The FaceBase Consortium

NIDCR Update

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A reminder of why we're here

- Facilitate craniofacial research by making large, integrated datasets available to the wider scientific community
- Develop the tools and datasets needed for a comprehensive, systems-wide, understanding of the developmental processes that create the face during embryogenesis and how those processes go awry.



FaceBase objectives

Data Collection and Technology Development: "-omics" datasets on midface development at the molecular, cellular, tissue and organism levels deposited into FaceBase database



Data Coordination, Integration and Sharing: Comparative analysis, gene regulatory networks, modeling



Expansion of FaceBase to new anatomic areas



Translational and clinical applications for prevention & intervention

Expanding FaceBase beyond the RFAs

- Today's webcast
 - This is our first webcast so please let the viewers out in television land know what types of data you'll be producing.
- Additional funding opportunities for:
 - Using FaceBase data
 - Software development through BD2K



Related FOA: PAR-13-178

- NIDCR Small Research Grants for Secondary Analysis of <u>FaceBase Data</u> (R03)
 - Intended to encourage "outside" use of the data that you have produced and to encourage analyses that tie together data from different FaceBase projects
 - Larger than normal R03 budget
 - Direct costs of up to \$200,000 per year are allowed with a maximum direct cost of \$300,000 for the entire two year funding period
 - Not intended for support of data generation
- Recipients of these grants are expected to share the results of their analyses through FaceBase



BD2K

- Funding announcements at bd2k.nih.gov
 - Including:
 - Software development and hardening
 - Development of analytical methods
 - Training
- The Data Discovery Index Coordination Consortium
 - http://biocaddie.org
- Software Discovery Index



Expanded NIH Genomic Data Sharing Policy

- NIH has expanded the scope of its "genomic" data sharing policy
 - All NIH-funded research generating <u>large-scale</u> human or nonhuman genomic data and the use of these data for subsequent research
 - Non-human data: Share through any widely used data repository
 - Human data: Share through an NIH-designated data repository
 - Examples of large-scale genomic data include, but are not limited to, GWAS, Single Nucleotide Polymorphism (SNP) arrays, and genome sequence, transcriptomic, epigenomic, and gene expression data
 - Applies to all funding mechanisms (grants, contracts, intramural support) with no minimum threshold for cost
 - Include genomic data sharing plan as part of application or proposal



Expanded NIH Genomic Data Sharing Policy

- Smaller-Scale Genomic Research Projects
 - Submissions may also be expected by the funding NIH Institute or Center (IC), based on:
 - State of the science
 - Programmatic priorities of the funding IC
 - Utility of the data for the research community
- More info at http://gds.nih.gov/index.html
 - See Supplemental Information for examples of large-scale genomic data
- Contact Emily Harris for NIDCR specific info at harrisel@nidcr.nih.gov



Introductions

- Susan Lowenthal grants management specialist for your FaceBase awards
 - The person to talk to about award/budget issues
- The "Scientific Leadership Group"
 - Helps the Consortium and NIDCR assess progress, solve problems, etc.
 - Mike Cherry, Stanford
 - Large data integration projects
 - Melissa Haendel, OHSU
 - Data curation and integration
 - Max Muenke, NHGRI/NIH
 - Genetics/genomics of normal and abnormal human development



