

Publications

1. **Hoyt, C. T.**, Konotopez, A., & Ebeling, C. (2017). PyBEL: a computational framework for Biological Expression Language. *Bioinformatics (Oxford, England)*, 34(4), 703–704.
2. **Hoyt, C. T.**, Domingo-Fernández, D., & Hofmann-Apitius, M. (2018). BEL Commons: an environment for exploration and analysis of networks encoded in Biological Expression Language. *Database*, 2018(3), 1–11.
3. **Hoyt, C. T.**, *et al.* (2019). Re-curation and rational enrichment of knowledge graphs in Biological Expression Language. *Database*, 2019, baz068.
4. **Hoyt, C. T.**, *et al.* (2019). Bio2BEL: Integration of Structured Knowledge Sources with Biological Expression Language. *BMC Bioinformatics*, submitted.
5. Gündel, M., **Hoyt, C. T.**, & Hofmann-Apitius, M. (2018). BEL2ABM: Agent-based simulation of static models in Biological Expression Language. *Bioinformatics*, 34(13), 2316–2318.
6. Muslu, Ö., **Hoyt, C. T.**, Hofmann-Apitius, M., & Fröhlich, H. (2019). Guilty-Targets: Prioritization of Novel Therapeutic Targets with Deep Network Representation Learning. *IEEE/ACM Trans. Comput. Biol. Bioinform.*, submitted.
7. **Hoyt, C. T.**, *et al.* (2018). A systematic approach for identifying shared mechanisms in epilepsy and its comorbidities. *Database*, 2018(1).