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, San Francisco, CA, 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

- 2017- OpenBEL Consortium
- 2016- Erasmus Student Network
- 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

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/