

QSAR/QSPR Modeling as a Cornerstone for Modern Molecular Design and Property Prediction

Aleksandar M. Veselinović^{1*}

¹Department of Chemistry, Faculty of Medicine, University of Niš, Serbia

*aleks.veselinovic@gmail.com; aveselinovic@medfak.ni.ac.rs

Quantitative Structure-Activity/Property Relationship (QSAR/QSPR) modeling is fundamental in modern chemistry, pharmaceutical sciences, and materials research. As the chemical space expands rapidly, the ability to predict molecular properties and biological activities directly from molecular structure has become crucial for effective research planning, prioritization, and risk assessment.

Various QSPR/QSAR strategies have been developed, including the use of SMILES-based descriptors in conjunction with Monte Carlo optimization, as well as alternative descriptor systems and machine learning approaches. These innovations enable the development of fragment-based models and generalized predictive frameworks capable of estimating a wide range of physicochemical, pharmacokinetic, and toxicological endpoints.

Today, QSAR/QSPR modeling is recognized as a validated, practical, and essential tool across multiple sectors. Its applications include regulatory toxicology (OECD, REACH), predictive ADME profiling, safety assessment, formulation development, and synthesis optimization. Additionally, these models help bridge experimental gaps where data may be limited or ethically constrained.

QSPR modeling supports property prediction in various areas of chemistry and materials science. By estimating key physicochemical parameters, these models facilitate rational design, expedite candidate selection, streamline synthesis planning, and reduce the experimental workload.

In drug discovery, QSAR and Computer-Aided Drug Design (CADD) techniques accelerate the early selection of compounds, enabling the swift identification of promising candidates while eliminating those with unfavorable pharmacokinetic or toxicological profiles. This targeted approach focuses experimental efforts on viable structures, reduces synthesis cycles, minimizes late-stage failures, and decreases the time and cost associated with drug development.

In addition to application-driven modeling, ongoing research continues to advance descriptor engineering, integrate graph-based and SMILES-based descriptors, and develop hybrid machine learning frameworks. These efforts present broad opportunities for synergistic experimental-computational research, particularly in interdisciplinary projects that combine chemistry, toxicology, pharmaceutical sciences, green chemistry, and sustainable synthesis.