# Charles Tapley Hoyt, Ph.D.

### Interests

Systems and Networks Biology, Cheminformatics, Machine Learning, Knowledge Graphs, Network Representation Learning, Proteochemometrics, Target Prioritization, Drug Repositioning, Pathway Analysis

## Work

- 2020- Computational Biologist, Enveda Therapeutics, Bonn, Germany.
- 2018–19 Lecturer, University of Bonn, Bonn, Germany.
- 2016–19 **Research Fellow**, *Fraunhofer SCAI*, Sankt Augustin, Germany.
- 2012–15 **Teaching Assistant**, *Northeastern University*, Boston, MA, USA.
  - 2014 in Silico Lead Discovery Co-op, Novartis, Cambridge, MA, USA.
- 2013-14 Molecular Informatics Internship, Pfizer, Cambridge, MA, USA.
  - 2013 **Post-Selection Chemistry Co-op**, *GlaxoSmithKline*, Waltham, MA, USA.
  - 2012 Research Assistant, Pollastri Laboratory, Northeastern University, Boston, MA, USA.

#### Education

- 2018–19 **Doctor of Philosophy**, Computational Life Sciences, University of Bonn, Germany.
- 2015–17 Master of Science, Life Science Informatics, University of Bonn, Germany.
- 2011–15 Bachelor of Science, Chemistry, Northeastern University, USA.

#### Affiliations

- 2020- CoronaWhy
- 2017- OpenBEL Consortium
- 2016- Erasmus Student Network
- 2011 American Chemical Society

## Spoken Languages

English (Native)

German (Limited working proficiency)

## Programming Languages

Python, Bash, Fish, R, Java, SQL, SPARQL, Cypher, Javascript, HTML, CSS, XPath

# Projects

PyBEL, An ecosystem for biological knowledge graphs in BEL.

https://github.com/pybel

BEL Commons, Interactive exploration and analysis of biological knowledge graphs.

https://github.com/bel-commons

**PyKEEN**, Learning, evaluation and applications of knowledge graph embeddings.

https://github.com/pykeen/pykeen/