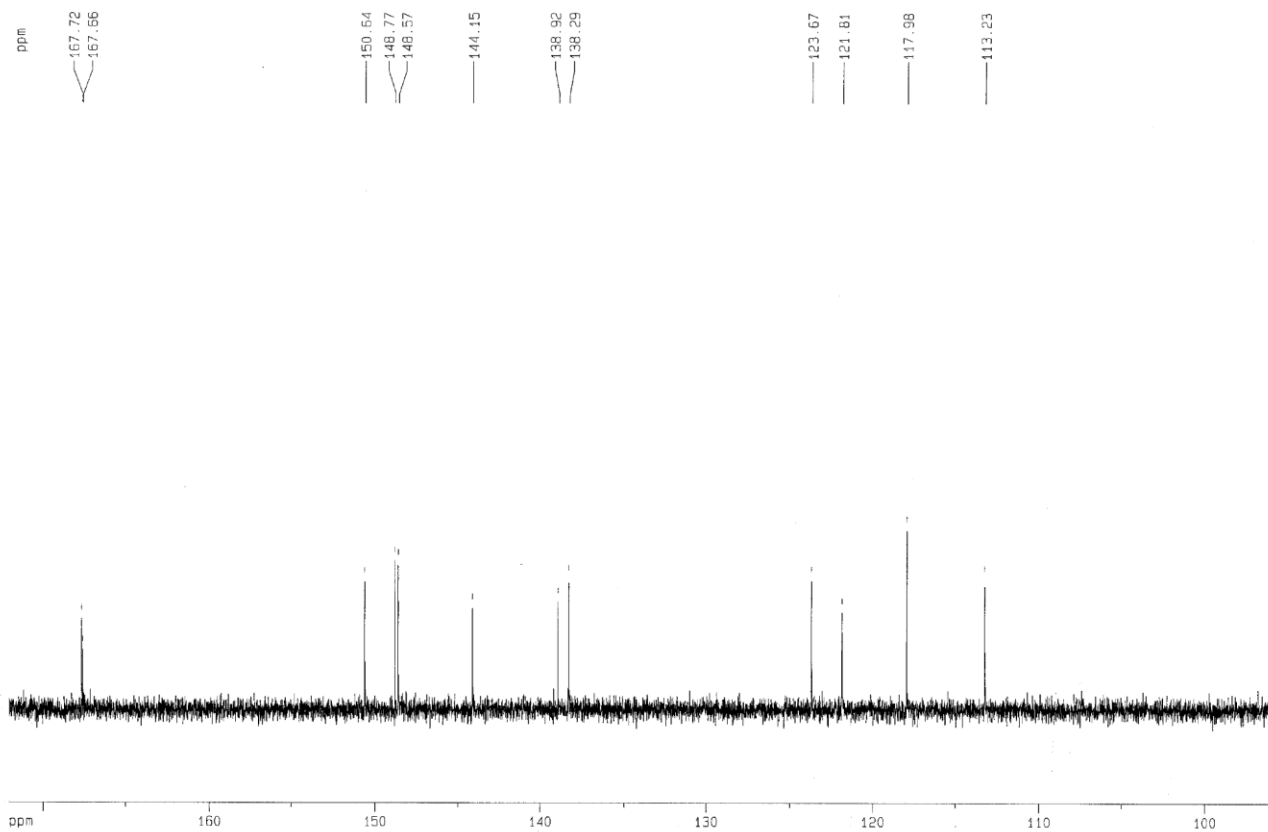
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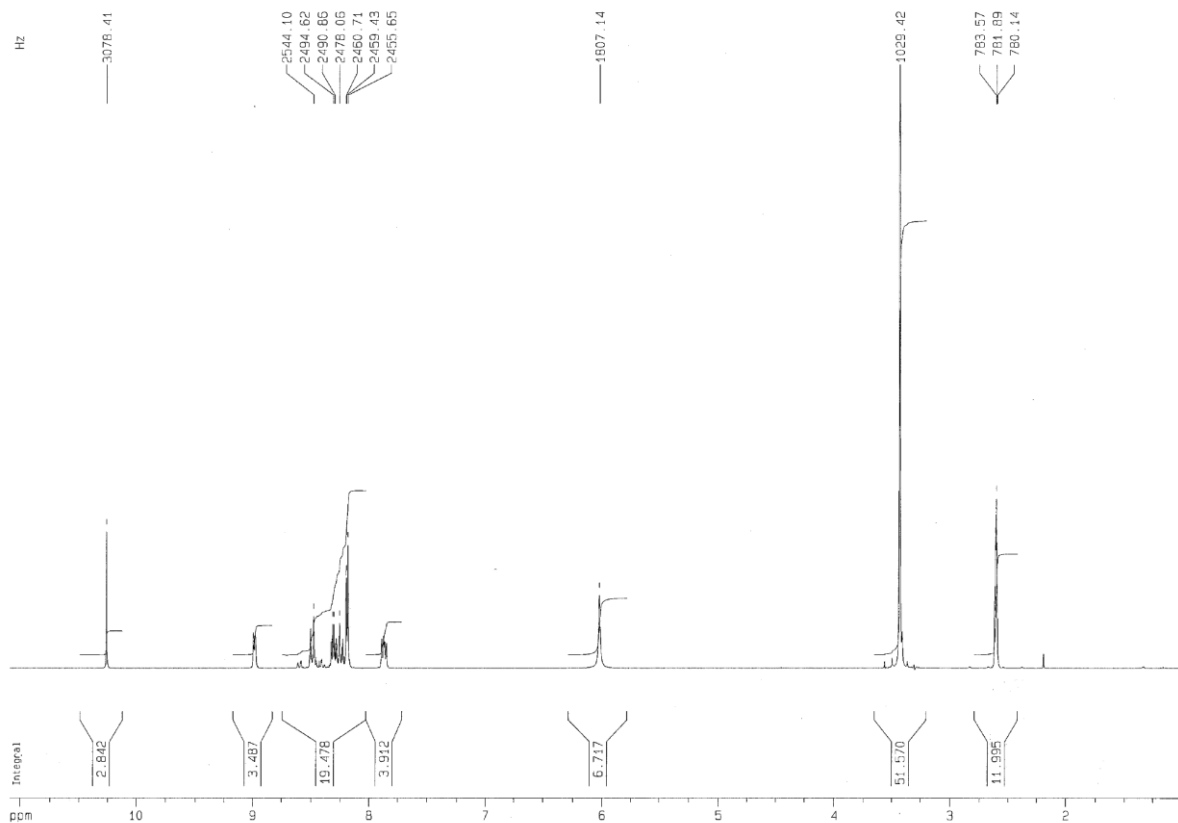
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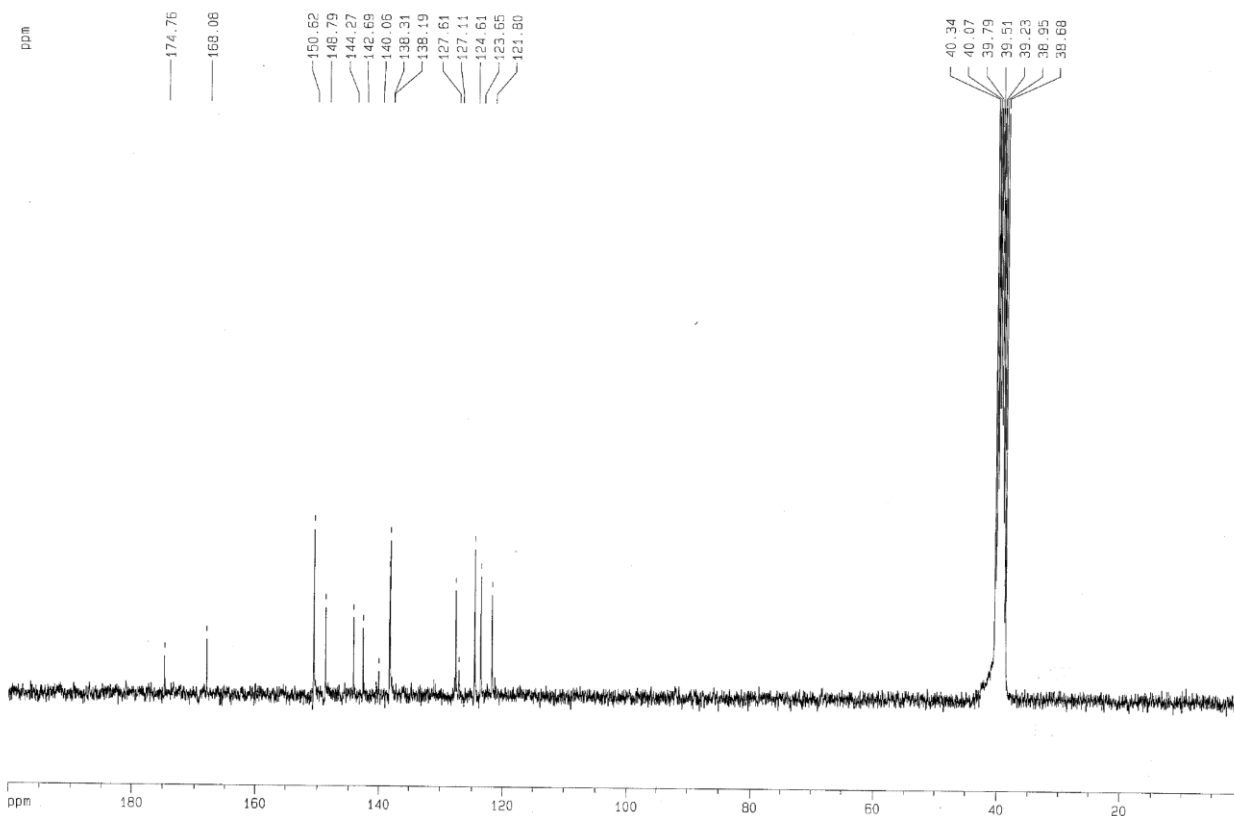
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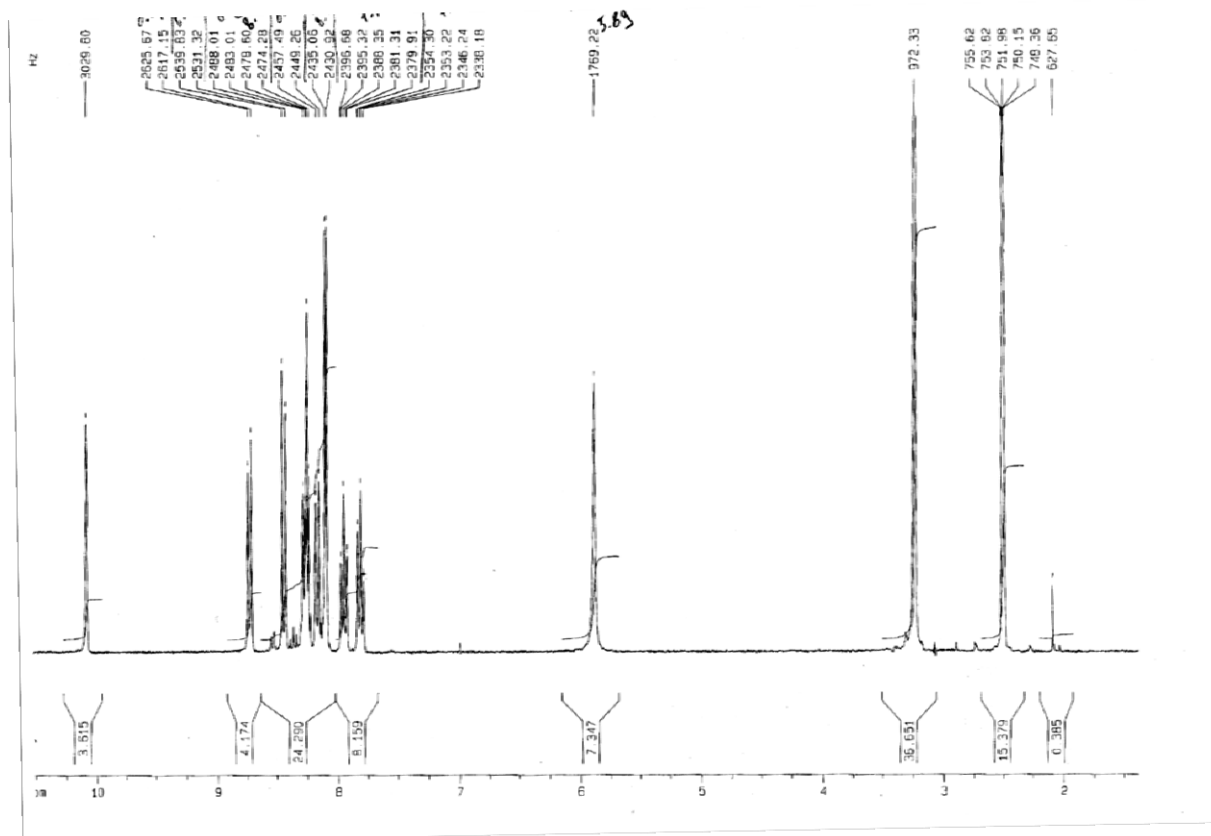
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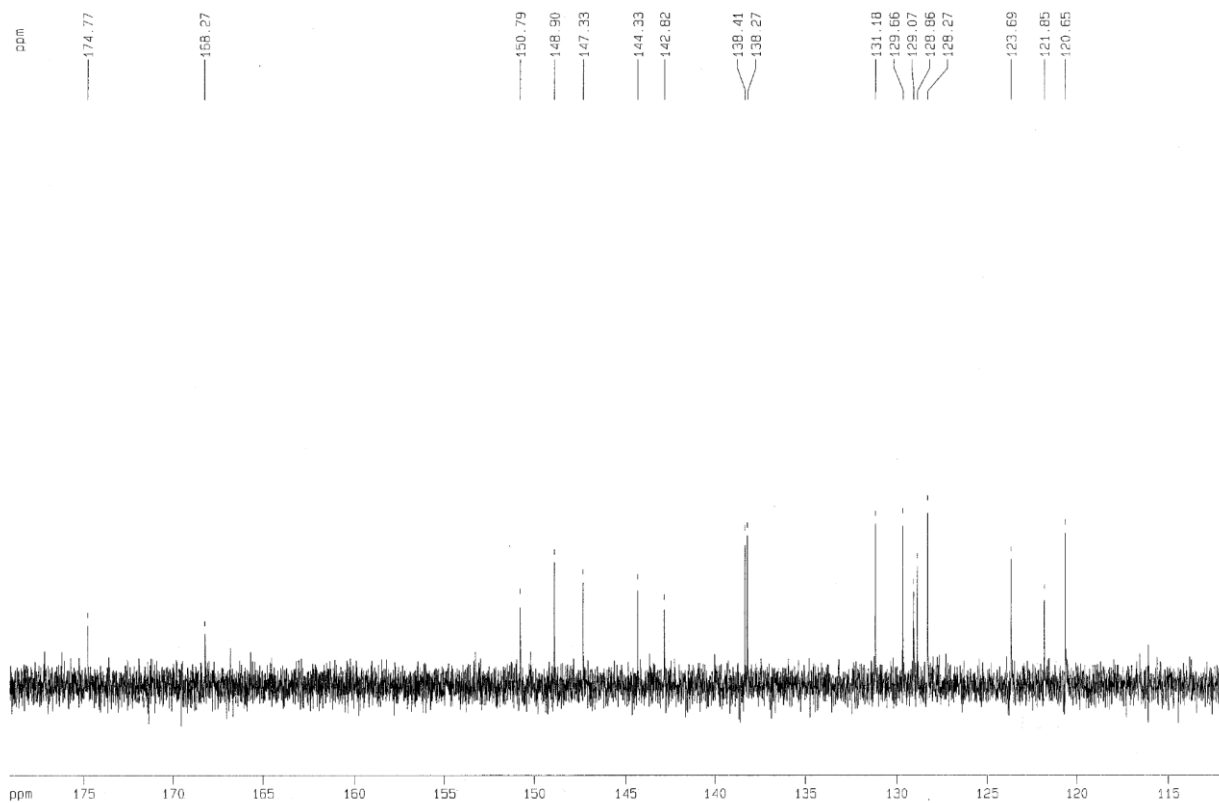
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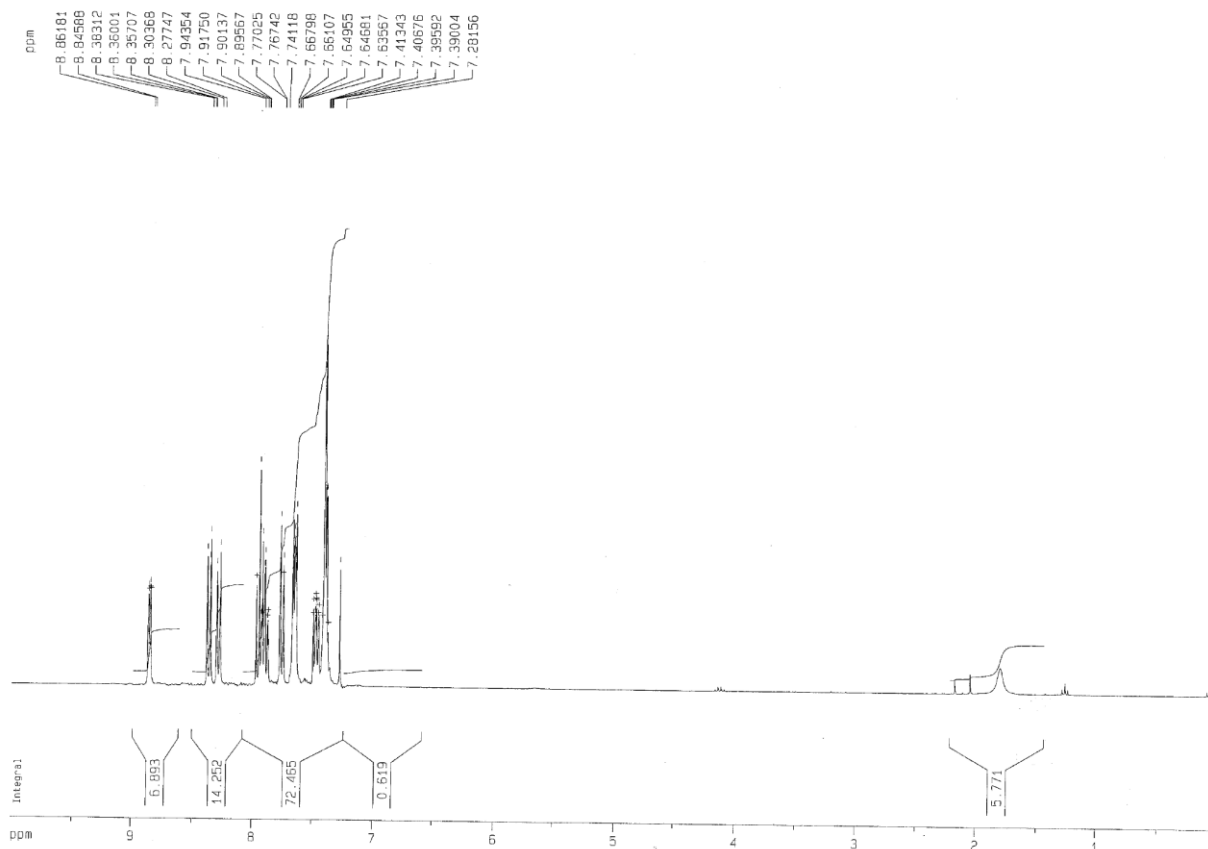
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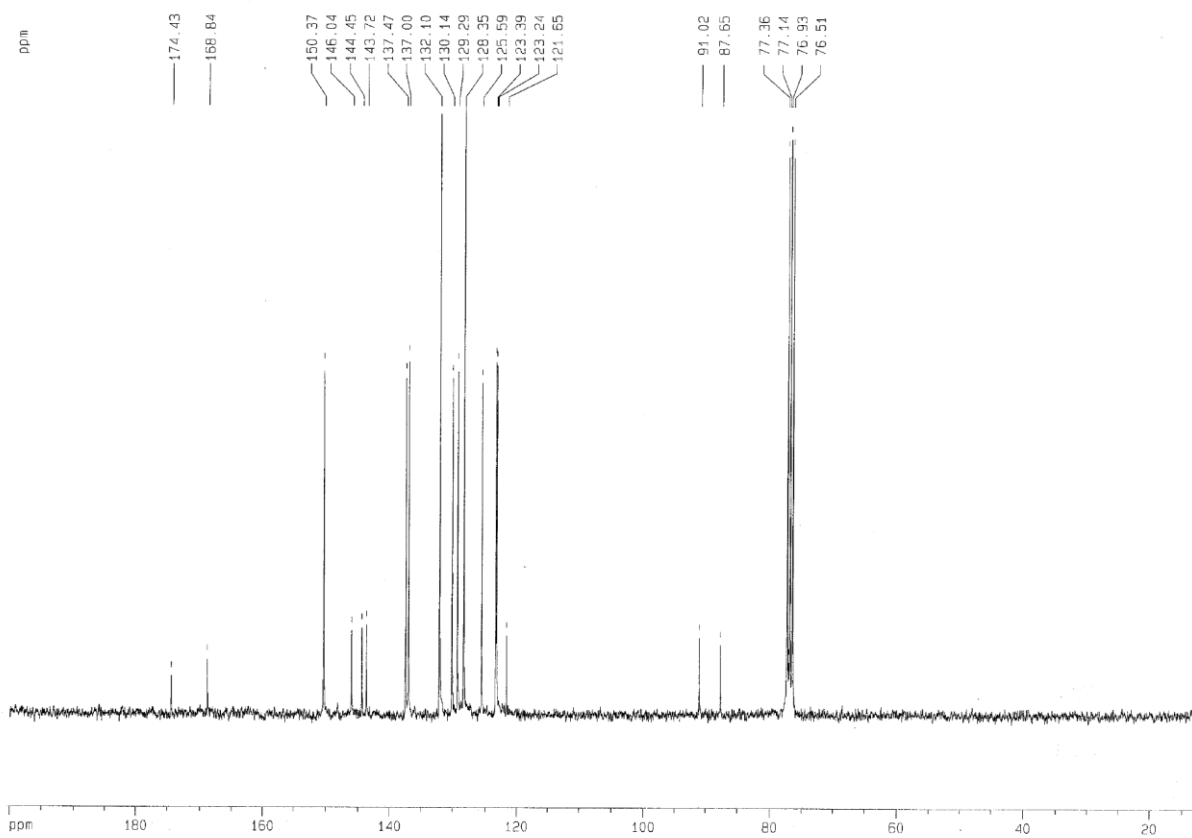
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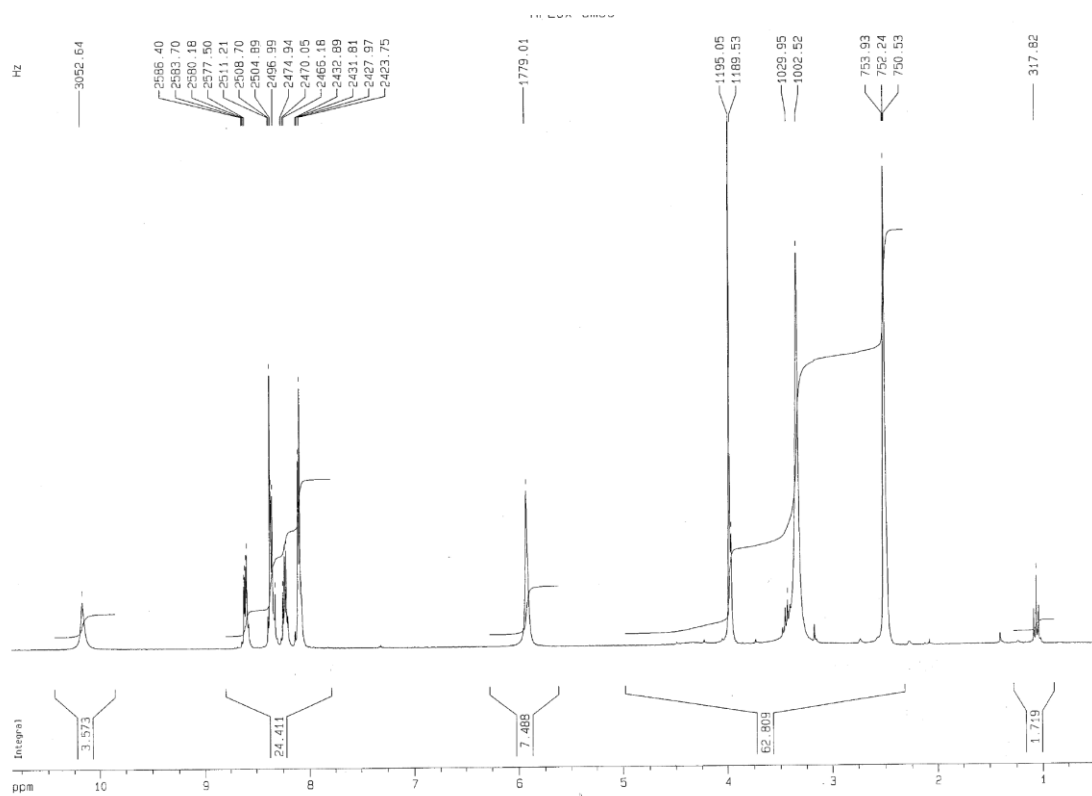
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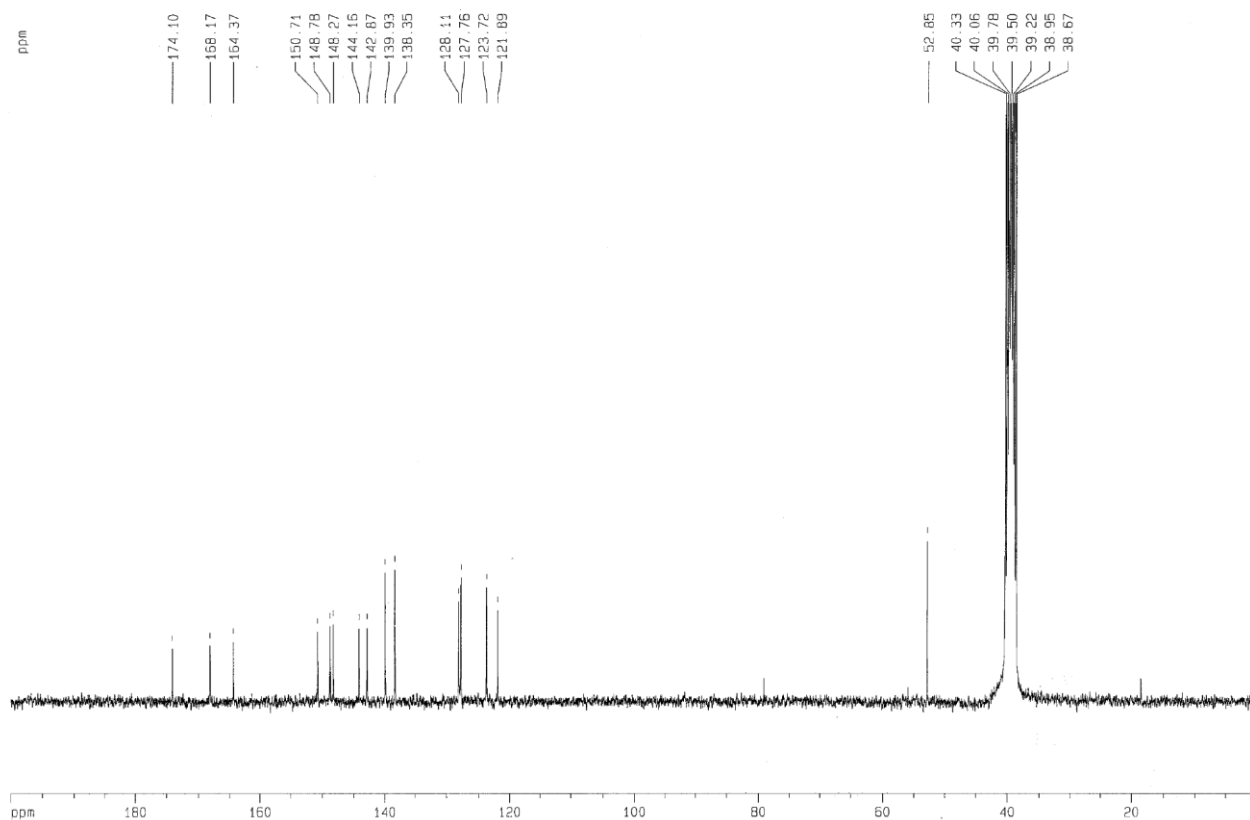
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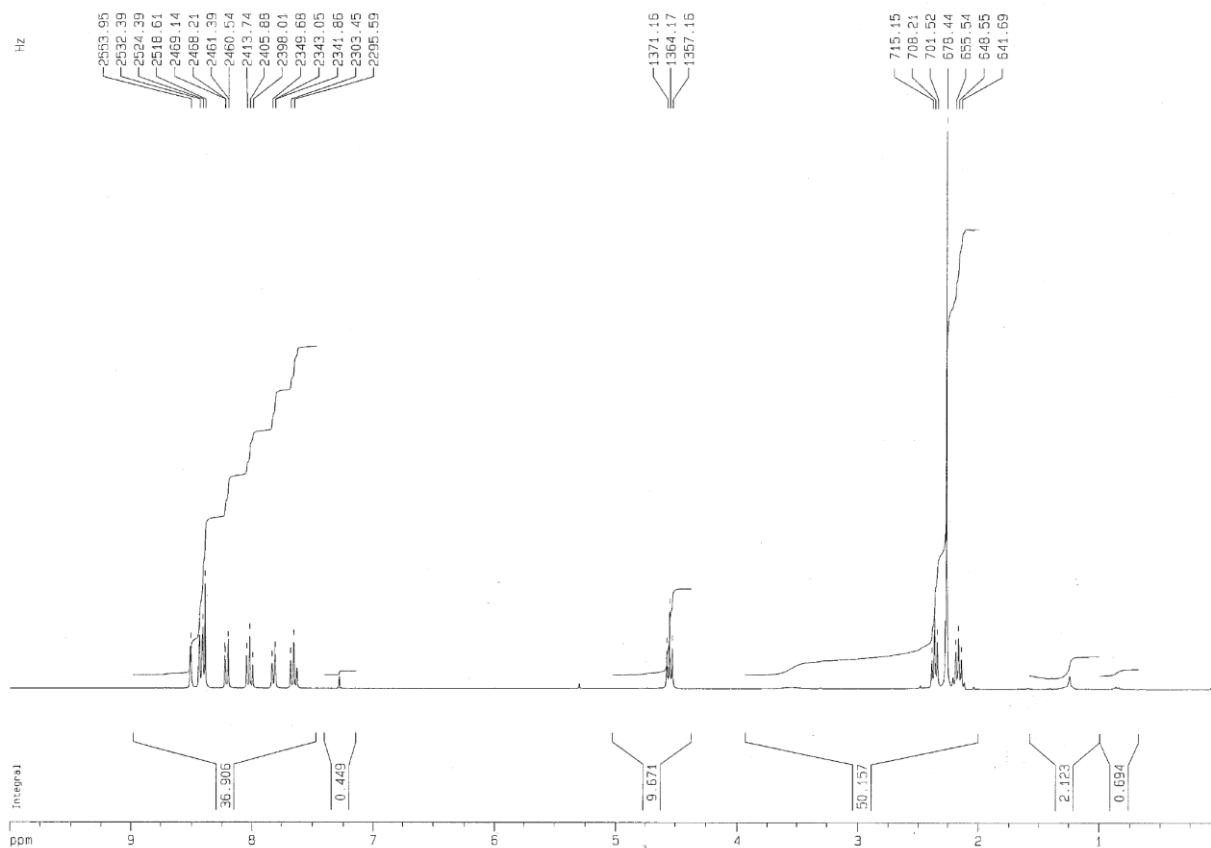
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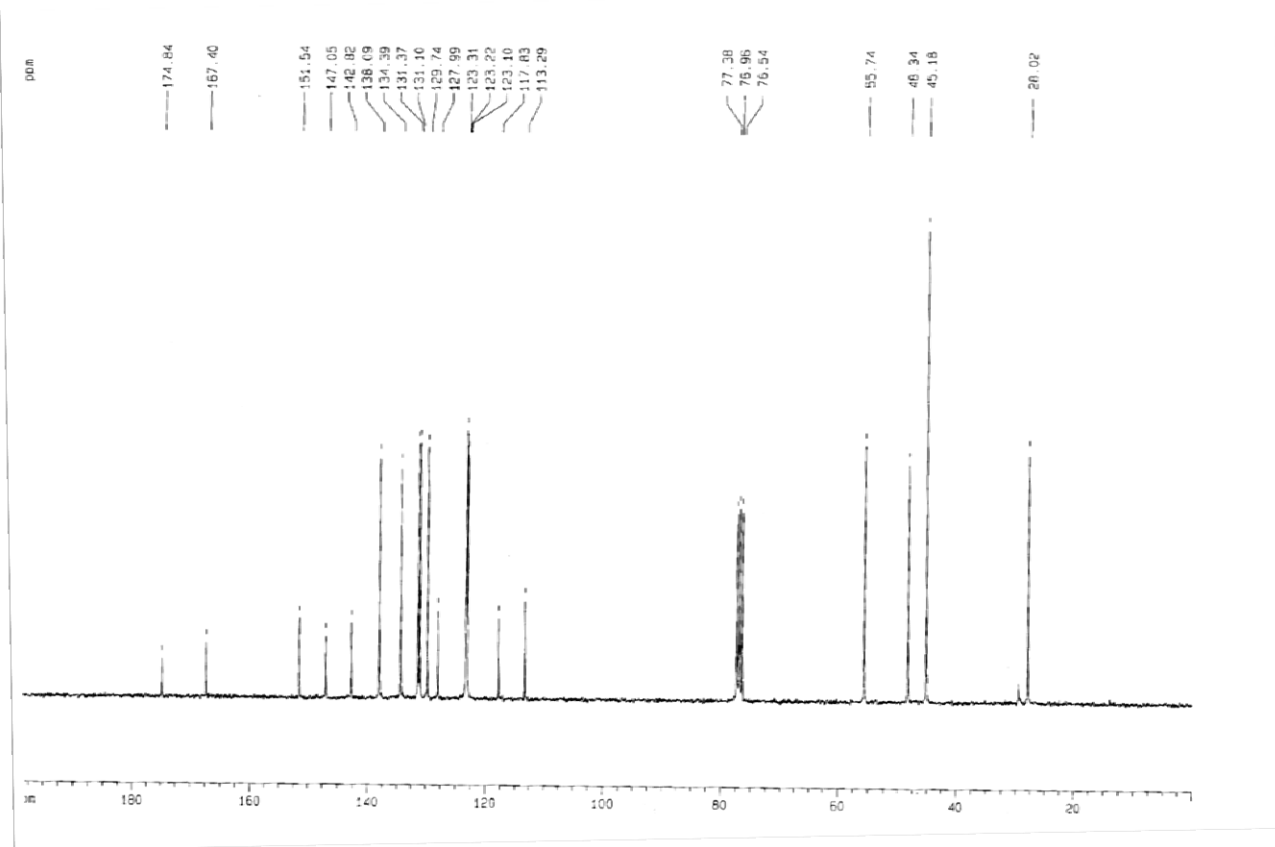
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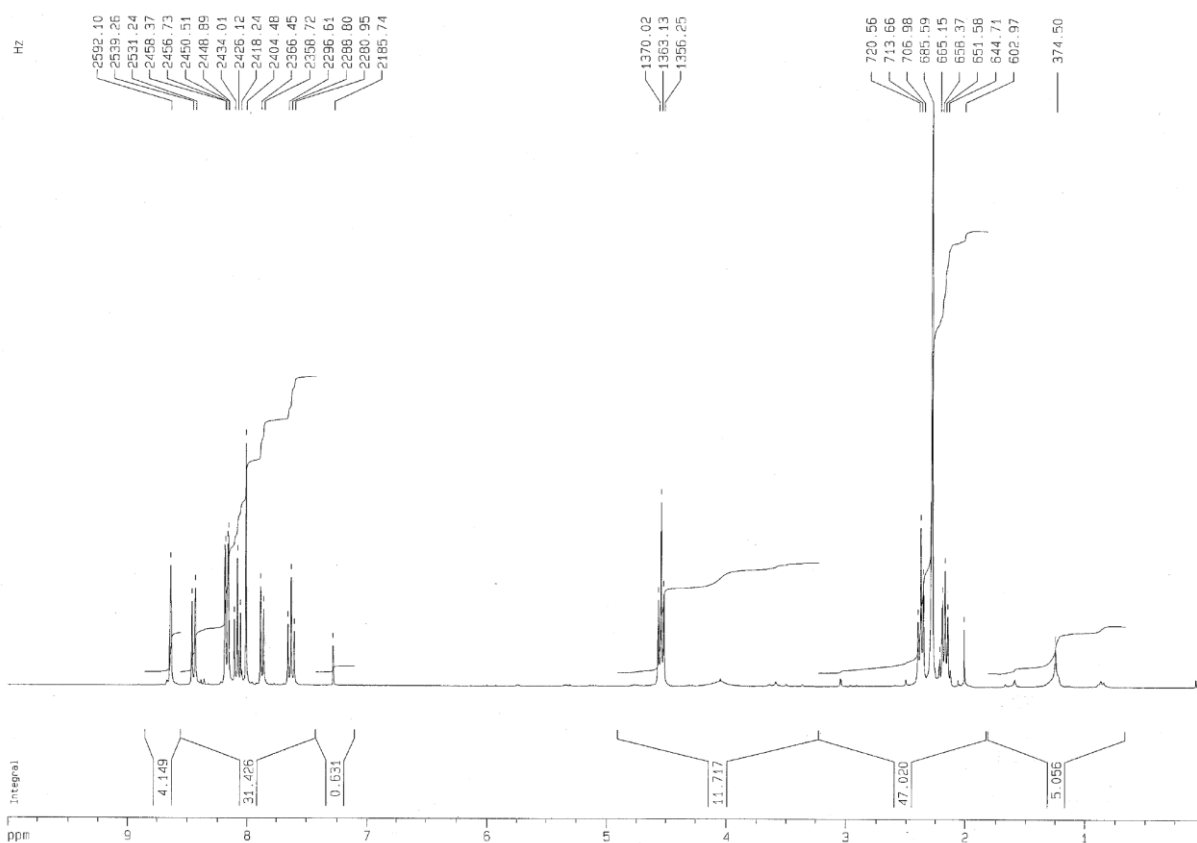
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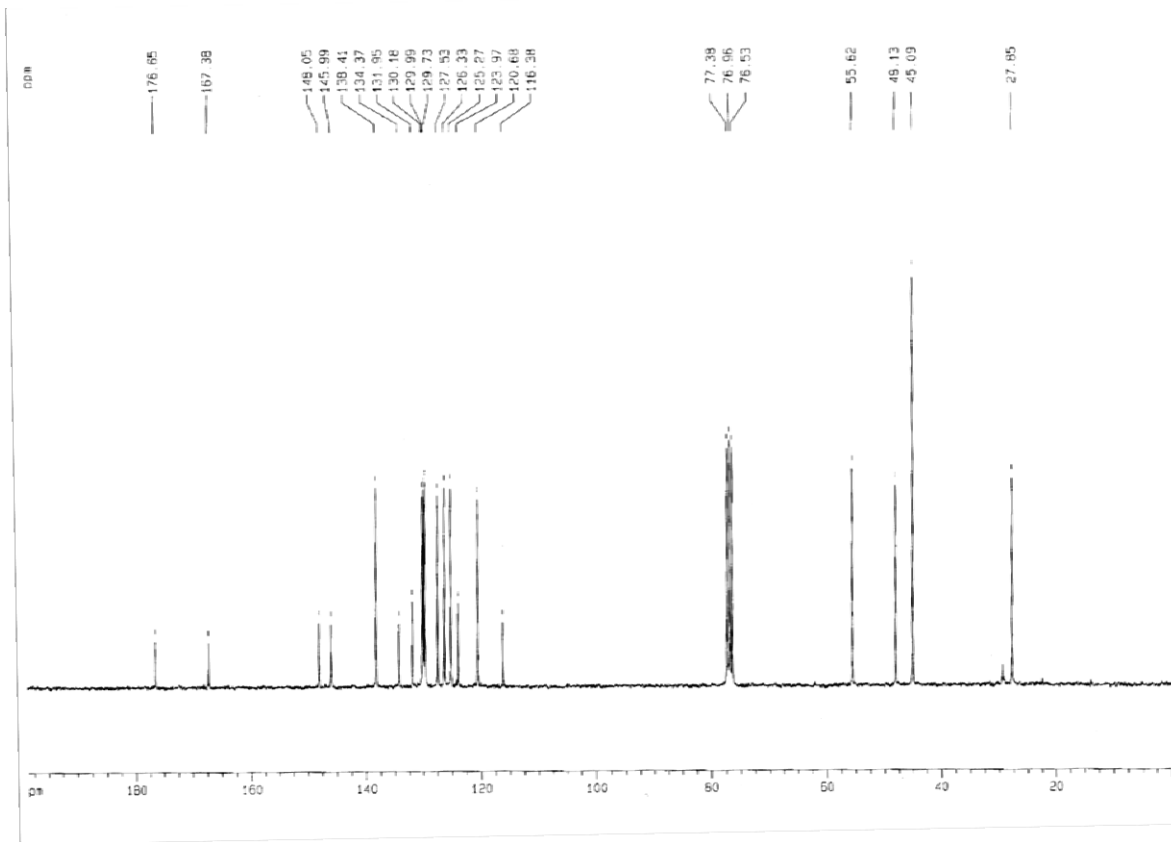
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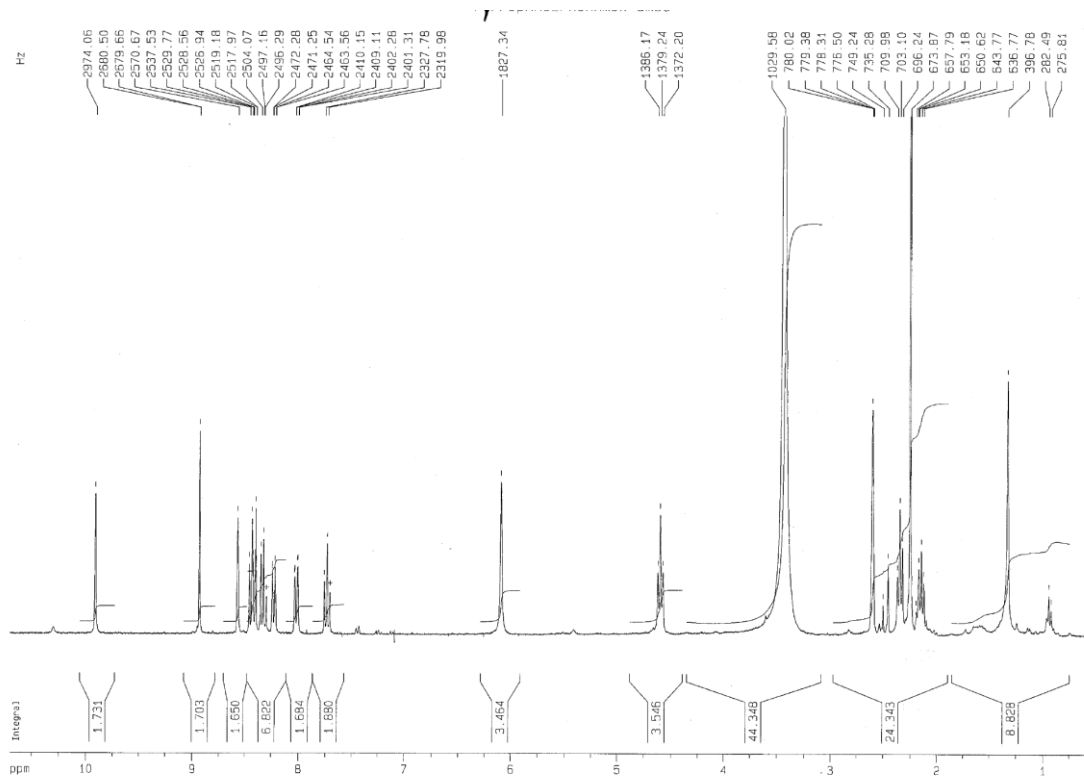
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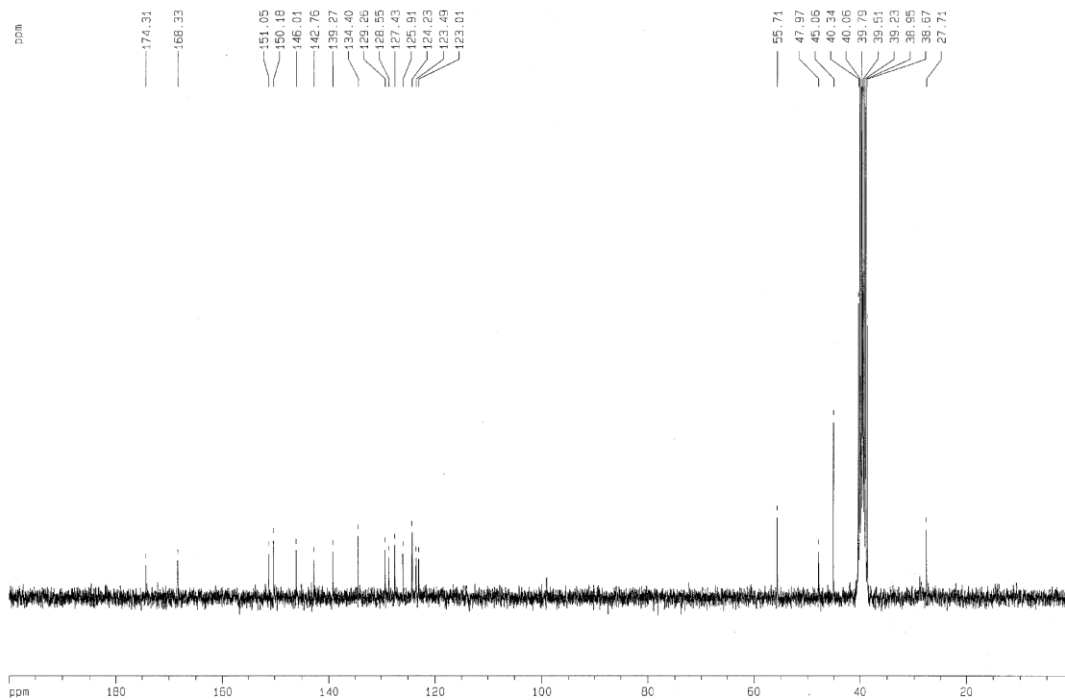
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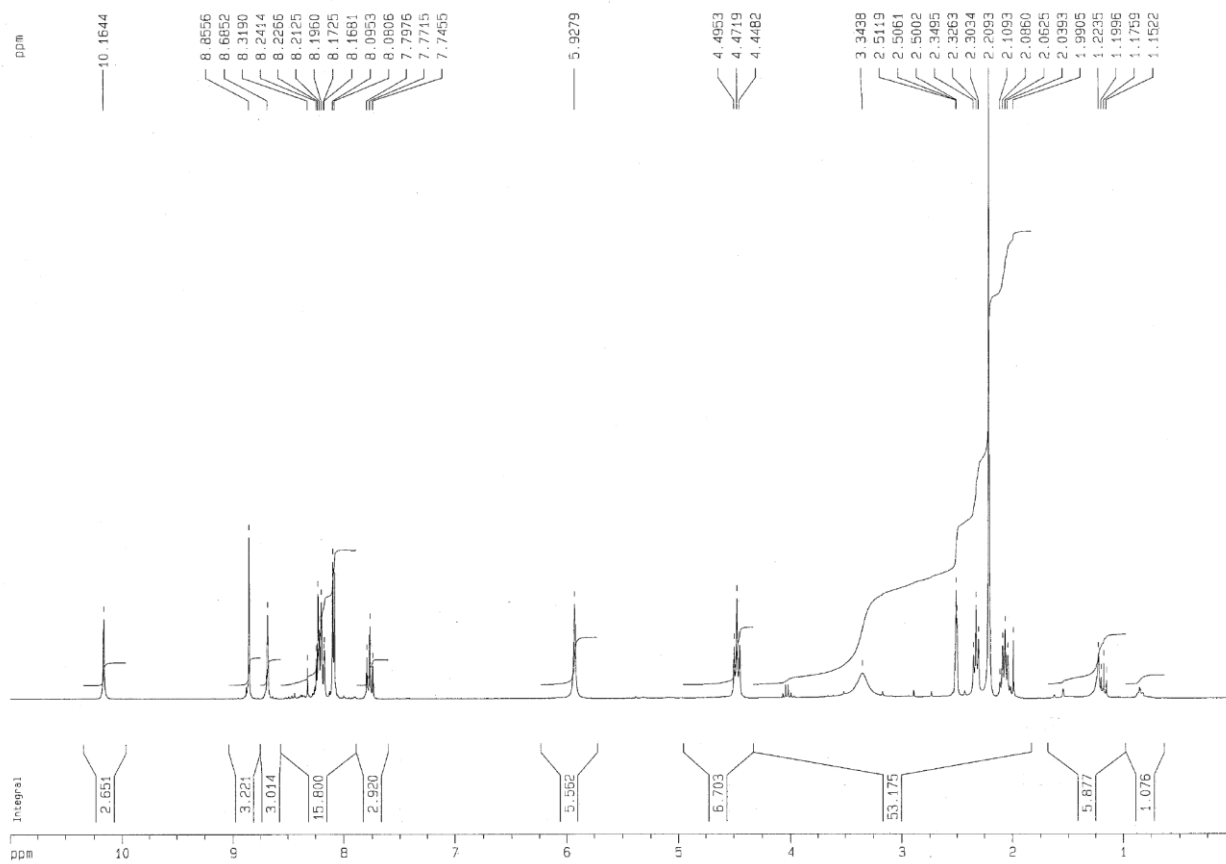
¹H-NMR – DMSO d₆: Fragment 35



¹³C-NMR – DMSO d6: Fragment 35



¹H-NMR – DMSO d6: Fragment 36



¹³C-NMR – DMSO d6: Fragment 36

