



Please find information regarding ENCODE representation at the ASHG 2023 Annual Symposium below. The ASHG 2023 Annual Symposium will be held at the Walter E. Washington Convention Center in Washington, D.C. and runs from November 1st to November 5th. All times below, reflect US – Eastern Standard Time

Wednesday, November 1, 2023

Time	Location	Presenting Author	Presenting Author's Site	Title of Presentation	Type
6:30pm	Ballroom ABC Level 3	Aman Patel	Stanford University	Reconstructing the cis-regulatory landscape of archaic hominids using deep learning	Plenary Talk

Thursday, November 2, 2023

Time	Location	Presenting Author	Presenting Author's Site	Title of Presentation	Type
3:00pm – 5:00pm	Exhibit & Poster Hall AB	Yunzhe Jiang	Yale University	Epigenetic characterization of pseudogenes across human tissues	Poster (#PB2261)

Friday, November 3, 2023

Time	Location	Presenting Author	Presenting Author's Site	Title of Presentation	Type
8:30am – 10:00am	Ballroom C Level 3	Tim Reddy, Carolyn Hutter, Barbara Wold, Jill Moore, Kushal Dey, Martin Hirst	Multiple	The ENCODE consortium: 2003 - 2023	Invited Session

3:00pm – 5:pm	Exhibit & Poster Hall AB	Kuei-Yueh Ko	Duke University	Functional characterization and network analysis of regulatory regions using genome-wide functional screens and chromatin interactions	Poster (#PB2274)
1:45pm	Room 146B Level 1	Sasha Boytsov	Altius Institute for Biomedical Sciences	Large-scale characterization of naturally occurring human regulatory genetic variation.	Platform Talk

Saturday, November 4, 2023

Time	Location	Presenting Author	Presenting Author's Site	Title of Presentation	Type
1:30pm	Ballroom C Level 3	Artem Kim	University of Southern California	Fine-mapping causal cell-types of human diseases and disease variants	Platform Talk
2:15pm – 4:15pm	Exhibit & Poster Hall AB	Giovanni Quinones-Valdez	University of California, Los Angeles	Long read RNA-sequencing demarcates cis and trans-directed alternative RNA splicing.	Poster (#PB3420)
2:15pm – 4:15pm	Exhibit & Poster Hall AB	Kristy Mualim	Stanford University	Genome-wide maps of regulatory enhancer-gene interactions across human cell types and tissues.	Poster (#PB4497)