References Cited

- [1] Davoud Torkamaneh, Jérôme Laroche, and François Belzile. "Genome-Wide SNP Calling from Genotyping by Sequencing (GBS) Data: A Comparison of Seven Pipelines and Two Sequencing Technologies". In: PLOS ONE 11.8 (Aug. 2016), pp. 1-14. DOI: 10.1371/journal.pone.0161333.
- [2] Corinne Watts et al. "DNA metabarcoding as a tool for invertebrate community monitoring: a case study comparison with conventional techniques". In: Austral Entomology 58.3 (2019), pp. 675-686. Doi: 10.1111/aen.12384.
- [3] Carla Hurt et al. "First Worldwide Molecular Phylogeny of the Morphologically and Ecologically Hyperdiversified Snapping Shrimp Genus Alpheus (Malacostraca: Decapoda)". In: Molecular Phylogenetics and Evolution (in press).
- [4] Ramesh Paudel, Timothy Muncy, and William Eberle. "Detecting DoS Attack in Smart Home IoT Devices Using a Graph-Based Approach". In: 2019 IEEE International Conference on Big Data (Big Data). 2019, pp. 5249-5258. DOI: 10.1109/BigData47090.2019. 9006156.
- [5] Ramesh Paudel, Peter Harlan, and William Eberle. "Detecting the Onset of a Network Layer DoS Attack with a Graph-Based Approach". In: International Conference of the Florida AI Research Society (FLAIRS). May 2019, pp. 38-43.
- [6] Ramesh Paudel et al. "Cognitive Health Prediction on the Elderly Using Sensor Data in Smart Homes". In: International Conference of the Florida AI Research Society (FLAIRS). May 2018, pp. 317-322.
- [7] Lenin Mookiah, Chris Dean, and William Eberle. "Graph-Based Anomaly Detection on Smart Grid Data". In: International Conference of the Florida AI Research Society (FLAIRS). May 2017, pp. 306-311.