# Virtual Workshop: Methylation Analysis

## Nov. 17-19, 2026
## 10:00 am - 2:00 pm EST

Ready to examine epigenetic modifications?
Join our Virtual Methylation Analysis workshop to gain hands-on experience analyzing conversion-based (e.g., EM-Seq) and direct detection-based (e.g., Oxford Nanopore) methylation data. You will learn how to generate methylation calls, map reads, assess data quality and perform quality control, quantify methylation levels, and perform differential methylation analyses using real experimental datasets on a university HPC. Perfect for researchers of all levels - no prior bioinformatics experience required!

**Cost**: $630 ($525 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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