



# Journal of Applicable Chemistry

2014, 3 (2): 622-629

(International Peer Reviewed Journal)



## Molecular Docking Studies Of Lawsone Derivatives With Tuberculosis Protein (PDB CODE: 1v0j)

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Accepted on 12<sup>th</sup> February 2014

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### ABSTRACT

*A series of seven numbers of lawsone derivatives are examined for the Insilico molecular docking studies with the tuberculosis protein (PDB CODE: 1v0j). The compound 6 shows the lowest inhibition constant 4.82  $\mu$ M and the compound 7 shows the highest inhibition constant 74.17  $\mu$ M.*

**Keywords:** Lawsone derivatives; Molecular docking; Tuberculosis protein; 1v0j.

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