

## 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.