

Supporting information for

Unfolding the Antibacterial Activity and Acetylcholinesterase Inhibition Potential of Benzofuran-Triazole Hybrids: Synthesis, Antibacterial, Acetylcholinesterase Inhibition, and Molecular Docking Studies

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¹H and ¹³C NMR spectra of compound **10a-e**

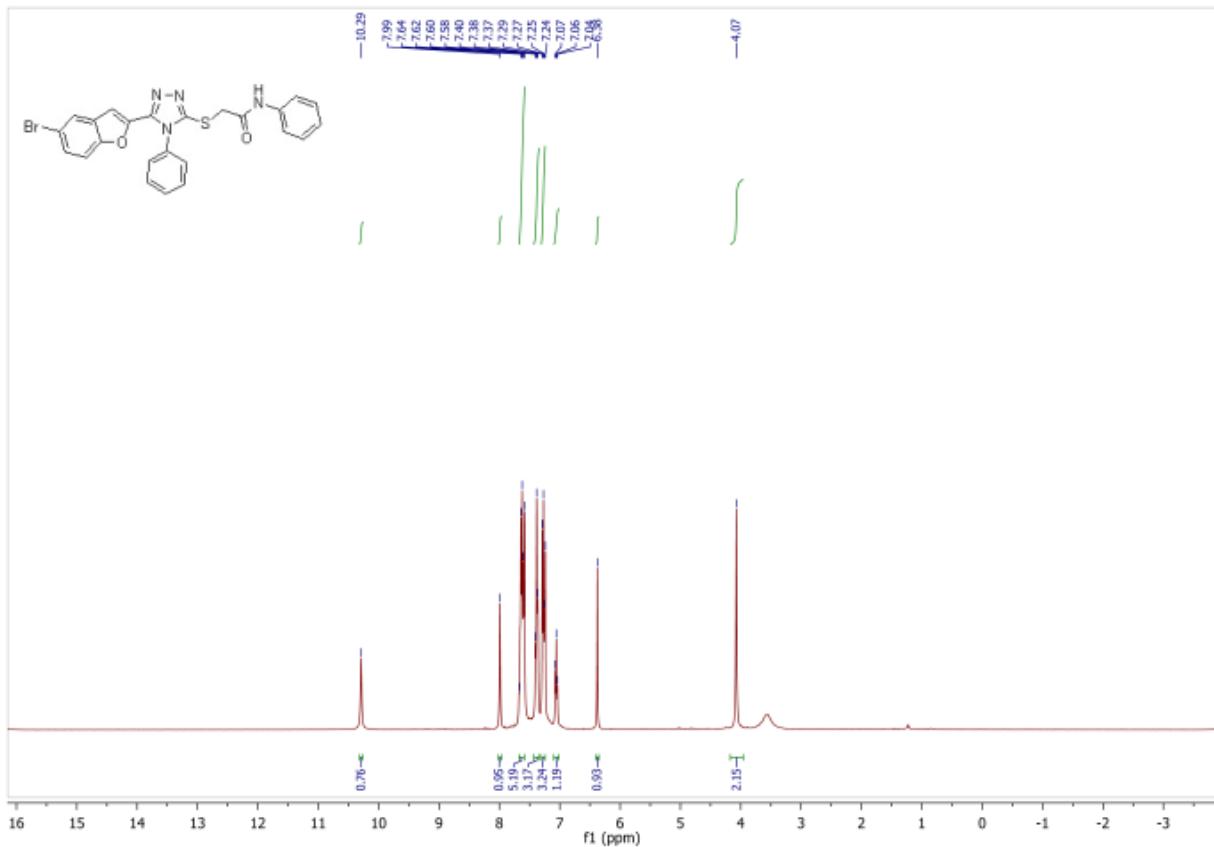


Figure S1: ¹H NMR spectrum of 2-((5-(5-bromobenzofuran-2-yl)-4-phenyl-4H-1,2,4-triazol-3-yl)thio)-N-phenylacetamide (**10a**).

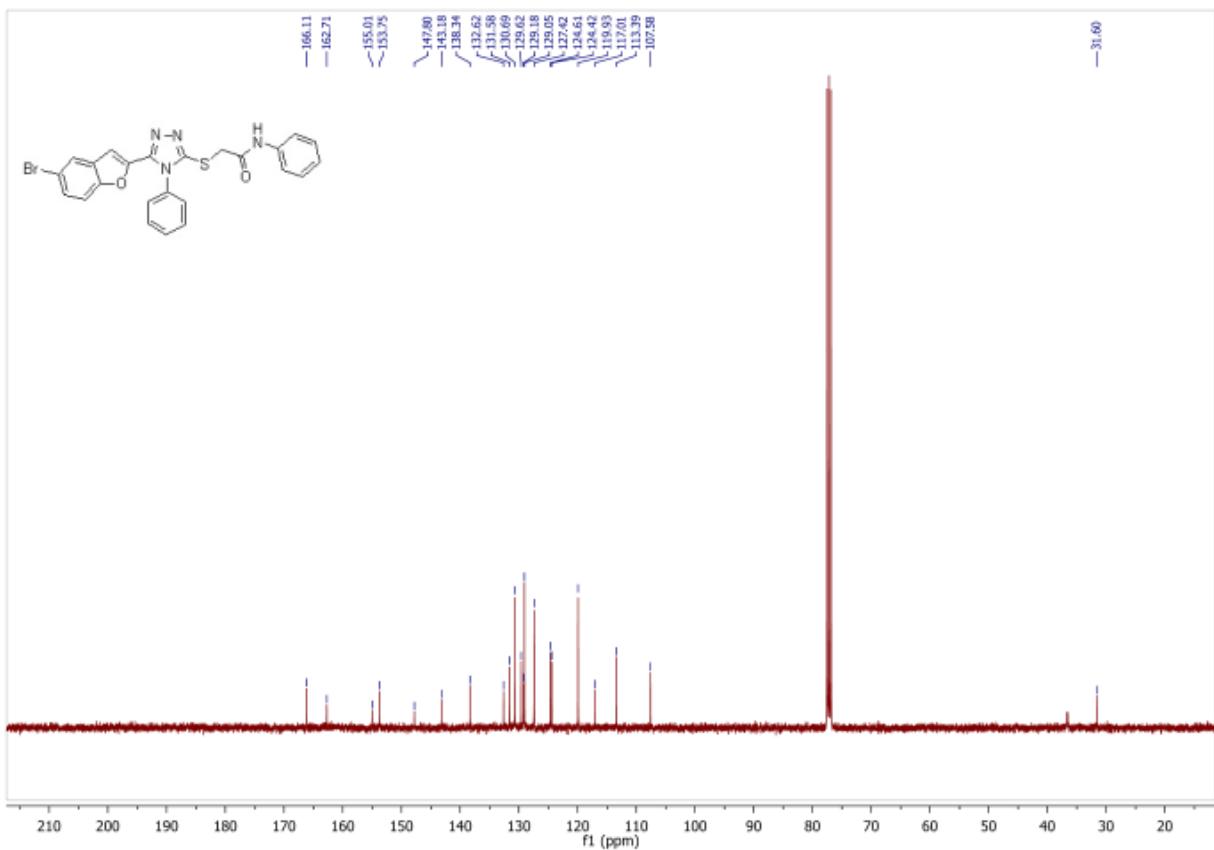


Figure S2: ^{13}C NMR spectrum of 2-((5-(5-bromobenzofuran-2-yl)-4-phenyl-4*H*-1,2,4-triazol-3-yl)thio)-*N*-phenylacetamide (**10a**).

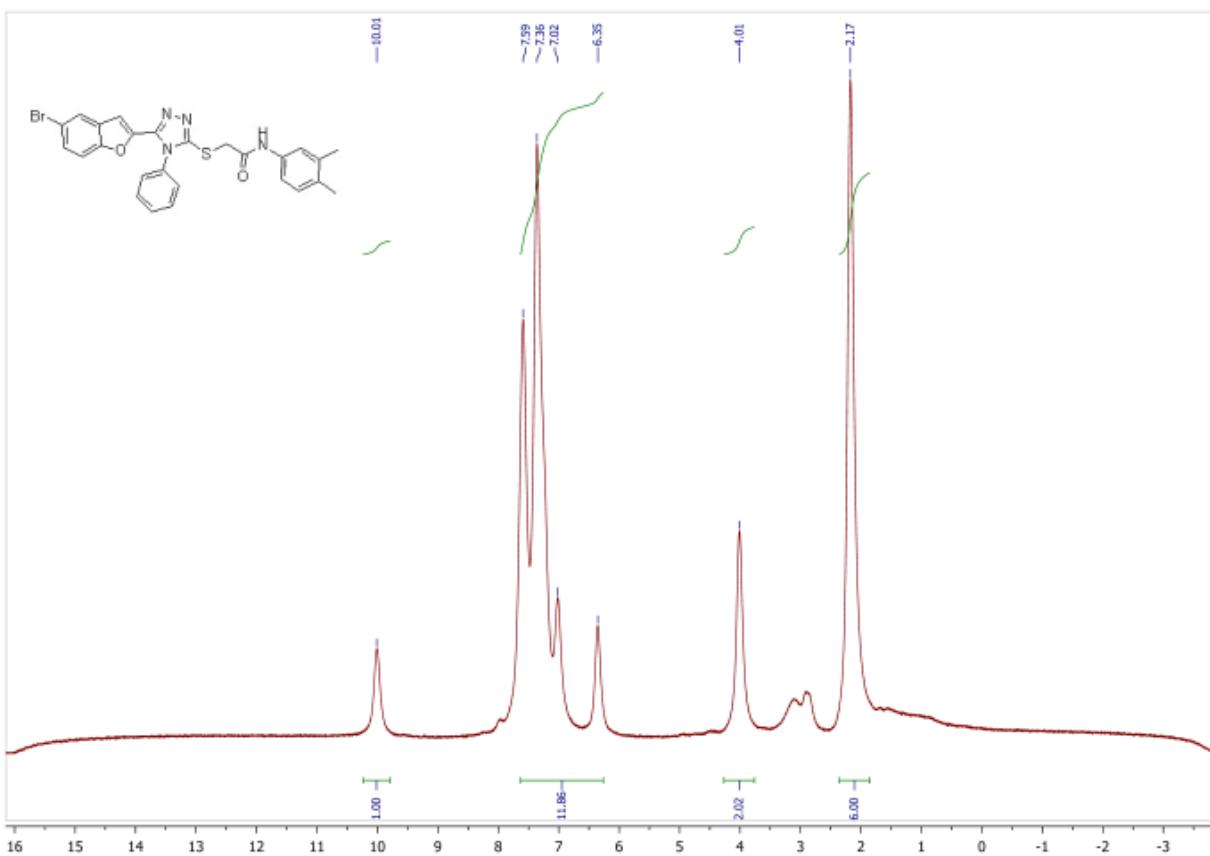


Figure S3: ^1H NMR spectrum of 2-((5-(5-bromobenzofuran-2-yl)-4-phenyl-4*H*-1,2,4-triazol-3-yl)thio)-*N*-(3,4-dimethylphenyl)acetamide (**10b**).

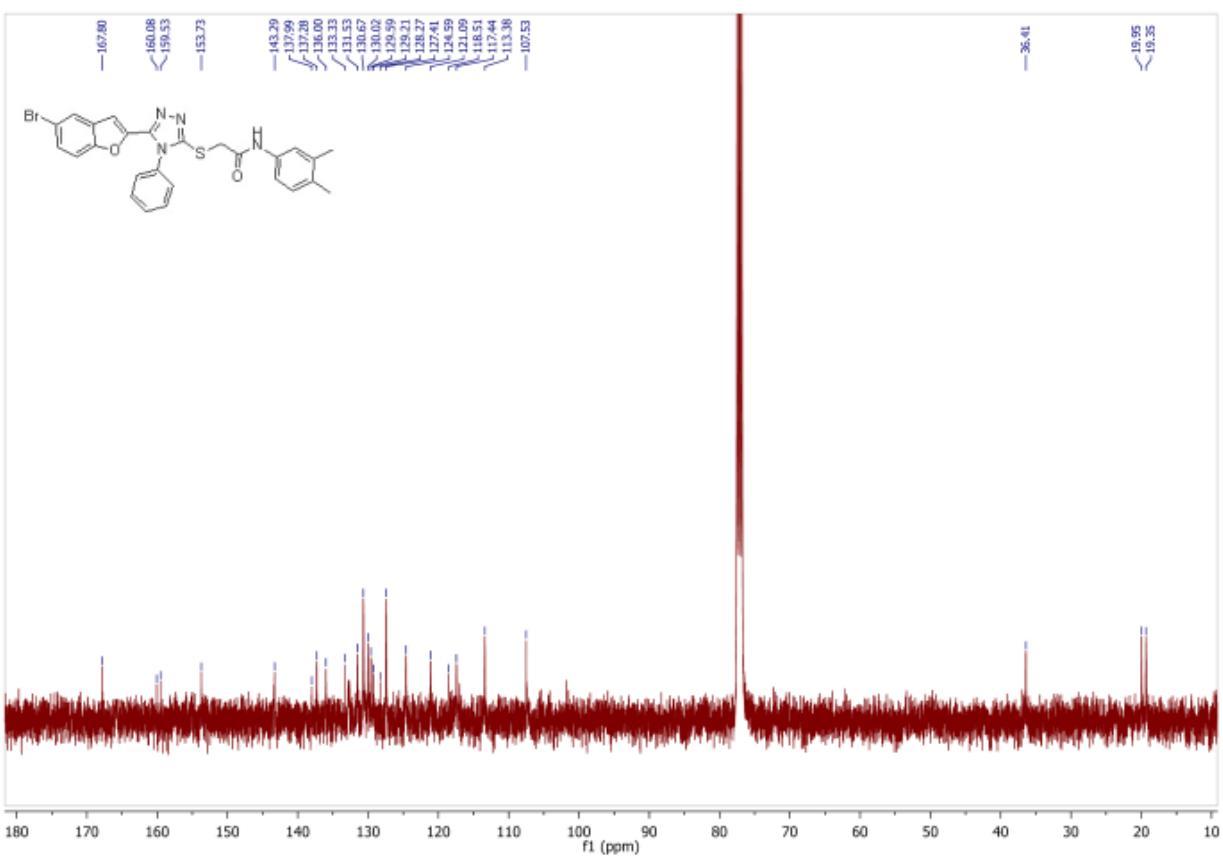


Figure S4: ^{13}C NMR spectrum of 2-((5-(5-bromobenzofuran-2-yl)-4-phenyl-4*H*-1,2,4-triazol-3-yl)thio)-*N*-(3,4-dimethylphenyl)acetamide (**10b**).

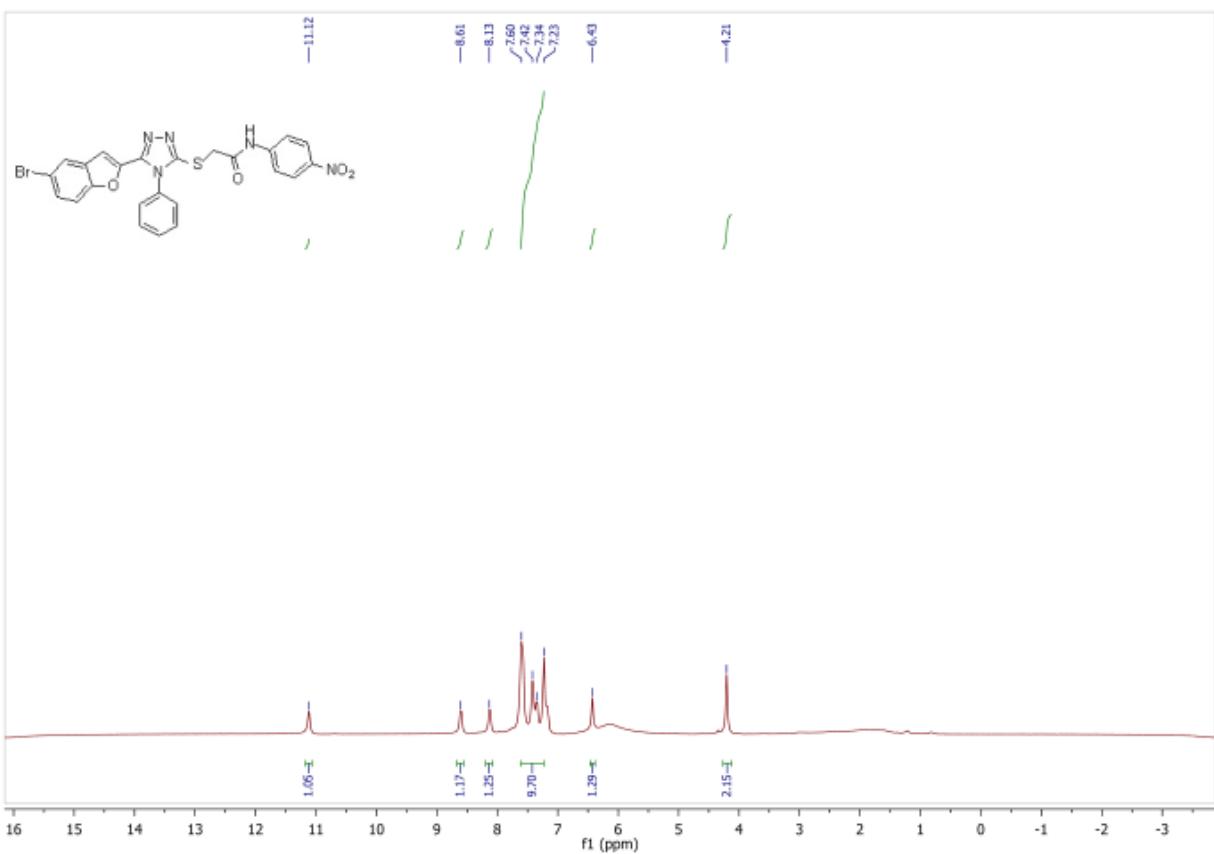


Figure S5: ^1H NMR spectrum of 2-((5-(5-bromobenzofuran-2-yl)-4-phenyl-4*H*-1,2,4-triazol-3-yl)thio)-*N*-(4-nitrophenyl)acetamide (**10c**).

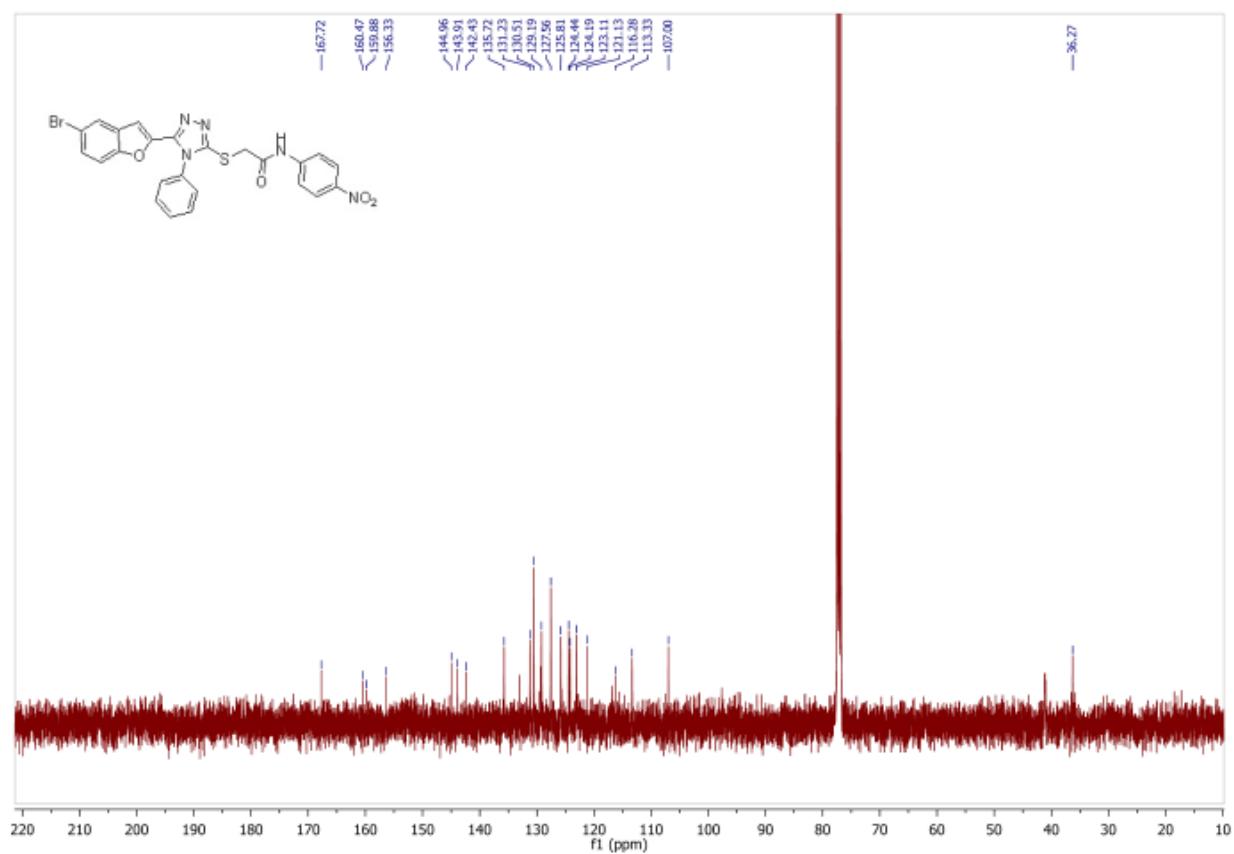


Figure S6: ^{13}C NMR spectrum of 2-((5-(5-bromobenzofuran-2-yl)-4-phenyl-4*H*-1,2,4-triazol-3-yl)thio)-*N*-(4-nitrophenyl)acetamide (**10c**).

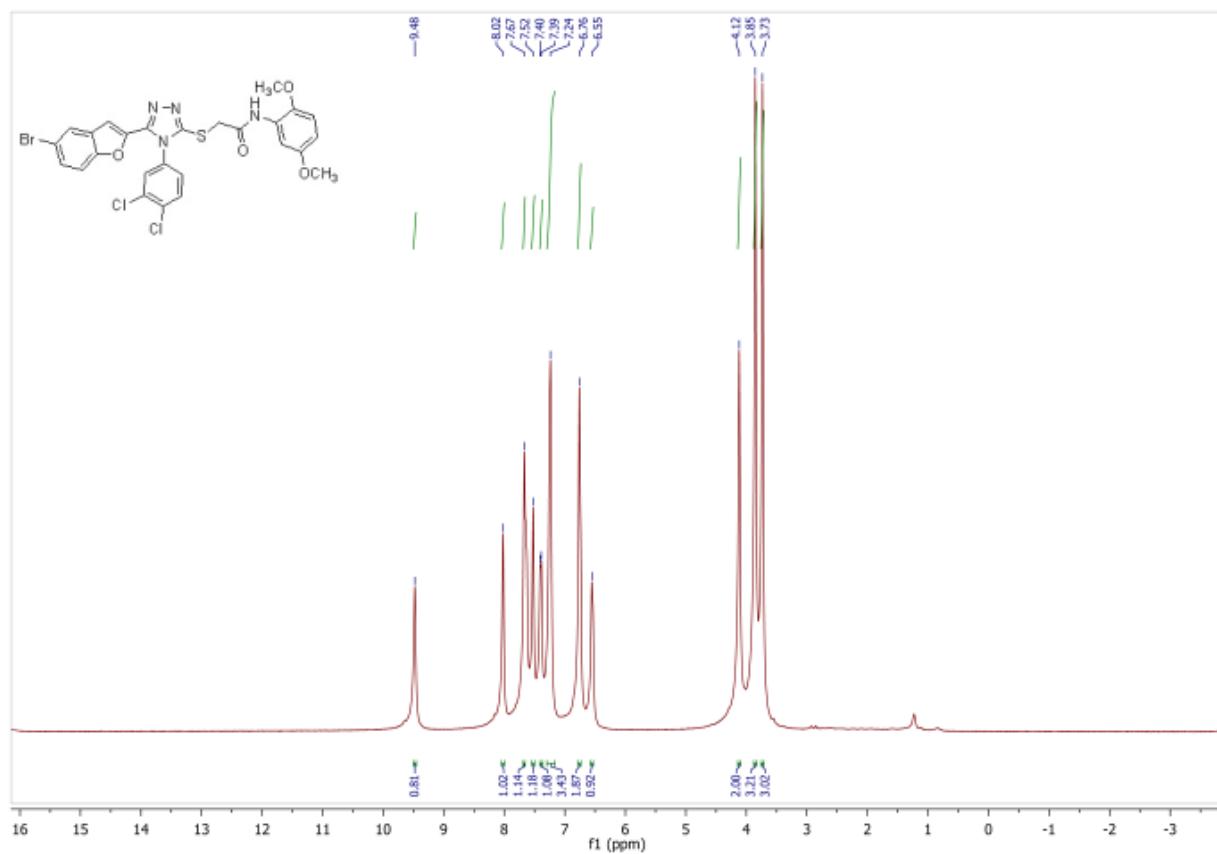


Figure S7: ¹H NMR spectrum of 2-((5-bromobenzofuran-2-yl)-4-(3,4-dichlorophenyl)-4*H*-1,2,4-triazol-3-yl)thio)-*N*-(2,5-dimethoxyphenyl)acetamide (**10d**).

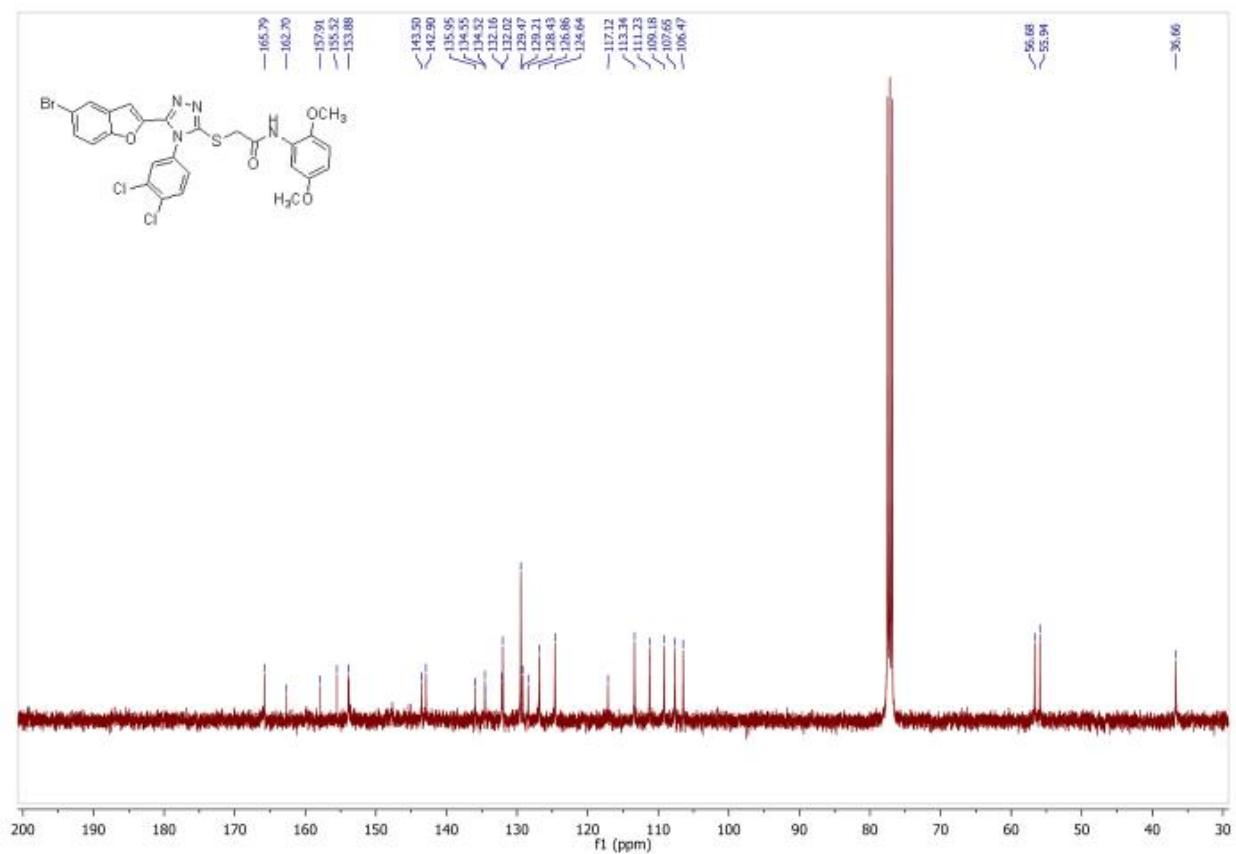


Figure S8: ^{13}C NMR spectrum of 2-((5-(5-bromobenzofuran-2-yl)-4-(3,4-dichlorophenyl)-4*H*-1,2,4-triazol-3-yl)thio)-*N*-(2,5-dimethoxyphenyl)acetamide (**10d**).

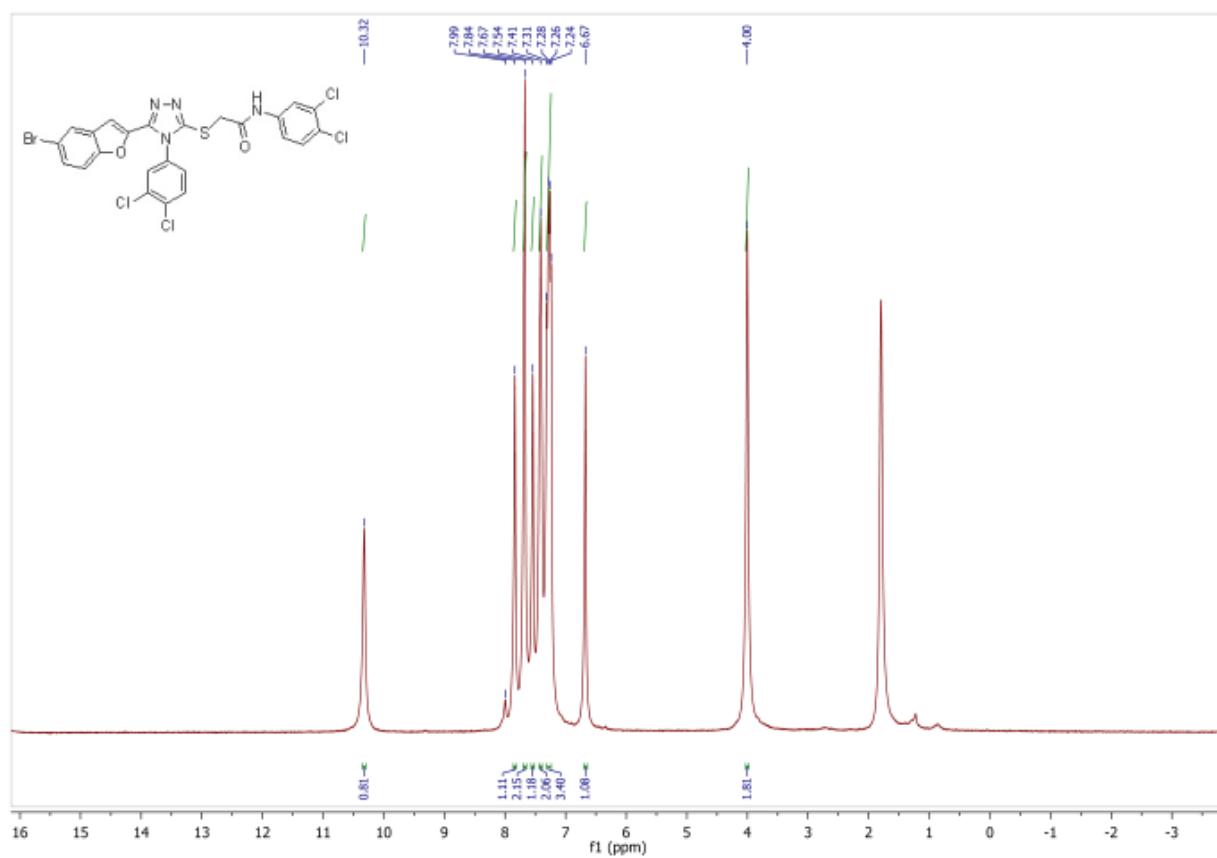


Figure S9: ^1H NMR spectrum of 2-((5-bromobenzofuran-2-yl)-4-(3,4-dichlorophenyl)-4*H*-1,2,4-triazol-3-yl)thio)-*N*-(3,4-dichlorophenyl)acetamide (**10e**).

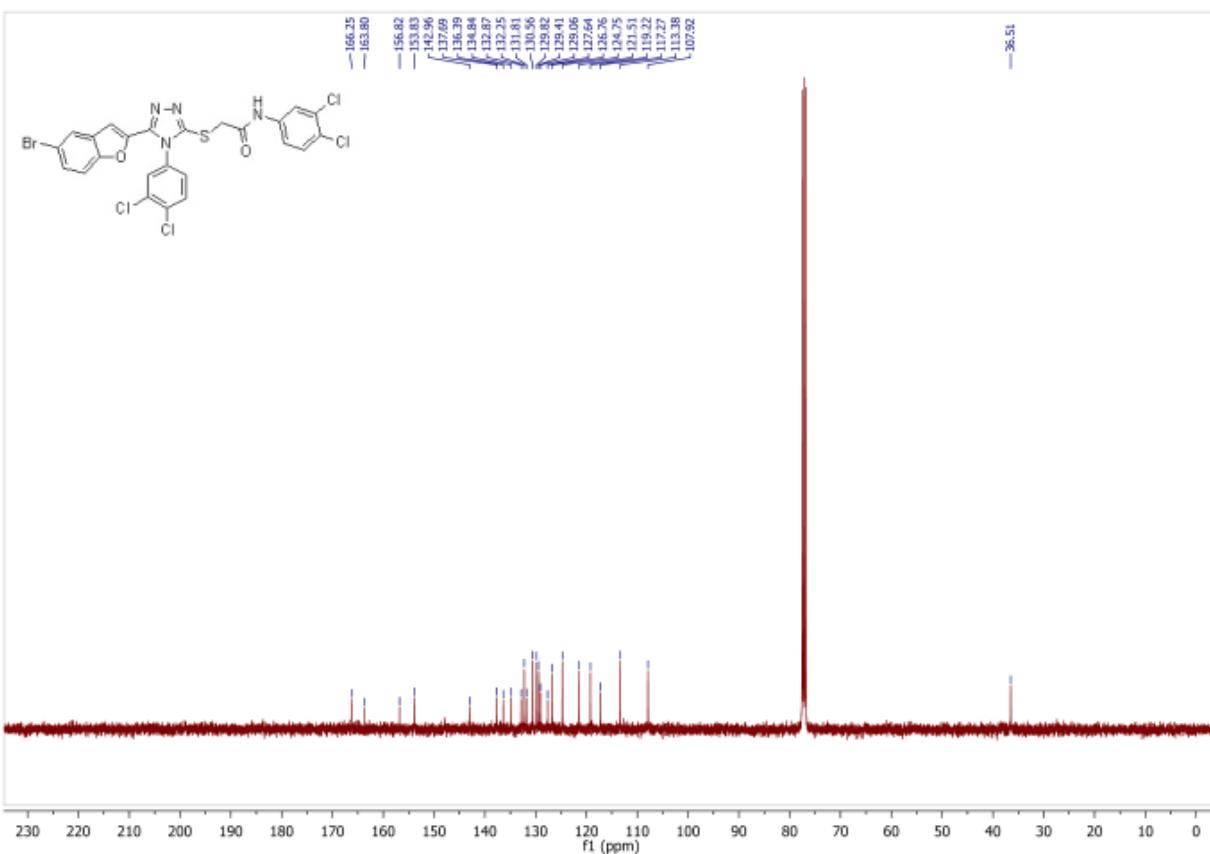


Figure S10: ^{13}C NMR spectrum of 2-((5-bromobenzofuran-2-yl)-4-(3,4-dichlorophenyl)-4*H*-1,2,4-triazol-3-yl)thio)-*N*-(3,4-dichlorophenyl)acetamide (**10e**).

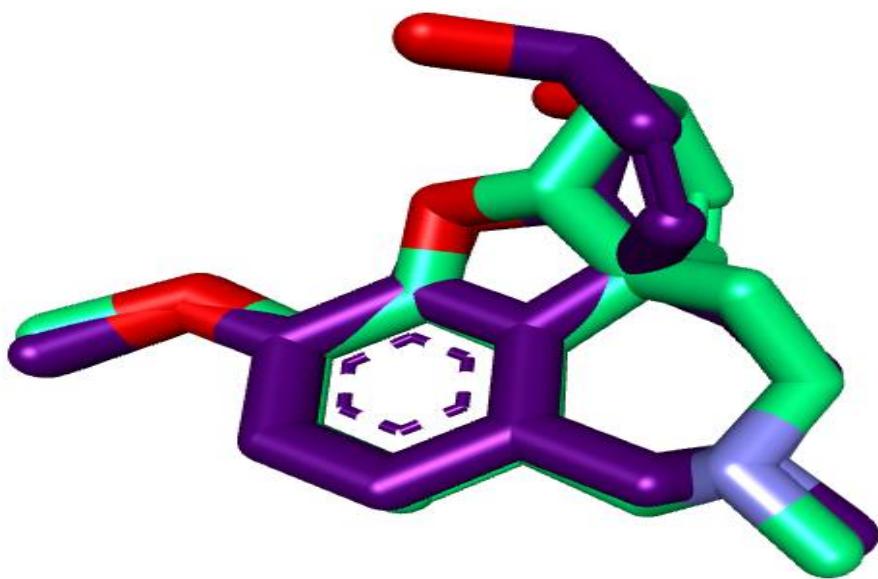


Figure S11. Cognate re-docking of co-crystallized ligand of AChE (4EY6) to validate IFD protocol; Co-crystallized apo-conformation (green) of native ligand aligned against its re-docked conformation reproduced by IFD protocol.