

3D Analysis of Normal Facial Variation: Data Repository and Genetics

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FaceBase Annual Meeting: May 1-3, 2013



Goal 1: To construct a web-based normative repository of 3D facial measurements and images

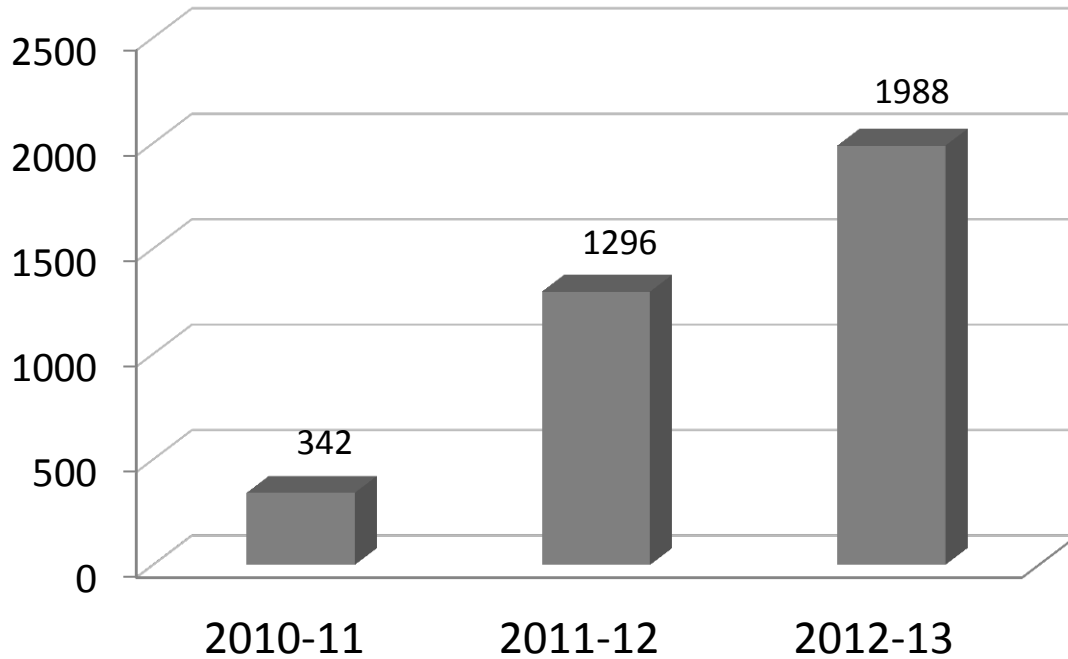
- 1a) Ascertain 3500 European Caucasian individuals – age 3 to 40 – from multiple sites: Pittsburgh, Seattle, and Houston (and Iowa!)
- 1b) Acquire 3D facial surface images, basic demographic descriptors and saliva samples from each participant
- 1c) Extract quantitative data from 3D surface images including landmarks and linear measurements
- 1d) Deposit clean data (including images and measures) into a web-based repository accessible through FaceBase.org (*3D Facial Norms*)

Goal 2: To investigate the genetic basis of variation in facial shape

- 2a) Submit saliva samples for genotyping (proposed year 5)
- 2b) Perform morphometric analyses on 3D data set to derive shape variation descriptors
- 2c) Identify genetic variants associated with normal facial shape variation
- 2d) Submit genotype data and results to FaceBase.org

Progress over the Past Year

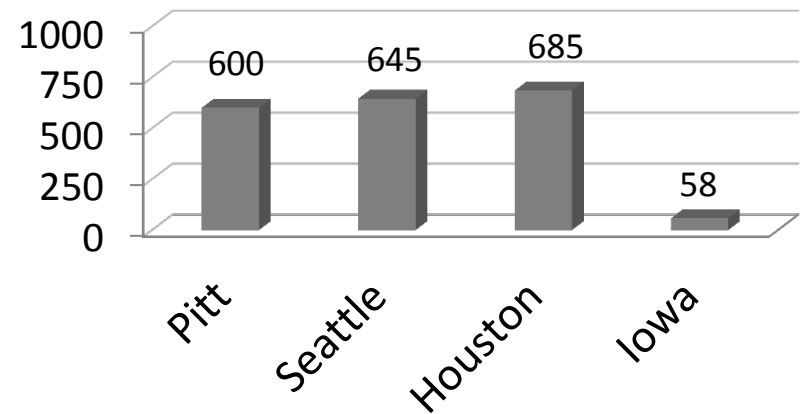
Overall Recruitment Effort



3D Facial Norms Repository

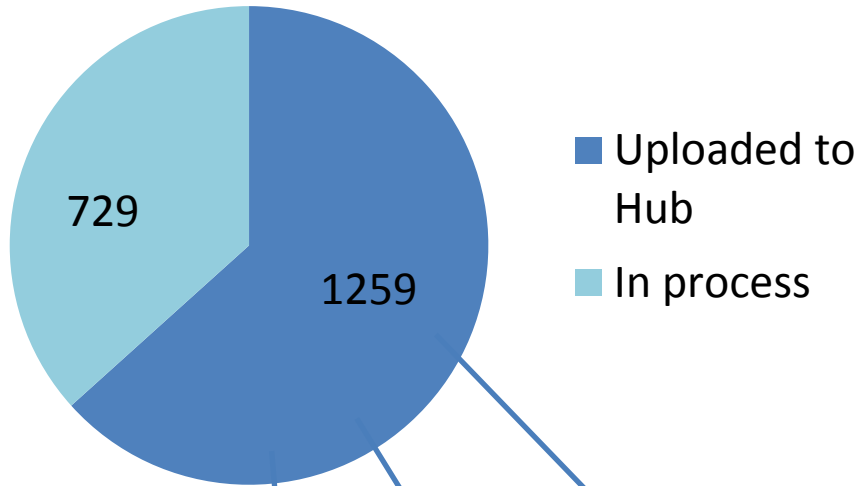
Recruitment Target:
3500 healthy unrelated
individuals age 3-40

Site Recruitment Figures



Progress over the Past Year

Phenotypic Data Collection

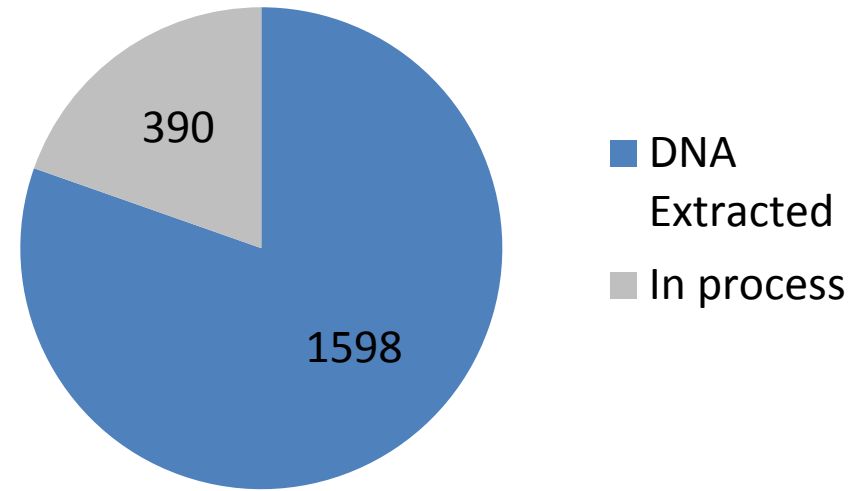


Demographics

3D Facial Surfaces

Landmark Coordinates and Measurements

Saliva Kits



3D Facial Norms Web-Interface



USERNAME OR E-MAIL * PASSWORD *

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DATA

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SEARCH

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YOU ARE HERE: [Home](#) » 3D Facial Norms Database

3D Facial Norms Database

Overview

[3D Database Search](#)

[Summary Statistics](#)

[Data Analysis](#)

[Tools](#)

[Technical Notes](#)



3D DATABASE SEARCH

Explore the 3D Norms Database through a customizable search interface. This is the main portal for querying and downloading individual-level phenotype and genotype data based on user-defined parameters. Phenotypic data include 3D facial landmark coordinates, anthropometric facial measurements, demographic descriptors, and 3D facial surface models.

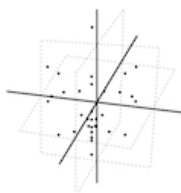
GO →



SUMMARY STATISTICS

View normative age- and sex-based averages for selected craniofacial measurements and 3D facial surface models.

GO →



DATA ANALYSIS

A compendium of results from past analyses based on the 3D Facial Norms dataset.

GO →



TOOLS

A collection of practical web applications for the clinical and research community.

GO →



TECHNICAL NOTES

Detailed background information on all aspects of the 3D Facial Norms Project.

GO →



ABSTRACT PAGE

GO →

 Analyses of human craniofacial variation and growth patterns

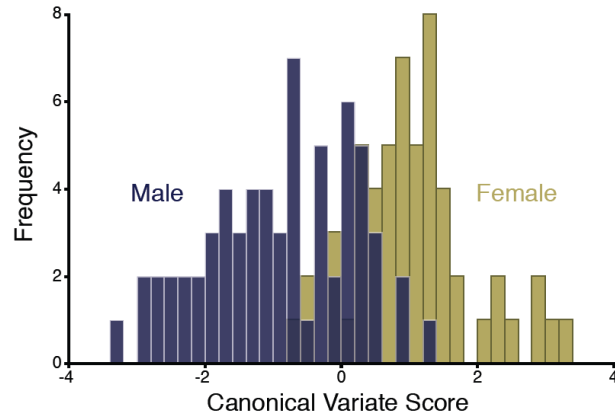
Normative control data for craniofacial comparisons

Development of novel image analysis methods

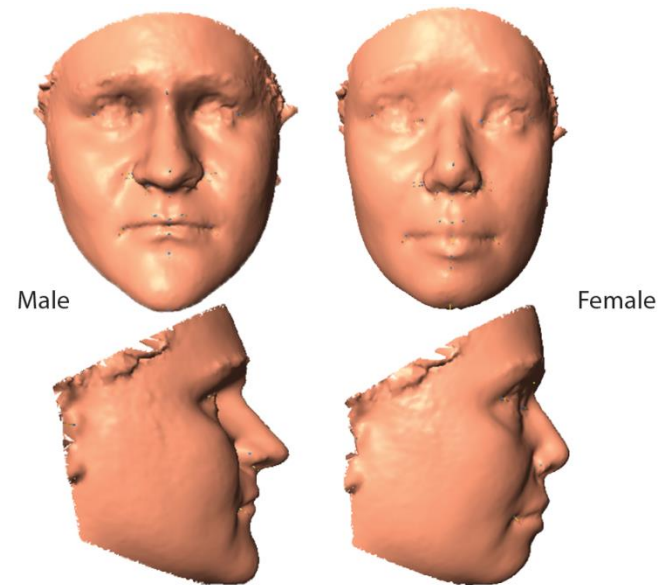
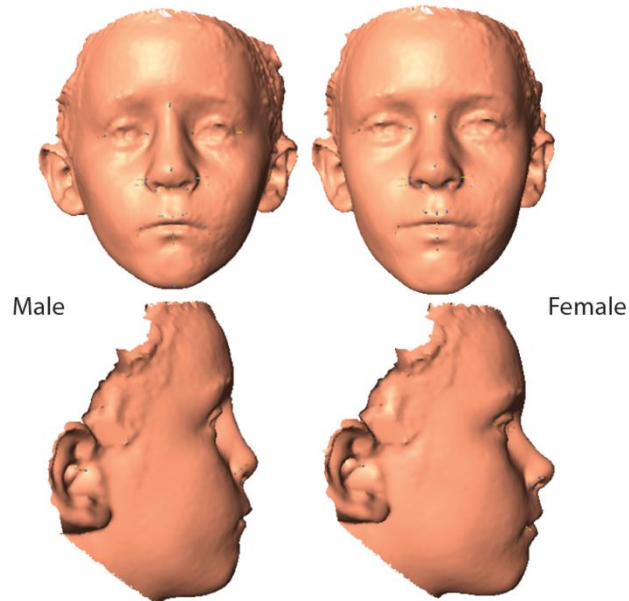
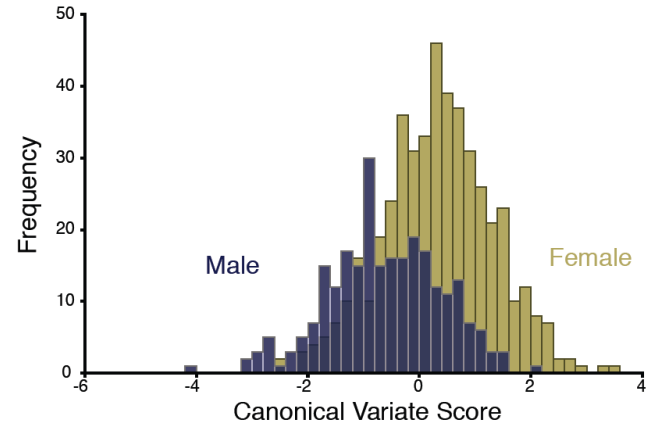
Genomic studies of craniofacial traits

Sexual Dimorphism in facial shape across the lifespan

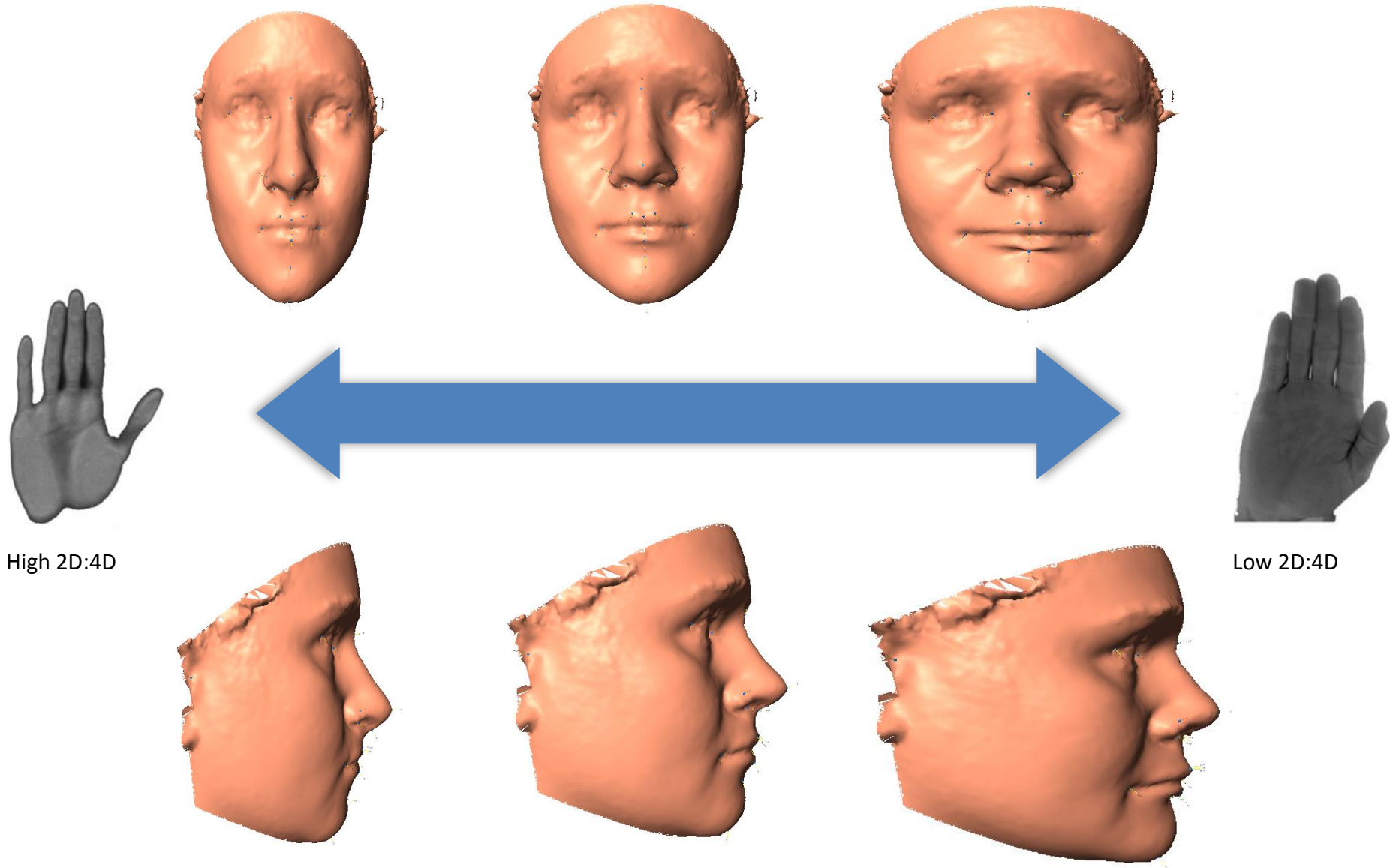
Juveniles 3-10 (n = 119)



Adults 18-40 (n = 722)



Relationship between facial shape and digit ratio in males



Using the 3D Norms Data

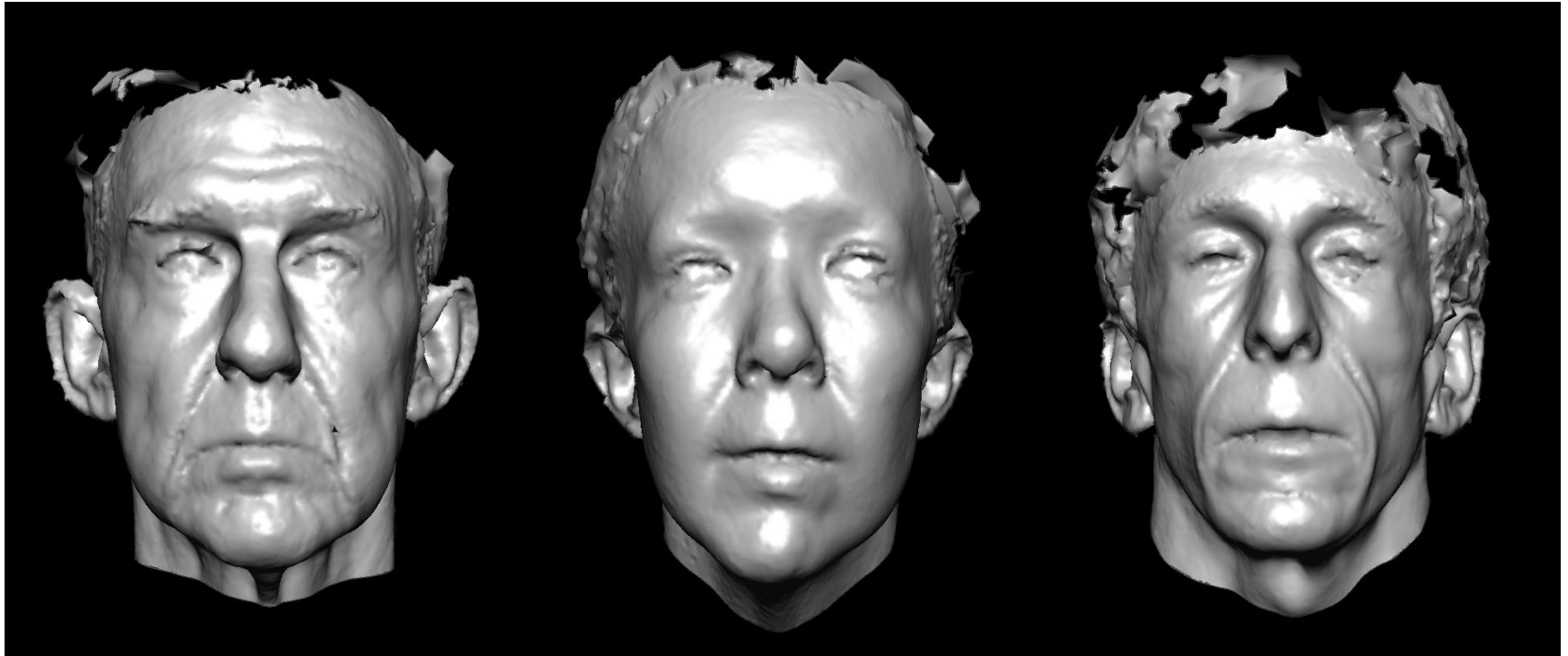
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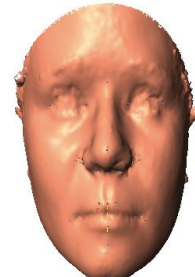
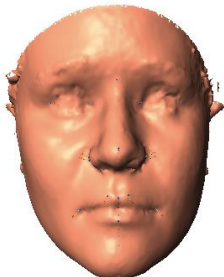
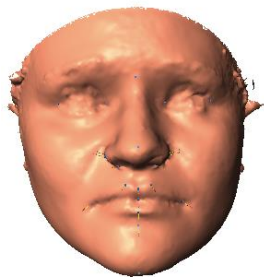
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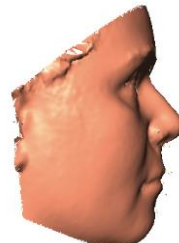
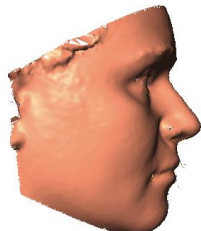
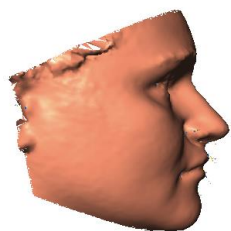
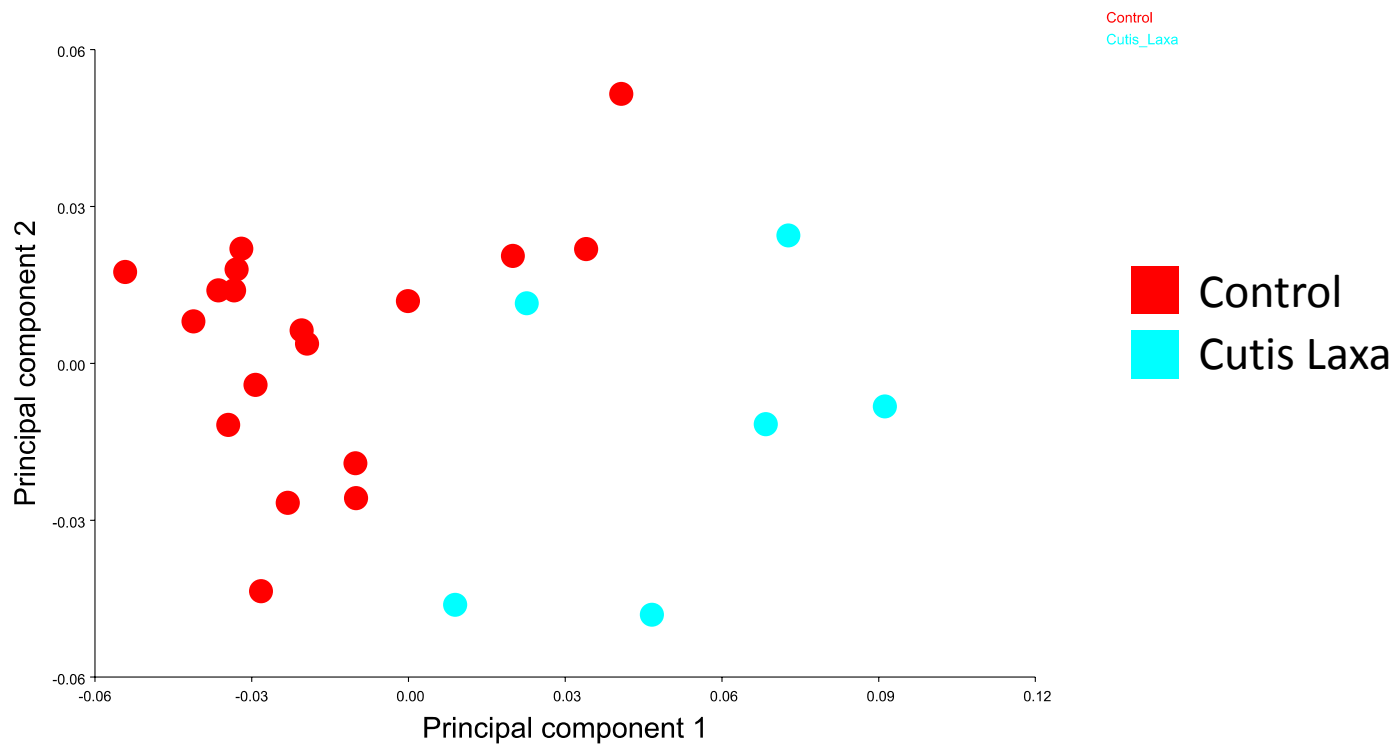
Assessment of craniofacial dysmorphology in Cutis Laxa



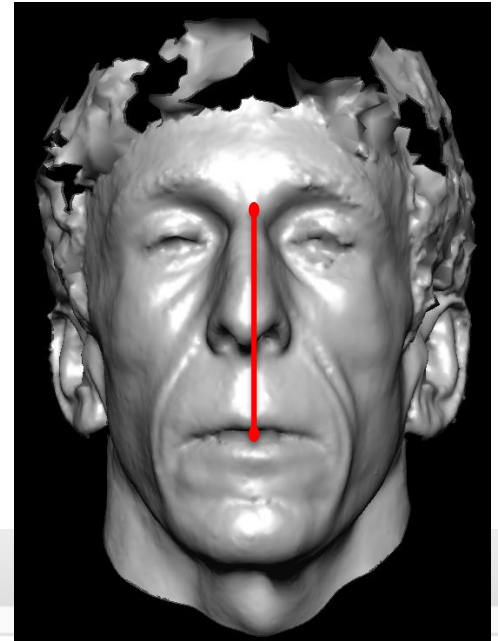
Control
Morph




Cutis Laxa
Morph



Using Z-Score Calculator Tool to compare upper facial height



MEASUREMENT	VALUE	Z-SCORE
Lower Facial Depth Left	<input type="text"/> mm	<input type="text"/> σ
Morphological Facial Height	<input type="text"/> mm	<input type="text"/> σ
Upper Facial Height	<input type="text" value="92.10"/> mm	<input type="text" value="2.02"/> σ



A normal distribution curve is shown with a horizontal axis ranging from -5 to 5. A blue dot is placed on the curve at the position corresponding to a z-score of 2.02.

Using the 3D Norms Data

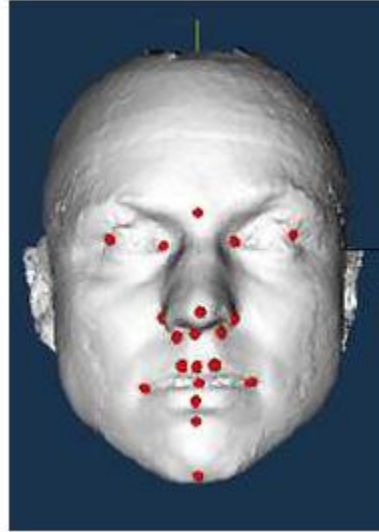
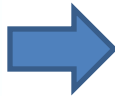
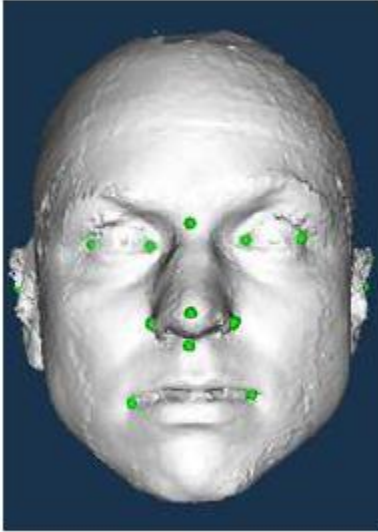
Analyses of human craniofacial variation and growth patterns

Normative control data for craniofacial comparisons

 **Development of novel image analysis methods**

Genomic studies of craniofacial traits

Automated landmarking of facial surfaces



Step 1: Initial Identification of facial landmarks using geometric method

Step 2: Deformable registration – improves fit and adds additional landmarks

Compared to manual landmark placement

Point Name	Geometric Method Average Distance(mm)	Deformable Method Average Distance(mm)
Nasion	—	2.92±1.62
Pronasale	1.29±0.68	1.59±0.81
Subnasale	2.35±2.16	2.45±0.80
Alare(R)	3.24±2.61	1.78±1.15
Alare(L)	3.14±2.41	3.07±1.15
Labiale Superius	—	2.27±1.15
Stomion	—	1.49±0.90
Labiale Inferius	—	2.27±1.41
Sublabiale	—	3.17±1.87
Subalare(R)	—	2.36±1.06
Subalare(L)	—	1.59±0.93
Crista Philtri(R)	—	2.31±1.27
Crista Philtri(L)	—	1.99±1.03
Chelion(R)	3.14±2.41	3.08±2.14
Chelion(L)	2.80±2.38	3.08±1.64
Gnathion	—	5.31±3.54
Endocanthion(R)	4.78±1.45	2.39±1.09
Endocanthion(L)	4.58±1.70	2.78±1.50
Exocanthion(R)	3.15±2.21	3.34±1.63
Exocanthion(L)	2.72±1.86	3.68±1.91

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Genotyping Strategy

- CIDR application planned for summer
- Goal is to have 2200-2500 samples in hand at the time of application
- Key factor in thinking about genotyping: **what is most useful for the FaceBase user community**
 - A genome-wide panel probably makes most sense for FaceBase users
 - Illumina Core Panel (300k SNPs) + Exome, or...
 - OmniExpress Panel (750k SNPs) + Exome, or...
 - Omni 2.5 Panel (2.5mil SNPs) + Exome
 - Additional custom panel for testing relevant SNPs from recent analyses
 - Replication and extension dataset
 - Meta-analysis/mega-analysis dataset

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