



















- NR Wheeler, P Benchek, BW Kunkle, KL Hamilton-Nelson, M Warfe, JR Fondran, JL Haines, WS Bush (2019). “Big data solutions for reproducibility and scalability of genomic sequencing studies.” *Pac Symp Biocomput.* To appear.
- Y Yao, S Ramsey (2019). “CERENKOV3: Clustering and molecular network-derived features improve computational prediction of functional noncoding SNPs.” *Pac Symp Biocomput.* To appear.