Supplementary Information

Novel Pyrimidine Linked Acyl Thiourea Derivatives as Potent α-Amylase and Proteinase

K Inhibitors: Design, Synthesis, Molecular Docking and ADME Studies

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Figure 1. ¹H-NMR Spectrum of compound (6a)





Figure 2.¹³C-NMR Spectrum of compound (6a)



Figure 3. FT-IR Spectrum of compound (6a)





Figure 4. ¹H-NMR Spectrum of compound (6b)



Figure 5.¹³C-NMR Spectrum of compound (6b)











Figure 7. ¹H-NMR Spectrum of compound (6c)



Figure 8. ¹³C-NMR Spectrum of compound (6c)

Figure 9. FT-IR Spectrum of compound (6c)



Figure 10. ¹H-NMR Spectrum of compound (6d)



Figure 11.¹³C-NMR Spectrum of compound (6d)



Figure 12. FT-IR Spectrum of compound (6d)



Figure 13. ¹H-NMR Spectrum of compound (6e)





Figure 14.¹³C-NMR Spectrum of compound (6e)



Figure 15. FT-IR Spectrum of compound (6e)





Figure 16. ¹H-NMR Spectrum of compound (6f)





Figure 17.¹³C-NMR Spectrum of compound (6f)



Figure 18. FT-IR Spectrum of compound (6f)





Figure 19. ¹H-NMR Spectrum of compound (6g)





Figure 20.¹³C-NMR Spectrum of compound (6g)

Figure 21. FT-IR Spectrum of compound (6g)





Figure 22. ¹H-NMR Spectrum of compound (6h)





Figure 23. ¹³C-NMR Spectrum of compound (6h)



Figure 24. FT-IR Spectrum of compound (6h)



Figure 25. FT-IR Spectrum of compound (6i)



Figure 26. FT-IR Spectrum of compound (6j)

6a





6c



6d

6b





6e





6f



6g



6h



6i



6j



Fig 27. 3D docking pose and 2D binding interaction of the synthesized compounds (6a-j) in the binding pocket of α -amylase

6a



6b



6c



6d





6f



6g





6i



6j



Fig 28. 3D docking pose and 2D binding interaction of the synthesized compounds (**6a-j**) in the binding pocket of Proteinase K