Supplementary Data

New 1,2,3-triazole-containing hybrids as antitumor candidates: Design, click-reaction synthesis, DFT calculations and molecular docking study

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I- DFT Calculations

Figure S1: Optimized structure of **4** and **5** derivatives at B3LYP /6-311++G(d,p) level of theory in the gas phase.





Figure S2: optimized structure of **6** and **7** derivatives at B3LYP/6-311++G(d,p) level of theory in the gas phase.



II- Molecular docking data



(A)

(B)

Figure S3. Docking of compound 6 within ATP bind site of EGFR. A) 3D biding mode. B) 2D binding mode.



Figure S4. Docking of compound 5 within ATP bind site of EGFR. A) 3D biding mode. B) 2D binding mode.



Figure S5. Docking of compound 4 within ATP bind site of EGFR. A) 3D biding mode. B) 2D binding mode.



Figure S6. Docking of compound 8 within ATP bind site of EGFR. A) 3D biding mode. B) 2D binding mode.



Figure S7. Docking of compound 9 within ATP bind site of EGFR. A) 3D biding mode. B) 2D binding mode.



Figure S8. Docking of compound 2 within ATP bind site of EGFR. A) 3D biding mode. B) 2D binding mode.



Figure S9. Docking of compound 3 within ATP bind site of EGFR. A) 3D biding mode. B) 2D binding mode.



Figure S10. ¹H NMR of compound 2







Figure S11. ¹H NMR of compound 3



Figure S12. ¹³C NMR of compound 3



Figure S13. 1H NMR of compound 4



Figure S14. ¹³C NMR of compound 4



Figure S15. ¹H NMR of compound 5







Figure S17. ¹H NMR of compound 6



Figure S18. ¹³C NMR of compound 6



Figure S19. ¹H NMR of compound 7



Figure S20. ¹H NMR of compound 8



Figure S21. ¹³C NMR of compound 8



Figure S22. ¹H NMR of compound 9