

N-Derivatives of (Z)-methyl 3-(4-oxo-2-thioxothiazolidin-5-ylidene) methyl)-1*H*-indole-2-carboxylates as antimicrobial agents. *In silico* and *in vitro* evaluation.

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Table S1. Prediction of organ toxicity and toxicity end points for compounds **1-17**.

| Comp. | Organ toxicity | | | Toxicity end points | | | |
|-----------|-----------------|-----------------|-----------------|---------------------|-----------------|-----------------|------------------|
| | Cardiotoxicity | Hepatotoxicity | Carcinogenicity | Immunotoxicity | Mutagenicity | Cytotoxicity | Phospholipidosis |
| 1 | Inactive (0.79) | Inactive (0.68) | Inactive (0.62) | Inactive(0.55) | Inactive (0.74) | Inactive(0.64) | Inactive (0.99) |
| 2 | Inactive (0.75) | Active (0.56) | Inactive (0.63) | Inactive (0.97) | Inactive (0.55) | Inactive (0.64) | Inactive (0.89) |
| 3 | Inactive (0.93) | Active (0.59) | Inactive (0.61) | Active (0.61) | Inactive (0.54) | Inactive(0.61) | Inactive (0.89) |
| 4 | Inactive (0.93) | Active (0.60) | Inactive (0.60) | Active(0.72) | Inactive (0.53) | Inactive(0.64) | Inactive (0.89) |
| 5 | Inactive (0.60) | Inactive (0.66) | Inactive (0.57) | Active (0.88) | Inactive (0.69) | Inactive (0.65) | Inactive (0.81) |
| 6 | Inactive (0.93) | Active (0.52) | Inactive (0.56) | Inactive (0.76) | Inactive (0.58) | Inactive (0.65) | Inactive (0.93) |
| 7 | Inactive (0.84) | Inactive (0.50) | Inactive (0.58) | Inactive (0.76) | Inactive (0.61) | Inactive (0.62) | Inactive (0.89) |
| 8 | Inactive (0.84) | Active (0.55) | Inactive (0.66) | Active (0.64) | Inactive (0.64) | Inactive (0.70) | Inactive (0.99) |
| 9 | Inactive (0.84) | Active (0.51) | Inactive (0.62) | Active (0.77) | Inactive (0.71) | Inactive (0.63) | Inactive (0.99) |
| 10 | Inactive (0.93) | Active (0.59) | Inactive (0.62) | Active (0.84) | Inactive (0.58) | Inactive (0.67) | Inactive (0.89) |
| 11 | Inactive (0.84) | Inactive (0.53) | Inactive (0.63) | Inactive (0.82) | Inactive (0.69) | Inactive (0.64) | Inactive (0.99) |
| 12 | Inactive (0.84) | Inactive (0.54) | Inactive (0.68) | Inactive (0.88) | Inactive (0.63) | Inactive (0.67) | Inactive (0.99) |
| 13 | Inactive (0.93) | Active (0.57) | Inactive (0.60) | Active (0.98) | Inactive (0.59) | Inactive (0.70) | Inactive (0.81) |
| 14 | Inactive (0.84) | Inactive (0.52) | Inactive (0.63) | Active (0.78) | Inactive (0.70) | Inactive (0.63) | Inactive (0.99) |
| 15 | Inactive (0.93) | Active (0.56) | Inactive (0.63) | Inactive (0.54) | Inactive (0.58) | Inactive (0.73) | Inactive (0.89) |
| 16 | Inactive (0.93) | Active (0.51) | Inactive (0.59) | Active (0.81) | Inactive (0.57) | Inactive (0.65) | Inactive (0.93) |
| 17 | Inactive (0.79) | Active (0.59) | Inactive (0.63) | Active (0.79) | Inactive (0.71) | Inactive (0.63) | Inactive (0.99) |

Number in brackets indicate possibilities. Cardiotoxicity is referred to the probability of inhibition of the inward rectifying voltage gated potassium channel in the heart encoded by the human ether-a-go-go related gene (hERG).

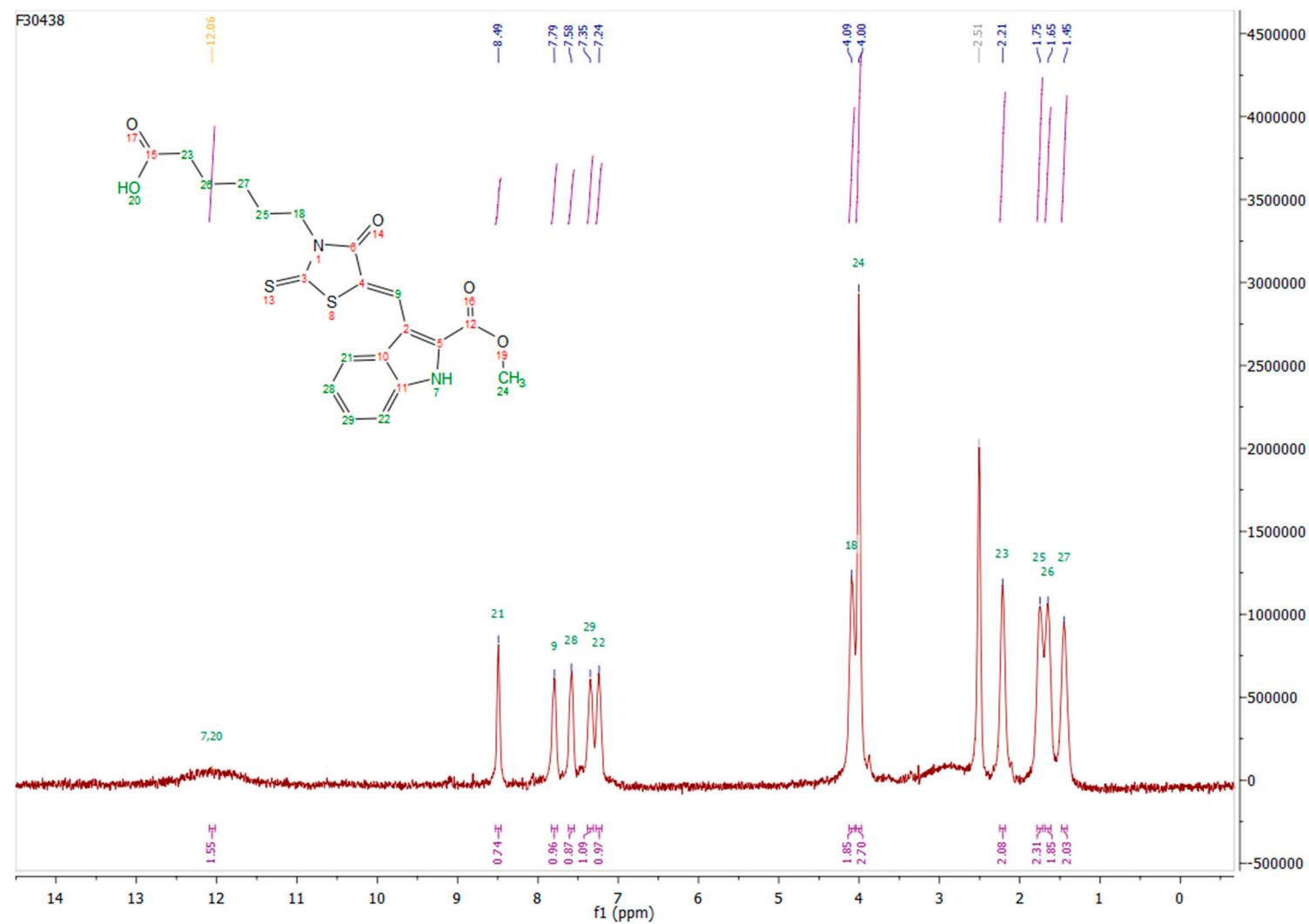
Table S2. Prediction of adverse outcomes according to TOX21 of compounds 1-17.

| Comp. | Tox21-Nuclear receptor signalling pathways | | | | | | | Tox21-Stress response pathways | | | | |
|-------|--|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------------------|--------------------|--------------------|--------------------|--------------------|
| | AhR | AR | AR-LBD | ARO | ER | ER-LBD | PPAR γ | NRF2/ARE | HSE | MMP | p53 | ATAD5 |
| 1 | Inactive (0.77) | Inactive (0.98) | Inactive (0.97) | Inactive (0.93) | Inactive (0.87) | Inactive (0.95) | Inactive (0.81) | Inactive (0.85) | Inactive (0.85) | Inactive (0.85) | Inactive (0.81) | Inactive (0.90) |
| 2 | Inactive (0.61) | Inactive (0.95) | Inactive (0.97) | Inactive (0.81) | Inactive (0.76) | Inactive (0.92) | Inactive (0.81) | Inactive (0.90) | Inactive (0.90) | Inactive (0.54) | Inactive (0.75) | Inactive (0.71) |
| 3 | Inactive (0.60) | Inactive (0.92) | Inactive (0.97) | Inactive (0.84) | Inactive (0.73) | Inactive (0.92) | Inactive (0.82) | Inactive (0.89) | Inactive (0.89) | Inactive (0.51) | Inactive (0.71) | Inactive (0.75) |
| 4 | Inactive (0.63) | Inactive (0.94) | Inactive (0.97) | Inactive (0.83) | Inactive (0.72) | Inactive (0.90) | Inactive (0.81) | Inactive (0.91) | Inactive (0.91) | Inactive (0.51) | Inactive (0.70) | Inactive (0.75) |
| 5 | Inactive (0.78) | Inactive (0.95) | Inactive (0.93) | Inactive (0.91) | Inactive (0.87) | Inactive (0.94) | Inactive (0.82) | Inactive (0.91) | Inactive (0.91) | Inactive (0.75) | Inactive (0.82) | Inactive (0.94) |
| 6 | Inactive (0.82) | Inactive (0.95) | Inactive (0.95) | Inactive (0.90) | Inactive (0.87) | Inactive (0.94) | Inactive (0.88) | Inactive (0.92) | Inactive (0.92) | Inactive (0.69) | Inactive (0.89) | Inactive (0.94) |
| 7 | Inactive (0.65) | Inactive (0.98) | Inactive (0.96) | Inactive (0.93) | Inactive (0.84) | Inactive (0.95) | Inactive (0.84) | Inactive (0.91) | Inactive (0.91) | Inactive (0.79) | Inactive (0.81) | Inactive (0.83) |
| 8 | Inactive (0.63) | Inactive (0.97) | Inactive (0.97) | Inactive (0.87) | Inactive (0.79) | Inactive (0.88) | Inactive (0.87) | Inactive (0.89) | Inactive (0.89) | Inactive (0.65) | Inactive (0.79) | Inactive (0.91) |
| 9 | Inactive (0.78) | Inactive (0.98) | Inactive (0.97) | Inactive (0.90) | Inactive (0.84) | Inactive (0.90) | Inactive (0.79) | Inactive (0.85) | Inactive (0.85) | Inactive (0.82) | Inactive (0.82) | Inactive (0.87) |
| 10 | Inactive (0.65) | Inactive (0.96) | Inactive (0.96) | Inactive (0.87) | Inactive (0.71) | Inactive (0.83) | Inactive (0.78) | Inactive (0.89) | Inactive (0.89) | Inactive (0.54) | Inactive (0.73) | Inactive (0.82) |
| 11 | Inactive (0.72) | Inactive (0.98) | Inactive (0.97) | Inactive (0.93) | Inactive (0.89) | Inactive (0.95) | Inactive (0.80) | Inactive (0.86) | Inactive (0.86) | Inactive (0.83) | Inactive (0.84) | Inactive (0.79) |
| 12 | Inactive (0.64) | Inactive (0.97) | Inactive (0.97) | Inactive (0.91) | Inactive (0.82) | Inactive (0.92) | Inactive (0.88) | Inactive (0.90) | Inactive (0.90) | Inactive (0.67) | Inactive (0.79) | Inactive (0.89) |
| 13 | Inactive (0.63) | Inactive (0.95) | Inactive (0.95) | Inactive (0.88) | Inactive (0.72) | Inactive (0.88) | Inactive (0.86) | Inactive (0.89) | Inactive (0.89) | Inactive (0.53) | Inactive (0.75) | Inactive (0.84) |
| 14 | Inactive (0.78) | Inactive (0.98) | Inactive (0.96) | Inactive (0.89) | Inactive (0.84) | Inactive (0.89) | Inactive (0.81) | Inactive (0.86) | Inactive (0.86) | Inactive (0.82) | Inactive (0.83) | Inactive (0.86) |
| 15 | Inactive (0.58) | Inactive (0.97) | Inactive (0.96) | Inactive (0.89) | Inactive (0.80) | Inactive (0.93) | Inactive (0.89) | Inactive (0.91) | Inactive (0.91) | Inactive (0.65) | Inactive (0.77) | Inactive (0.88) |
| 16 | Inactive (0.84) | Inactive (0.97) | Inactive (0.93) | Inactive (0.91) | Inactive (0.80) | Inactive (0.90) | Inactive (0.90) | Inactive (0.91) | Inactive (0.91) | Inactive (0.73) | Inactive (0.89) | Inactive (0.93) |
| 17 | Inactive (0.79) | Inactive (0.98) | Inactive (0.96) | Inactive (0.94) | Inactive (0.89) | Inactive (0.95) | Inactive (0.80) | Inactive (0.89) | Inactive (0.89) | Inactive (0.85) | Inactive (0.81) | Inactive (0.84) |

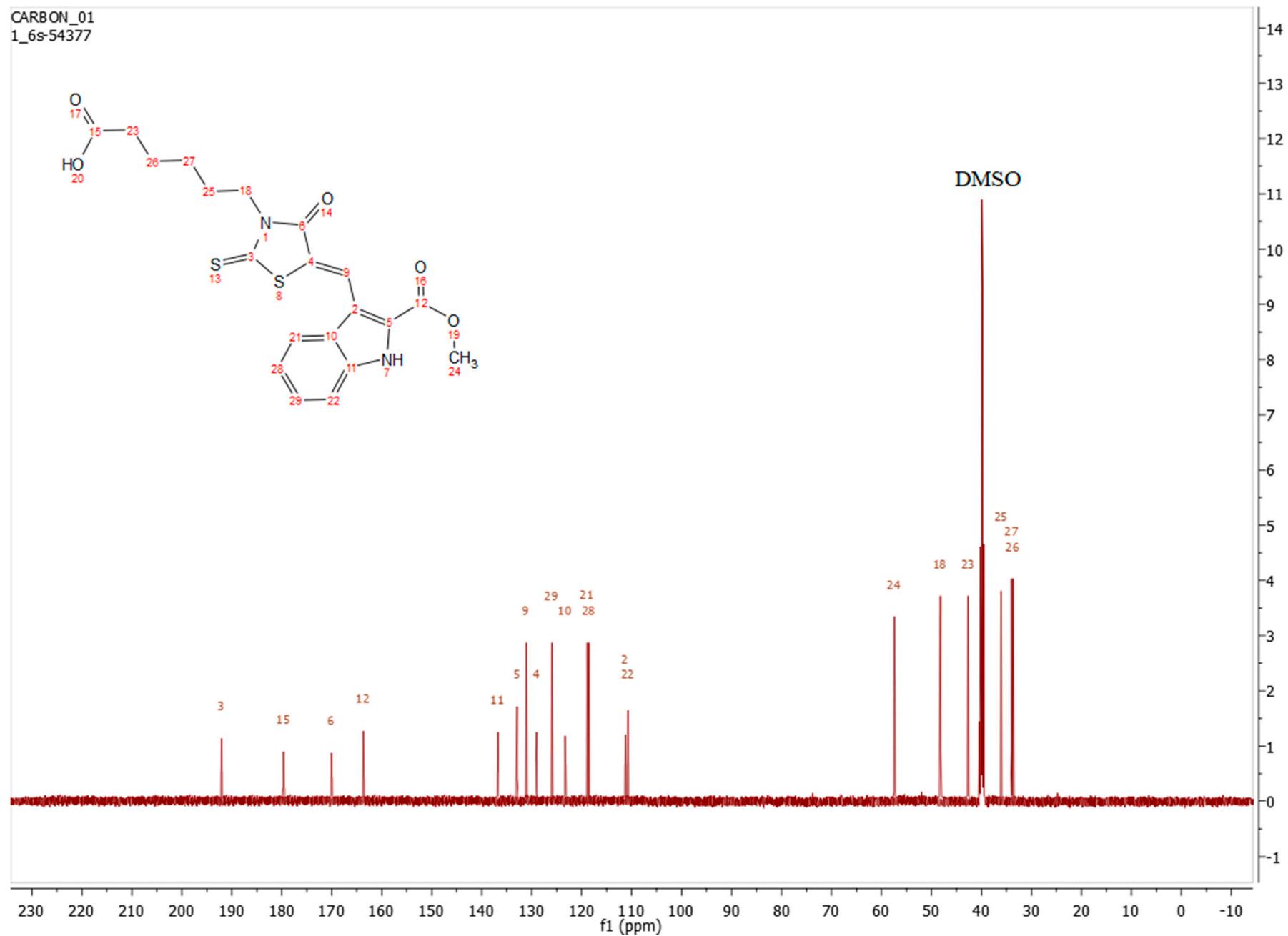
(1) Aryl hydrocarbon Receptor, (2) Androgen Receptor, (3) Androgen Receptor Ligand Binding Domain, (4) Aromatase, (5) Estrogen Receptor Alpha, (6) Estrogen Receptor Ligand Binding Domain, (7) Peroxisome Proliferator Activated Receptor Gamma, (8) Nuclear factor (erythroid-derived 2)-like 2/antioxidant responsive element, (9) Heat shock factor response element, (10) Heat shock factor response element, (11) Phosphoprotein (Tumor Suppressor) p53, (12) ATPase family AAA domain-containing protein 5

¹H-NMR and ¹³C-NMR of compounds

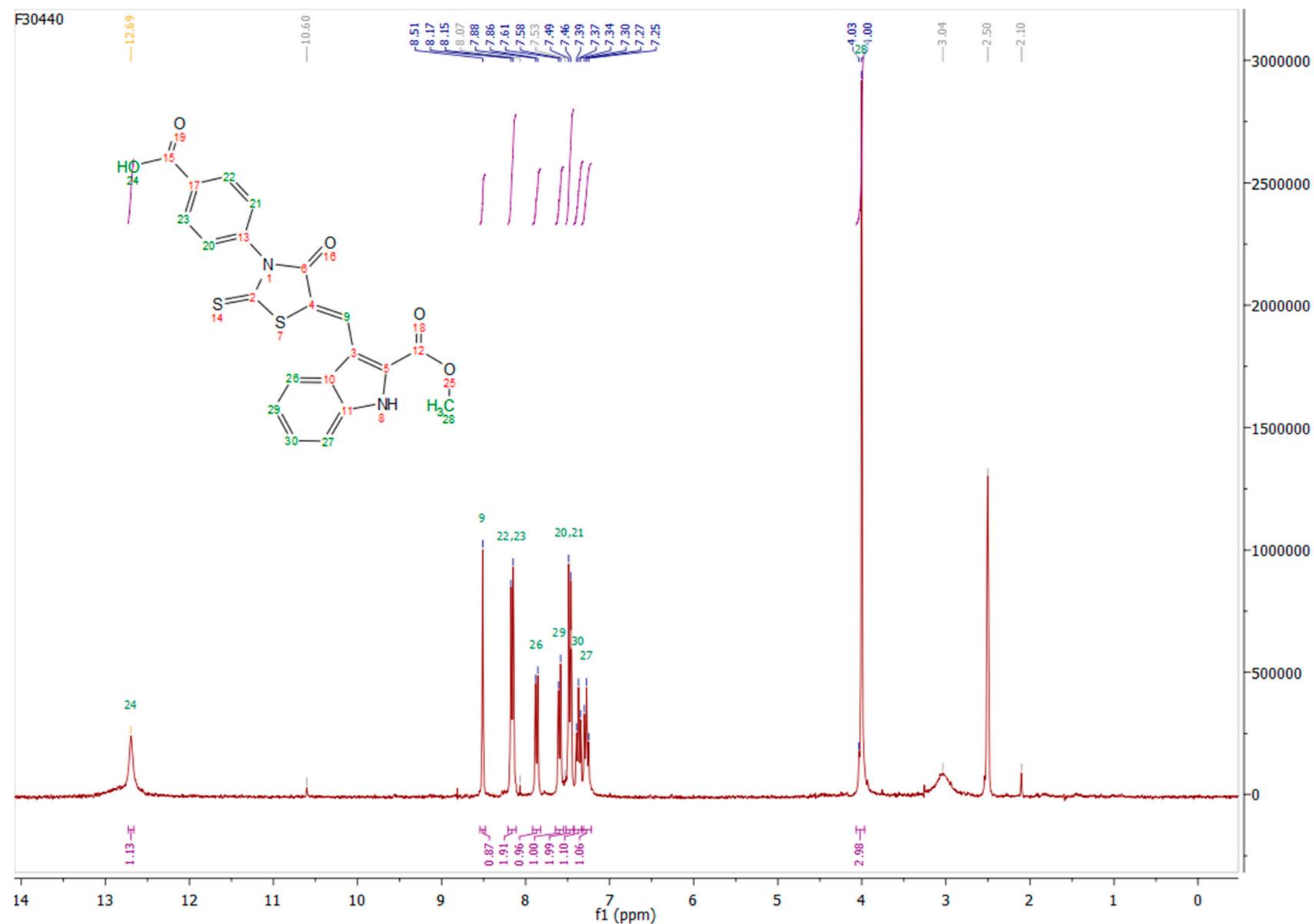
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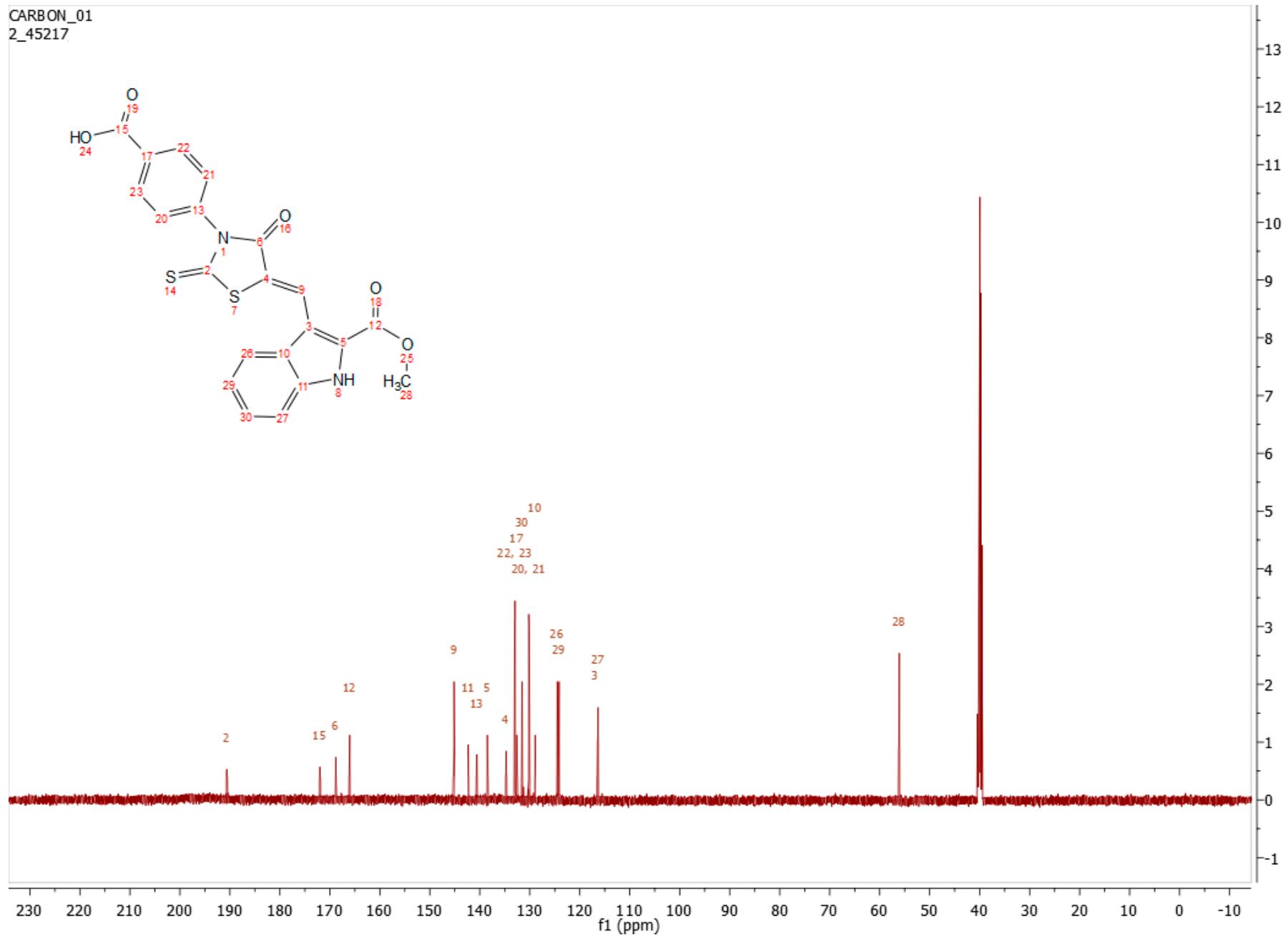
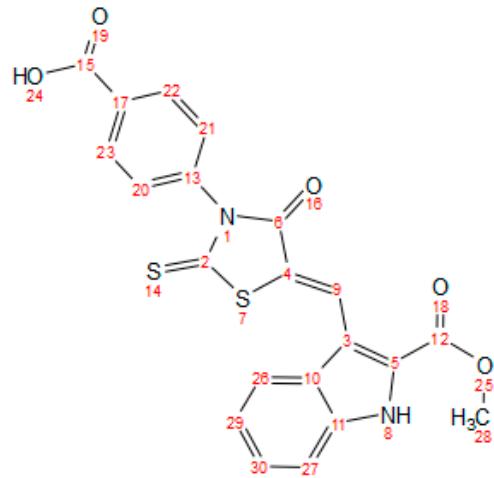
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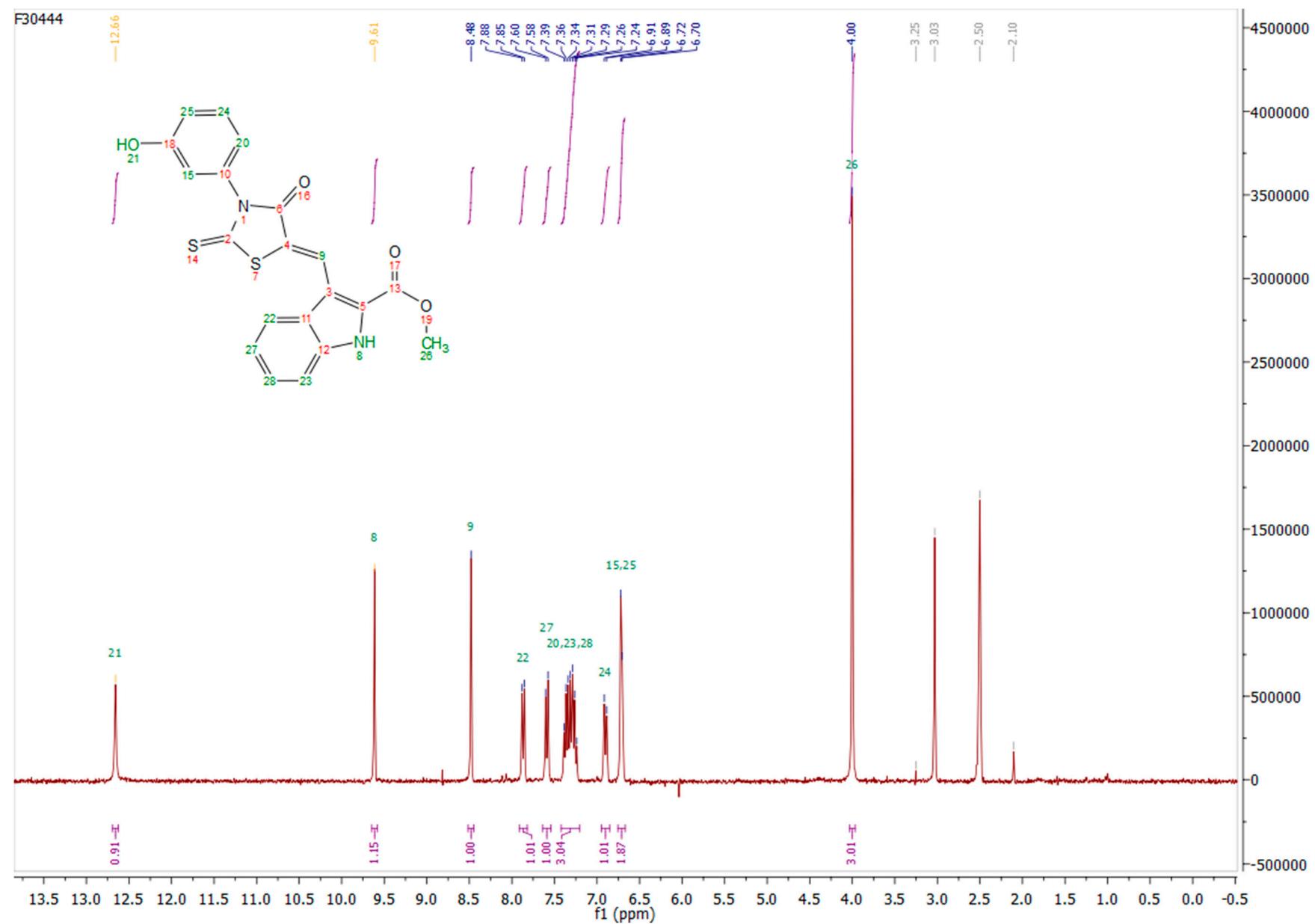
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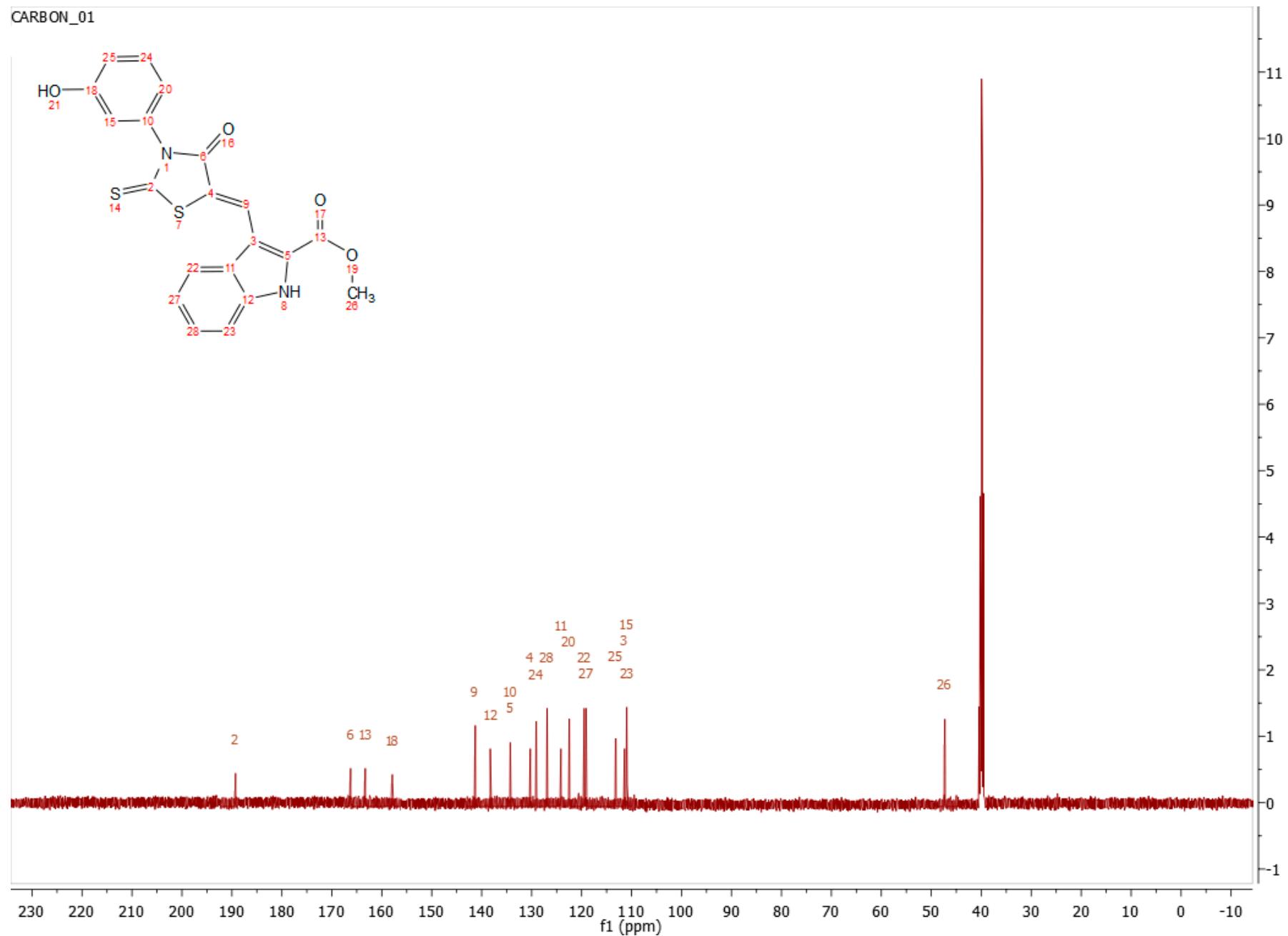
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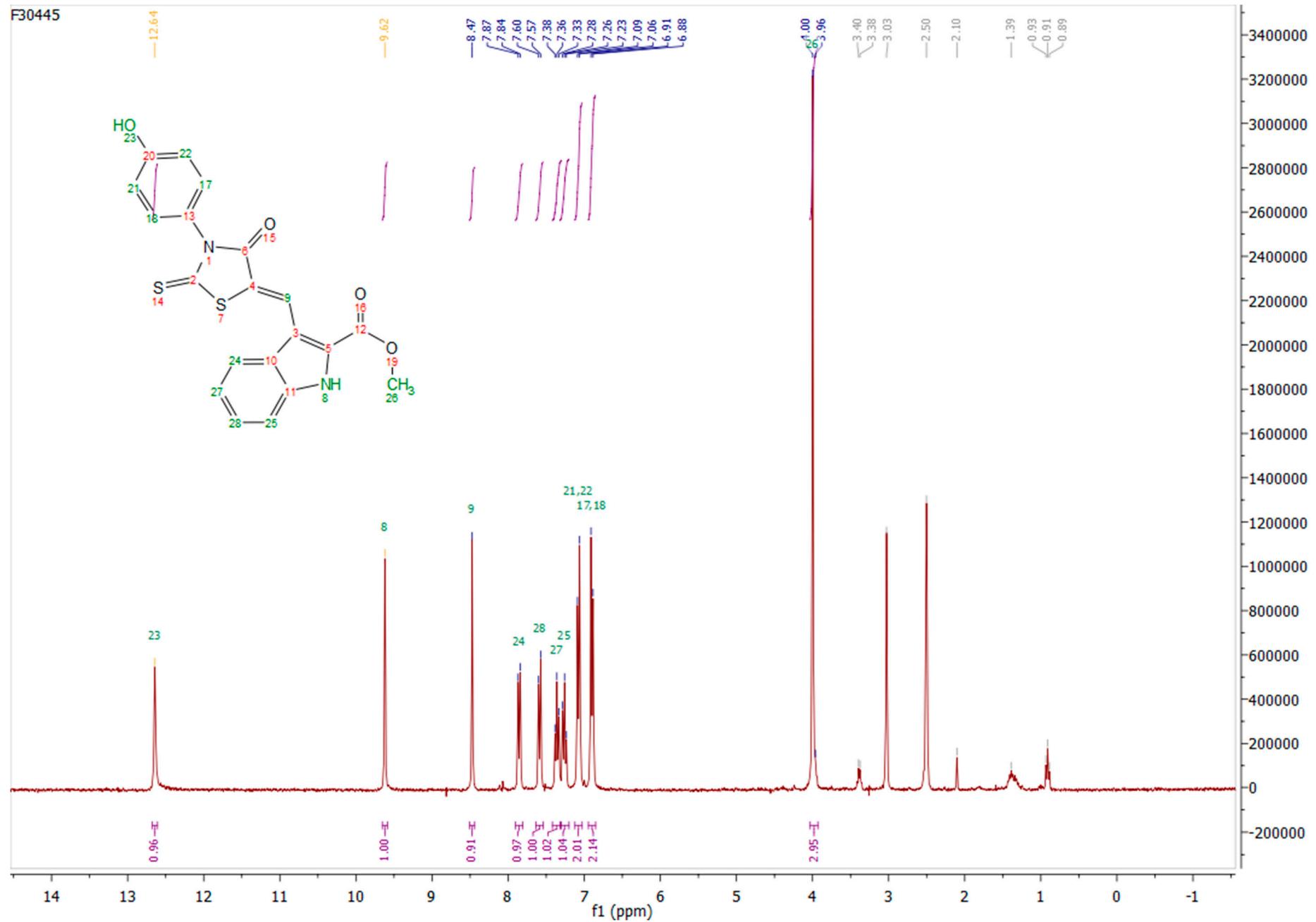
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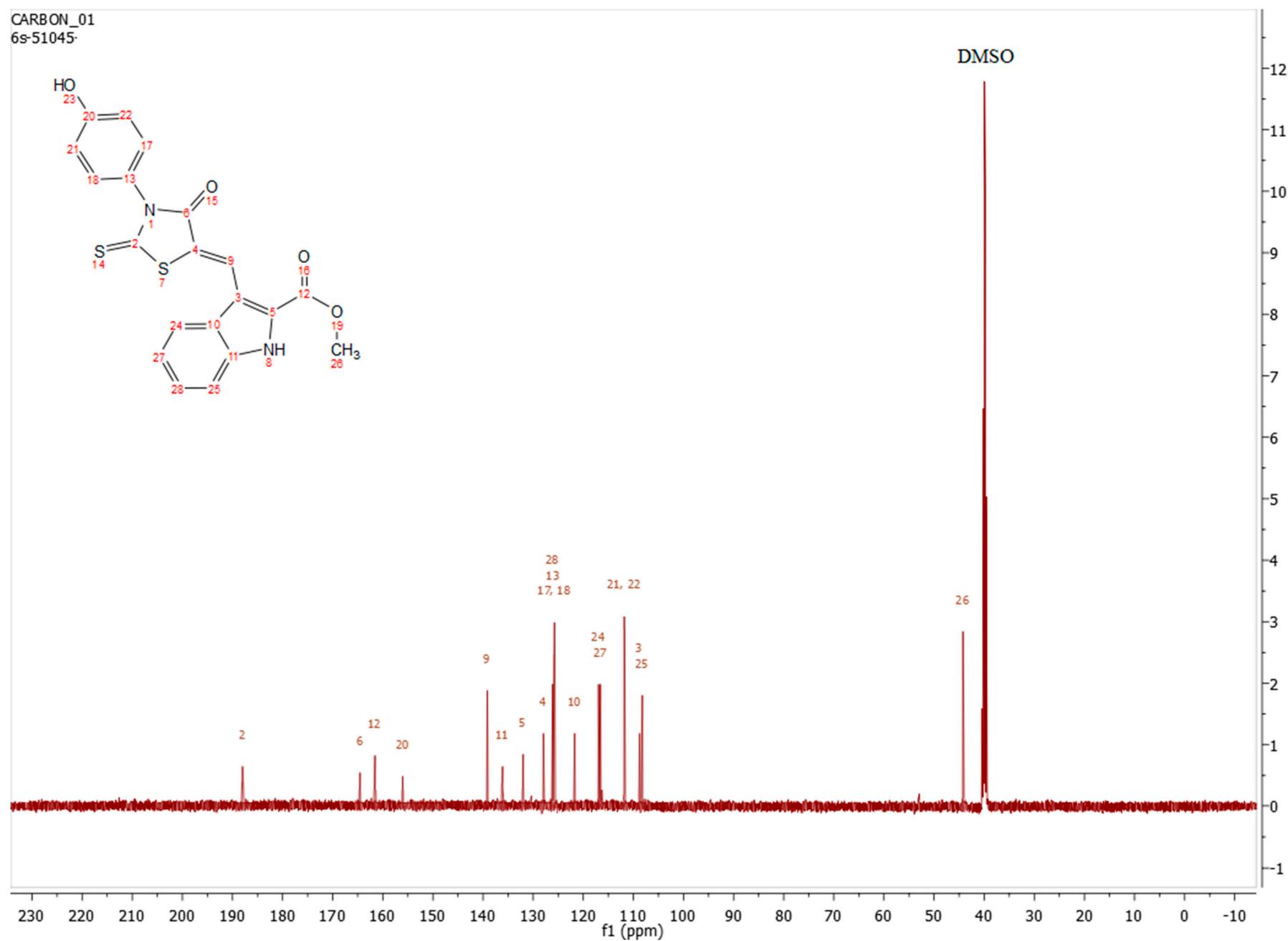
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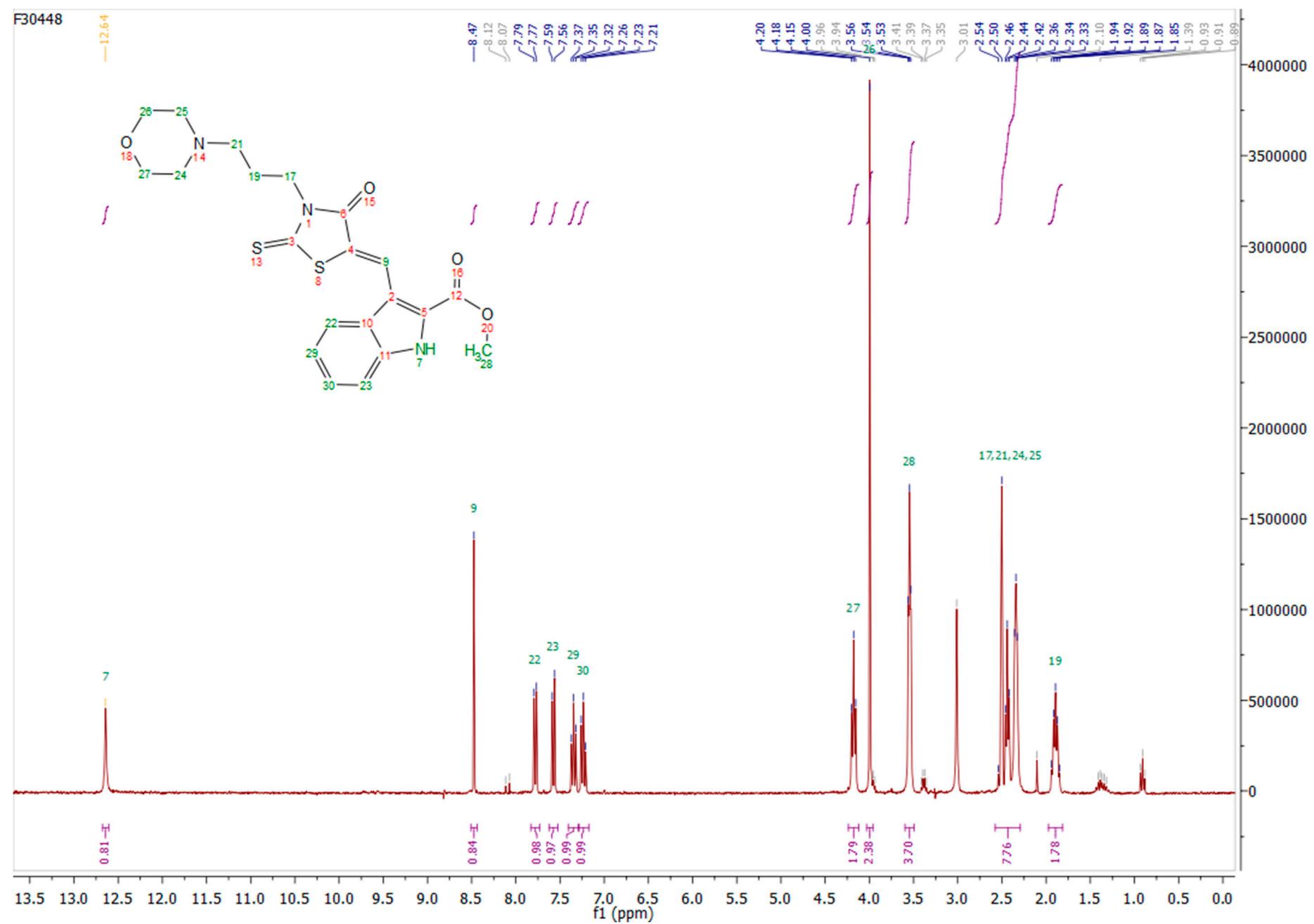
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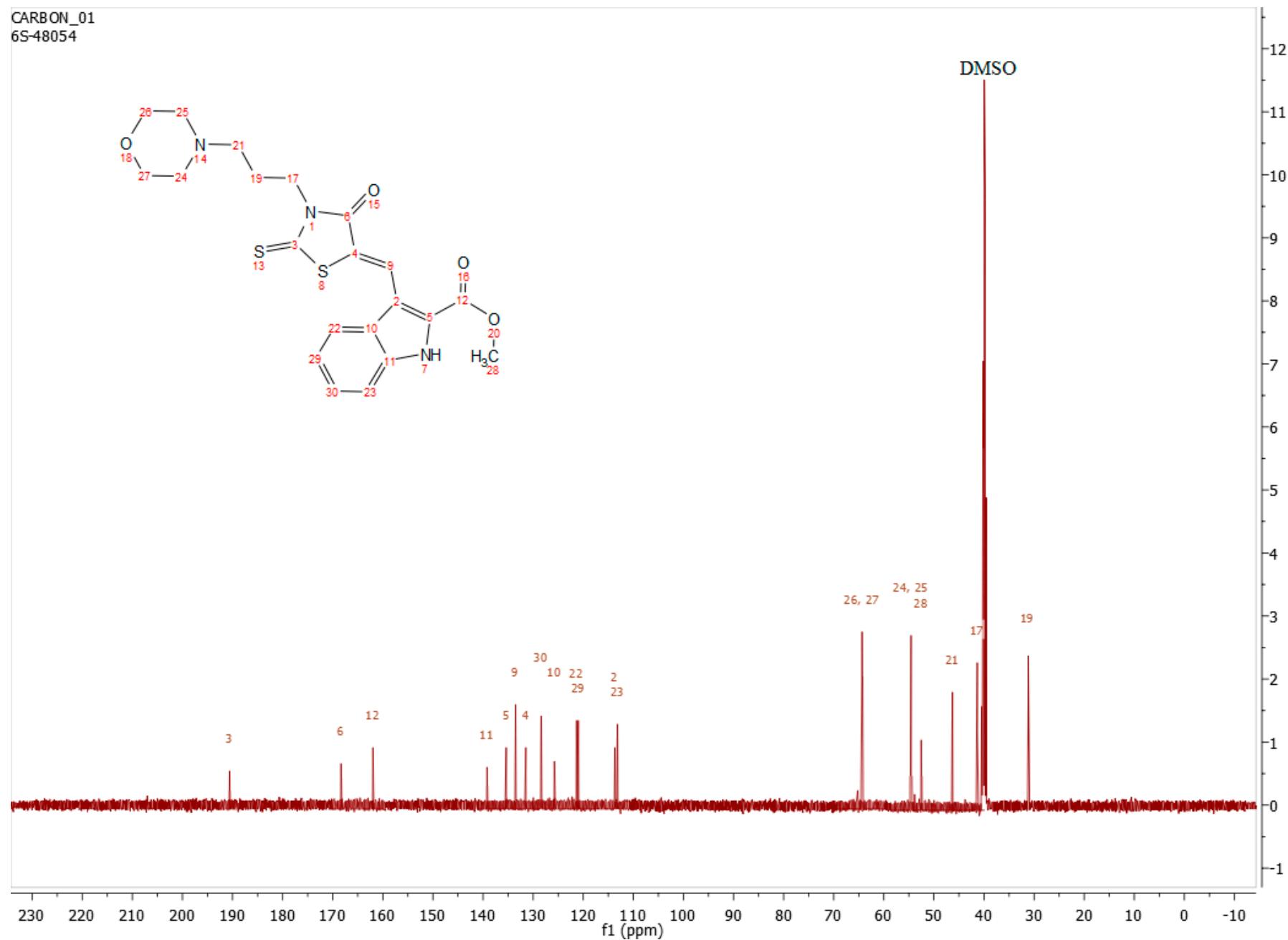
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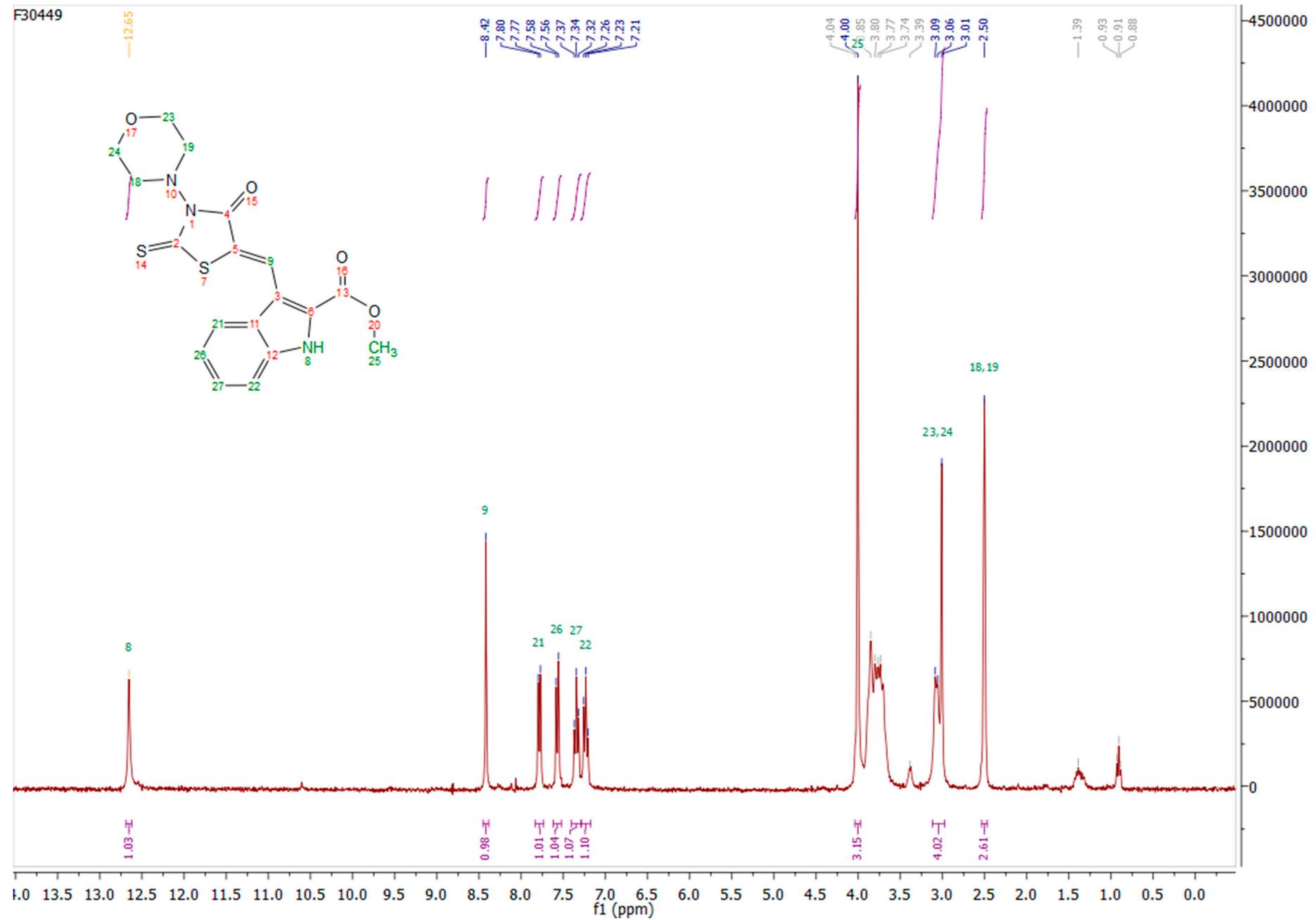
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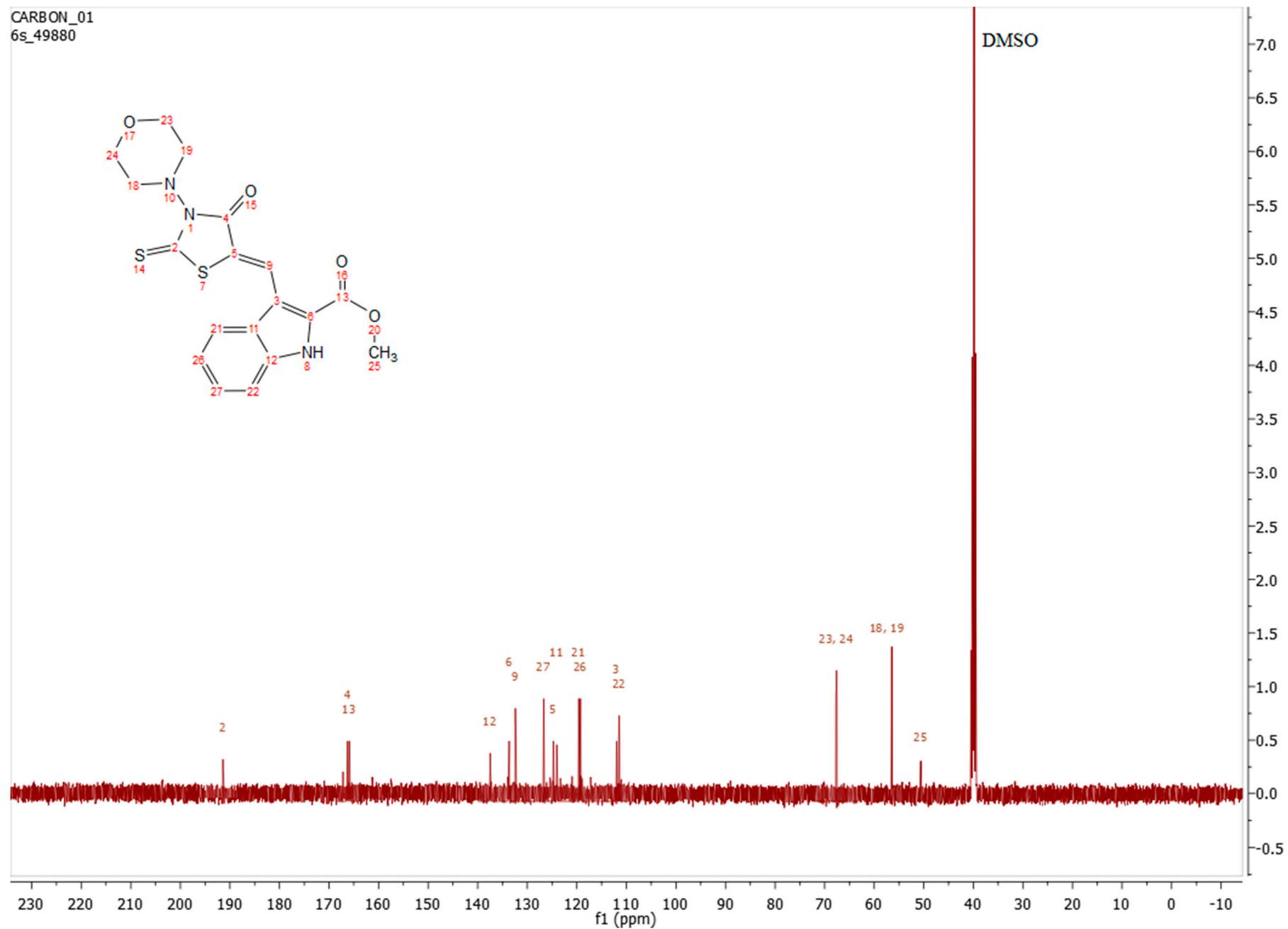
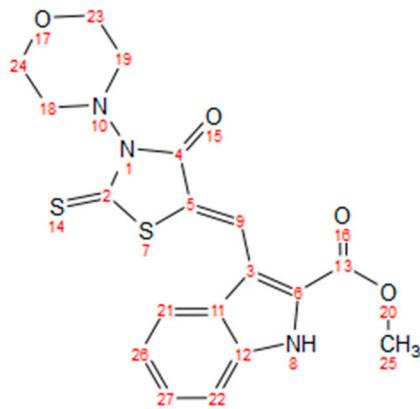
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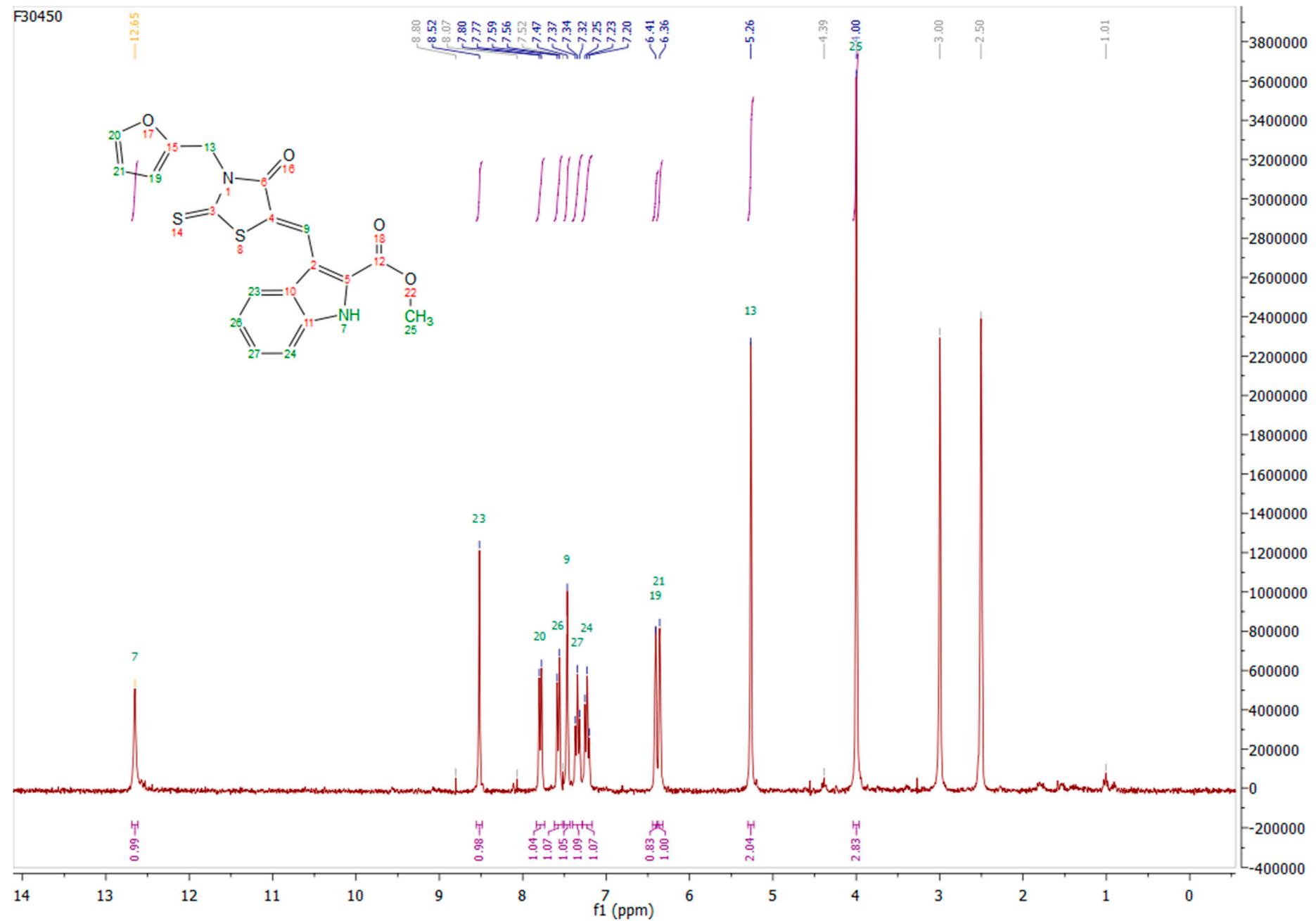
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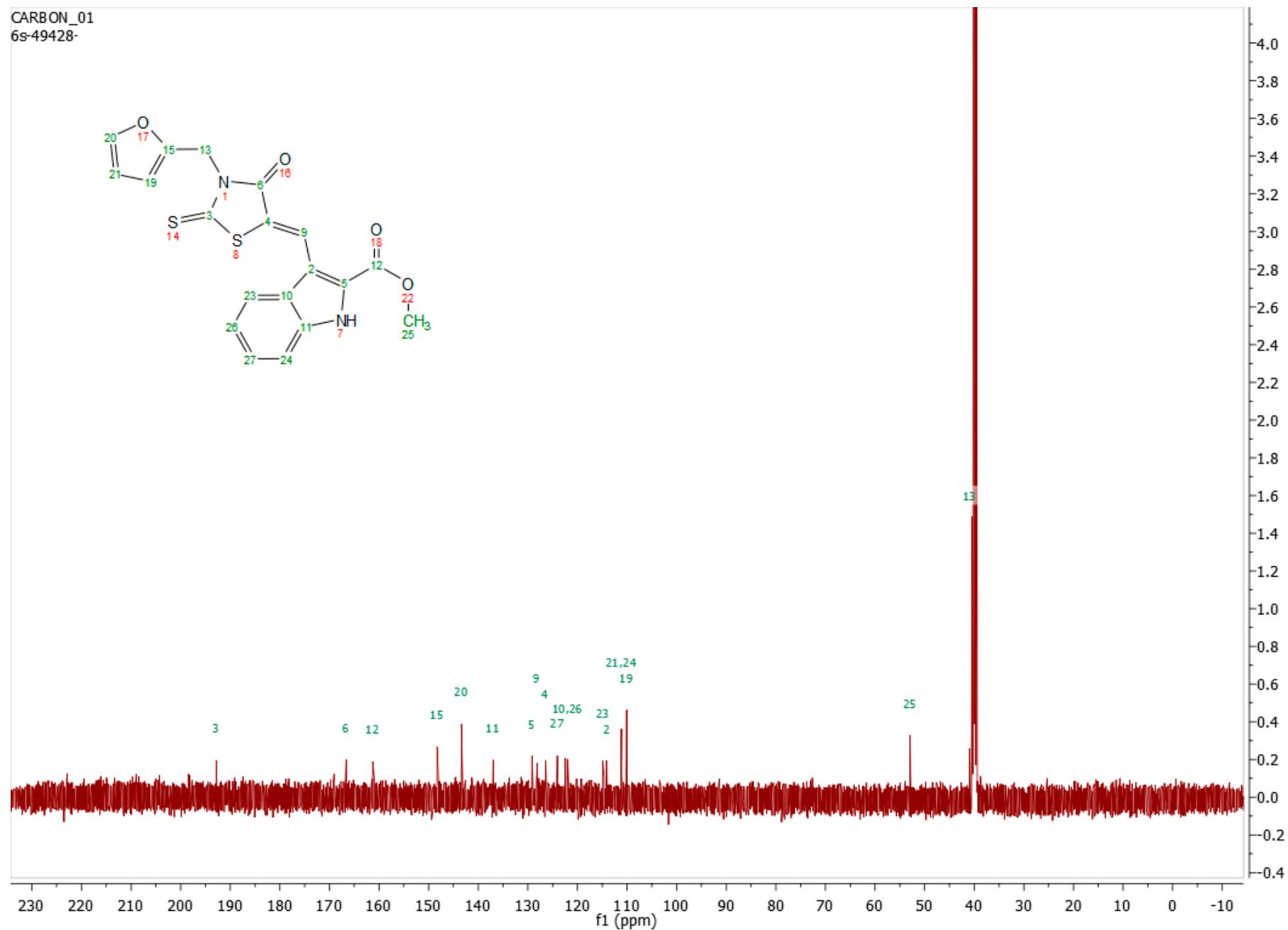
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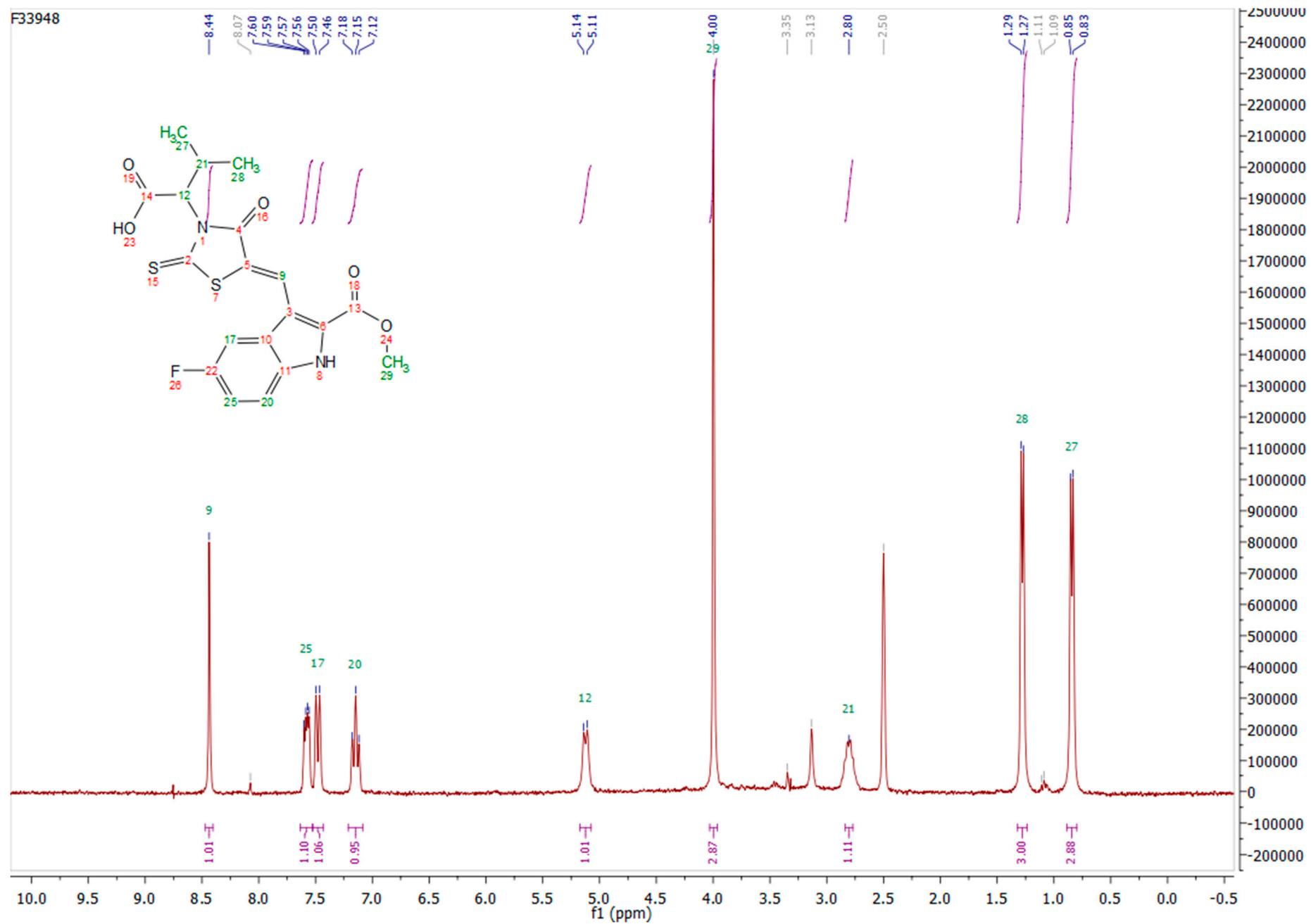
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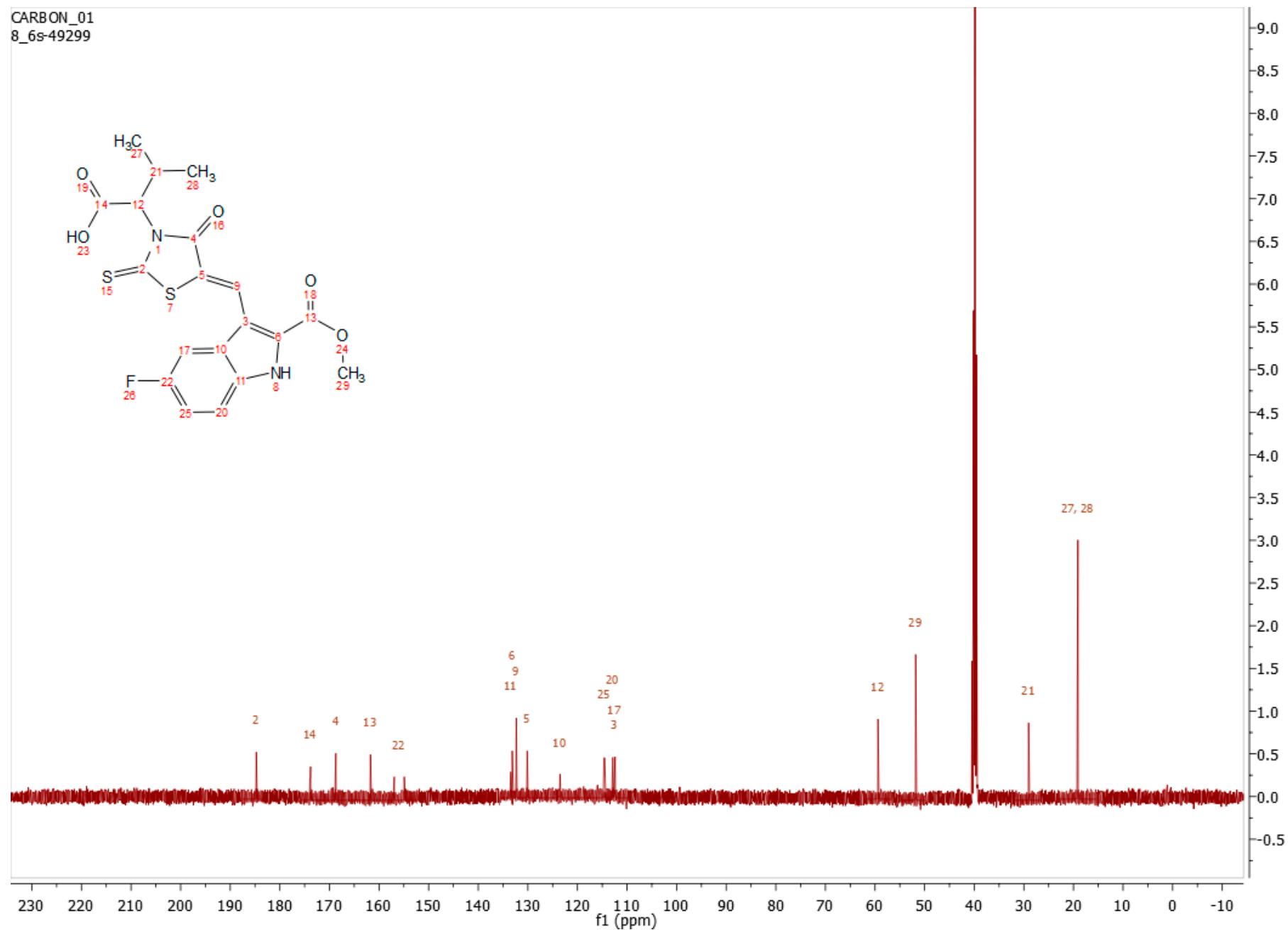
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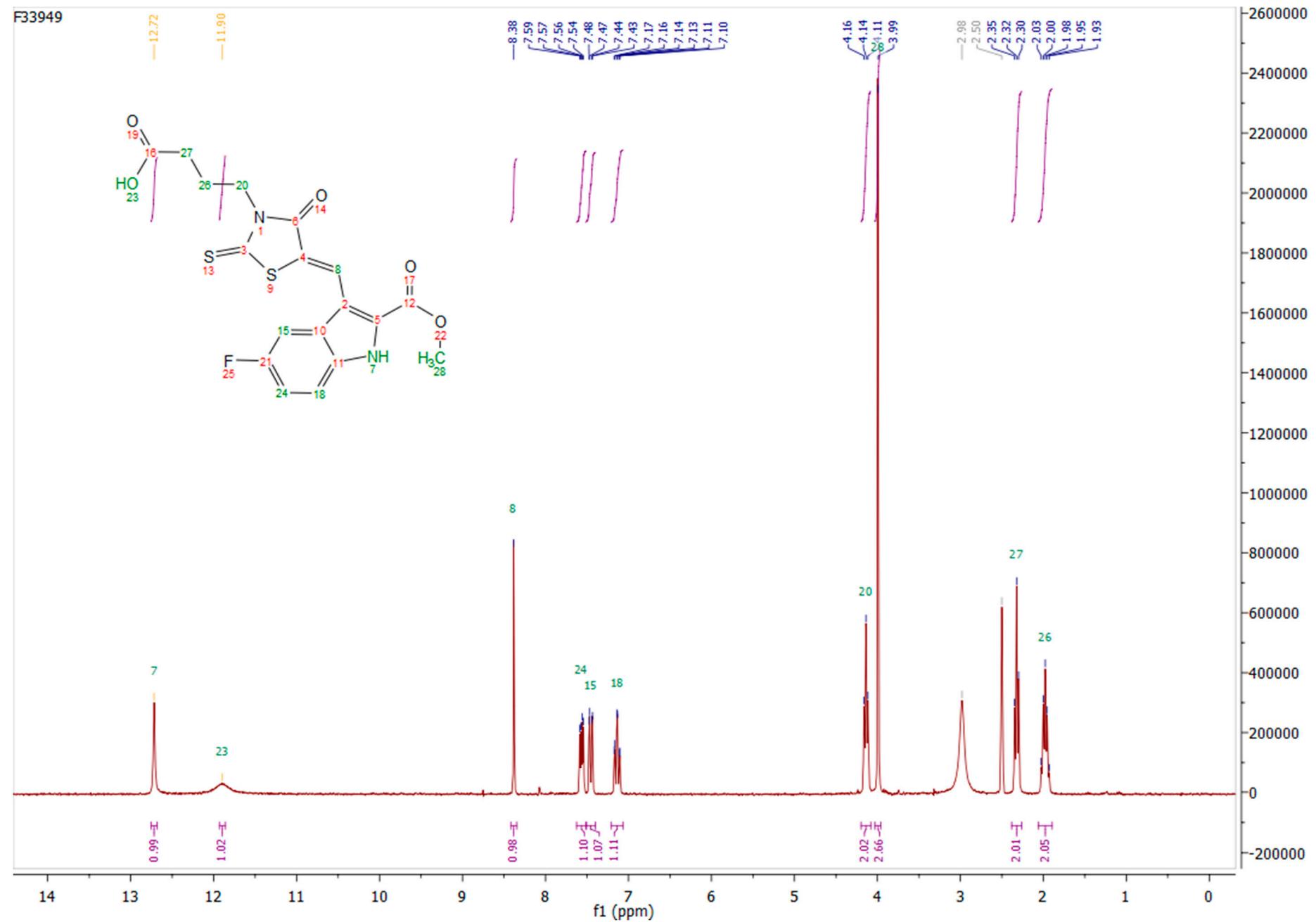
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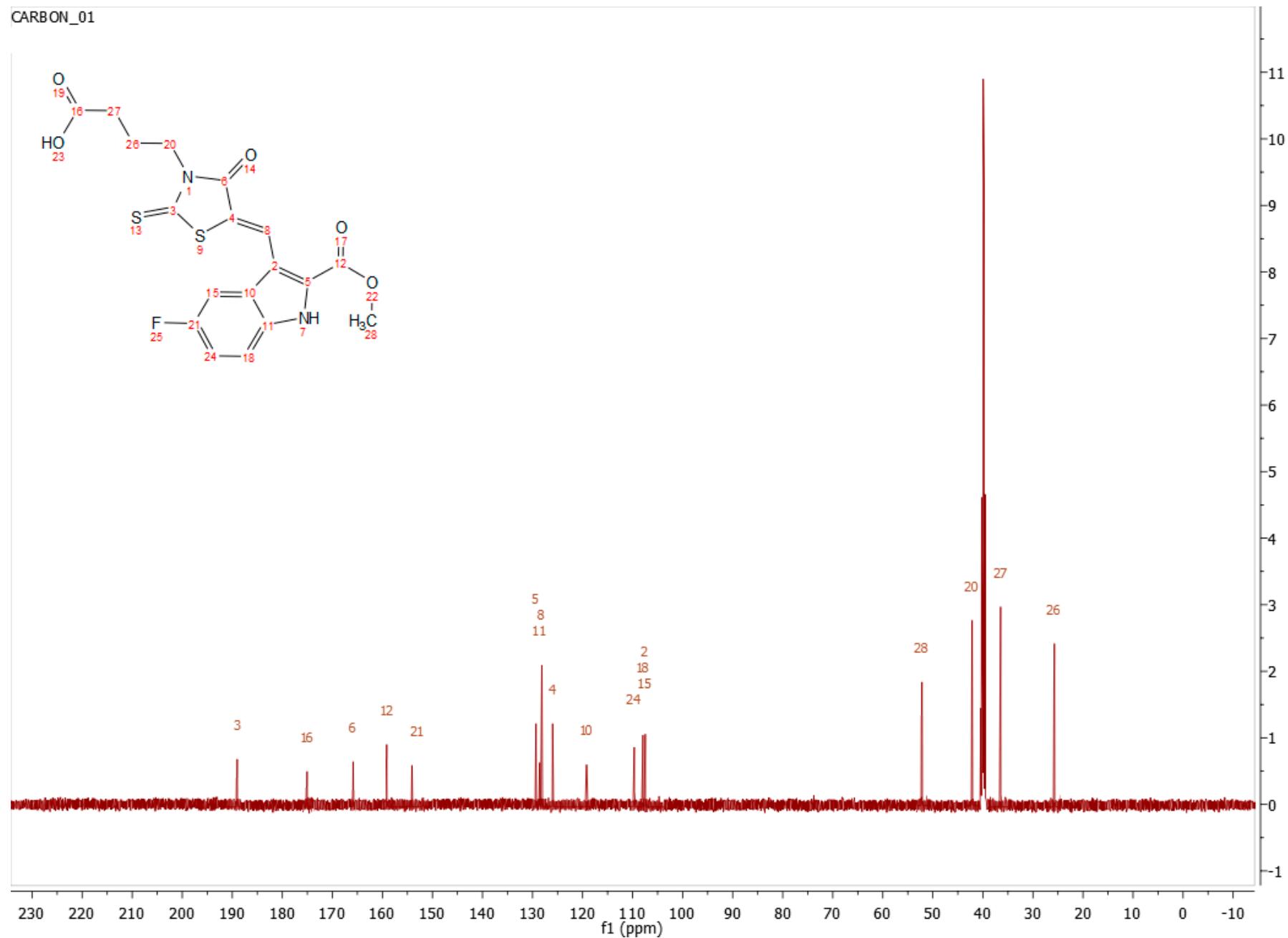
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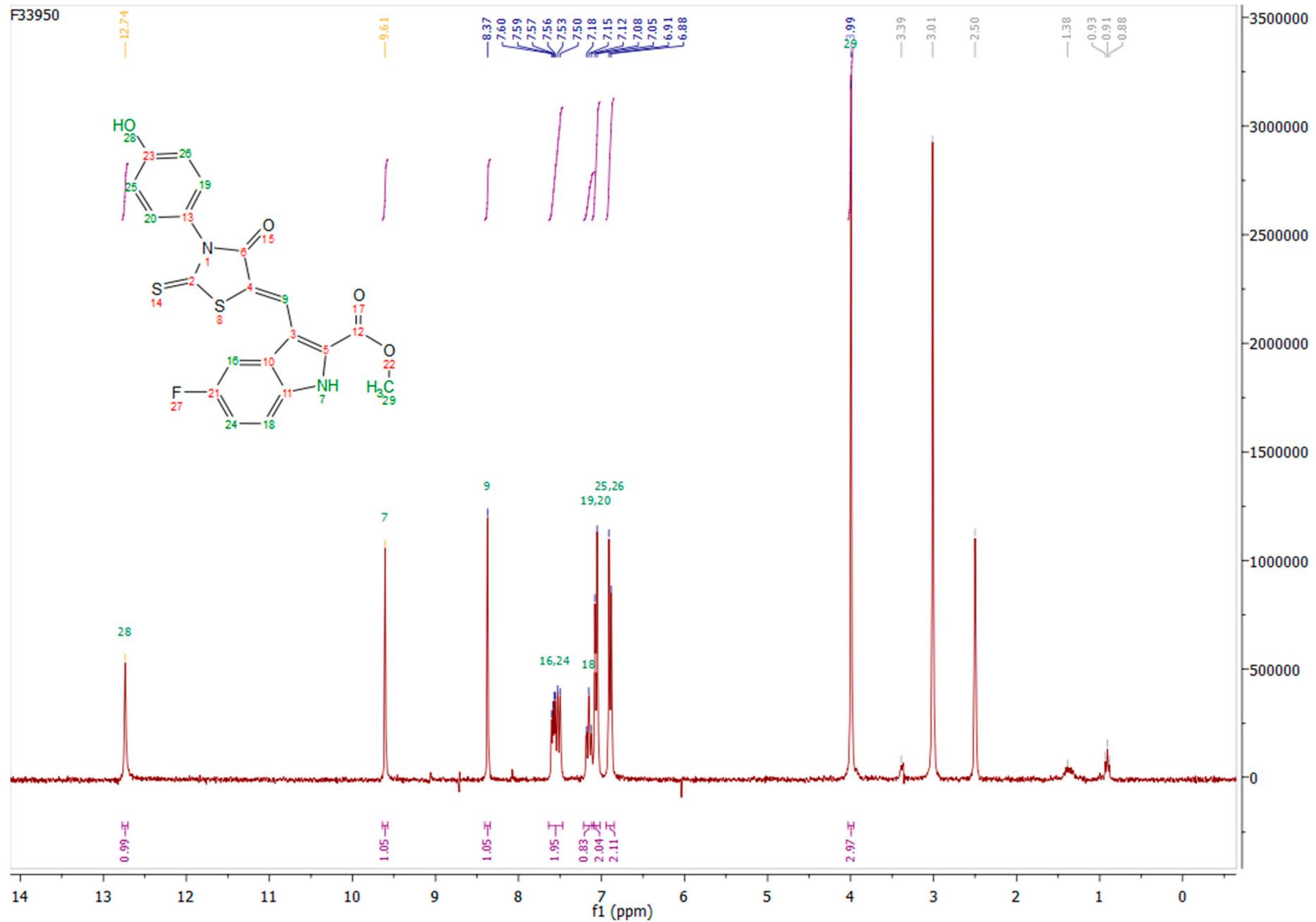
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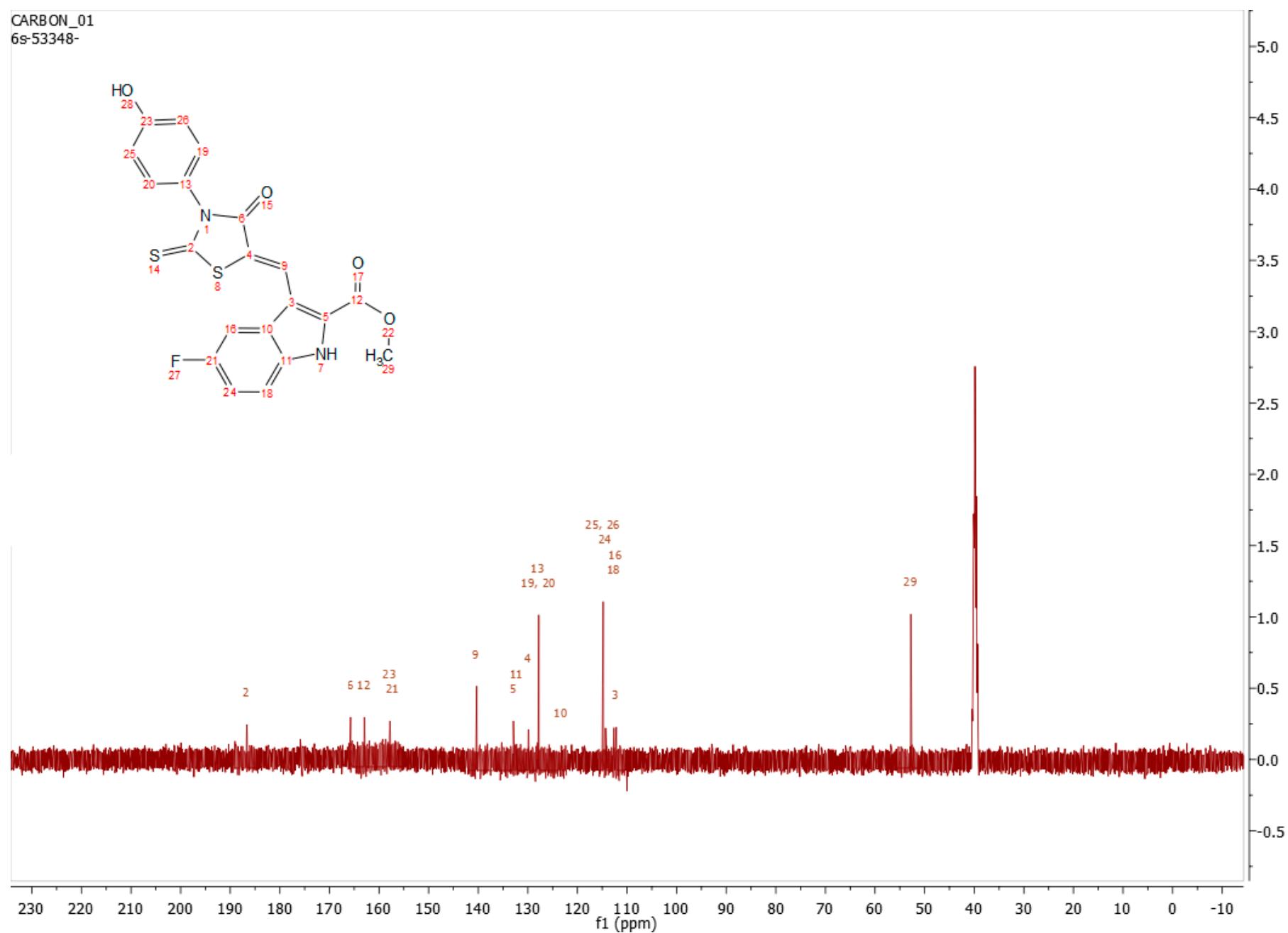
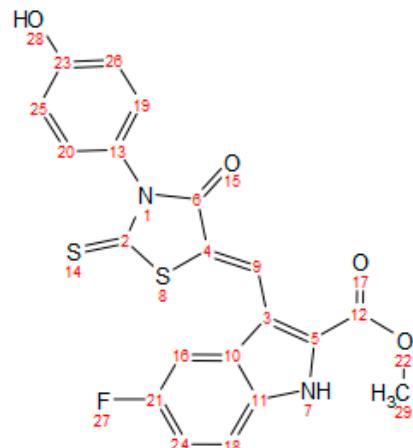
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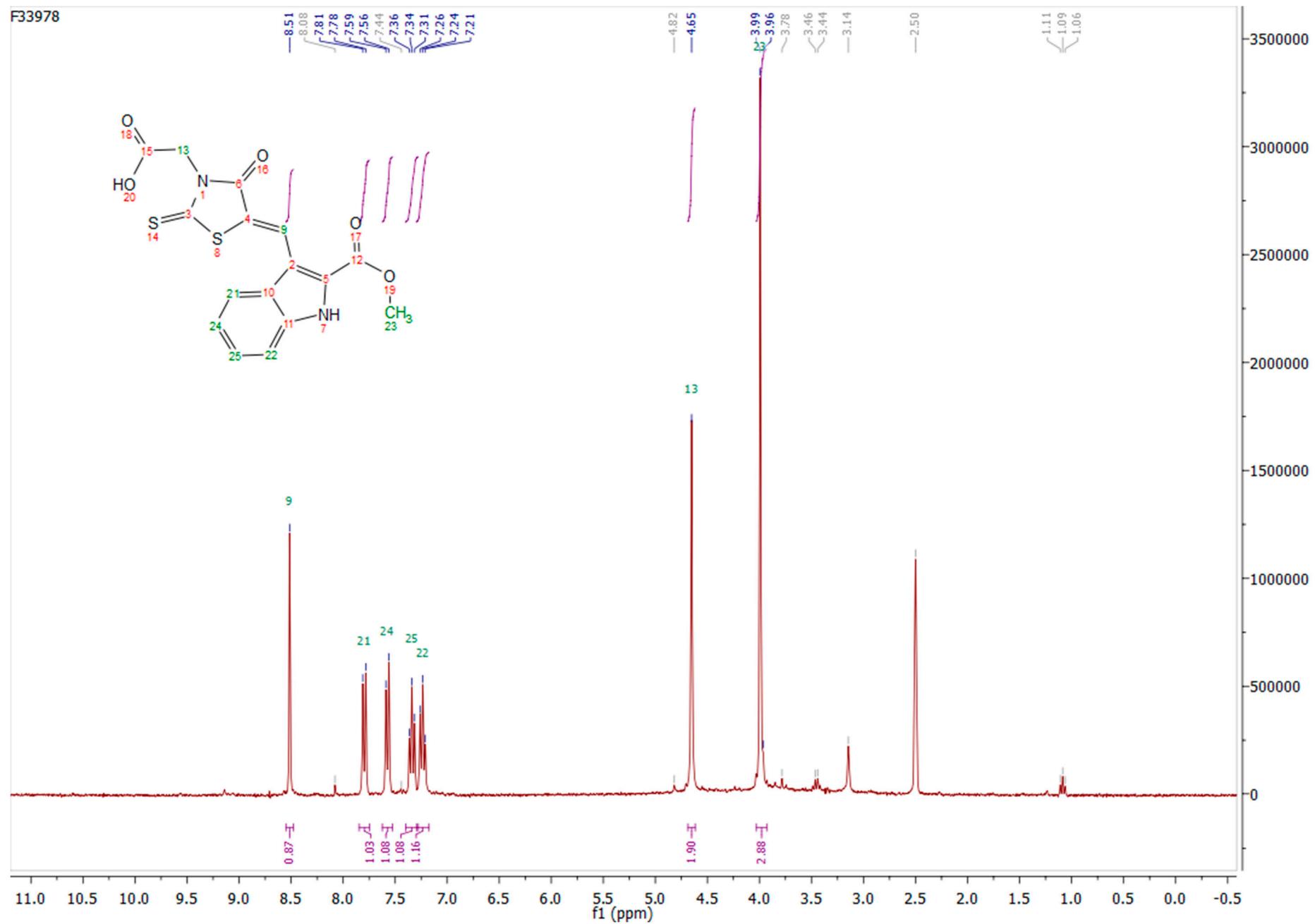
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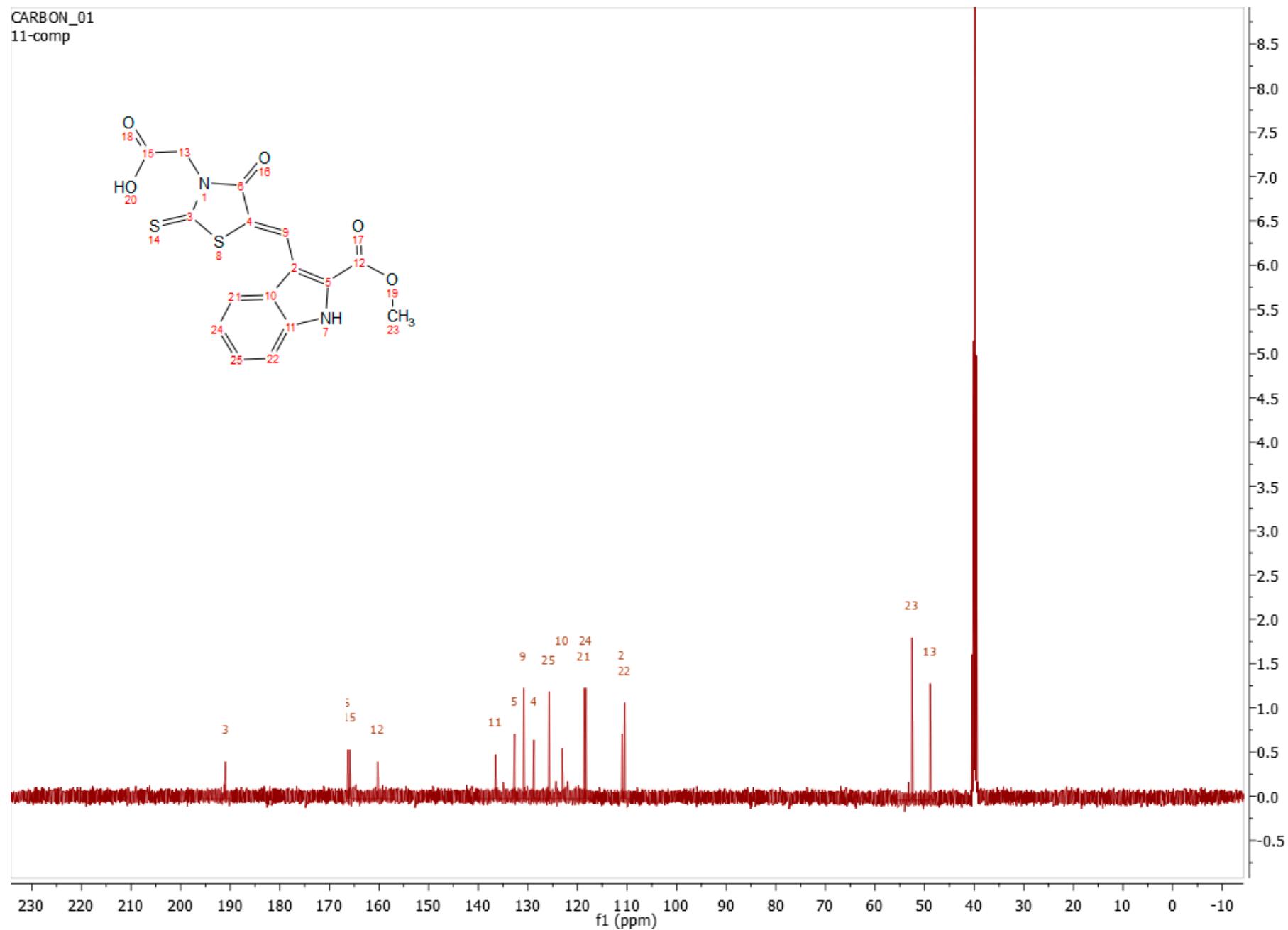
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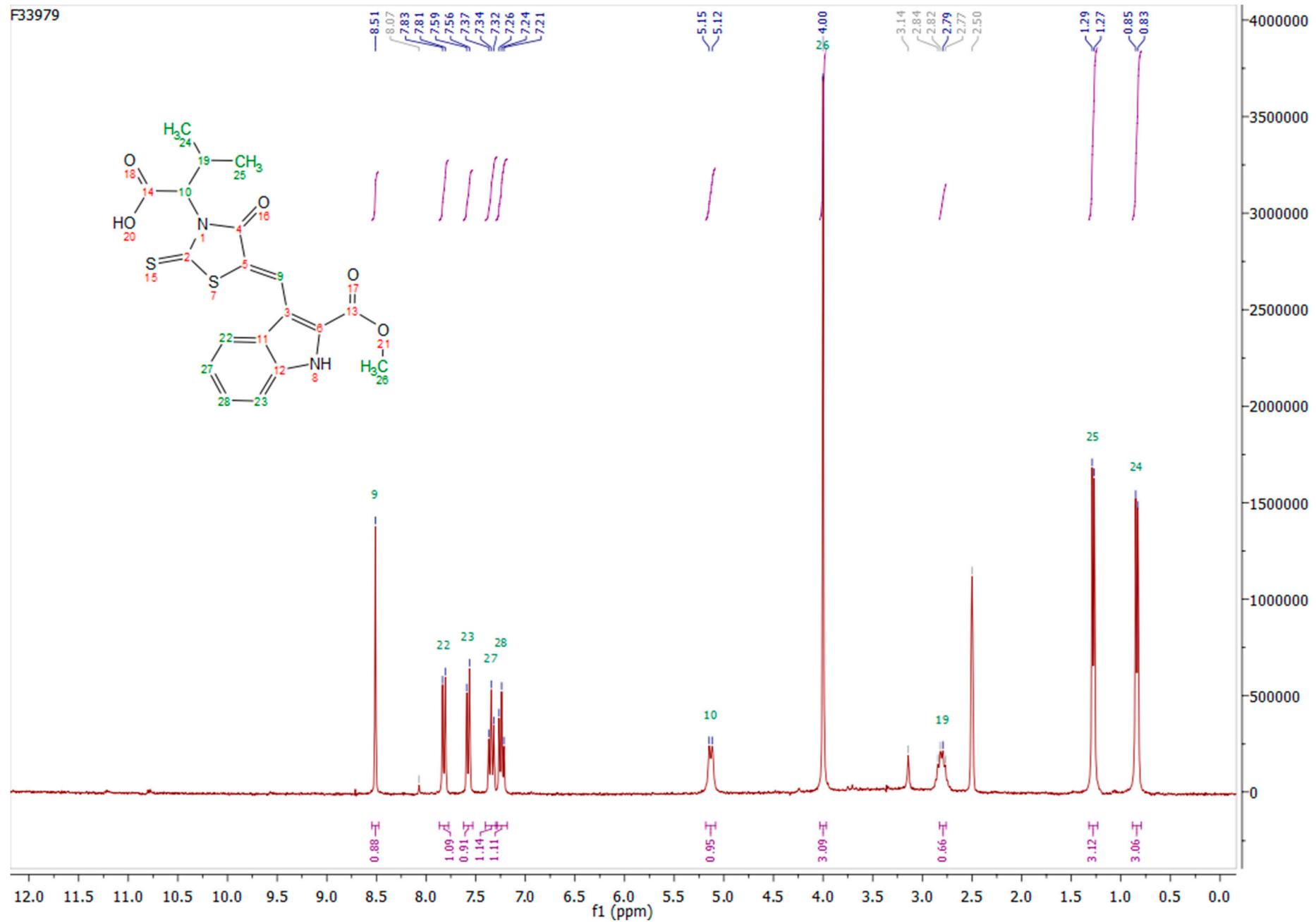
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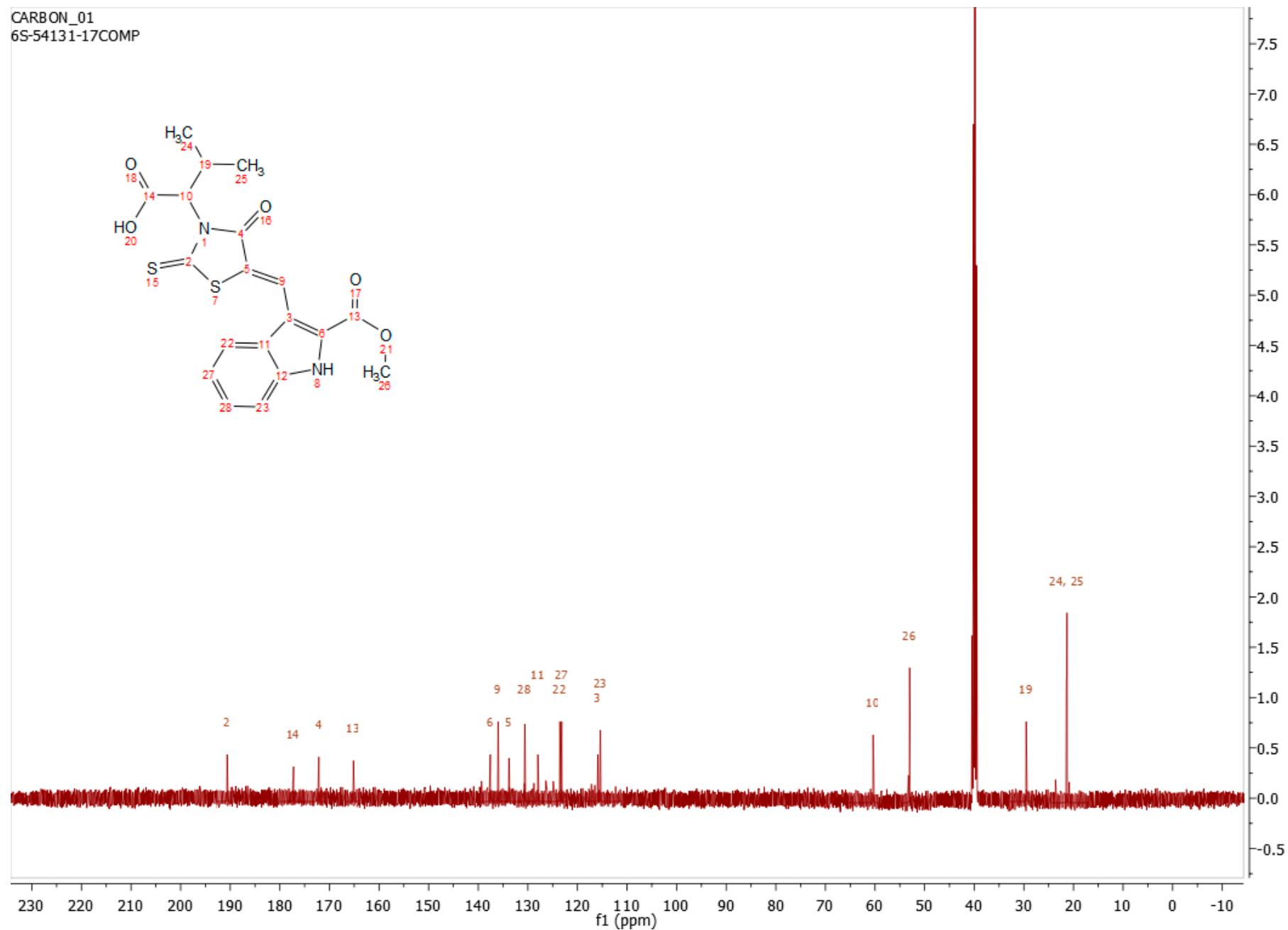
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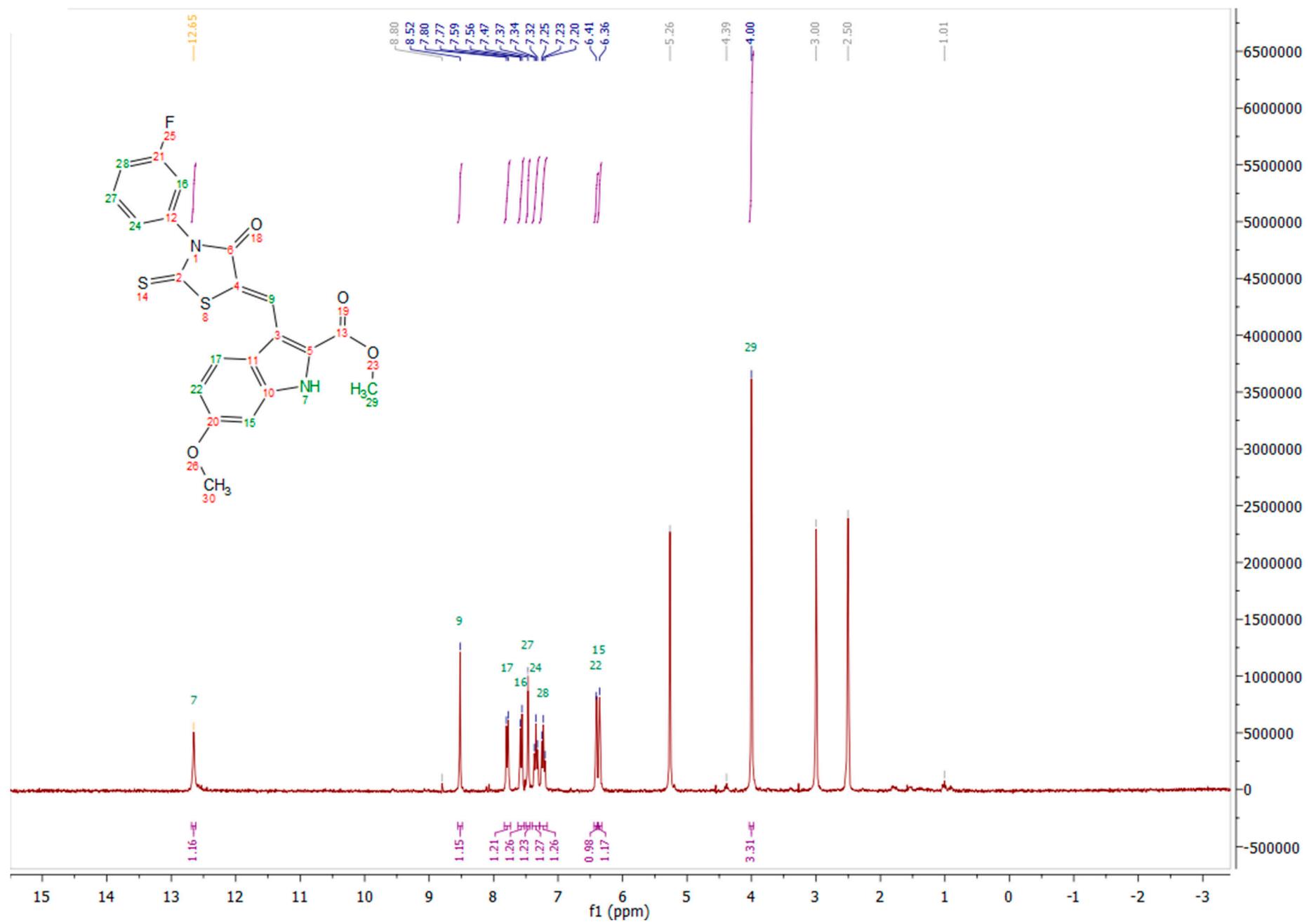
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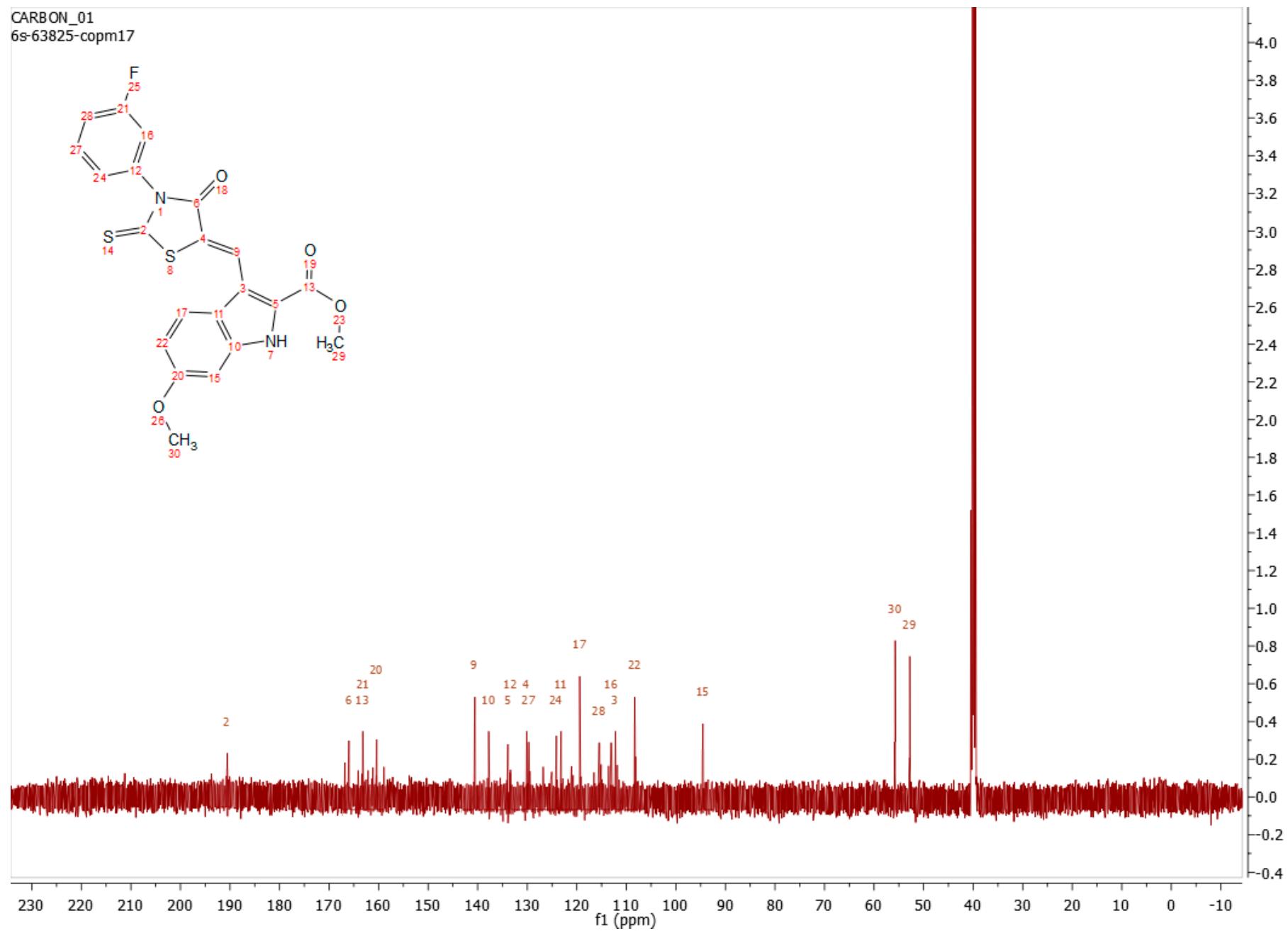
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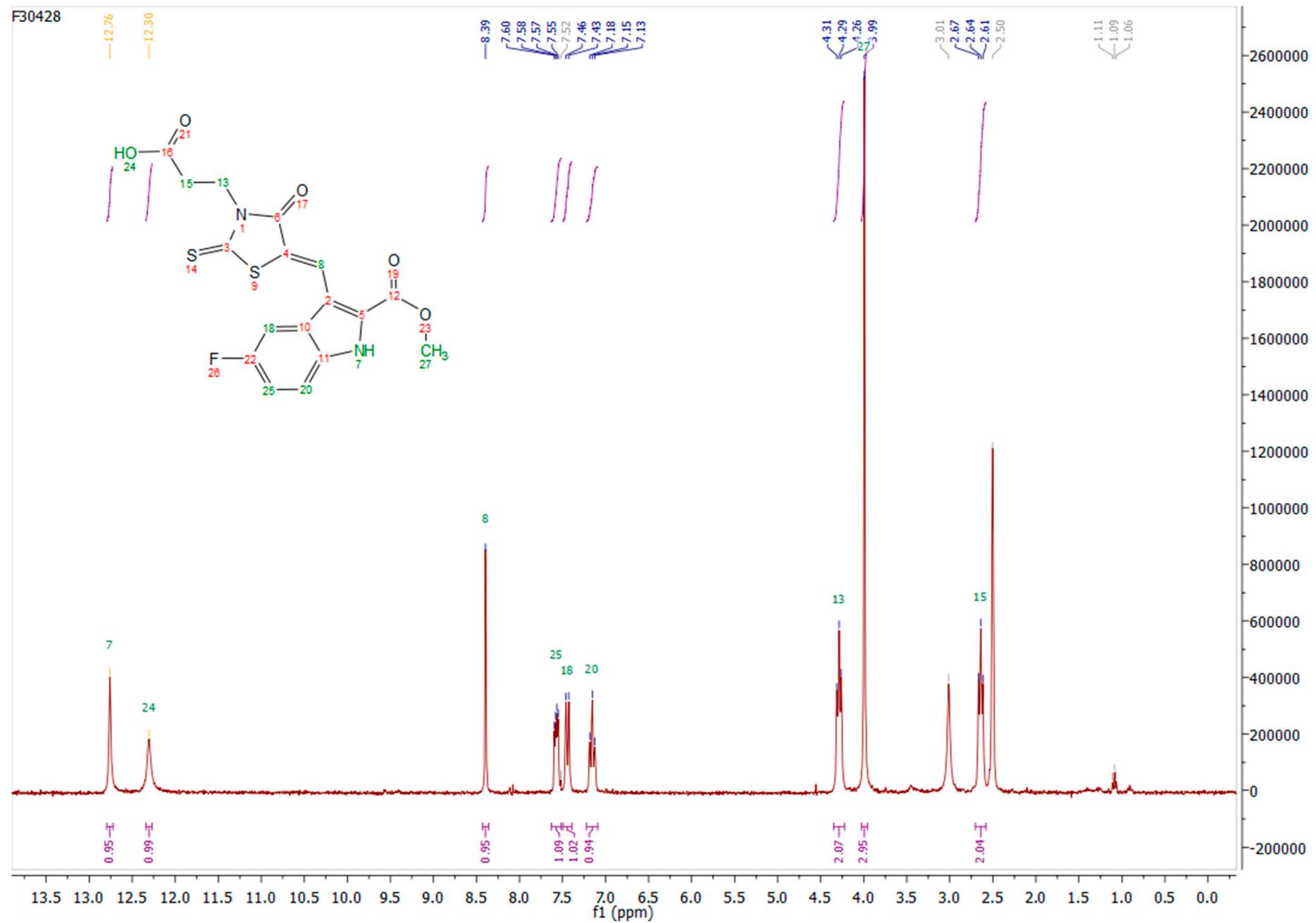
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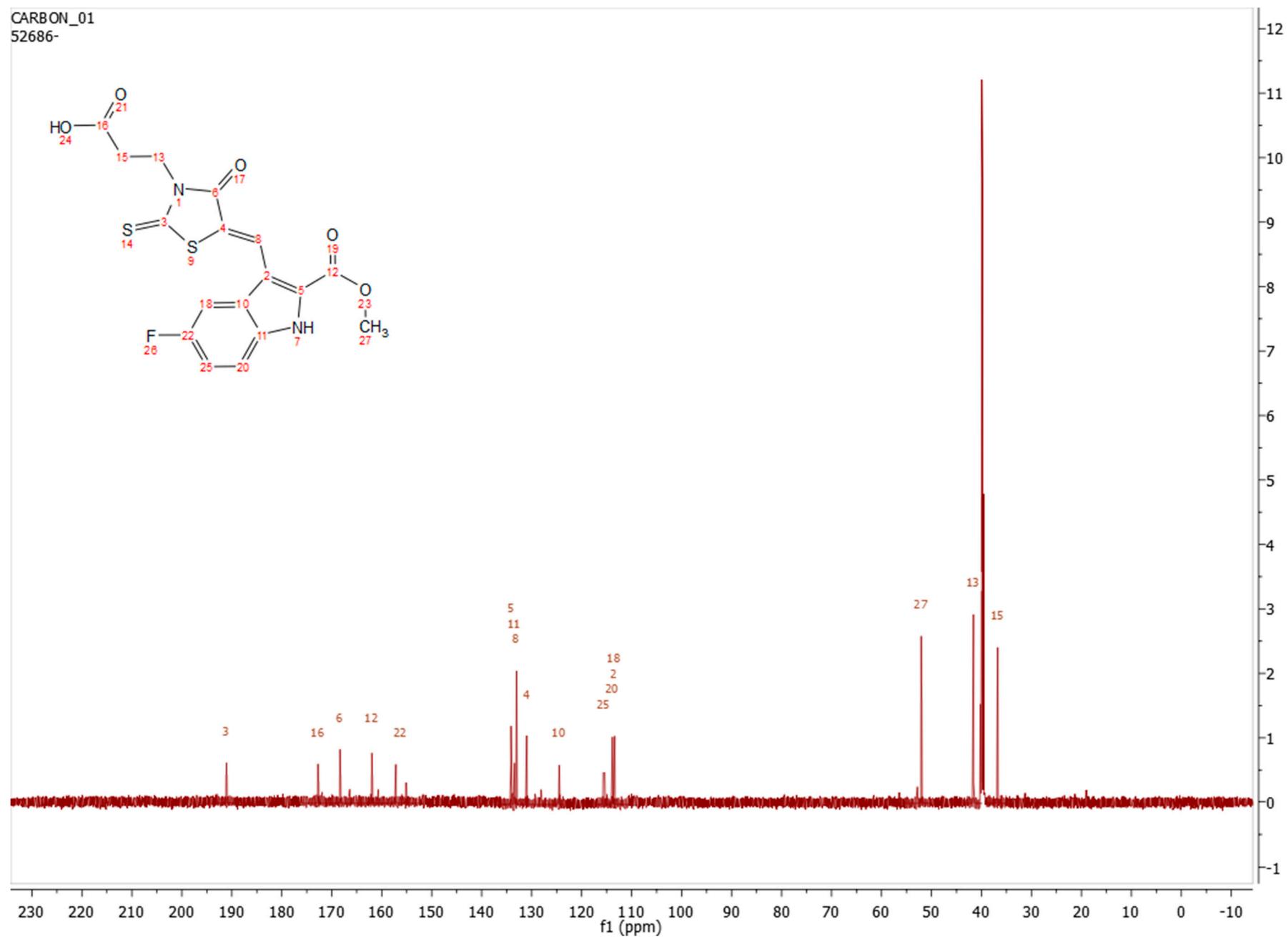
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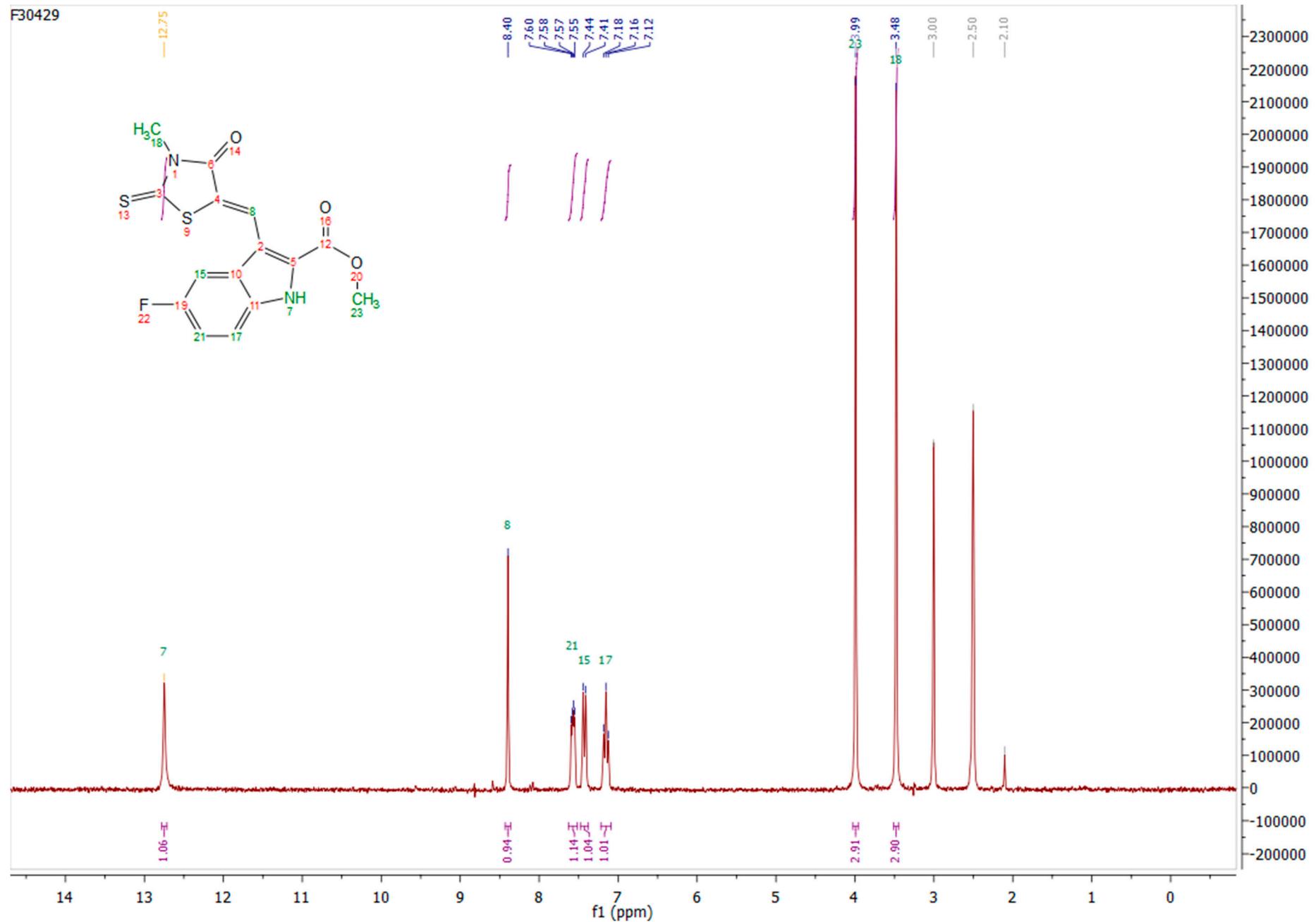
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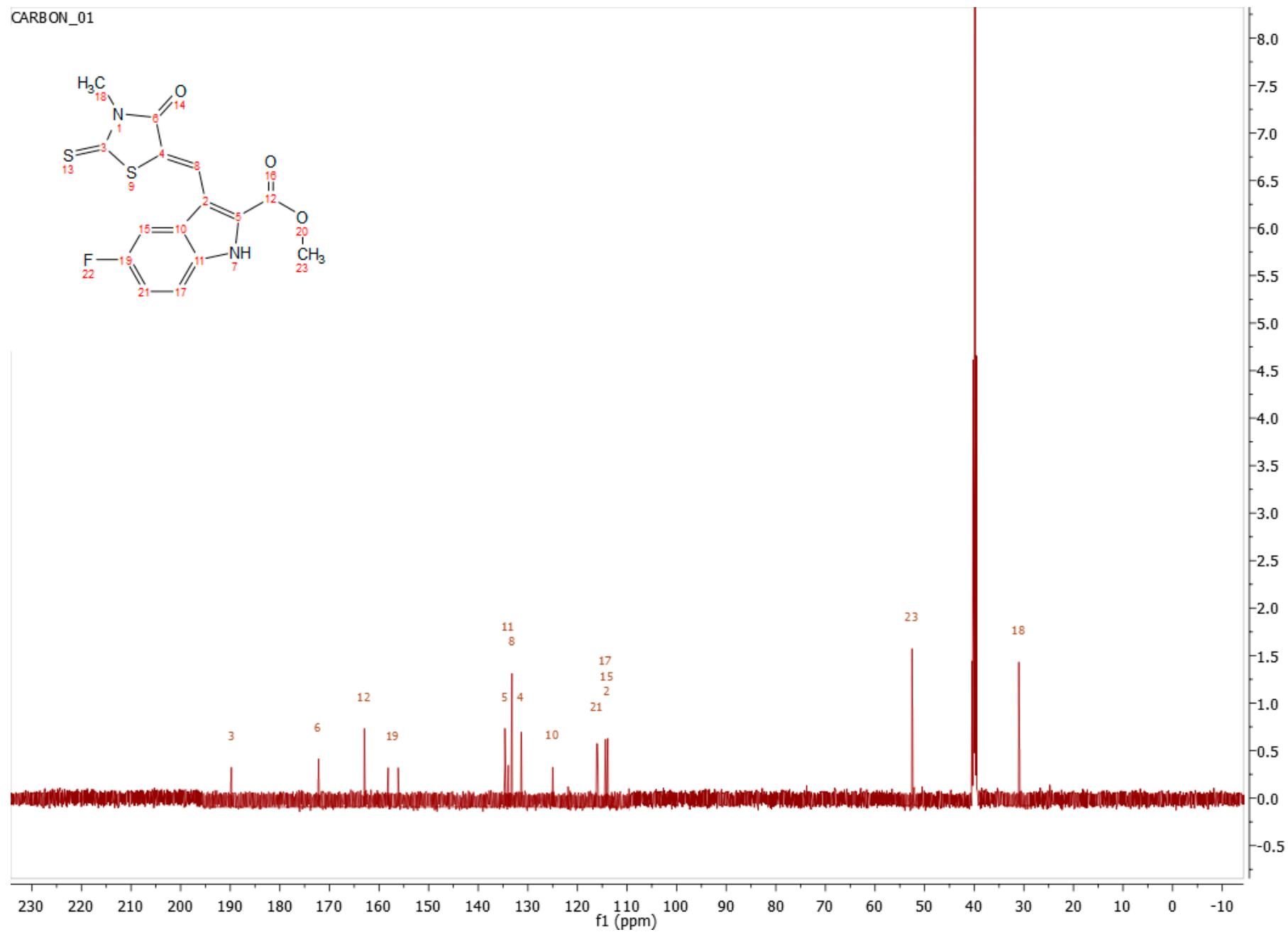
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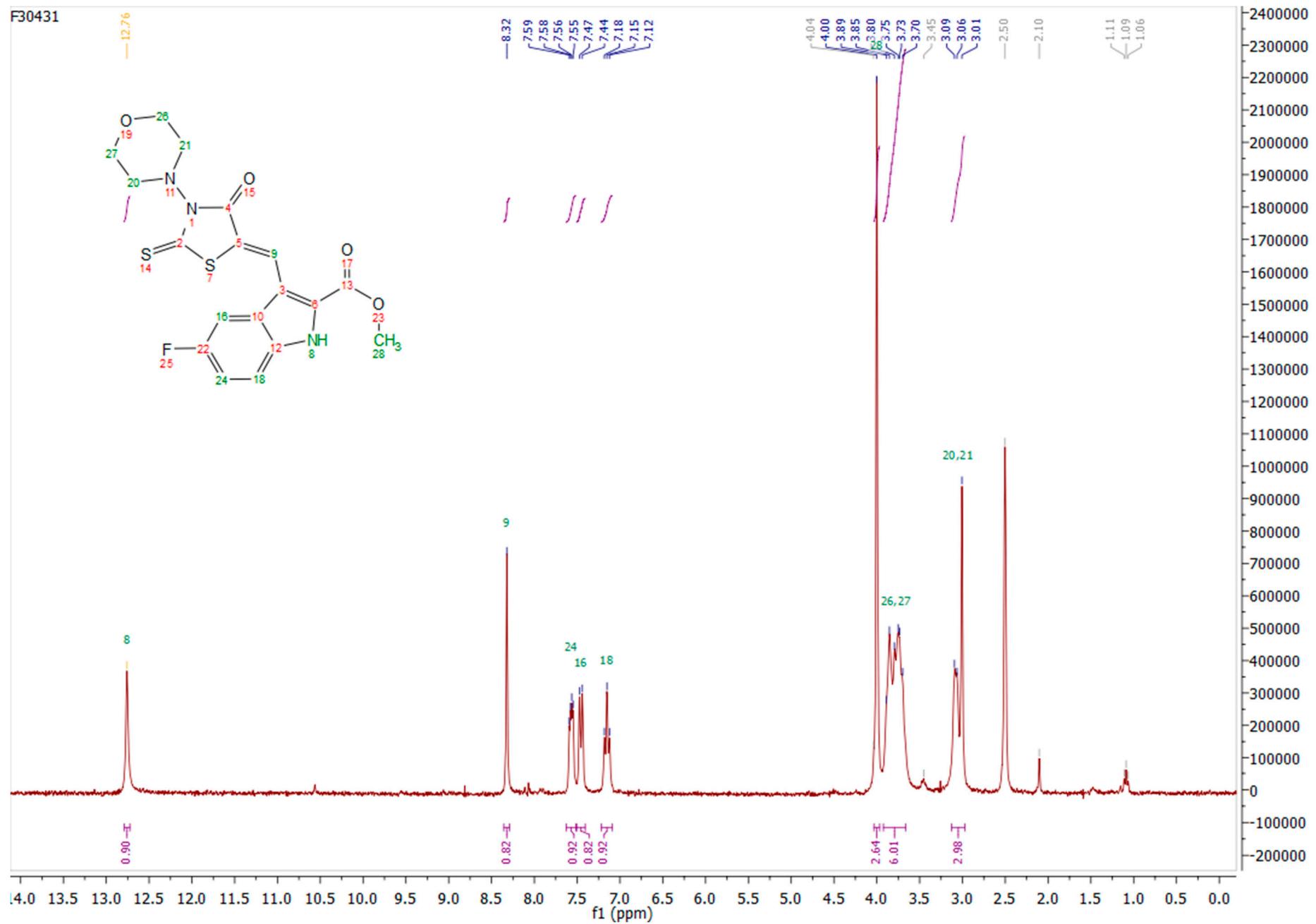
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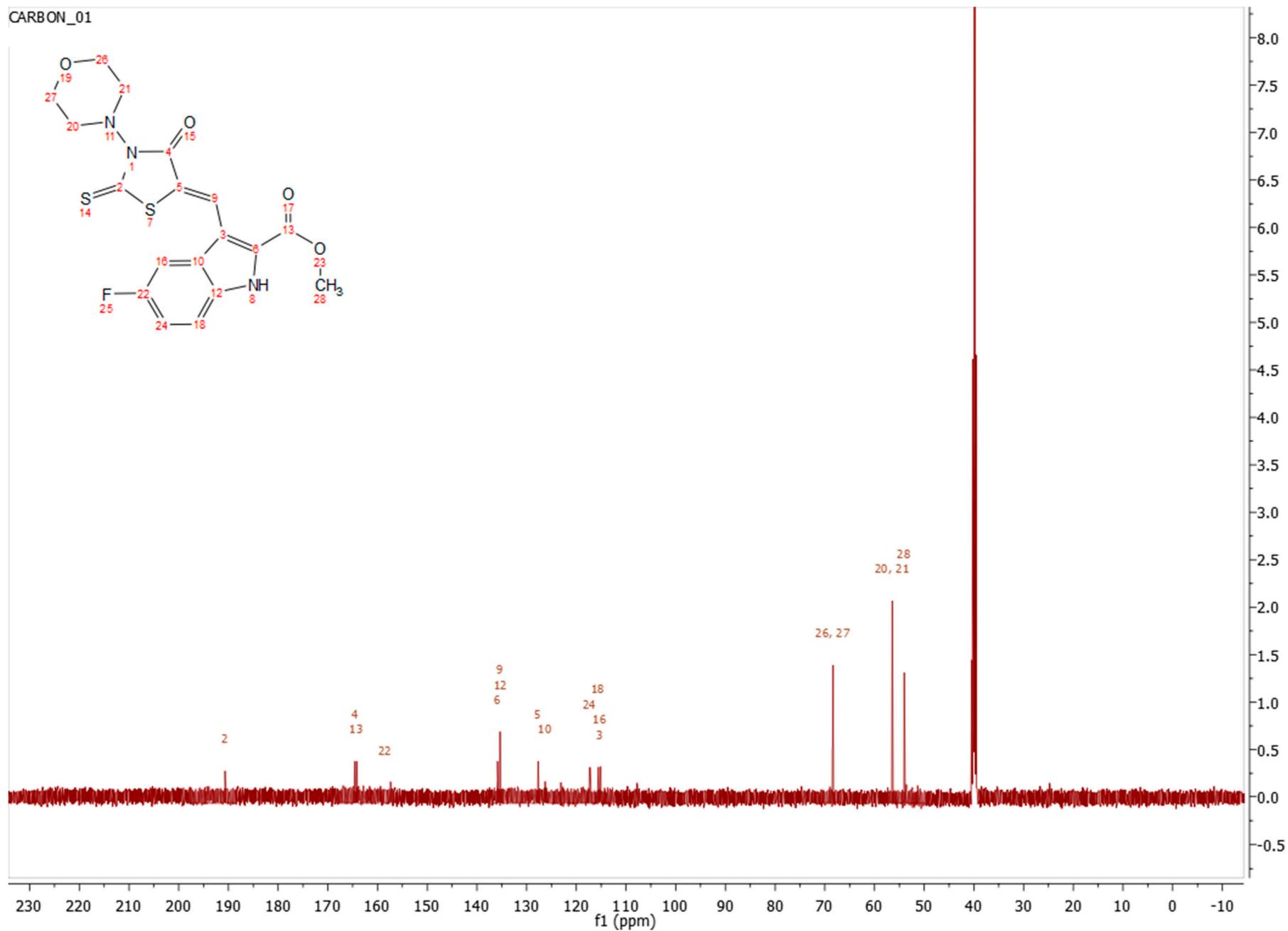
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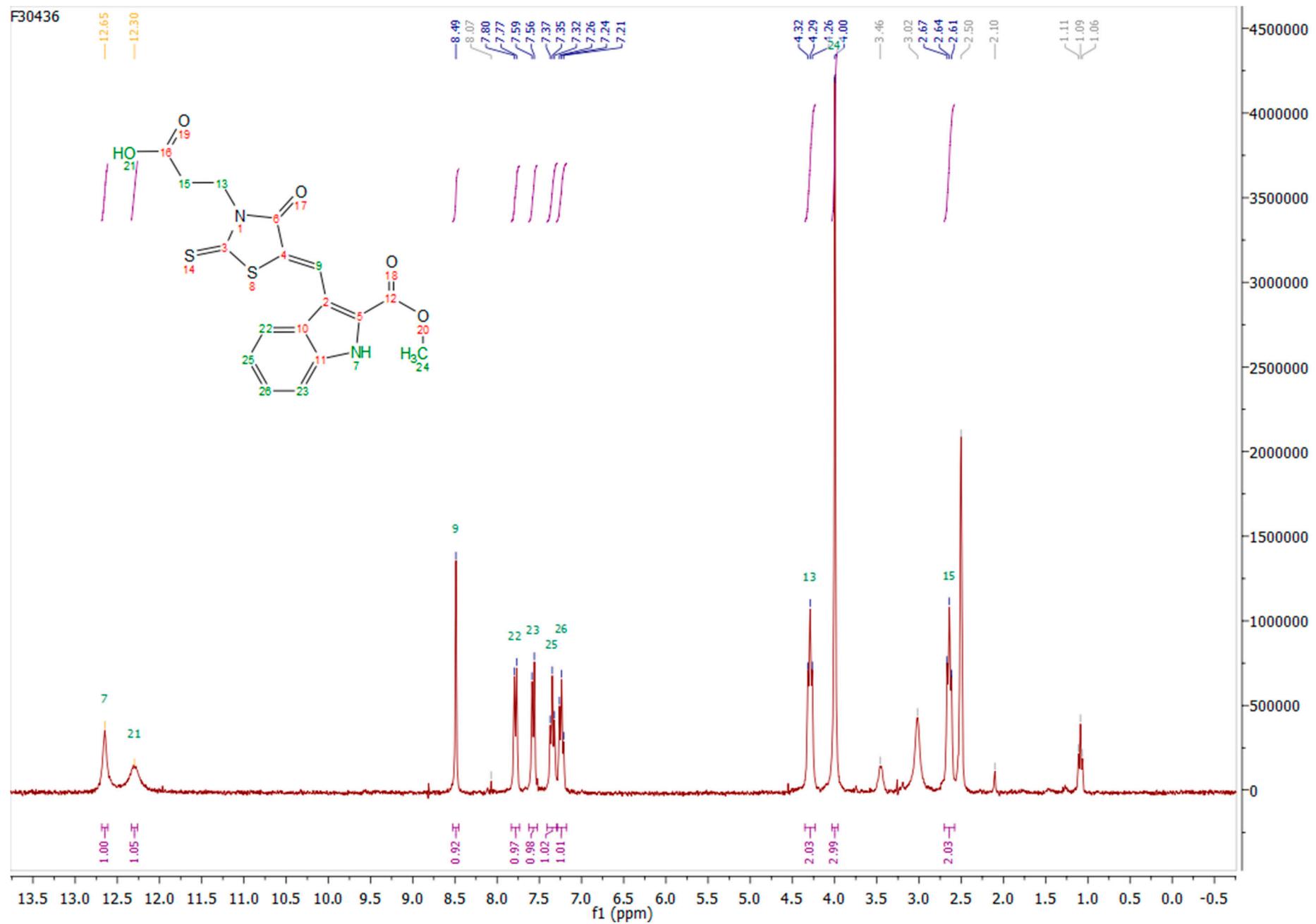
Compound 16



CARBON_01



Compound 17



CARBON_01

