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DADNP: Dual Antibacterial Drug-Nanoparticle Systems Machine learning Approach

Karel Diéguez-Santana ^{a,b}

^a Department of Organic and Inorganic Chemistry, University of Basque Country UPV/EHU, 48940 Leioa, Spain

^b Departamento Ciencias de la Vida, Universidad Estatal Amazónica, Paso Lateral Vía Napo, km 2.5, 160150, Puyo, Pastaza, Ecuador

Graphical Abstract



Abstract.

The rise of new infectious diseases, combined with an increase in antibiotic resistance among bacterial pathogens, poses a significant health danger to humans. This ever-increasing threat of bacterial resistance necessitates the development of novel techniques to overcome this barrier. One of the successful strategies for combating antibiotic resistance has been the conjugation of nanoparticles (NPs) with antimicrobial moieties such as antibiotics, peptides, or other biomolecules. However, Dual Antibacterial Drug-Nanoparticle (DADNP) discovery is a slow

process due to the high number of combinations of NP vs. AD compounds, assays, etc. Artificial Intelligence/Machine Learning (AI/ML) algorithms may speed up it if they predict which putative DADNP systems should be short listed for assay. Nevertheless, the low amount of DADNP activity indicates that AI/ML analysis is tough. To solve this problem in an additive manner, the IFPTML = Information Fusion (IF) + Perturbation-Theory (PT) + Machine Learning (ML) technique was applied. Two datasets were combined (>165000 ChEMBL AD experiments with 300 NP assays) against multiple bacteria species. Eleven non-linear ML algorithms were developed using the Waikato Environment for Knowledge Analysis (WEKA) and STATISTICA. The analysis of the values of all the IFPTML models (Training/Validation Series) presents good performances (Accuracy global of 88.8-98.3%), Similarly, AUROC values are high (92-99%) in most cases. In the analysis and comparison of the algorithms used, ANN, RF, and KNN models stand out as having the highest $Sn \approx Sp \approx 88.5\% - 99.0\%$ and $AUROC \approx 0.94 - 0.99$ in both series. These results suggest that the IFPTML models may become a useful tool in the design of DADNP systems for antibacterial therapy against multidrug-resistant microbiological infections.

Introduction

The main bibliographic sources used in this document are listed below [1-15].

References

1. Diéguez-Santana, K.; Casañola-Martin, G.M.; Green, J.R.; Rasulev, B.; González-Díaz, H. Predicting Metabolic Reaction Networks with Perturbation-Theory Machine Learning (PTML) Models. *Current Topics in Medicinal Chemistry* **2021**, *21*, 819-827, doi:10.2174/1568026621666210331161144.
2. Diéguez-Santana, K.; González-Díaz, H. Towards machine learning discovery of dual antibacterial drug–nanoparticle systems. *Nanoscale* **2021**, doi:10.1039/d1nr04178a.
3. Diéguez-Santana, K.; Rivera-Borroto, O.M.; Puris, A.; Pham-The, H.; Le-Thi-Thu, H.; Rasulev, B.; Casañola-Martin, G.M. Beyond Model Interpretability using LDA and Decision Trees for α -Amylase and α -Glucosidase Inhibitor Classification Studies. *Chemical Biology & Drug Design* **2019**, doi:10.1111/cbdd.13518.

4. Hanczar, B.; Hua, J.; Sima, C.; Weinstein, J.; Bittner, M.; Dougherty, E.R. Small-sample precision of ROC-related estimates. *Bioinformatics* **2010**, *26*, 822-830, doi:10.1093/bioinformatics/btq037.
5. Hill, T.; Lewicki, P. *Statistics: Methods and Applications*, 1st edition ed.; StatSoft, Inc.: 2005; p. 800.
6. Jelinkova, P.; Mazumdar, A.; Sur, V.P.; Kociova, S.; Dolezelikova, K.; Jimenez, A.M.J.; Koudelkova, Z.; Mishra, P.K.; Smerkova, K.; Heger, Z.; et al. Nanoparticle-drug conjugates treating bacterial infections. *Journal of Controlled Release* **2019**, *307*, 166-185, doi:10.1016/j.jconrel.2019.06.013.
7. Kleandrova, V.V.; Luan, F.; Gonzalez-Diaz, H.; Russo, J.M.; Speck-Planche, A.; Cordeiro, M.N. Computational tool for risk assessment of nanomaterials: novel QSTR-perturbation model for simultaneous prediction of ecotoxicity and cytotoxicity of uncoated and coated nanoparticles under multiple experimental conditions. *Environ Sci Technol* **2014**, *48*, 14686-14694, doi:10.1021/es503861x.
8. Kollef, M.H.; Golan, Y.; Micek, S.T.; Shorr, A.F.; Restrepo, M.I. Appraising contemporary strategies to combat multidrug resistant gram-negative bacterial infections--proceedings and data from the Gram-Negative Resistance Summit. *Clin Infect Dis* **2011**, *53 Suppl 2*, S33-55; quiz S56-38, doi:10.1093/cid/cir475.
9. Luan, F.; Kleandrova, V.V.; González-Díaz, H.; Russo, J.M.; Melo, A.; Speck-Planche, A.; Cordeiro, M.N.D.S. Computer-aided nanotoxicology: Assessing cytotoxicity of nanoparticles under diverse experimental conditions by using a novel QSTR-perturbation approach. *Nanoscale* **2014**, *6*, 10623-10630, doi:10.1039/c4nr01285b.
10. Nocedo-Mena, D.; Cornelio, C.; Camacho-Corona, M.D.R.; Garza-Gonzalez, E.; Waksman de Torres, N.; Arrasate, S.; Sotomayor, N.; Lete, E.; Gonzalez-Diaz, H. Modeling Antibacterial Activity with Machine Learning and Fusion of Chemical Structure Information with Microorganism Metabolic Networks. *J Chem Inf Model* **2019**, *59*, 1109-1120, doi:10.1021/acs.jcim.9b00034.
11. Puzyn, T.; Rasulev, B.; Gajewicz, A.; Hu, X.; Dasari, T.P.; Michalkova, A.; Hwang, H.M.; Toropov, A.; Leszczynska, D.; Leszczynski, J. Using nano-QSAR to predict the cytotoxicity of metal oxide nanoparticles. *Nat Nanotechnol* **2011**, *6*, 175-178, doi:10.1038/nnano.2011.10.
12. Santana, R.; Zuluaga, R.; Ganan, P.; Arrasate, S.; Onieva, E.; Montemore, M.M.; Gonzalez-Diaz, H. PTML Model for Selection of Nanoparticles, Anticancer Drugs, and Vitamins in the Design of Drug-Vitamin Nanoparticle Release Systems for Cancer Cotherapy. *Mol Pharm* **2020**, *17*, 2612-2627, doi:10.1021/acs.molpharmaceut.0c00308.
13. Santana, R.; Zuluaga, R.; Gañán, P.; Arrasate, S.; Onieva, E.; González-Díaz, H. Designing nanoparticle release systems for drug-vitamin cancer co-therapy with multiplicative perturbation-theory machine learning (PTML) models. *Nanoscale* **2019**, *11*, 21811-21823, doi:10.1039/c9nr05070a.
14. Speck-Planche, A.; Kleandrova, V.V.; Luan, F.; Cordeiro, M.N. Computational modeling in nanomedicine: prediction of multiple antibacterial profiles of nanoparticles using a quantitative structure-activity relationship perturbation model. *Nanomedicine (Lond)* **2015**, *10*, 193-204, doi:10.2217/nnm.14.96.
15. Urista, D.V.; Carrue, D.B.; Otero, I.; Arrasate, S.; Quevedo-Tumailli, V.F.; Gestal, M.; Gonzalez-Diaz, H.; Munteanu, C.R. Prediction of Antimalarial Drug-Decorated Nanoparticle Delivery Systems with Random Forest Models. *Biology (Basel)* **2020**, *9*, doi:10.3390/biology9080198.