



FaceBase2 Final Recap Webinar Agenda

Monday, May 20th

All times are Eastern Time Zone

Project talks: 20 min, Q&A 10 min	
11:00 AM	Opening Remarks – <i>NIDCR Staff</i>
11:05 AM	FB2 Hub Update - <i>Carl Kesselman, USC</i>
11:35 AM	Integrated Research of Functional Genomics and CR Morphogenesis <i>Yang Chai, USC</i>
12:05 PM	Rapid Identification and Validation of Human CR Developing Genes <i>Richard Maas, Eric Liao, Harvard</i>
12:35 PM	Anatomical Atlas, Transgenic Tools for Late Skull Formation in Zebrafish <i>Shannon Fisher, Boston Univ and Matthew Harris, Harvard</i>
1:05 PM	<i>Lunch – 30 minutes</i>
1:35 PM	Transcriptome Atlases of the Craniofacial Sutures <i>Ethylin Wang Jabs, Mt. Sinai</i>
2:05 PM	Genomic and Transgenic Resources for Craniofacial Enhancer Studies <i>Axel Visel, Berkeley</i>
2:35 PM	Human Genomics Analysis Interface for Facebase 2 <i>Mary Marazita, Univ Pittsburgh</i>
3:05 PM	<i>Break – 15 minutes</i>
3:20 PM	Epigenetic Landscapes and Regulatory Divergence of Human CR Traits <i>Joanna Wysocka, UCSF</i>
3:50 PM	RNA Dynamics in the Developing Mouse Face <i>Joan Hooper, UC Denver</i>
4:20 PM	Developing 3D Craniofacial Morphometry Data and Tools <i>Richard Spritz, UC Denver</i>
4:50 PM	Ontology of Craniofacial Development and Malformation <i>Jim Brinkley via slides/movie</i>
5:05 PM	Closing remarks – <i>Carl Kesselman, USC</i>
5:20 PM	Adjourn