Challenges and Advances in Computational Chemistry and Physics 33 Series Editor: Jerzy Leszczynski

Alla P. Toropova Andrey A. Toropov *Editors*

QSPR/QSAR Analysis Using SMILES and Quasi-SMILES





Challenges and Advances in Computational Chemistry and Physics

Volume 33

Series Editor

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This book series provides reviews on the most recent developments in computational chemistry and physics. It covers both the method developments and their applications. Each volume consists of chapters devoted to the one research area. The series highlights the most notable advances in applications of the computational methods. The volumes include nanotechnology, material sciences, molecular biology, structures and bonding in molecular complexes, and atmospheric chemistry. The authors are recruited from among the most prominent researchers in their research areas. As computational chemistry and physics is one of the most rapidly advancing scientific areas such timely overviews are desired by chemists, physicists, molecular biologists and material scientists. The books are intended for graduate students and researchers.

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QSPR/QSAR Analysis Using SMILES and Quasi-SMILES



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Preface

Who is this book for intended? Primarily for students who are planning their carrier. Ph.D. students can also get valuable ideas for their careers if they are sure that their scientific activity somehow connects with chemistry, biology, medicine, informatics, and mathematical chemistry. The author's team contains specialists in different directions of chemistry, biochemistry, and medicinal chemistry. The geography of the authors is vast enough: USA, Canada, Iran, India, China, Uzbekistan, Czech Republic, Portugal and Italy.

It seems that recognizing the differences in the paths of transition of randomness into regularity or, conversely, the ways of randomness into stable chaos may be of interest to everyone since this task affects any area of human activity. In fact, this book describes attempts to solve the mentioned problem concerning development processes QSPR/QSAR and nano-QSPR/QSAR.

The curious intrigue of the proposed book demonstrates the ability of randomness to provide patterns through variational autoencoders (VAEs) defined over SMILES string and molecular graph, the Monte Carlo technique, and using so-called quasi-SMILES (i.e., traditional SMILES extended via special symbols which are reflecting experimental conditions). However, the philosophic principle "nothing is the only" should make the reader sure that every model should be validated as much as possible, i.e., checked up under a diversity of experimental conditions.

Thus, there is the probability that the book can become curiously and attractive to various "random" readers (professors, engineers, players) who are capable of curios and wonder relevant to the process of building up models for different phenomena.

Milan, Italy

Alla P. Toropova Andrey A. Toropov

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Abbreviations

AAD	Average absolute deviation
ACE	Angiotensin-converting enzymes
AD	Applicability domain
AFM	Atomic force microscopy
ANFIS	Adaptive neuro-fuzzy inference system
ANN	Artificial neural networks
AZI	Augmented Zagreb index
BET	Brunauer, Emmett and Teller
CCC	Concordance correlation coefficient
CII	Correlation intensity index
CORAL	Correlation and logic
Cp	Isobaric heat capacity
CW	Correlation weights
DCW	Descriptor of correlation weights
DHFR	Dihydrofolate reductase
DLS	Dynamic light scattering
DTR	Decision tree regression
EDX	Energy dispersive X-ray spectrometry
EG	Ethylene glycol
EM	Electronic microscopy
EP	Endpoint
ESEM	Environmental scanning electron microscopy
F	Fischer ratio
FF-ANNs	Feed-forward artificial neural networks
FFF	Field flow filtration
FMO	Frontier molecular orbital theory
GBR	Gradient boosting regression
GNPs	Gold nanoparticles
GRNNs	Generalized regression neural networks
GRUs	Gated recurrent units
HOMO	Highest occupied molecular orbital

HSG	Hydrogen-suppressed molecular graphs
ICP-MS	Inductively coupled plasma mass spectrometry
ICPOES	
	Inductively coupled plasma emission spectroscopy
IIC	Index ideality of correlation
ILs	Ionic liquids
LC	Liquid chromatography
LDM	Liquid drop model
logP	Decimal logarithm of octanol-water partition coefficient
LSSVM	Least square support vector machine
LSTM	Long short-term memory
LUMO	Lowest unoccupied molecular orbital
MAE	Mean absolute error
MLP	Multilayer perceptron
MLR	Multiple regression analysis
MO-NPs	Metal oxide nanoparticles
MoRSE	3D-Molecular representation of structures based on electron
	diffraction
MVC	Multivariate characterization
MW	Molecular weight
MWCNTs	Multiwalls carbon nanotubes
NPs	Nanoparticles
OECD	Organization of Economic Co-operation and Development
PCA	Principal component analyses
PLS	Partial least-squares regression analysis
PPs	Principal properties
Q^2	Leave-one-out cross-validated correlation coefficient
QED	Quantitative estimate of drug-likeness
QSAR	Quantitative structure-activity relationship
QSGFEAR	Gibb's free energy of activation relationship
QSPR	Quantitative structure-property relationship
Quasi-SMILES	Quasi-simplified molecular input-line entry-system
R^2	Determination coefficient (or squared correlation coefficient)
RBF	Radial basis function
RF	Random forest
RMSE	Root-mean-square error
RNNs	Recurrent neural networks
SA	SMILES attributes
SADT	Self-accelerating decomposition temperature
SFS	Sequential forward selection
SMILES	Simplified molecular input-line entry-system
SNN	Siamese neural network
SVM	Support vector machine
SVR	Support vector inachine Support vector regression
SWCNTs	Single-wall carbon nanotubes
TC	Thermal conductivity
IC IC	i nemiai conductivity

TEM	Transmission electron microscopy
TF	Target function
TMACC	Topological maximum cross-correlation
VAEs	Variational autoencoders
VIF	Variation inflation factor
WW	Hyper-Wiener index
ΔG ‡	Gibb's activation free energy

Greek Symbols

ρ Density

 φ Volume fraction of nanoparticle (%)

Subscripts

- bf Base fluid
- nf Nanofluid
- p Nanoparticle
- v Volume fraction

Chemical Formulas

Ag	Silver
Al_2O_3	Aluminum oxide
AlN	Aluminum nitride
Au	Gold
Bi ₂ O ₃	Bismuth (III) oxide
CeO_2	Cerium (IV) oxide
Co ₃ O ₄	Cobalt (II,III) oxide
Cr_2O_3	Chromium (III) oxide
Cu	Copper
CuO	Copper oxide
Dy_2O_3	Dysprosium (III) oxide
Fe	Iron
Fe ₂ O ₃	Iron (III) oxide
Fe ₃ O ₄	Iron (II,III) oxide
Gd_2O_3	Gadolinium (III) oxide
HfO ₂	Hafnium (IV) oxide

Chapter 5 SMILES-Based Bioactivity Descriptors to Model the Anti-dengue Virus Activity: A Case Study



Soumya Mitra, Sumit Nandi, Amit Kumar Halder, and M. Natalia D. S. Cordeiro

Abstract The present work aims to demonstrate the significance of the newly suggested bioactivity descriptors (so-called signaturizers) towards developing predictive 2D-OSAR models. As a case study, we examined the development of 2D-QSAR models based on a dataset containing 77 compounds with inhibitory activity reported in a DENV2ProHeLa assay, which is basically a cell-based assay that estimates the Dengivirus-2 (DENV-2) protease inhibitory potential within cellular atmosphere. Indeed, though dengue is a well-known neglected tropical disease, its global incidence has risen sharply in recent years. Moreover, DENV infections may lead to serious and life-threatening diseases such as haemorrhagic fever and dengue shock syndrome. Inhibition of the DENV protease may therefore be a potential target for discovering anti-DENV agents. Interestingly, our initial attempts to set up QSAR models based solely on a number of chemicals descriptors coming from a range of different software packages/programs completely failed, since none of these yielded satisfactory statistical results. Hybrid QSAR models were generated also by combining both chemical and biological descriptors. Noteworthy is that the predictive quality of the 2D-QSAR models significantly improved by resorting instead to solely bioactivity descriptors or those combined with chemical descriptors. The comparison analysis carried out in this work certainly shows that bioactivity descriptors can be useful for setting up predictive models to characterise complex

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