# 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

- 2021 Postdoctoral Research Fellow, Harvard Medical School, Boston, MA, USA (remote).
  - 2020 Computational Biologist, Enveda Biosciences, Boulder, CO, USA (remote).
- 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

- 2021 International Society of Biocuration
- 2017- OpenBEL Consortium
- 2011 American Chemical Society
- 2020 CoronaWhy

## Spoken Languages

English (Native), German (Limited working proficiency)

## Programming Languages

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

### Projects

**Bioregistry**, An integrative registry of biological databases, ontologies, and nomenclatures. https://github.com/biopragmatics/bioregistry

**PyBEL**, An ecosystem for biological knowledge graphs in BEL.

https://github.com/pybel

**PyKEEN**, Learning, evaluation and applications of knowledge graph embeddings. https://github.com/pykeen/pykeen/