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Synthesis of novel coumarin-hydrazone hybrids as α -glucosidase inhibitors and their molecular docking studies

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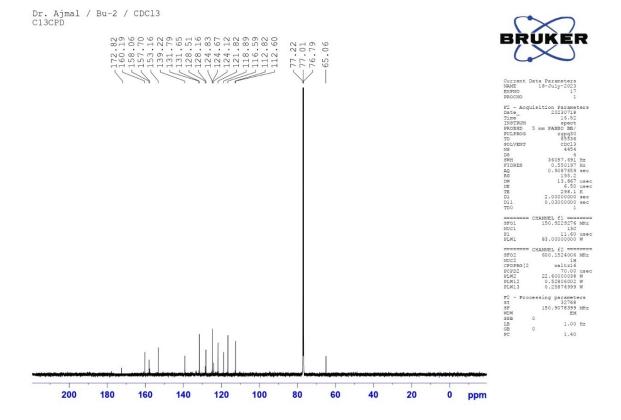
Both authors contributed equally

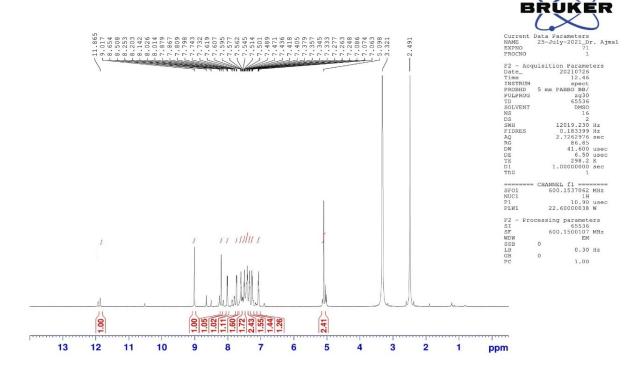
* Address for Correspondence

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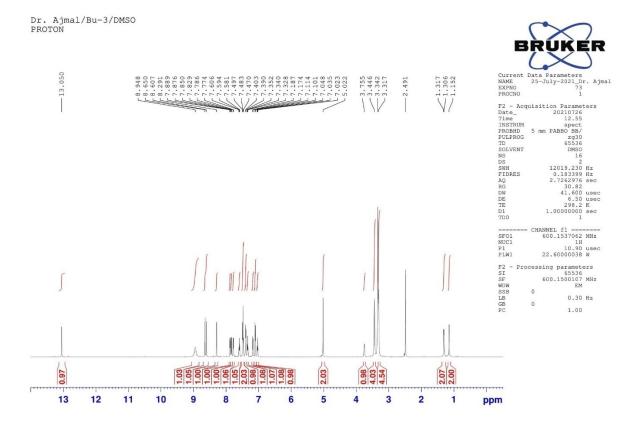
Prof. Dr. Ahmed Al-Harrasi, aharrasi@unizwa.edu.om

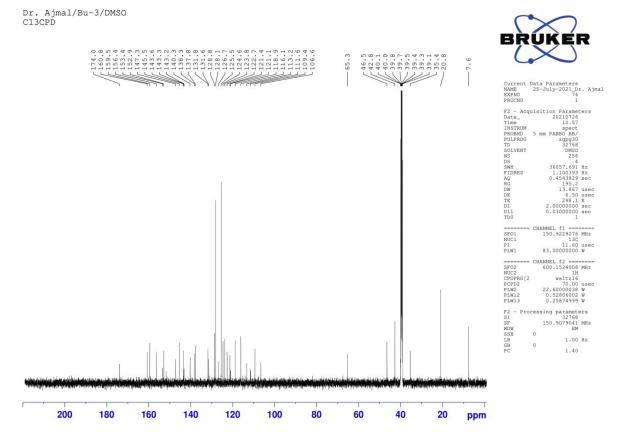
¹H-NMR and ¹³C-NMR Spectra of 7a



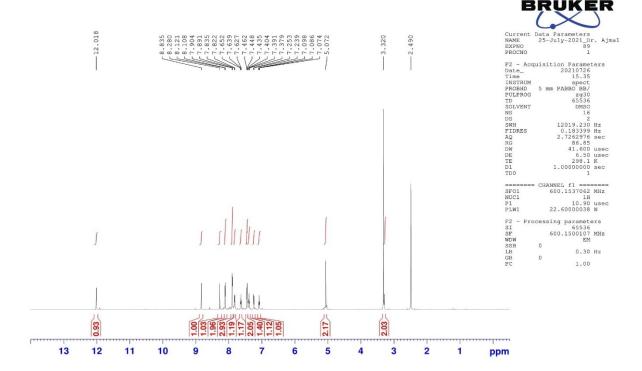


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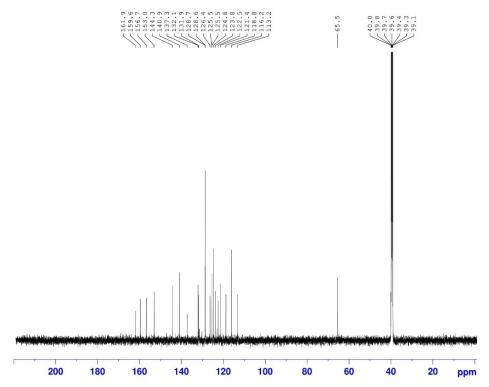




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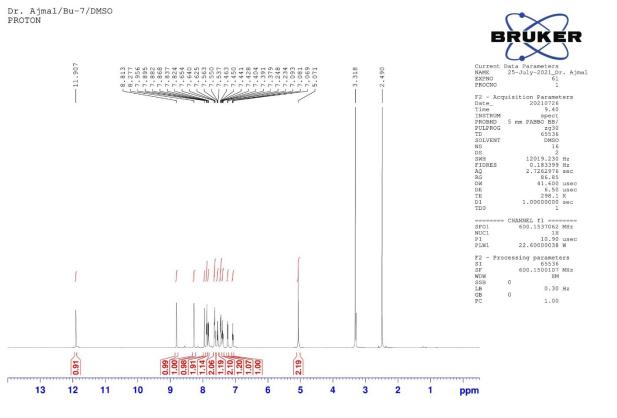
Dr. Ajmal/Bu-4/DMSO C13CPD



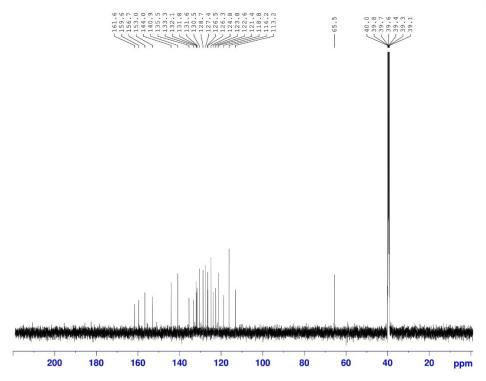


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¹H-NMR and ¹³C-NMR Spectra of 7d

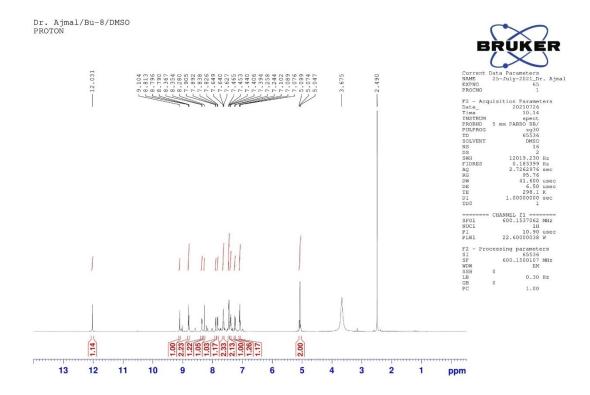


Dr. Ajmal/Bu-7/DMSO C13CPD

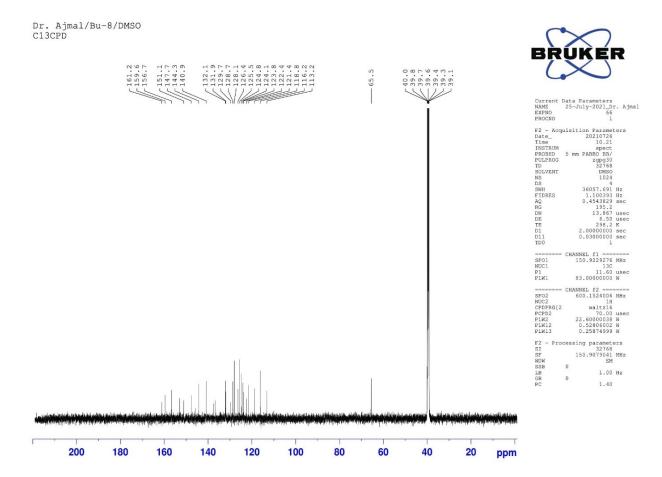




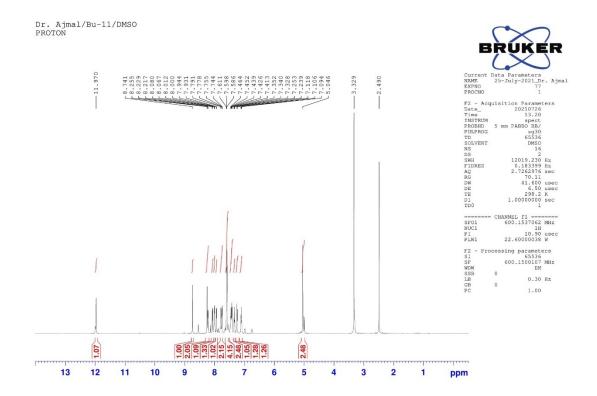
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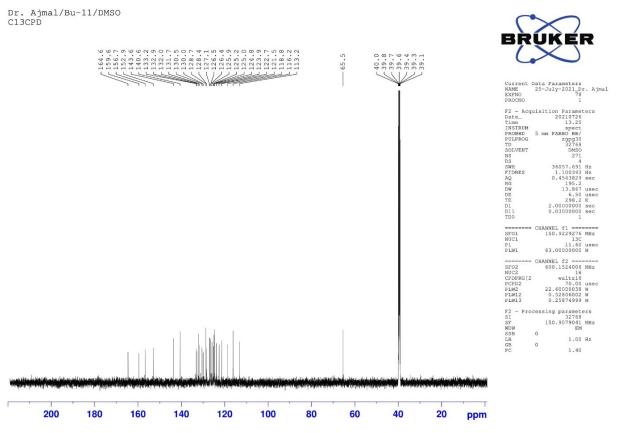


¹H-NMR and ¹³C-NMR Spectra of 7e

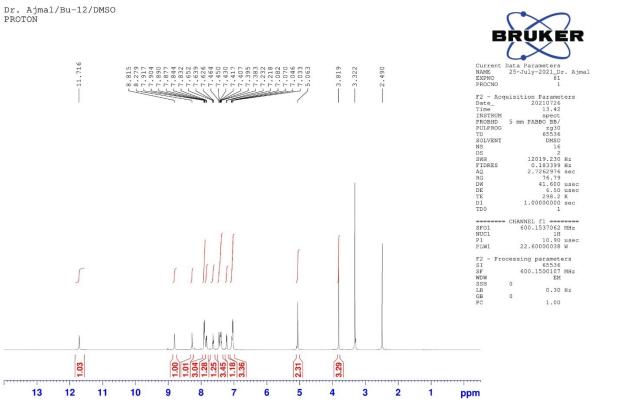


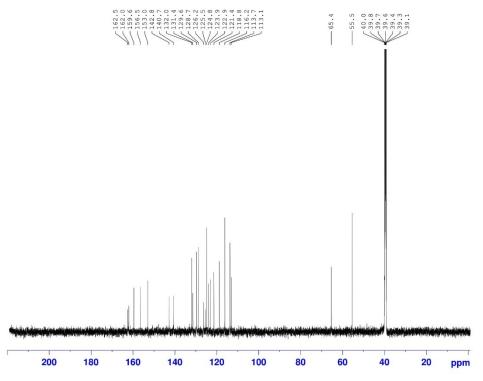
¹H-NMR and ¹³C-NMR Spectra of 7f





¹H-NMR and ¹³C-NMR Spectra of 7g

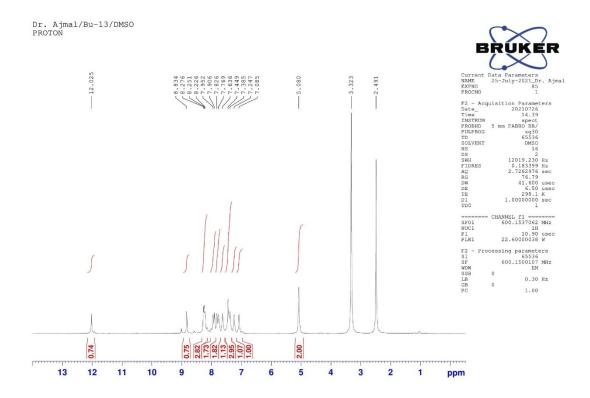




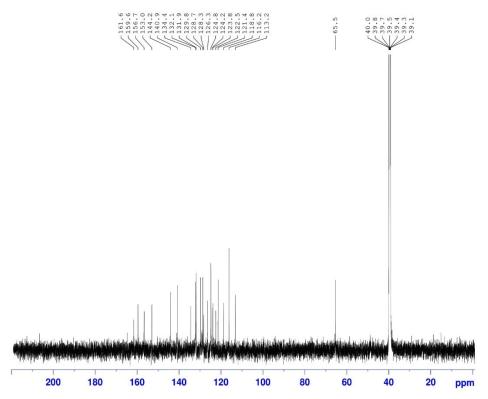


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NUC1 P1	13C
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¹H-NMR and ¹³C-NMR Spectra of 7h

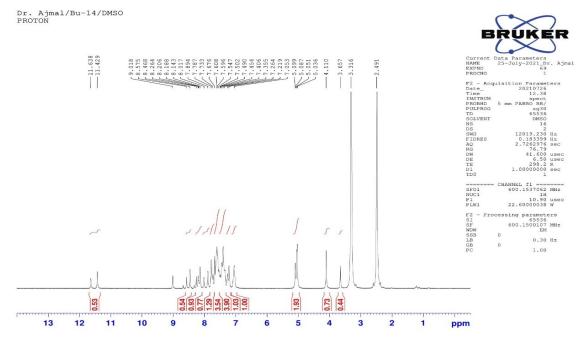


Dr. Ajmal/Bu-13/DMSO C13CPD

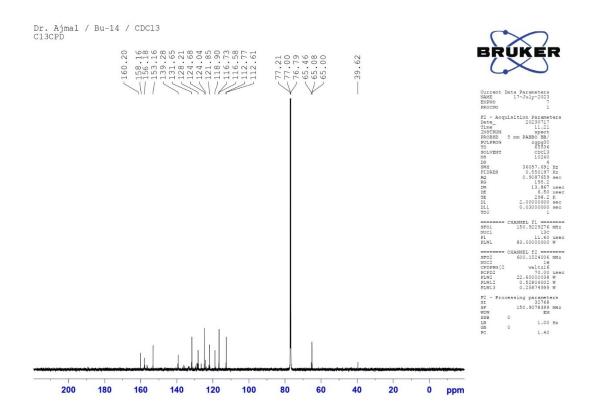




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¹H-NMR and ¹³C-NMR Spectra of 7i



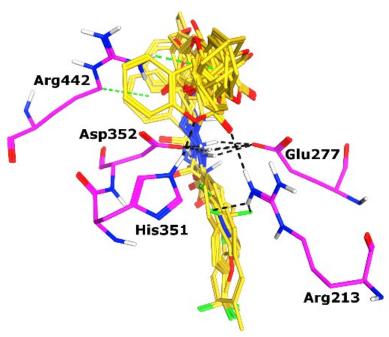


Figure S1. The docked conformations of all the compounds (presented in yellow sticks) are shown in the active site of α -glucosidase. The binding residues in the active site are presented in magenta sticks, H-bonds and hydrophobic bonds are depicted in black and green dotted lines, respectively.