

Juan A Castillo-Garit

List of Publications by Year in descending order

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Version: 2024-02-01

50
papers

1,227
citations

331259

21
h-index

377514

34
g-index

50
all docs

50
docs citations

50
times ranked

799
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Atom, atom-type and total molecular linear indices as a promising approach for bioorganic and medicinal chemistry: theoretical and experimental assessment of a novel method for virtual screening and rational design of new lead anthelmintic. <i>Bioorganic and Medicinal Chemistry</i> , 2005, 13, 1005-1020. | 1.4 | 97 |
| 2 | Markovian chemicals "in silico" design (MARCH-INSIDE), a promising approach for computer-aided molecular design I: discovery of anticancer compounds. <i>Journal of Molecular Modeling</i> , 2003, 9, 395-407. | 0.8 | 87 |
| 3 | Estimation of ADME Properties in Drug Discovery: Predicting Caco-2 Cell Permeability Using Atom-Based Stochastic and Non-stochastic Linear Indices. <i>Journal of Pharmaceutical Sciences</i> , 2008, 97, 1946-1976. | 1.6 | 72 |
| 4 | Protein linear indices of the ϵ -macromolecular pseudograph $\hat{\pm}$ -carbon atom adjacency matrix $\hat{\epsilon}$ ™ in bioinformatics. Part 1: Prediction of protein stability effects of a complete set of alanine substitutions in Arc repressor. <i>Bioorganic and Medicinal Chemistry</i> , 2005, 13, 3003-3015. | 1.4 | 70 |
| 5 | Atom, Atom-Type, and Total Linear Indices of the ϵ -Molecular Pseudograph $\hat{\epsilon}$ ™s Atom Adjacency Matrix $\hat{\epsilon}$ ™: Application to QSPR/QSAR Studies of Organic Compounds. <i>Molecules</i> , 2004, 9, 1100-1123. | 1.7 | 64 |
| 6 | Tomocomd-Cardd, a novel approach for computer-aided ? rational? drug design: I. Theoretical and experimental assessment of a promising method for computational screening and in silico design of new anthelmintic compounds. <i>Journal of Computer-Aided Molecular Design</i> , 2004, 18, 615-634. | 1.3 | 62 |
| 7 | Predicting antitrichomonal activity: A computational screening using atom-based bilinear indices and experimental proofs. <i>Bioorganic and Medicinal Chemistry</i> , 2006, 14, 6502-6524. | 1.4 | 53 |
| 8 | A novel approach to predict aquatic toxicity from molecular structure. <i>Chemosphere</i> , 2008, 73, 415-427. | 4.2 | 50 |
| 9 | Atom-based stochastic and non-stochastic 3D-chiral bilinear indices and their applications to central chirality codification. <i>Journal of Molecular Graphics and Modelling</i> , 2007, 26, 32-47. | 1.3 | 45 |
| 10 | Linear indices of the ϵ -macromolecular graph $\hat{\epsilon}$ ™s nucleotides adjacency matrix $\hat{\epsilon}$ ™ as a promising approach for bioinformatics studies. Part 1: Prediction of paromomycin $\hat{\epsilon}$ ™s affinity constant with HIV-1 $\hat{\nu}$ -RNA packaging region. <i>Bioorganic and Medicinal Chemistry</i> , 2005, 13, 3397-3404. | 1.4 | 44 |
| 11 | Tyrosinase Enzyme: 1. An Overview on a Pharmacological Target. <i>Current Topics in Medicinal Chemistry</i> , 2014, 14, 1494-1501. | 1.0 | 38 |
| 12 | In Silico Assessment of ADME Properties: Advances in Caco-2 Cell Monolayer Permeability Modeling. <i>Current Topics in Medicinal Chemistry</i> , 2019, 18, 2209-2229. | 1.0 | 38 |
| 13 | 3D-chiral Atom, Atom-type, and Total Non-stochastic and Stochastic Molecular Linear Indices and their Applications to Central Chirality Codification. <i>Journal of Computer-Aided Molecular Design</i> , 2005, 19, 369-383. | 1.3 | 35 |
| 14 | Atom-based 3D-chiral quadratic indices. Part 2: Prediction of the corticosteroid-binding globulinbinding affinity of the 31 benchmark steroids data set. <i>Bioorganic and Medicinal Chemistry</i> , 2006, 14, 2398-2408. | 1.4 | 34 |
| 15 | Prediction of acute toxicity of phenol derivatives using multiple linear regression approach for <i>Tetrahymena pyriformis</i> contaminant identification in a median-size database. <i>Chemosphere</i> , 2016, 165, 434-441. | 4.2 | 28 |
| 16 | A Review of QSAR studies to Discover New Drug-like Compounds Actives Against Leishmaniasis and Trypanosomiasis. <i>Current Topics in Medicinal Chemistry</i> , 2012, 12, 852-865. | 1.0 | 27 |
| 17 | Bond $\hat{\epsilon}$ -based 3D $\hat{\epsilon}$ -chiral linear indices: Theory and QSAR applications to central chirality codification. <i>Journal of Computational Chemistry</i> , 2008, 29, 2500-2512. | 1.5 | 26 |
| 18 | Computational discovery of novel trypanosomicidal drug-like chemicals by using bond-based non-stochastic and stochastic quadratic maps and linear discriminant analysis. <i>European Journal of Pharmaceutical Sciences</i> , 2010, 39, 30-36. | 1.9 | 26 |

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|----|---|-----|-----------|
| 19 | A Simple Method to Predict Blood-Brain Barrier Permeability of Drug- Like Compounds Using Classification Trees. <i>Medicinal Chemistry</i> , 2017, 13, 664-669. | 0.7 | 25 |
| 20 | 3D-chiral (2.5) atom-based TOMOCOMD-CARDD descriptors: theory and QSAR applications to central chirality codification. <i>Journal of Mathematical Chemistry</i> , 2008, 44, 755-786. | 0.7 | 23 |
| 21 | Atom-based non-stochastic and stochastic bilinear indices: Application to QSPR/QSAR studies of organic compounds. <i>Chemical Physics Letters</i> , 2008, 464, 107-112. | 1.2 | 23 |
| 22 | Ligand-based discovery of novel trypanosomicidal drug-like compounds: In silico identification and experimental support. <i>European Journal of Medicinal Chemistry</i> , 2011, 46, 3324-3330. | 2.6 | 19 |
| 23 | Comparative study to predict toxic modes of action of phenols from molecular structures. SAR and QSAR in <i>Environmental Research</i> , 2013, 24, 235-251. | 1.0 | 19 |
| 24 | An approach to identify new antihypertensive agents using Thermolysin as model: In silico study based on QSARINS and docking. <i>Arabian Journal of Chemistry</i> , 2019, 12, 4861-4877. | 2.3 | 19 |
| 25 | Bond-based bilinear indices for computational discovery of novel trypanosomicidal drug-like compounds through virtual screening. <i>European Journal of Medicinal Chemistry</i> , 2015, 96, 238-244. | 2.6 | 16 |
| 26 | In silicoAntibacterial Activity Modeling Based on the TOMOCOMD-CARDD Approach. <i>Journal of the Brazilian Chemical Society</i> , 2015, , . | 0.6 | 16 |
| 27 | Bond-based linear indices of the non-stochastic and stochastic edge-adjacency matrix. 1. Theory and modeling of ChemPhys properties of organic molecules. <i>Molecular Diversity</i> , 2010, 14, 731-753. | 2.1 | 15 |
| 28 | Identification <i>In Silico</i> and <i>In Vitro</i> of Novel Trypanosomicidal Drug-Like Compounds. <i>Chemical Biology and Drug Design</i> , 2012, 80, 38-45. | 1.5 | 14 |
| 29 | Machine learning-based models to predict modes of toxic action of phenols to <i>Tetrahymena pyriformis</i> . SAR and QSAR in <i>Environmental Research</i> , 2017, 28, 735-747. | 1.0 | 14 |
| 30 | Discrete Fourier Transform-Based Multivariate Image Analysis: Application to Modeling of Aromatase Inhibitory Activity. <i>ACS Combinatorial Science</i> , 2018, 20, 75-81. | 3.8 | 14 |
| 31 | Computational Identification of Chemical Compounds with Potential Activity against <i>Leishmania amazonensis</i> using Nonlinear Machine Learning Techniques. <i>Current Topics in Medicinal Chemistry</i> , 2019, 18, 2347-2354. | 1.0 | 12 |
| 32 | Applications of Bond-Based 3D-Chiral Quadratic Indices in QSAR Studies Related to Central Chirality Codification. <i>QSAR and Combinatorial Science</i> , 2009, 28, 1465-1477. | 1.5 | 11 |
| 33 | Prediction of aquatic toxicity of benzene derivatives using molecular descriptor from atomic weighted vectors. <i>Environmental Toxicology and Pharmacology</i> , 2017, 56, 314-321. | 2.0 | 11 |
| 34 | Prediction of Aquatic Toxicity of Benzene Derivatives to <i>Tetrahymena pyriformis</i> According to OECD Principles. <i>Current Pharmaceutical Design</i> , 2016, 22, 5085-5094. | 0.9 | 10 |
| 35 | Prediction of Caco-2 Cell Permeability Using Bilinear Indices and Multiple Linear Regression. <i>Letters in Drug Design and Discovery</i> , 2015, 13, 161-169. | 0.4 | 9 |
| 36 | State of the Art Review and Report of New Tool for Drug Discovery. <i>Current Topics in Medicinal Chemistry</i> , 2017, 17, 2957-2976. | 1.0 | 8 |

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|----|--|-----|-----------|
| 37 | Larvicidal activity prediction against mosquito using computational tools. Journal of Vector Borne Diseases, 2017, 54, 164-171. | 0.1 | 8 |
| 38 | Analysis of Proteasome Inhibition Prediction Using Atom-Based Quadratic Indices Enhanced by Machine Learning Classification Techniques. Letters in Drug Design and Discovery, 2014, 11, 705-711. | 0.4 | 7 |
| 39 | Undersampling: case studies of flaviviral inhibitory activities. Journal of Computer-Aided Molecular Design, 2019, 33, 997-1008. | 1.3 | 6 |
| 40 | Thorough evaluation of OECD principles in modelling of 1-[(2-hydroxyethoxy)methyl]-6-(phenylthio)thymine derivatives using QSARINS. SAR and QSAR in Environmental Research, 2020, 31, 741-759. | 1.0 | 6 |
| 41 | Learning from Multiple Classifier Systems: Perspectives for Improving Decision Making of QSAR Models in Medicinal Chemistry. Current Topics in Medicinal Chemistry, 2018, 17, 3269-3288. | 1.0 | 5 |
| 42 | Evolutionary algorithm-based generation of optimum peptide sequences with dengue virus inhibitory activity. Future Medicinal Chemistry, 2021, 13, 993-1000. | 1.1 | 4 |
| 43 | Ligand-based discovery of new potential acetylcholinesterase inhibitors for Alzheimer's disease treatment. SAR and QSAR in Environmental Research, 2022, 33, 49-61. | 1.0 | 4 |
| 44 | Machine learning approach to discovery of small molecules with potential inhibitory action against vasoactive metalloproteases. Molecular Diversity, 2022, 26, 1383-1397. | 2.1 | 3 |
| 45 | Computational identification of chemical compounds with potential anti-Chagas activity using a classification tree. SAR and QSAR in Environmental Research, 2021, 32, 71-83. | 1.0 | 3 |
| 46 | A Review of Computational Approaches Targeting SARS-CoV-2 Main Protease to the Discovery of New Potential Antiviral Compounds. Current Topics in Medicinal Chemistry, 2023, 23, 3-16. | 1.0 | 3 |
| 47 | <i>Dry</i> selection and <i>wet</i> evaluation for the <i>rational</i> discovery of new anthelmintics. Molecular Physics, 2017, 115, 2300-2313. | 0.8 | 2 |
| 48 | Atom based linear index descriptors in QSAR-machine learning classifiers for the prediction of ubiquitin-proteasome pathway activity. Medicinal Chemistry Research, 2018, 27, 695-704. | 1.1 | 1 |
| 49 | Computational approach to the discovery of potential neprilysin inhibitors compounds for cardiovascular diseases treatment. Medicinal Chemistry Research, 2020, 29, 897-909. | 1.1 | 1 |
| 50 | Computational Modeling of Aldose Reductase Inhibitory Activity of Flavonoids Derivatives for Diabetic Complications Treatment. Letters in Drug Design and Discovery, 2021, 18, . | 0.4 | 0 |