

Synthesis, DFT and Molecular docking study of novel bis 1,2,3-triazole derivatives of 2-hydroxyquinoline-4-carboxylate as antimicrobial agents

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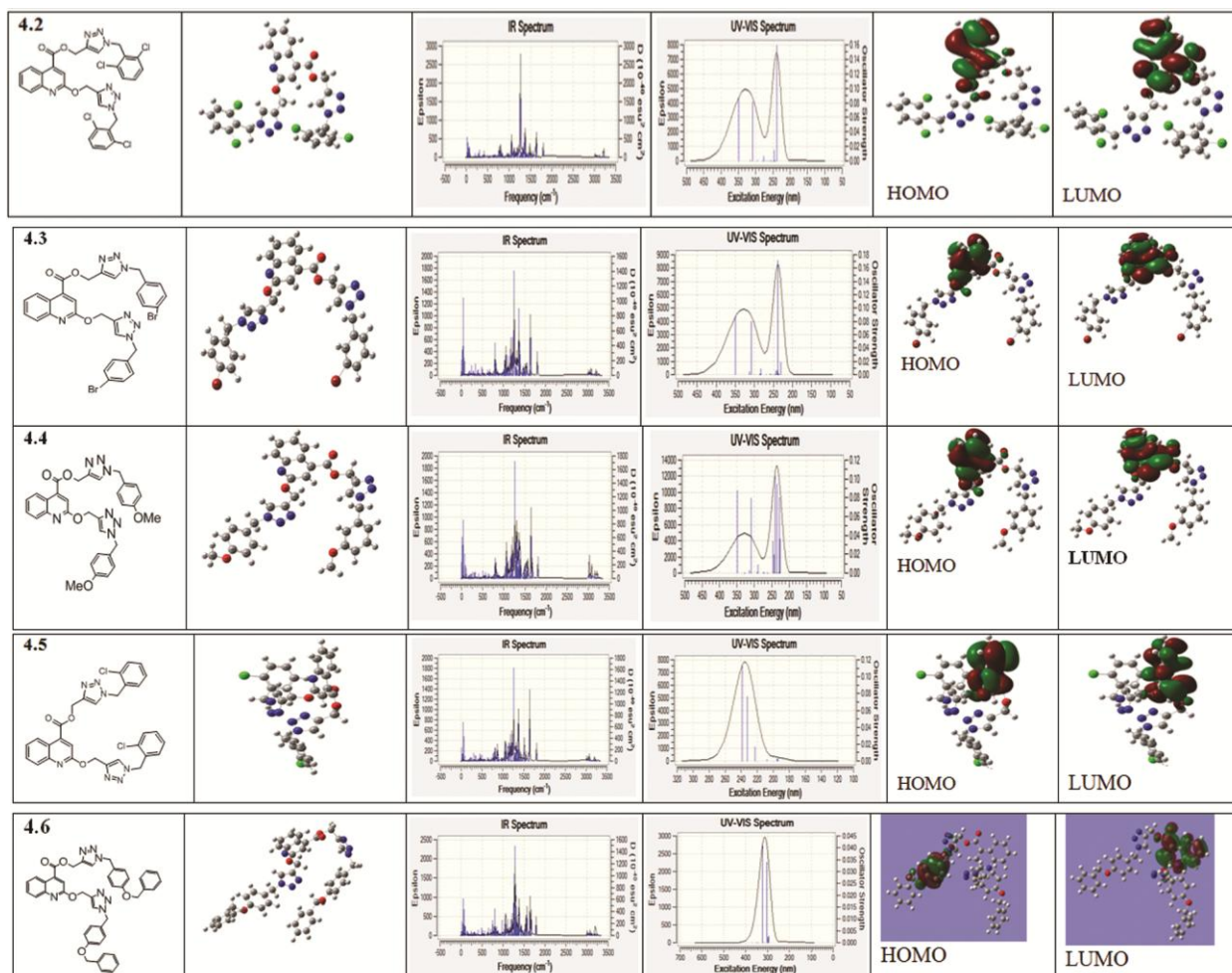
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Supplementary Data

Figures



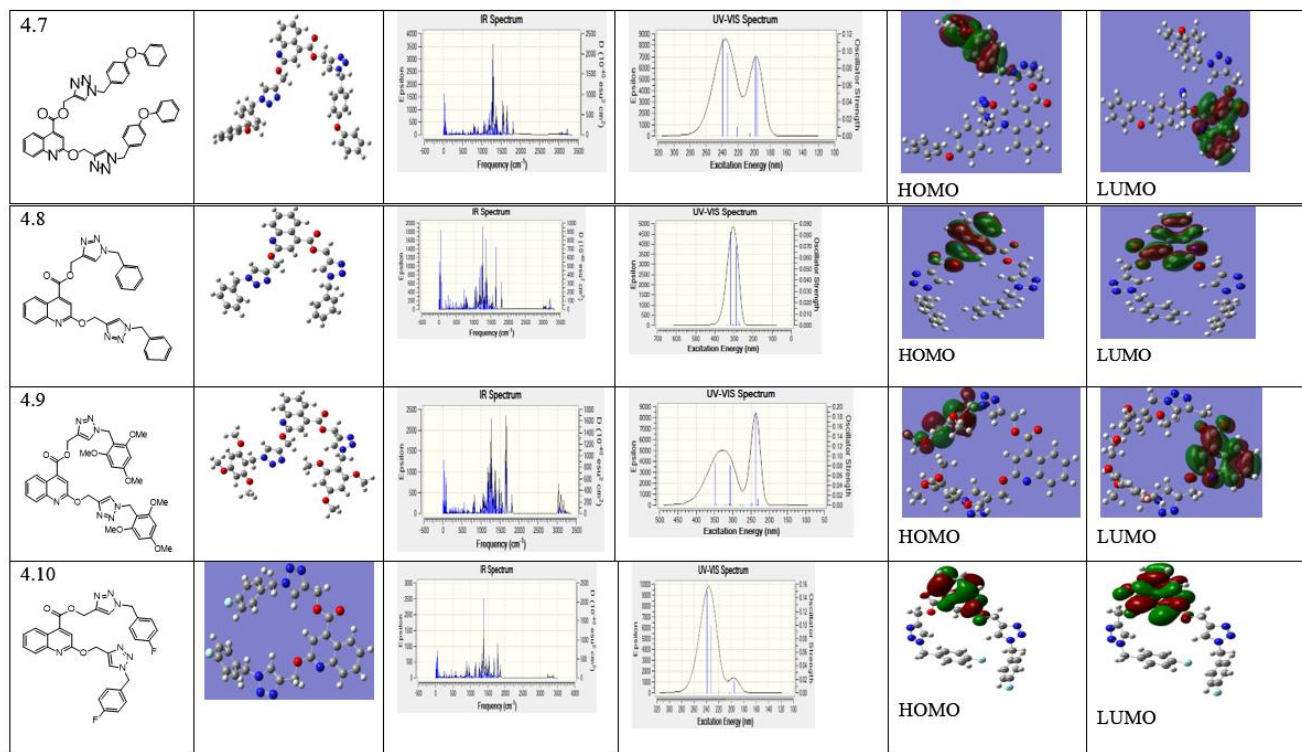


Fig.1 — Display of Chemdraw Structures, Ball and Stick models, UV and IR spectrum and Molecular orbital energy diagrams HOMO and LUMO of Compounds 4.2 to 4.10 using DFT

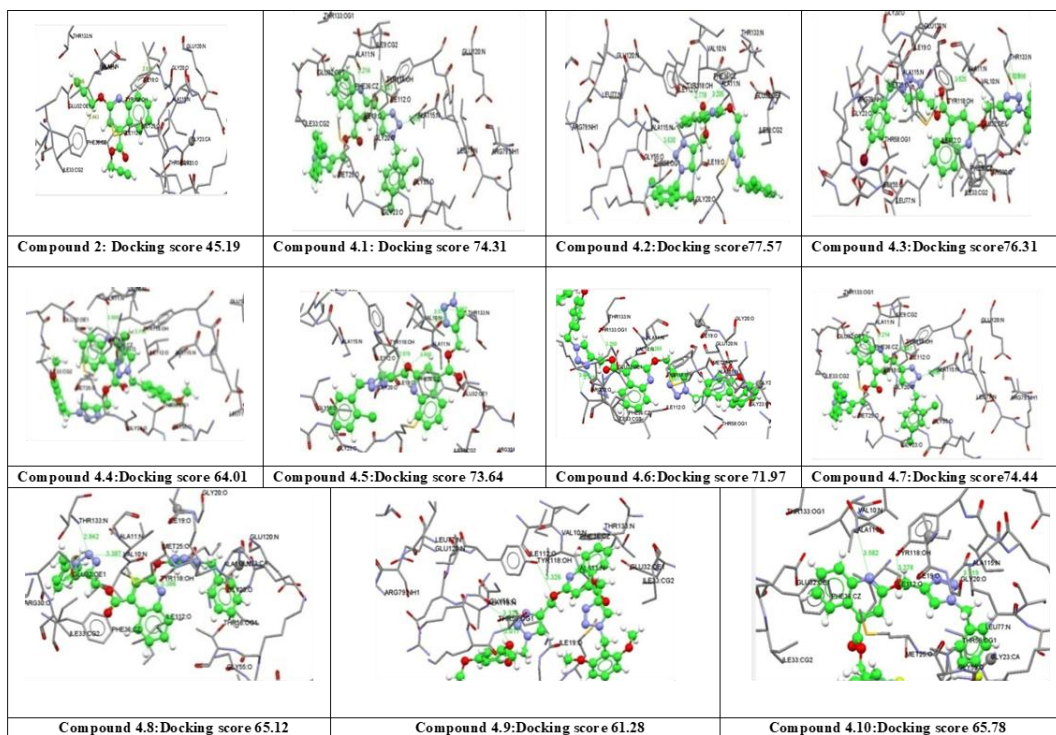
Molecular Docking Studies:

Fig. 2 — Binding orientations of 1,2,3-Triazole derivatives 2, 4.1, 4.2 and 4.3 4.4, 4.5, 4.6, 4.7, 4.8, 4.9, 4.10 in the active site of 1M78

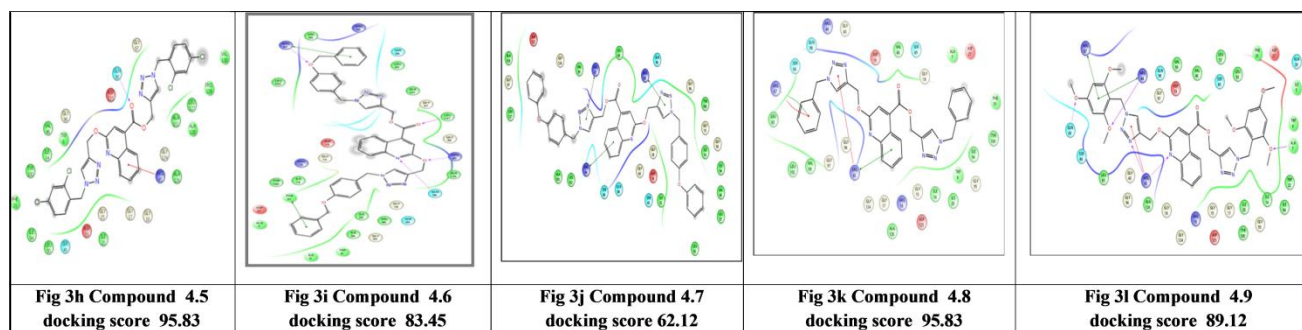
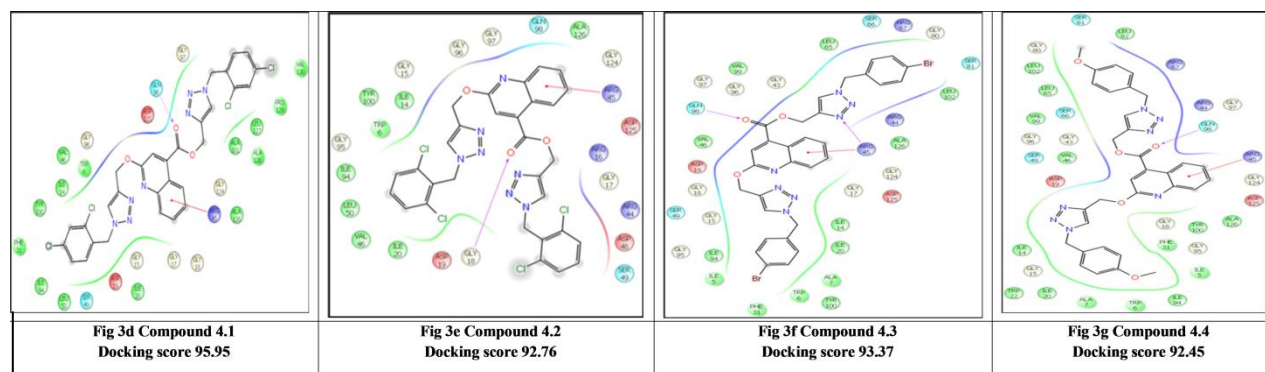
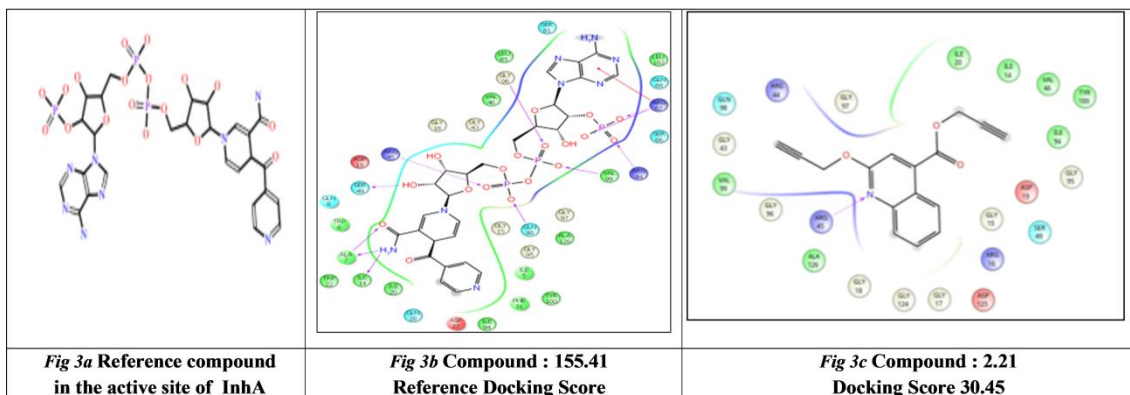
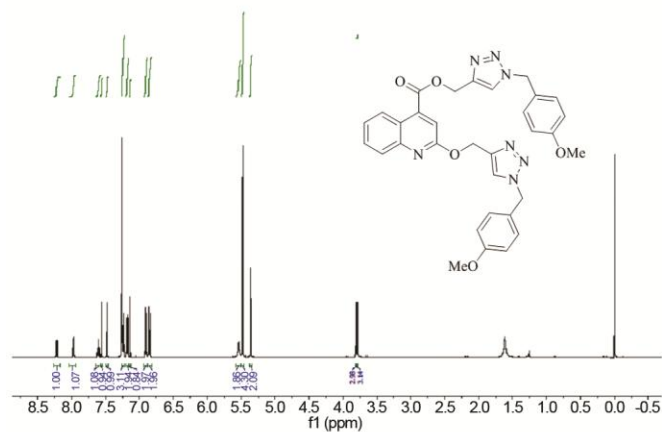


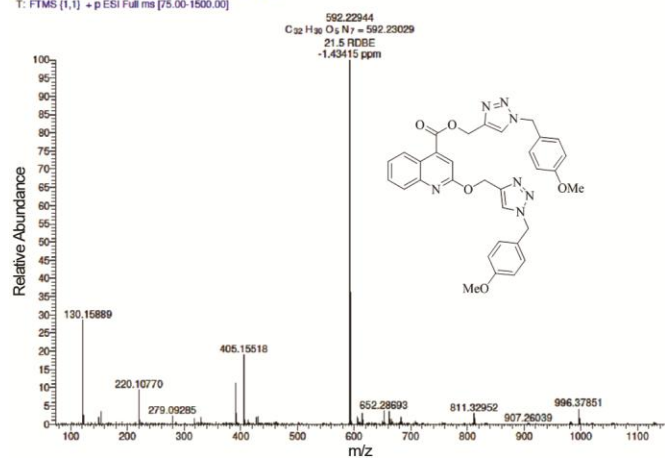
Fig 3a-c — Binding orientations of Reference compound and compound 2 in the active site of InhA; and 3d-l — Binding orientations of 1,2,3-Triazole derivatives 4.1, 4.2, 4.3, 4.4, 4.5, 4.6, 4.7, 4.8, 4.9, 4.10 in the active site of InhA

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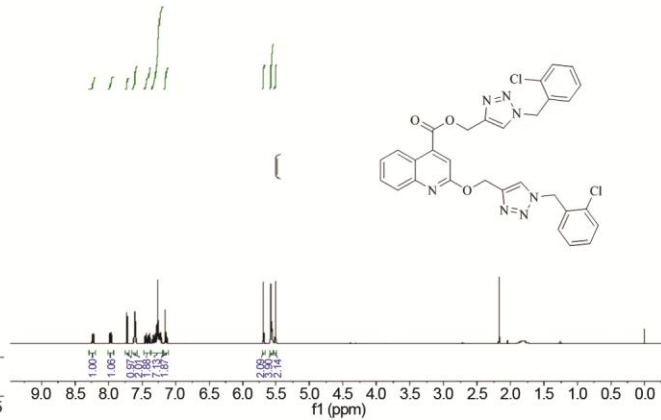


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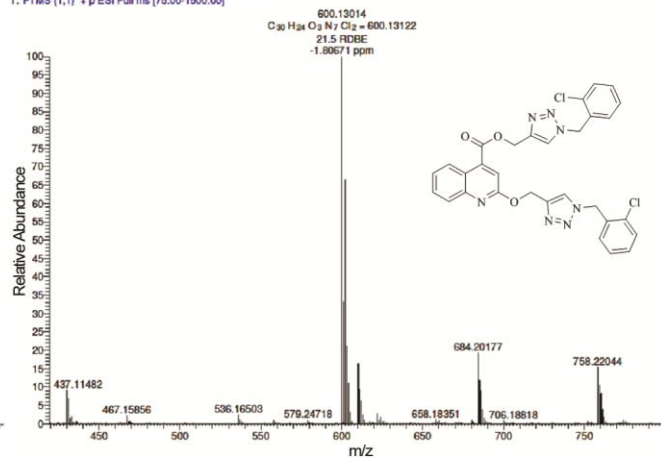


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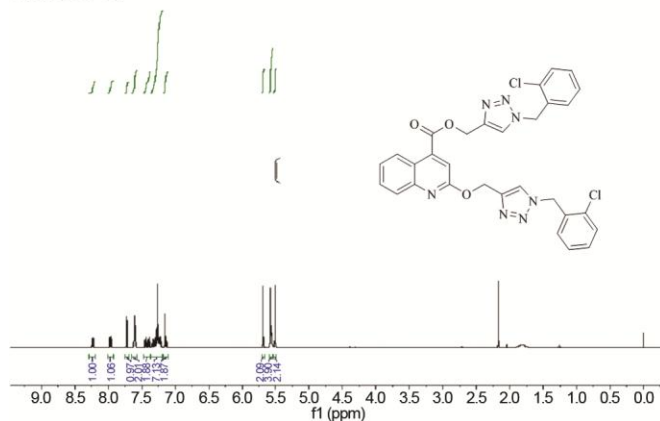


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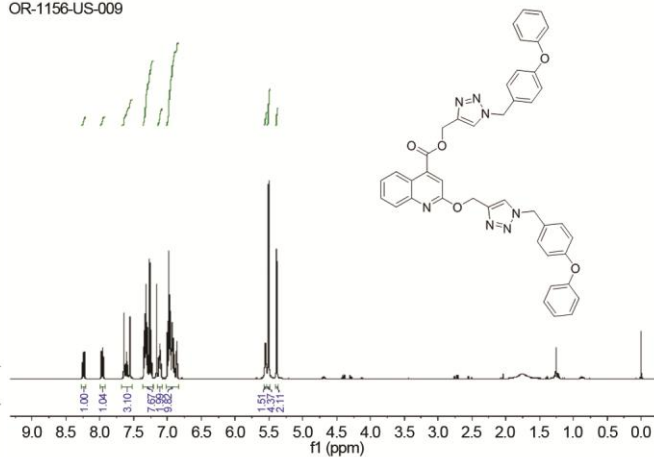
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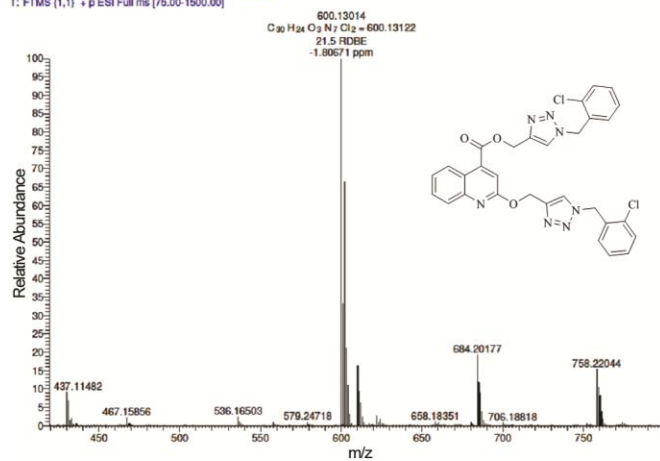


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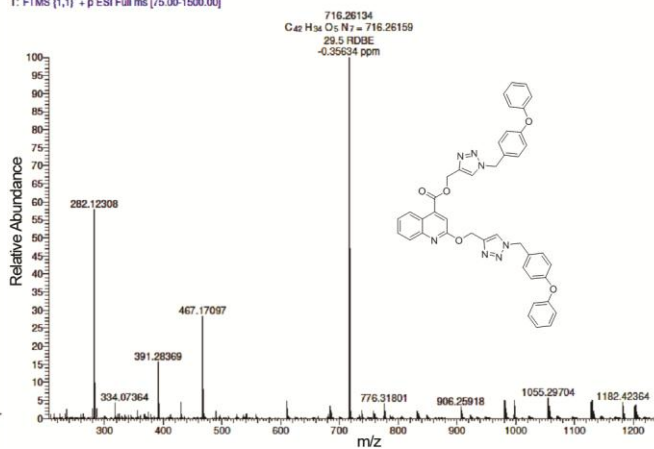
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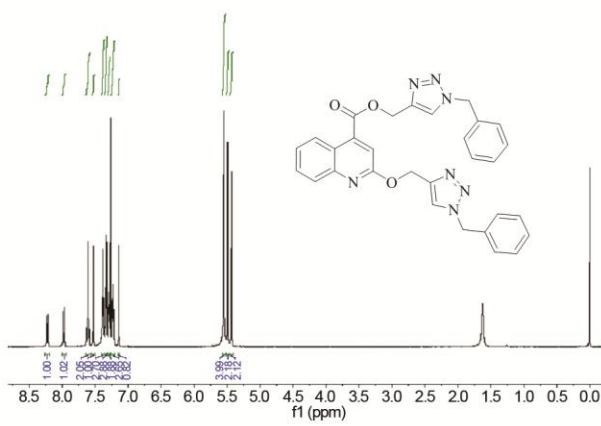


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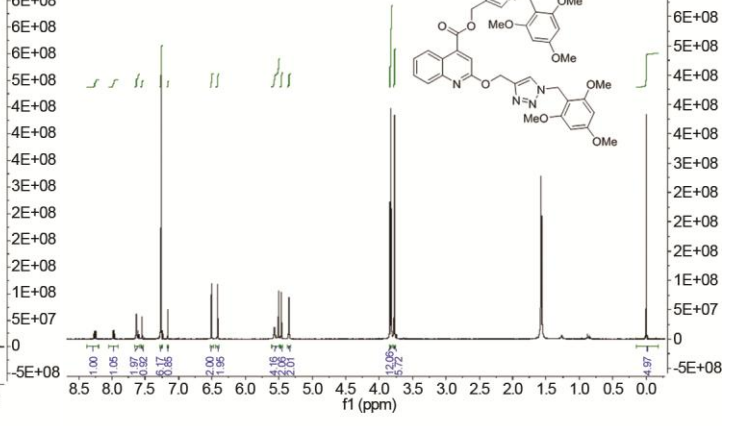
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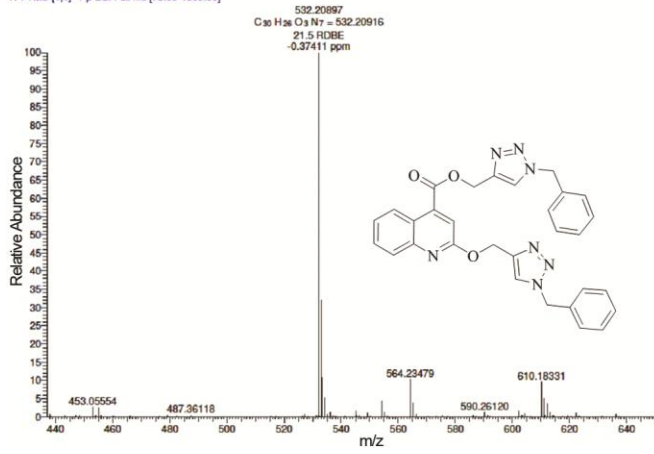


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