

# Alessio Valentini, Ph.D.

I am a passionate researcher who lives at the interface between computer science and computational chemistry. I built my career delivering high quality software to solve chemistry problems. I am an eager learner who enjoys new challenges and tinkers with all things computer science.

## Work Experience:

- **Research Associate/PostDoc at Stanford University.** Project leader and main developer of the Retropaths package, a new software that automates exploration of chemical synthesis channels using graph theory, funded by a multimillion MURI DOD ONR grant (*California 2020-present*).
- **PostDoc at University of Liège.** Project leader and main developer of a package to perform and analyze quantum dynamics for attosecond time-scale events (*Belgium 2016-2019*).
- **PostDoc at University of Siena.** Developed tools in the OpenMolcas quantum chemistry suite for semiclassical dynamics and automatic generation of hybrid quantum mechanics / molecular mechanics protein models (*Italy 2015-2016*).

## Skills:

- Developed several Python (**11** years) and Haskell (**3** years) packages for molecular dynamics, reaction discovery, data analysis and data wrangling pipelines.
- Proficiency in developing high performance codes in collaborative settings, as proven by my work in the OpenMolcas suite and in the Martínez group at Stanford University.
- Installed and managed two university clusters for scientific calculations (*2009-2019*).
- Effective communication skills, with a strong passion for visual effects and 3D rendering, as proven by my 8 presentations and 2 awards at international scientific conferences.

## Career achievements:

- Published 22 articles with more than 2400 total citations in peer-reviewed journals in the fields of photochemistry, photobiology, reaction discovery and chemical pedagogy tools.

## Education:

- **Ph.D. in Computational Chemistry** on *Semiclassical molecular dynamics*. University of Alcalá, (*Spain 2011-2015*).
- **M.Sc. in Chemistry** on *Automatic generation of QM/MM models for photoactive proteins*. University of Siena, (*Italy 2008-2011*).

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## Python libraries:

Cython  
NetworkX  
Numpy/Scikit  
Pandas  
Matplotlib  
Rdkit/OpenBabel  
Mkdocs

## Secondary languages:

CUDA  
C/C++  
Fortran  
React/Javascript

## Daily tools:

Bash  
VScode  
JupyterLab  
Docker  
CI/CD

## Misc:

Blender  
Gimp  
Inkscape  
Latex  
d3.js

## Real life languages:

Italian  
English  
Spanish

