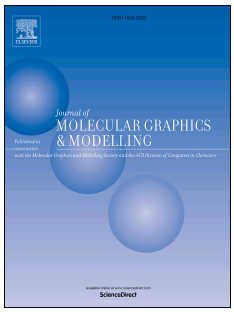
Accepted Manuscript



Identification of potent and selective COX-2 inhibitors via structure-based virtual screening and molecular dynamics simulation

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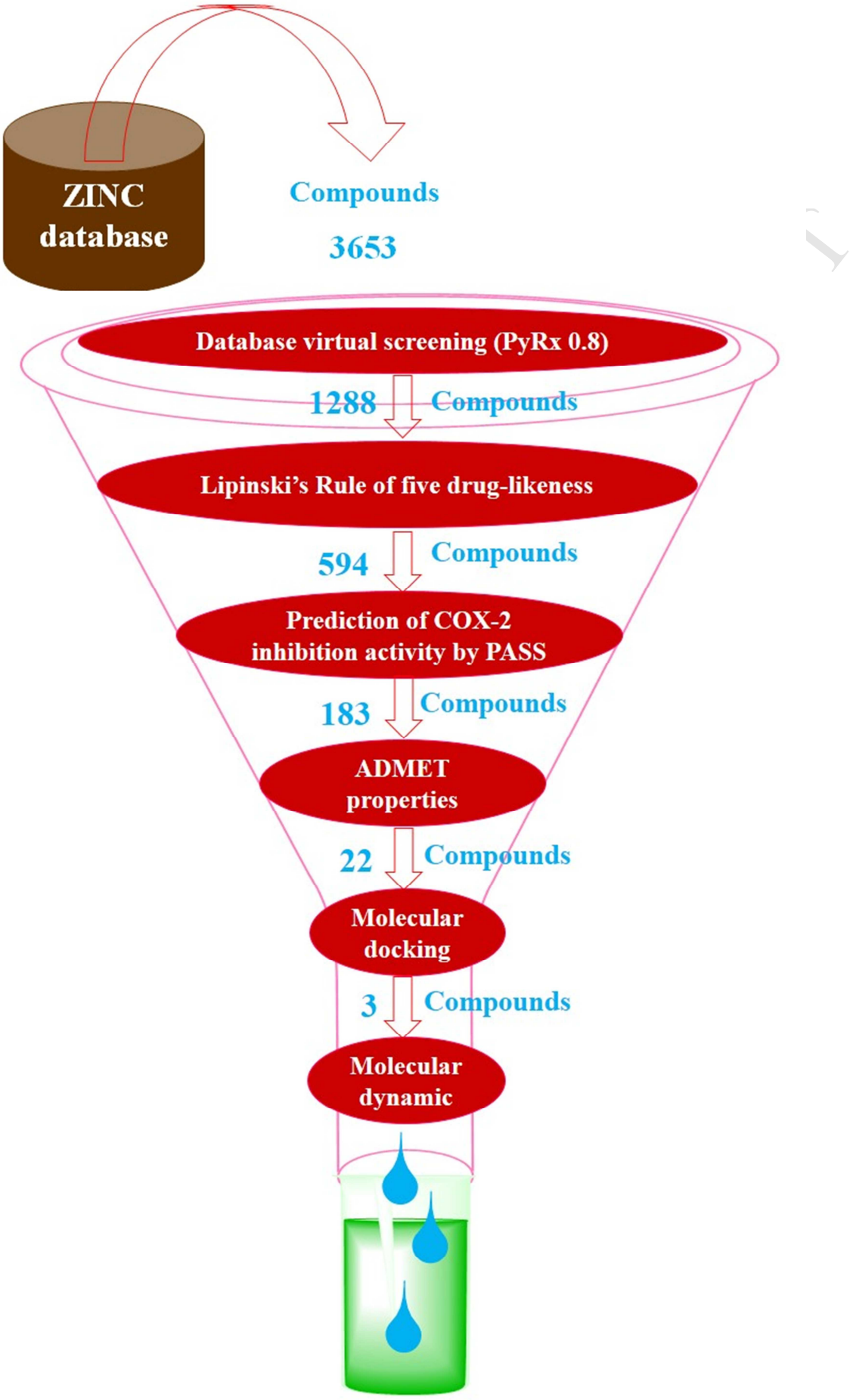
ZINC database
Compounds
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