

Virtual Workshop: Variant Detection



January 27-29, 2026
10:00 am - 2:00 pm EST

Ready to dive into the world of variant detection? Join our virtual workshop, where we cover Linux and high-performance computing basics, high-throughput sequencing, reference alignment and quality control, variant calling models and pipelines (GATK, Freebayes, BCFtools), and variant call filtering, comparison, and annotation. Designed for beginners; no prior bioinformatics experience required.



Cost: \$500 (\$400 for UConn affiliates)

Where: Live Sessions via MS Teams | Recordings distributed

Register now: <https://bioinformatics.uconn.edu/cbc-workshops/>

Questions? Email cbcsupport@helpspotmail.com



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