

CURRICULUM VITAE

PERSONAL INFORMATION

Name: Antonio
Surname: de la Vega de León
Birth Date: 18 September 1986
ID Number: 53615149-X
Address: Clemensstrasse 57, 53225 Bonn, NRW, Germany
Telephone Number: +49-176-47951874
Email: antoniovega86@gmail.com / vega@bit.uni-bonn.de

ACADEMIC FORMATION

- 2010 - 2012:** M. Sc. Life Science Informatics from the Bonn-Aachen International Center for Information Technology, part of the Rheinische Friedrich-Wilhelms-Universität Bonn, Germany. Grade: 1.2 ($\approx 9.5/10$)
- 2004 - 2010:** *Licenciado* in Biology from the *Universidad Complutense de Madrid*, Spain. Grade: 8.67/10
- 2002 - 2004:** *Bachillerato* in Life Science from *Colegio Everest*, Madrid, Spain. Grade: *Matrícula de Honor* (9.1/10)

PROFESSIONAL EXPERIENCE

- November 2016 – Present:** Postdoctoral research assistant (Marie-Curie fellow) at the group of Prof. Gillet (University of Sheffield) for the European project Diagnostic and Drug Discovery Initiative for Alzheimer's Disease (FP7-PEOPLE-2013-IAPP)
- November 2012 – October 2016:** Research assistant on the workgroup from Prof. Bajorath (University of Bonn), doing my PhD Thesis on the topic of applications of MMPs and visualization techniques for computational compound optimization
- May 2012 – November 2012:** Student assistant (SHK) on the workgroup from Prof. Bajorath (University of Bonn), developing a new multi-target activity landscape as the Master Thesis for the Life Science Informatics Masters
- October 2011 – February 2012:** Student assistant (SHK) in the University of Bonn, tutoring the exercise part of the lecture Foundations of Information Management WS 11/12 for the Life Science Informatics Masters

October 2009 - May 2010: Assistant on a bioinformatics project assembling and annotating the genomes of *Rhodococcus ruber* and *Gordonia cholesteroivorans* under supervision of Prof. Julian Perera, Biochemistry and Molecular Biology department of the Universidad Complutense de Madrid

July - September 2009: Internship on the molecular genetics laboratory for diagnostic GENYCA INNOVA.

October 2008 - Februar 2009: Assistant of Professor Tomás Naranjo Pompa on the cytogenetic laboratory of the Universidad Complutense de Madrid

October 2006 - June 2007: Small experiment on the population genetics laboratory of the University

September 2006: Laboratory assistant in Barcelona on the company Ecofloat S.A.

July - August 2006: Costumer service of Iberia Equipajes S.A., working on phone service.

JOURNAL PUBLICATIONS

Anighoro A, **de la Vega de León A** & Bajorath J. Predicting bioactive conformations and binding modes of macrocycles. J Comput-Aided Mol Des, in press. [dx.doi.org/10.1007/s10822-016-9973-5](https://doi.org/10.1007/s10822-016-9973-5)

de la Vega de León A & Bajorath J. Chemical space visualization: transforming multi-dimensional chemical spaces into similarity-based molecular networks. Future Med Chem 8, 1769-1778, 2016. [dx.doi.org/10.4155/fmc-2016-0023](https://doi.org/10.4155/fmc-2016-0023)

Horvath D, Marcou G, Varnek A, Kayastha S, **de la Vega de León A** & Bajorath J. Prediction of activity cliffs using condensed graphs of reaction representations, descriptor recombination, support vector machine classification, and support vector regression. J Chem Inf Model, in press. [dx.doi.org/10.1021/acs.jcim.6b00359](https://doi.org/10.1021/acs.jcim.6b00359)

Shanmugasundaram V, Zhang L, Kayastha S, **de la Vega de León A**, Dimova D & Bajorath J. Monitoring the progression of structure-activity relationship information during lead optimization. J Med Chem 59, 4235-4244, 2016. [dx.doi.org/10.1021/acs.jmedchem.5b01428](https://doi.org/10.1021/acs.jmedchem.5b01428)

de la Vega de León A, Kayastha S, Dimova D, Schultz T & Bajorath J. Visualization of multi-property landscapes for compound selection and optimization. J Comput-Aided Mol Des 29, 695-705, 2015. [dx.doi.org/10.1007/s10822-015-9862-3](https://doi.org/10.1007/s10822-015-9862-3)

Hameed A, Khan K, Zehra S, Ahmed R, Shafiq Z, Bakht S, Yaqub M, Hussain M, **de la Vega de León A**, Furtmann N, Bajorath J, Ahmad H, Tahir M & Iqbal J. Synthesis, biological evaluation and molecular docking of N-phenyl thiosemicarbazones as urease inhibitors. Bioorg Chem 61, 51-57, 2015. [dx.doi.org/10.1016/j.bioorg.2015.06.004](https://doi.org/10.1016/j.bioorg.2015.06.004)

Kayastha S, **de la Vega de León A**, Dimova D & Bajorath J. Target-based analysis of ionization states of bioactive compounds. Med Chem Commun 6, 1030-1035, 2015. [dx.doi.org/10.1039/C5MD00051C](https://doi.org/10.1039/C5MD00051C)

- de la Vega de León A** & Bajorath J. Prediction of compound potency changes in matched molecular pairs using support vector regression. *J Chem Inf Model* 54, 2654-2663, 2014. [dx.doi.org/10.1021/ci5003944](https://doi.org/10.1021/ci5003944)
- de la Vega de León A**, Hu Y & Bajorath J. Systematic identification of matching molecular series and mapping of screening hits. *Mol Inf* 33, 257-263, 2014. [dx.doi.org/10.1002/minf.201400017](https://doi.org/10.1002/minf.201400017)
- Stumpfe D, **de la Vega de León A**, Dimova D & Bajorath J. Advancing the activity cliff concept, part II [v1; ref status: indexed, f1000r.es/34p] *F1000Research* 3:75, 2014. [dx.doi.org/10.12688/f1000research.4057](https://doi.org/10.12688/f1000research.4057)
- de la Vega de León A** & Bajorath J. Formation of activity cliffs is accompanied by systematic increases in ligand efficiency from lowly to highly potent compounds. *AAPS J* 16, 335-341, 2014. [dx.doi.org/10.1208/s12248-014-9567-x](https://doi.org/10.1208/s12248-014-9567-x)
- Hu Y, **de la Vega de León A**, Zhang & Bajorath J. Matched molecular pair-based data sets for computer-aided medicinal chemistry [v2; ref status: indexed, f1000r.es/2w9] *F1000Research* 3:36, 2014. [dx.doi.org/10.12688/f1000research.3-36.v2](https://doi.org/10.12688/f1000research.3-36.v2)
- de la Vega de León A** & Bajorath J. Matched molecular pairs derived by retrosynthetic fragmentation. *Med Chem Commun* 5, 64-67, 2014. [dx.doi.org/10.1039/C3MD00259D](https://doi.org/10.1039/C3MD00259D)
- Fernandez de las Heras L, Alonso S, **de la Vega de León A**, Xavier D, Perera J & Navarro Llorens JM. Draft genome sequence of the steroid degrader *Rhodococcus ruber* Strain Chol-4. *Genome Announc* 1:e00215-13, 2013. [dx.doi.org/10.1128/genomeA.00215-13](https://doi.org/10.1128/genomeA.00215-13)
- de la Vega de León A** & Bajorath J. Compound optimization through data set-dependent chemical transformations. *J Chem Inf Model* 53, 1263-1271, 2013. [dx.doi.org/10.1021/ci400165a](https://doi.org/10.1021/ci400165a)
- de la Vega de León A** & Bajorath J. Design of a three-dimensional multi-target activity landscape. *J Chem Inf Model* 52, 2876-2883, 2012. [dx.doi.org/10.1021/ci300444p](https://doi.org/10.1021/ci300444p)

OTHER SCIENTIFIC WORK

- Shanmugasundaram V, Liying Z, Kayastha S, **de la Vega de León A**, Dimova D & Bajorath J. Data sets for SAR progression analysis. Freely available data set. [dx.doi.org/10.5281/zenodo.32794](https://doi.org/10.5281/zenodo.32794)
- de la Vega de León A**, Kayastha S, Dimova D, Schultz T & Bajorath J. ChEMBL20 data sets for multi-property landscape analysis. Freely available data set. [dx.doi.org/10.5281/zenodo.21782](https://doi.org/10.5281/zenodo.21782)
- Hu Y, **de la Vega de León A**, Zhang B & Bajorath J. Detailed data sets of MMP-cliffs, SAR transfer series, RECAP-MMPs and compound activities. Freely available data set. [dx.doi.org/10.5281/zenodo.8418](https://doi.org/10.5281/zenodo.8418)
- Kayastha S, **de la Vega de León A**, Dimova D, Schultz T & Bajorath J. Visualization of multi-property landscapes for compound selection and optimization. Poster at Chemoinformatics Strasbourg Summer School 2016
- Dimova D, Kayastha S, **de la Vega de León A** & Bajorath J. Monitoring the progression of structure-activity relationship information during lead optimization. Poster at 2016 Frontiers in Medicinal Chemistry

Zhang L, Starr J, Dimova D, Iyer P, Gupta-Ostermann D, **de la Vega de León A** & Bajorath J, Shanmugasundaram V. Novel applications of SAR matrices in pharmaceutical research. Poster at the Spring 2014 Dallas ACS National Meeting

Müller G, Benningshof J, van Meurs P, Stumpfe D, **de la Vega de León A**, Furtmann N, Dimova D & Bajorath J. Synthetic and cheminformatic exploration of macrocyclic and peptidomimetic medicinal chemistry space. Poster at XXIII International Symposium on Medicinal Chemistry (EFMC-ISMIC 2014)

de la Vega de León A & Bajorath J. Compound optimization through data set-dependent chemical transformations. Poster at 9th German Conference on Chemoinformatics (GCC 2013)

de la Vega de León A, Lounkine E, Vogt M & Bajorath J. Design of diverse and focused compound libraries. In: Tutorials in Cheminformatics. Eds: Varnek A. (in preparation)

Vogt M, **de la Vega de León A** & Bajorath J. Algorithmic chemoinformatics. In: Tutorials in Cheminformatics. Eds: Varnek A. (in preparation)

TEACHING EXPERIENCE

SS2016: Assistant in the Python-based Programming Lab Course II

WS2015/16: Two theoretical lectures for Chemoinformatics; assistant in the exercises of Chemoinformatics and Structural Bioinformatics lectures

SS2015: Assistant in the Python-based Programming Lab Course II

WS2014/15: Two theoretical lectures for Bridging Course Chemistry; assistant in the Research Practical invitational lab course

SS2014: Three theoretical lectures for Bioinformatics II; assistant in the Python-based Lab Course II

WS2013/14: One theoretical lecture for Bridging Course Chemistry; assistant in the Java-based Chemoinformatics Lab Course

SS2013: Assistant in the MOE-based Molecular Modeling and Drug Design Lab Course

WS2012/13: Assistant in the Java-based Chemoinformatics Lab Course

WS2011/12: Assistant in the Foundations of Information Management Exercises

LANGUAGES

Spanish: Native level

English: High level
Certificate in Advanced English (CAE) from Cambridge University
Internet based TOEFL test (iBT) score: 107
12 years of studies in a bilingual school

German: Medium/high level
Zentrale Mittelstufenprüfung (ZMP) Certificate from Goethe Institut
Advanced level from the *Escuela Oficial de Idiomas*
More than six years living in Germany

SKILLS

Programming: Java, Python, R

Scientific Software: MOE, AutoDock, KNIME

Computer Software: Microsoft Office, Inkscape, Latex