XXXI Symposium on Bioinformatics and Computer-Aided Drug Discovery

AmlActive (AIA): A Large-scale QSAR Based Target Fishing and Polypharmacology Predictive Web Tool

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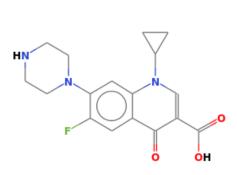


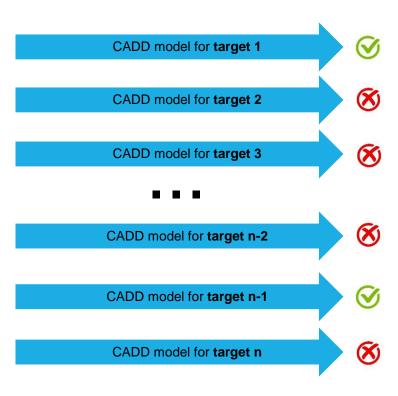






Reverse virtual screening (target fishing)







Introduction

Web tools for reverse virtual screening











Web tools for reverse virtual screening





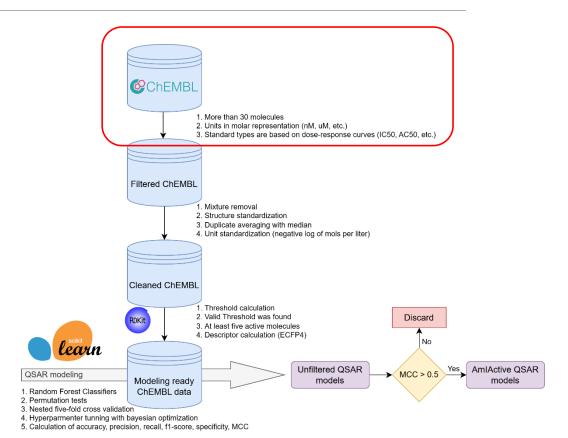


PLATO r35

Materials and methods



- Data was extracted from ChEMBL v30
- Targets where either:
 - Single proteins
 - Protein Families
 - Protein Complexes
 - Organisms
 - Cell-lines
 - Tissues
- Assay units where in some form of molar representation
- Assay metric was based on dose-response curves
- All filters considered, the final dataset had at least 30 molecules

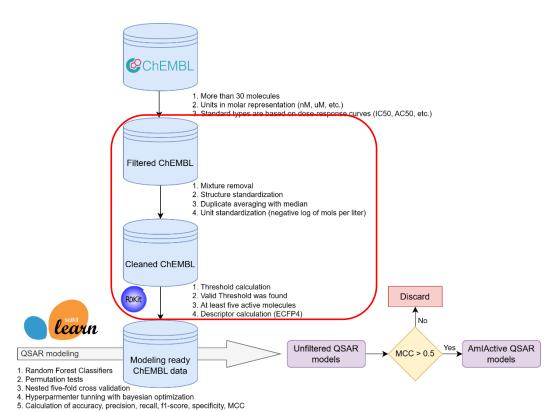




Materials and methods



- Mixtures were removed
- Structures were standardized
- Duplicate activities where averaged (median)
- Activity values where converted to a negative logarithm of moles per liter form
- Classifier activity thresholds were stablished
- Extended connectivity fingerprints (ECFP4)

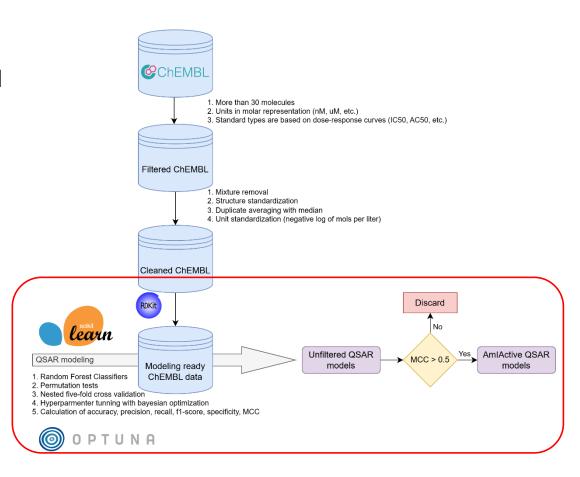








- Random forest algorithm
- Hyperparemeter tunning with bayesian optimization and nested five-fold cross-validation
- Calculated classifier metrics:
 - Acurracy
 - Precision
 - Recall
 - F1-Score
 - Specificity
 - Matthews Correlation Coefficient (MCC)
- Models that had a MCC > 0.5 were selected as the models of the system





- After all the filtering and cleaning procedures, we obtained 3239 QSAR models.
- Of these 3239 models, 2277 corresponded to distinct targets of interest.
- The target types distribution was:
 - Single proteins (≈69%)
 - Cell lines (≈15%)
 - Organisms (≈6%)
 - Protein complexes (≈4%)
 - Protein Families (≈4%)
 - Tissues (≈0.6%)



	Threshold	Accuracy	F1-score	мсс	Precision	Recall	Specificity
Cell-line	5.40	0.82	0.81	0.65	0.84	0.81	0.83
Organism	5.47	0.82	0.81	0.65	0.83	0.81	0.82
Protein Complex	6.64	0.82	0.82	0.65	0.82	0.83	0.80
Protein Family	6.25	0.82	0.82	0.65	0.83	0.83	0.81
Single Protein	6.43	0.82	0.82	0.65	0.83	0.83	0.81
Tissue	5.49	0.82	0.82	0.65	0.83	0.82	0.82





https://amiactive.ccen.ufpb.br/



AmlActive

AmlActive is a target fishing web tool that allows the scientific community to screen thousands of biological activities by clicking a single button.



Use the tool



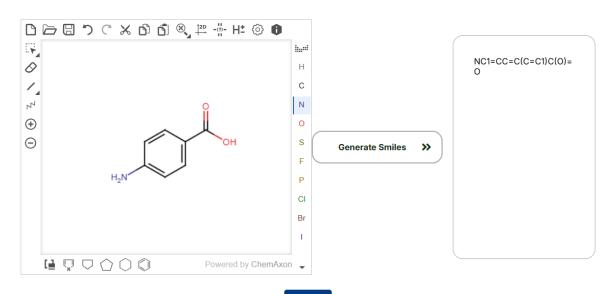






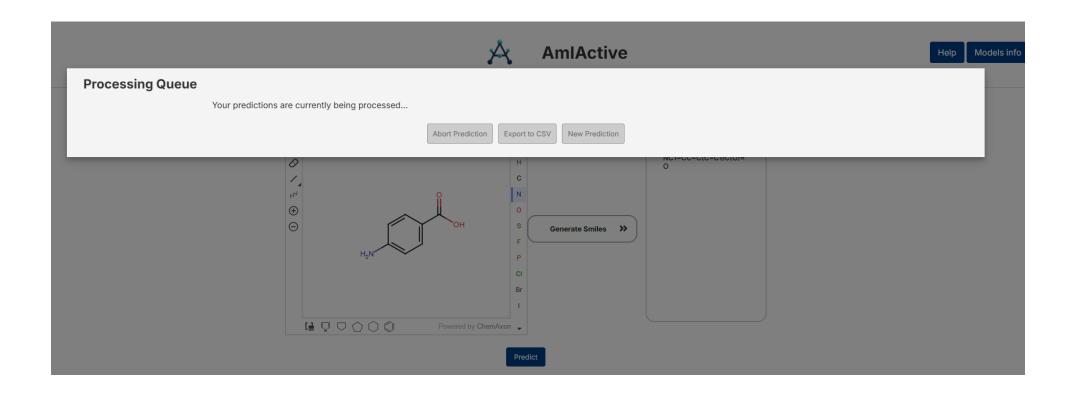


Help Models info

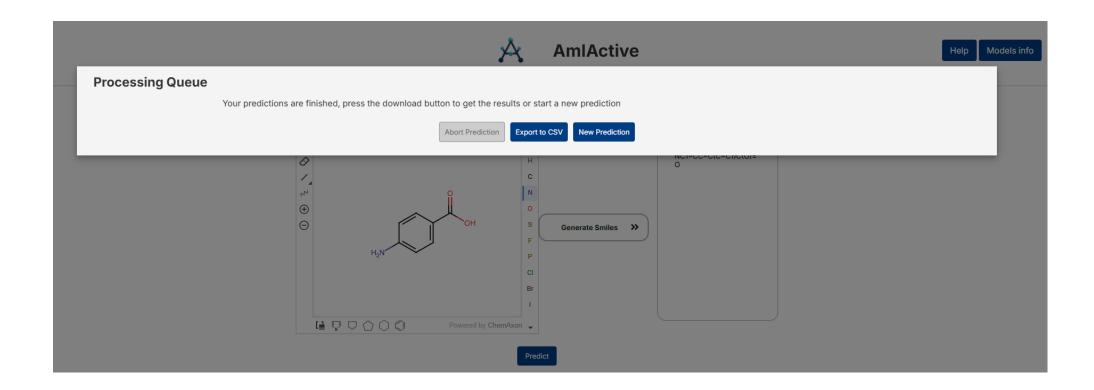


Predict











Conclusion

- AmlActive (AIA) is a brazilian QSAR based reverse screening webtool
- The AIA system has 3,239 models which correspond to 2,277 distinct targets
- The AIA system targets are either: Single proteins, Protein families, Protein complexes, organisms, cell-lines or tissues
- The AIA system can be accessed at: https://amiactive.ccen.ufpb.br/

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Thank you!

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