

Development of Web and Mobile Applications for Chemical Toxicity Prediction

Vinicius M. Alves,^{a,b} Rodolpho C. Braga,^a Eugene Muratov^{b,c} and Carolina H. Andrade^{*,a}

^aLaboratório de Planejamento de Fármacos e Modelagem Molecular, LabMol, Faculdade de Farmácia, Universidade Federal de Goiás, 74605-170 Goiânia-GO, Brazil

^bLaboratory for Molecular Modeling, Division of Chemical Biology and Medicinal Chemistry, UNC Eshelman School of Pharmacy, University of North Carolina, 27599 Chapel Hill, NC, USA

^cDepartment of Chemical Technology, Odessa National Polytechnic University, 65000 Odessa, Ukraine

Computational tools are recognized to provide high-quality predictions for the assessment of chemical toxicity. In the recent years, mobile devices have become ubiquitous, allowing for the development of innovative and useful models implemented as chemical software applications. Here, we will briefly discuss this recent uptick in the development of web-based and mobile applications for chemical problems, focusing on best practices, development, usage and interpretation. As an example, we also describe two innovative apps (Pred-hERG and Pred-Skin) for chemical toxicity prediction developed in our laboratory. These applications are based on predictive quantitative structure-activity relationships (QSAR) models developed using the largest publicly available datasets of structurally diverse compounds. The developed tools ensure both highly accurate predictions and easy interpretation of the models, allowing users to discriminate potential toxicants and to purpose structural modifications to design safer chemicals.

Keywords: web app, mobile, toxicity prediction, QSAR, Pred-hERG, Pred-Skin

1. Introduction

Chemical safety assessment is a fundamental step in development and regulation of drugs, cosmetics and other chemicals.¹ In the last few decades, society has less tolerated animal testing.² As a consequence, legislations, guidelines, and practice of animal experiments have been implementing the principles to reduce, refine, and replace animals used in laboratory. As a prime example, *in vivo* evaluation for cosmetic products is forbidden in Europe since 2003³ and the sale of cosmetics and ingredients tested on animals are banned since 2013.⁴ In Brazil, since 2015, alternative methods recognized in the National Regulatory Agency (ANVISA) are sufficient for the approval of chemicals⁵ and animal testing will not be allowed for endpoints with approved alternative methods (e.g., eye irritation, skin irritation, acute toxicity, etc.) starting from 2019.⁶

Computational methods have become an effective alternative method for the evaluation of new or untested

*e-mail: carolina@ufg.br, carolhandrade@gmail.com

compounds, such as drug candidates, cosmetics and pesticides, providing high-quality predictions at low-cost and reasonable time.^{7,8} Toxicity prediction often relies on structural alerts,⁹ read-across,¹⁰ and quantitative structure-activity/property relationships (QSAR/QSPR) models.^{11,12} Structural alerts are molecular substructures that are associated with a particular adverse biological effect.¹³ Read-across is a technique that extrapolates data based on structural similarity (usually using structural alerts) to previously measured compound(s) for those lacking experimental data.¹⁴ This method has earned prominence from a regulatory perspective^{15,16} due to its simplicity and transparency.¹⁷

Scientific applications implemented in a web interface (web apps) are an interesting alternative to standalone software applications, since they are ready to use, fast, and they usually present an intuitive interface. Several web apps to solve chemical and molecular modeling problems have been proposed through the years, such as the Platform for Unified Molecular Analysis (PUMA),¹⁸ SWISS-MODEL,^{19,20} Chembench,²¹ Pred-hERG,^{22,23} Pred-Skin,²⁴ among others. Conversely, to a lesser extent,

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The Green Solvents mobile app is one of the first chemistry apps. It is based on a dataset of solvents containing several informations to serve as a solvent guide for chemists.²⁸ The TB Mobile app provides information on over 700 molecules screened *versus Mycobacterium tuberculosis* from the Collaborative Drug Discovery (CDD) database. The app also performs similarity search, which can infer potential targets from similar structures.²⁹ The Lead Designer app³⁰ provides a tool for drawing compounds and predict permeability through blood-brain barrier membrane, CYP3A4 binding potency, and some physical-chemical properties.

Recently, our group has developed and implemented predictive QSAR models in two distinctive tools available as a web app (Pred-hERG)²³ and web and mobile apps (Pred-Skin),²⁴ for the fast prediction of hERG cardiac toxicity and skin sensitization, respectively. Here, we aim to describe the details behind the development of these two innovative and impactful web-based and mobile applications for chemical toxicity prediction.

2. Chemical Toxicity Prediction Using Apps

Chemical toxicity prediction usually relies on chemical similarity, by using structural alerts and read-across, as well as QSAR models.³¹ Structural alerts are molecular fragments that are associated with a positive response for

a particular endpoint.⁹ Chemical read-across uses chemical similarity to assess the effect of untested chemicals.^{10,32} Lastly, QSAR models are a major computational approach that uses statistical or machine learning algorithms to establish a correlation of molecular representation to activity and can be used to provide statistically significant prediction of chemicals.³³

A reliable mobile or web server for chemical toxicity prediction should be based on highly predictive and externally accurate models. As we and collaborators have recently shown,³¹ structural alerts are oversensitive, disproportionally flagging chemicals as toxic. Therefore, the use of statistically validated QSAR models are preferred. The best practices for development and validation of QSAR models have been recommended by the Organization for Economic Co-operation and Development (OECD)³⁴ and elaborated by Tropsha.35 Briefly, any QSAR model should have "(i) a defined endpoint; (ii) an unambiguous algorithm; (*iii*) a defined applicability domain; (*iv*) appropriate measures of goodness-of-fit, robustness, and predictivity; (v) and, if possible, a mechanistic interpretation." In addition, the reproducibility crisis has questioned the quality of experimental data published in peer-reviewed journals.36 For this reason, the curation of the chemical and biological data is of upmost importance.37-39

The workflow we have adopted for the development of the web apps is shown on Figure 1. First, we compile the biggest dataset available for the particular endpoint, following by the curation protocol proposed by Fourches *et al.*³⁷⁻³⁹ Then, models are developed following the best practices for model development and validiation.³⁵ Web-based and mobile apps are then developed (see details



Figure 1. General workflow for development of QSAR models and their implementation for web and mobile applications for toxicity prediction of chemical compounds.

in next section). Finally, apps are made available for use of the scientific community to predict new compounds using an intuitive tool with highly confident predictions and graphical interpretation.

One of the advantages of building apps based on QSAR models is the wide applicability of these models through different industrial chemical classes. As recently reported,⁴⁰ QSAR models generated for cosmetics, drugs and pesticides can be used interchangeably, i.e., a model developed using mainly drugs and drug-like compounds, can be used also to evaluate cosmetics and pesticides since a QSAR model built mainly using cosmetics compounds usually shares the same chemical space of most drugs or even pesticides. This is of high value for endpoints such as skin sensitization, as many industrial products (e.g., cosmetics, drugs, pesticides, food additives, etc.) could be in touch with the skin and cause allergic reactions.⁴¹ Despite the wide applicability of models generated using structurally diverse chemical datasets, one should be aware of the applicability domain of the model to evaluate the confidence of the prediction.⁴²⁻⁴⁴ This is especially true because chemical space has faced a substantial growth in the last decade. For instance, the Chemical Abstract Services has reached 100 million chemical substances in 2015,45 and the current version of the ZINC database46 contains over 35 million purchasable compounds.

3. Development of Web-Based and Mobile Applications

The development of integrative web-based and mobile applications running machine learning routines written in Python is possible by using Flask.⁴⁷ Flask is a small framework for creating web microframeworks. Flask was created from scratch as an extensible framework with a solid core of essential services and easy integration with Python extensions. The template support of Flask is provided by Jinja2, while the Web Server Gateway Interface (WSGI) subsystems is based on the Werkzeug toolkit. Usually web applications features such as validation of webforms, user authentication, and access to databases are employed through extensions. The main drawback of Flask is the lack of support for projects dealing with a large source file, leading the structuration of the app entirely to the developer.

The implementation of enormously complex workflows for cheminformatics applications written in Python with Flask is possible by integrating RDKit⁴⁸ and scikit-learn. RDKit is an open-source cheminformatics application programming interface (API) developed by Gregory Landrum at Novartis Rational Discovery for building predictive models for pharmacokinetics properties, toxicity, and biological activity. RDKit can be used to calculate molecular descriptors, such as Morgan fingerprints (extended circular fingerprints like),^{49,50} and to enable the generation of appropriate molecular representations for developing QSAR models using machine learning algorithms. The scikit-learn API, a combination of NumPy, SciPy, and matplotlib packages, can run machine learning scripts, perform datasets splits, data curation, model generation, and calculation of the appropriate statistical measures in conjunction with 5-fold external validation process, allowing an appropriate evaluation of the generated models.

Python routines integrating developed machine learning models using RDKit and scikit-learn works in the back-end of the app (Figure 2).^{51,52} All the functionalities are orchestrated by Flask (back-front-end), which is responsible to call the individual modules in the back-end and interact with the user in the front-end, using the responsive web-based templates in HTML5 and JavaScript (front-end). The features described here were used to develop the two innovative web apps, named Pred-hERG, to identify possible hERG blockers and non-blockers, and Pred-Skin, for assessment of skin sensitization potential of chemicals (see Pred-hERG and Pred-Skin sections).

4. Pred-hERG

The Pred-hERG is a web app that allows users to predict blockers and non-blockers of the hERG channels, and important drug anti-target associated with lethal cardiac arrhythmia.⁵³ This app has a fast and intuitive interface. We implemented binary (blocker *vs.* non-blocker) and multi-classification models, which are able to distinguish weak/moderate and strong/extreme blockers. We also implemented the probability maps of atomic contribution as predicted by the models, allowing users to interpret the results and propose structural modifications for the predicted compound. The current version of the app (v. 4.0) was developed using ChEMBL⁵⁴ version 23, containing 8,134 compounds with hERG blockage data after curation. This app is publicly available at the website.⁵⁵

5. Pred-Skin

The Pred-Skin²⁴ is an open source web-based and mobile application for evaluation of skin sensitization potential using externally validated QSAR models based on human and animal (local lymph node assay, LLNA) data. The app represents a benchmark for the prediction of skin sensitization, since it is the first tool to provide O S F Cl Br I P X

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Figure 2. Pred-Skin front-back-end technology (full-stack).⁵² (1) front-end using the responsive web-based templates in HTML5 and JavaScript for the user to draw/paste the chemical; (2) RDKit ECFP-4 fingerprint (Morgan) representation is calculated on the back-end; (3) scikit-learn API machine learning model runs on the back-end; (4) the front-end responsive framework templates based on Flask is rendered with the models predictions and the probability maps.⁵¹

predictions from models based on human data. Predictions for a single compound are produced within a few seconds. The following outputs are provided: (*i*) binary predictions of human and murine skin sensitization; (*ii*) multiclass predictions of murine skin sensitization potency; and (*iii*) probability maps illustrating the predicted contribution of chemical fragments. This app is freely available to the public at the website⁵⁶ and at the App Store.

6. Usage and Interpretation of Apps for Toxicity Prediction

There is a strong need for the development of userfriendly tools that would allow non-experts to usually predict their compounds of interest and also visualize important structural features related to increase or decrease the activity/toxicity of these compounds. QSAR models have been commonly referenced as a non-interpretable approach.⁵⁷ This common sense has led experimentalists and regulatory agencies to prefer the use of more simple and transparent approaches, such as structural alerts or read-accross.⁵⁸ However, many studies have shown that QSAR models can be interpreted and structural features associated with activity or toxicity identified.⁵⁹⁻⁶⁴ These substructures derived from the interpretation of QSAR models can be used to design novel compounds with improved toxicity profile.⁶⁵

Willing to provide predictive and interpretative tools for toxicity prediction, we have been working to implement highly predictive QSAR models in web and mobile apps with intuitive usage and interpretation. Both Pred-hERG and Pred-Skin allows the user to make predictions by pasting SMILES strings or drawing molecules in the JSME⁶⁶ molecule editor. Alternatively, the user can load .sdf or .mol files. After hitting the "Predict" button, the user will receive the QSAR predictions on the computer/ mobile screen.

In addition, we have implemented the probability maps proposed by Riniker and Landrum⁵¹ in our apps. This feature provides a graphical visualization of the predicted fragment contribution, allowing the user to interpret the prediction and to design safer compounds. The atoms are highlighted according to their predicted activity contribution. Atoms and molecular fragments highlighted in green represent a positive contribution (increase in the toxicity potential); whereas, pink fragments represent a negative contribution (decrease in toxicity potential). The gray fragments do not contribute to toxicity potential and gray isolines define the frontier between the positive (green) and the negative (pink) contributions.

7. Final Remarks

Despite the progress made in the last decade to predict toxicity endpoints, we observe that these efforts still do not fully guarantee that all new chemicals do not induce toxicity or harzard to human health. Therefore, the development of alternative methods, including *in silico* or computational methods for addressing toxicity potential of chemicals is of upmost importance. Machine learning methods, such as modern QSAR approaches have become more powerful due to the rapid expansion of bioactivity and toxicity data available in chemical databases such as TOXNET,67,68 ChEMBL,69 and PubChem.70,71 In this paper, we have discussed and reviewed the development of web-based and mobile applications for chemical problems, focusing on best practices, development, usage, and application. We have also described two innovative and freely accessible web-based and mobile applications for evaluation of potential hERG blockers and skin sensitizers developed in our laboratory. These applications are based on robust and predictive QSAR models developed using the largest publicly available data sets of structurally diverse compounds. The freely available Pred-hERG and Pred-Skin web apps ensure both highly accurate predictions and easy interpretation of the models, allowing users to perform rapid screening of large libraries of virtual compounds to discriminate potential toxicants as to purpose structural modifications to design safer chemicals.

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Vinicius M. Alves received his PhD in Pharmaceutical Sciences (2017) from the Federal University of Goiás, Brazil. He has experience in the development and implementation of innovative cheminformatics and molecular modeling approaches;

quantitative structure-activity relationships (QSAR) modeling; complex data analysis; and application of predictive models in screening large libraries of virtual compounds for prioritizing hits to be tested experimentally. He received in 2014 the Chemical Information Scholarship for Scientific Excellence awarded by the American Chemical Society. He also visited the University of Florida (USA) as a Research Intern (2011) and attended the University of North Carolina at Chapel Hill (USA) in 2012 and in 2015-2016 as a visiting scholar.



Rodolpho C. Braga is a Data Scientist and Drug Discovery Specialist working at Altox Alternative Toxicology Inc. Worked 9 months on the Lead Optimization Latin America project (LOLA) by Drugs for Neglected

Diseases initiative (DNDi), as a cheminformatician and computational chemistry scientist. Also, he was a Visiting Professor in Medicinal Chemistry at University of Turin (UniTO) (2015-2016). He was awarded with CINF Scholarship for Scientific Excellence of the American Chemical Society (ACS-2014). A Research Scientist with experience conducting and overseeing a variety of successful computational chemistry, cheminformatics, project management, molecular modeling, and complex data analysis research projects. Academic qualifications, including a Postdoctoral (2016), a Doctor of Philosophy (2015) and a Master of Science (2012) in Medicinal Chemistry from the Federal University of Goiás.



Eugene N. Muratov is a Research Assistant Professor and Associate Director of Laboratory for Molecular Modeling at the UNC Eshelman School of Pharmacy, University of North Carolina, Chapel Hill. He received his MSc in technology of

organic substances from Odessa National Polytechnic University in 2000 and PhD in organic chemistry in 2004 from the A. V. Bogatsky Physical-Chemical Institute. Before obtaining research assistant professor position in UNC in 2012, he spent several years as a researcher in Odessa, Jackson, Strasbourg, and Chapel Hill. In 2014-2017 he was visiting Federal University of Goiás, Goiânia, Brazil, as Invited Professor. His research interests are in the areas of cheminformatics (especially QSAR), computer-assisted drug design, antiviral research, computational toxicology, and medicinal chemistry. He has co-authored more than 100 peer-reviewed publications, and edited two books.



Carolina H. Andrade is Associate Professor at Federal University of Goiás. Graduated in Pharmacy (2004) and Doctor of Philosophy (2009) in Drugs and Medicines from University of São Paulo. Her research focuses on Computer-Aided

Drug Design, aiming at discovering new drug candidates for Neglected Tropical Diseases and Cancer. Moreover, cheminformatics tools and QSAR for the prediction of toxicity properties of chemical compounds are also her field of expertise. She is CNPq productivity research fellow since 2012. In 2014, she was awarded with the "For Women in Science" award from L'Oréal-ABC-UNESCO and in 2015 she received the "International Rising Talents" from L'Oréal-UNESCO. In 2016, she was elected Affiliated member of the Brazilian Academy of Sciences. Currently, she is vice-director of the Medicinal Chemistry Division of the Brazilian Chemical Society.

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