

New D2R partial agonist candidates: an in silico approach from statistical models, molecular docking, and ADME/Tox properties

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Structural Chemistry

2021; 32(5):2019-2033

ARTICLE IDENTIFIERS

DOI: 10.1007/s11224-021-01742-w

PMID: unavailable

PMCID: not available

JOURNAL IDENTIFIERS

LCCN: not available

pISSN: 1040-0400

eISSN: not available

OCLC ID: not available

CONS ID: not available

US National Library of Medicine ID: not available

This article was identified from a query of the SafetyLit database.