

Identification of new leads for human IGFBP-2: a therapeutic target for cardiovascular diseases

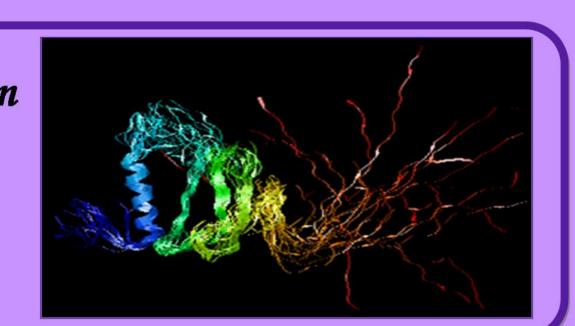
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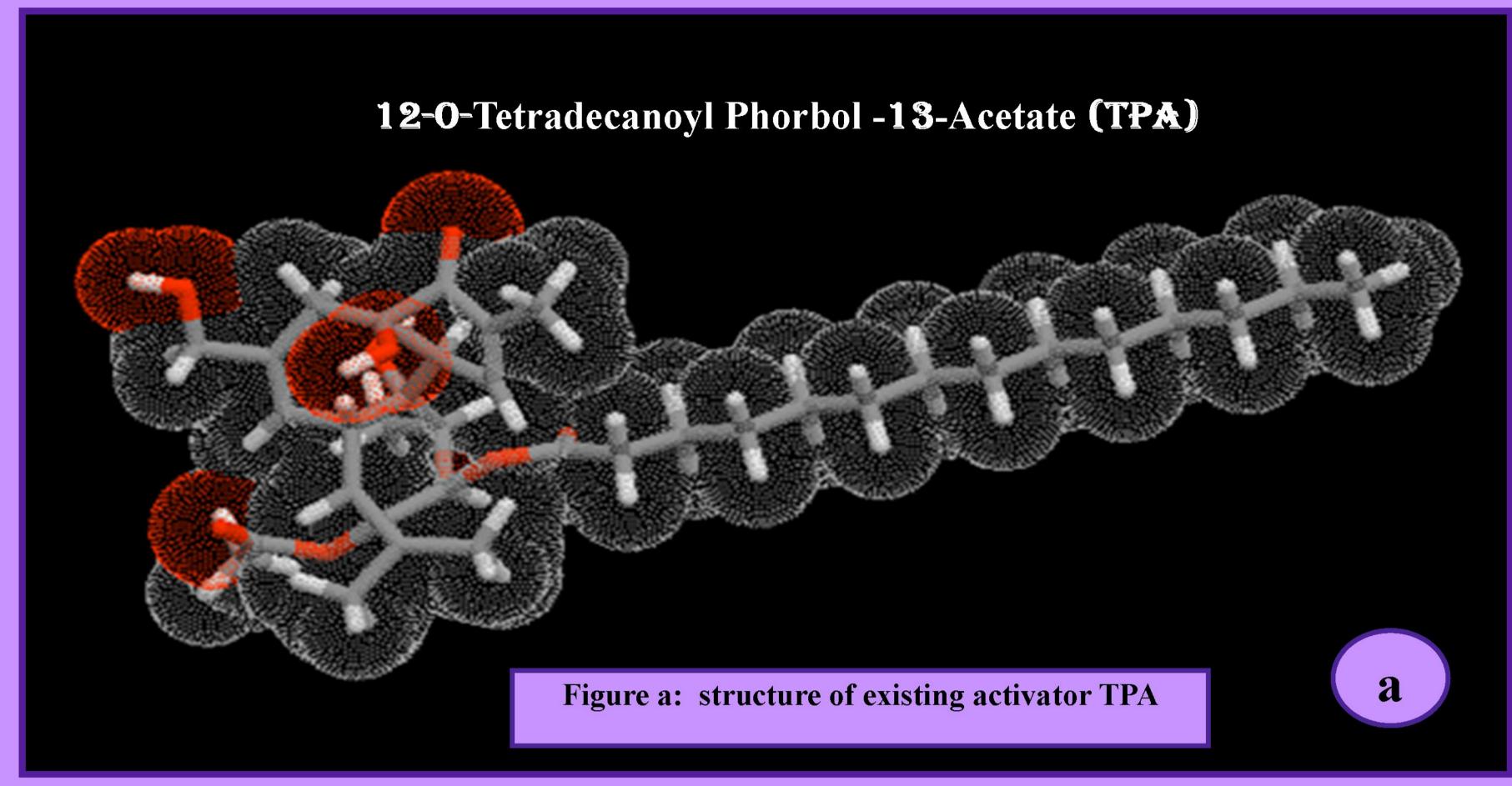
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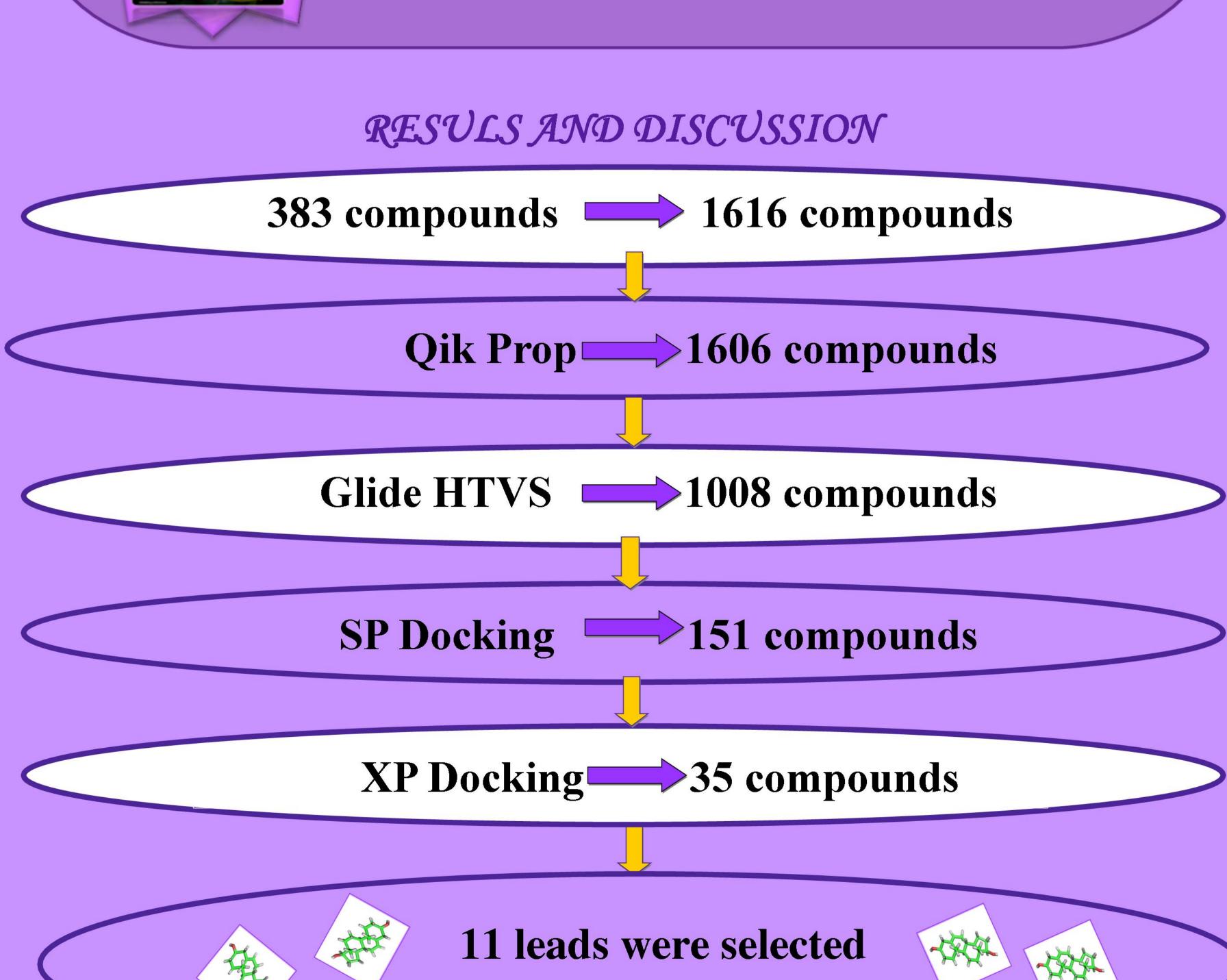


INTRODUCTION: Human insulin like growth factor binding protein-2 is the largest member of insulin-like growth factor binding protein family. Human IGFBP-2 is present in circulation where it acts to prolong half life of circulating IGFs, peripheral tissues and also in biological fluids. Human IGFBP-2 is down regulated in various diseases like obesity, type II diabetes mellitus, which upon aging leads to heart stroke. Therefore, human IGFBP-2 was selected as a drug target for cardiovascular disease therapy. In the present study high-throughput virtual screening and molecular docking studies were used to identify novel leads that up regulate the activity of human IGFBP-2.





MATERIALS AND METHODS **Retrieval of** V-198, R-201, I-202, M-205, R-206, L-207, P-208, N-209, R-211, G-215, H-216, L-217, Y-218, S-219, L-220, H-221, I-222, K-237, M-238, S-239, CAST L-240, N-241, Q-243, R-244, G-245, Glu-246, K-256, I-258, G-260, P-262, T-263, G-282, V-283, H-284, T-285, Q-286, R-287 and M-289. LIGAND BINDING SITES IN-HOUSE LIBRARY Ligand.info 383 structural analogues of TPA Docking analysis **Docking complex**



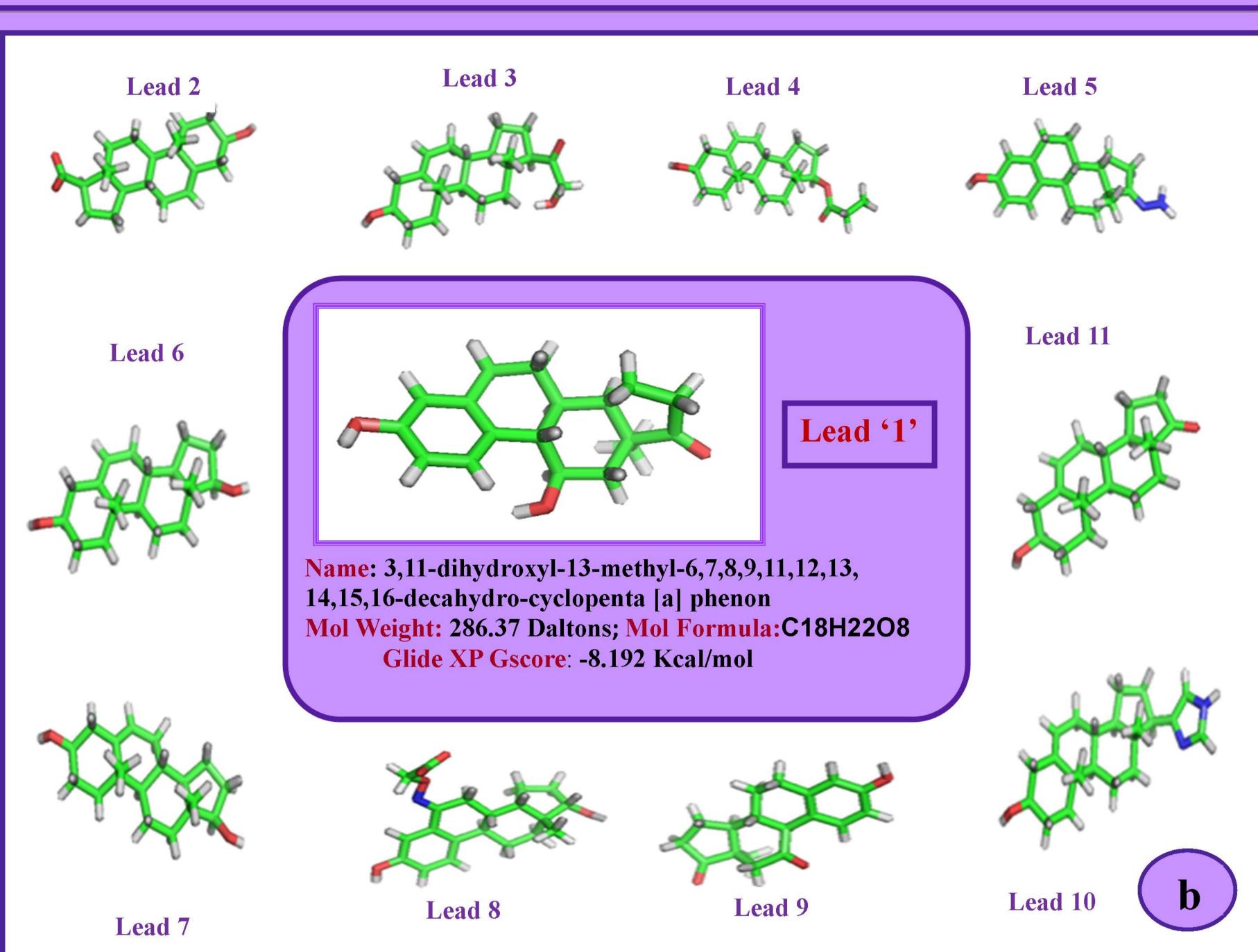


Figure b: Structures of proposed leads

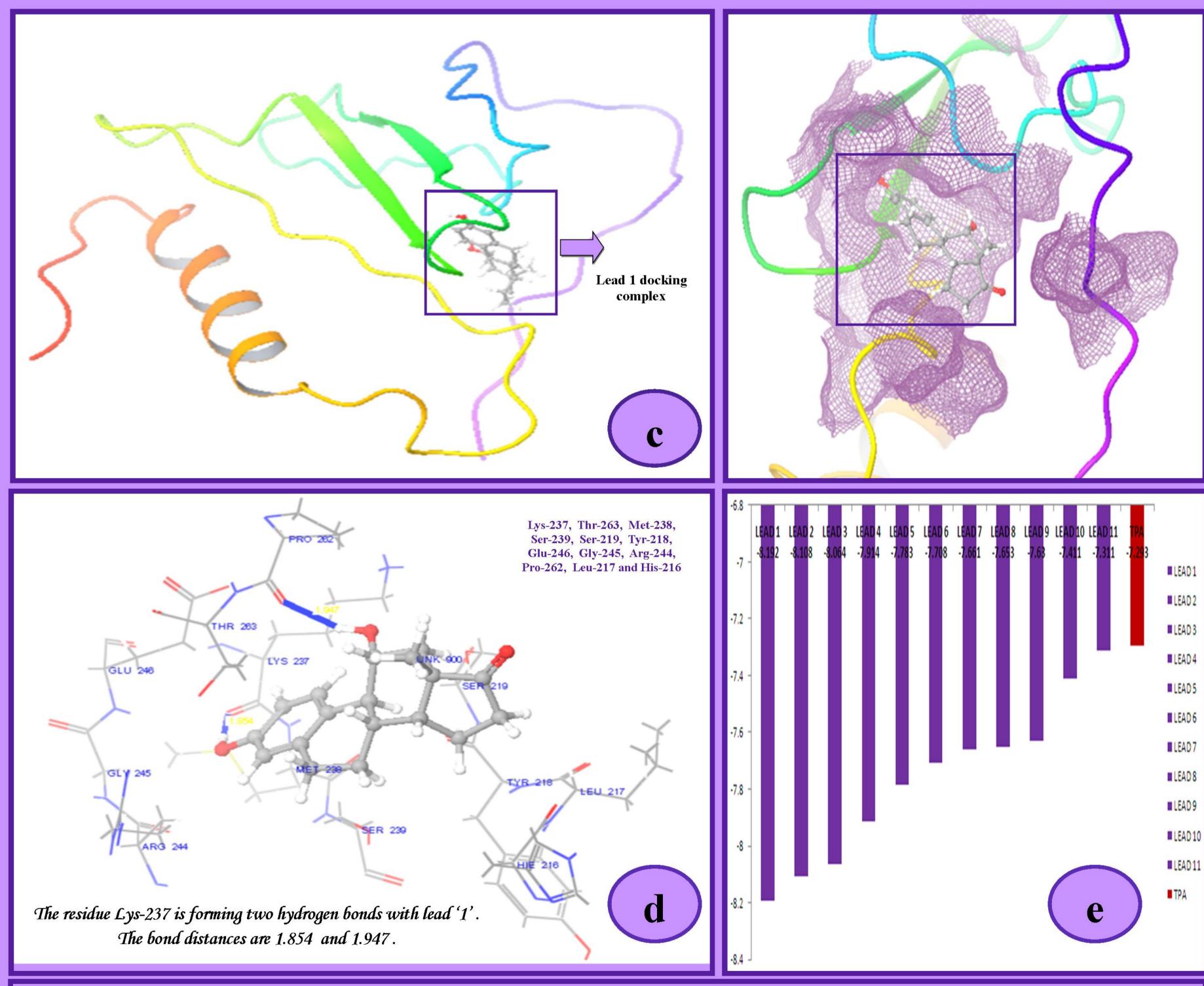


Figure c: Docking complex of lead '1' with human IGFBP-2. . Figure d: Hydrogen bond network between lead'1' L human IGFBP-2. . Figure e: Comparison of docking scores between lead molecules L TPA activator

CONCLUSION:

Low circulating levels of human IGFBP-2 leads to heart stroke.. Lead '1' 3,11-dihydroxyl-13-methyl-6,7,8,9,11,12,13,14,15,16-decahydro-cyclopenta [a] phenon is having good pharmacological properties, better docking score and good binding affinity to human IGFBP-2 protein than the existing activator (TPA). Hence, lead '1' is proposed as a promising lead to elevate the activity of Human IGFBP-2 protein if synthesized and validated in animal models.

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