# **Computer-aided facial recognition of individuals with FG (Opitz-**Kaveggia) syndrome caused by p.Arg961Trp mutation in MED12

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# FG (Opitz-Kaveggia) syndrome is an X-linked recessive syndrome characterized by relatively large

head, frontal upsweep, hypertelorism, ptosis, small ears, broad and flat thumbs, imperforate anus, hypotonia, agenesis of the corpus callosum and moderate intellectual disability. The classical form of FG syndrome is caused by a recurrent p.Arg961Trp mutation in the MED12 gene. FG syndrome is misdiagnosed and recent literature confirmed that several patients with similar features had microdeletion/microduplication syndromes and syndromes caused by FLNA mutations.

With the advent of novel automatic face analysis techniques, our ability to analyze facial morphology from photographs has improved significantly. In this study we examined whether a computer-based dysmorphological analysis can be used in order to discern between FG patients and non-FG patients that present superficially similar facial features. For this, we used a collection of 18 genetically verified FG cases and 18 non-FG cases that were clinically suspected to be FG but do not carry mutations in the MED12 gene.

A modern face analysis system that was developed specifically for dysmorphological analysis was used. The system is fully automatic and starts by detecting the face in the image. Then, 130 facial fiducial points are localized and various measurements are taken. The final classification is based on these measurements as well as on a global "gestalt" detector that estimate the probability of the subject having FG based on the appearance of the entire facial image. A statistical technique called cross validation was used to estimate the recognition capability of the computer system. At each one of 20 rounds, the data was split randomly to training and testing data, each comprising of 50% of the samples.

The system was trained to separate between the two classes using the train dataset, and evaluated on the test data. The statistical power of the test in correct recognition of FG patients with the p.Arg961Trp mutation was estimated using the conventional recognition metric of average area under the ROC curve and was 90%, which is considered very high. The gestalt of the face as captured by multiple local patterns of facial texture contributed significantly to the correct

# recognition of the individuals with FG syndrome.

# Overall, we have demonstrated that computer-based analysis can be successfully used in supporting experts for the correct recognition of patients with FG syndrome.

# **Computer-aided facial recognition of individuals with FG (Opitz-Kaveggia)** syndrome caused by p.Arg961Trp mutation in MED12

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FG (Opitz-Kaveggia) syndrome is an X-linked recessive syndrome characterized by relatively large head, frontal upsweep, hypertelorism, ptosis, small ears, broad and flat thumbs, imperforate anus, hypotonia, agenesis of the corpus callosum and moderate intellectual disability. The classical form of FG syndrome is caused by a recurrent p.Arg961Trp mutation in the MED12 gene. FG syndrome is often misdiagnosed and recent literature confirmed that several patients with similar features had microdeletion/microduplication syndromes and syndromes caused by FLNA mutations.

With the advent of novel automatic face analysis techniques, our ability to analyze facial morphology from photographs has improved significantly. In this study we examined whether a computer-based dysmorphological analysis can be used in order to discern between FG patients and non-FG patients that present superficially similar facial features.

Greenwood Genetic Center BEILINSON HOSPITAL

1. Detect

2. Localize facial feature points

3. Represent

4. Build models

F - are the observed image data

L - are the decision variables

 $(L_1^n)$   $(L_2^n) \cdots (L_m^n)$ 

A - models the noise

R - are the output correlation variables

 $i=1,\ldots,k_n$ 

are

5. Evaluate

**Positive samples:** 

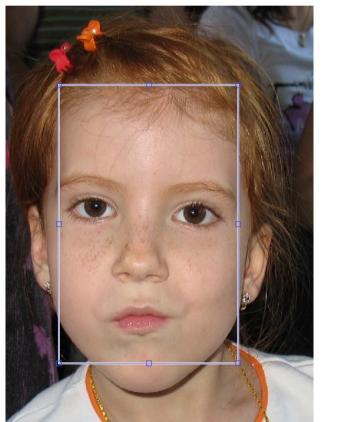


A collection of 18 genetically verified FG cases

**Negative samples:** 



A collection of 18 non-FG cases that were clinically suspected to have FG, but do not carry mutations in the MED12 gene.





Using a probabilistic model the frame of the face is located

Within this frame 130 fiducial points are using located local

Various local properties such as ratios of distances and local image descriptors are estimated

P.,

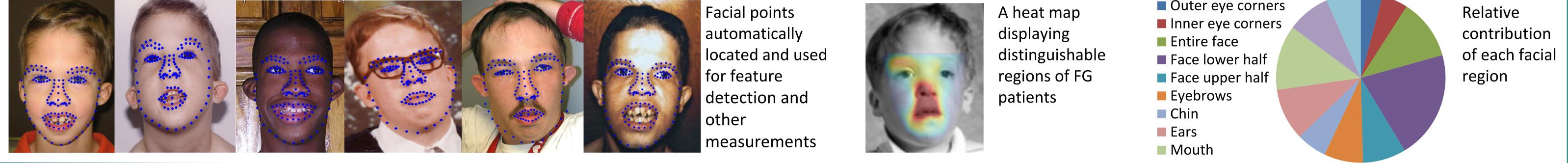
n = 1, 2, 3, .Statistical models called Bayesian networks constructed to measure FG dysmorphic feature and probabilities

Every specific image is placed on the curves measuring its FG feature probability ("empirical pvalue")

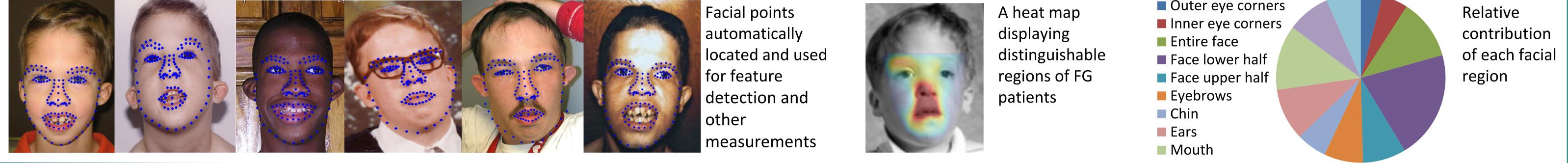
## **Testing the System**

Automatic facial contours analysis is used to extract relative measurements, fiducial points and dysmorphic feature evaluation

image detections



A "gestalt" description of the face is used to evaluate the entire images at once

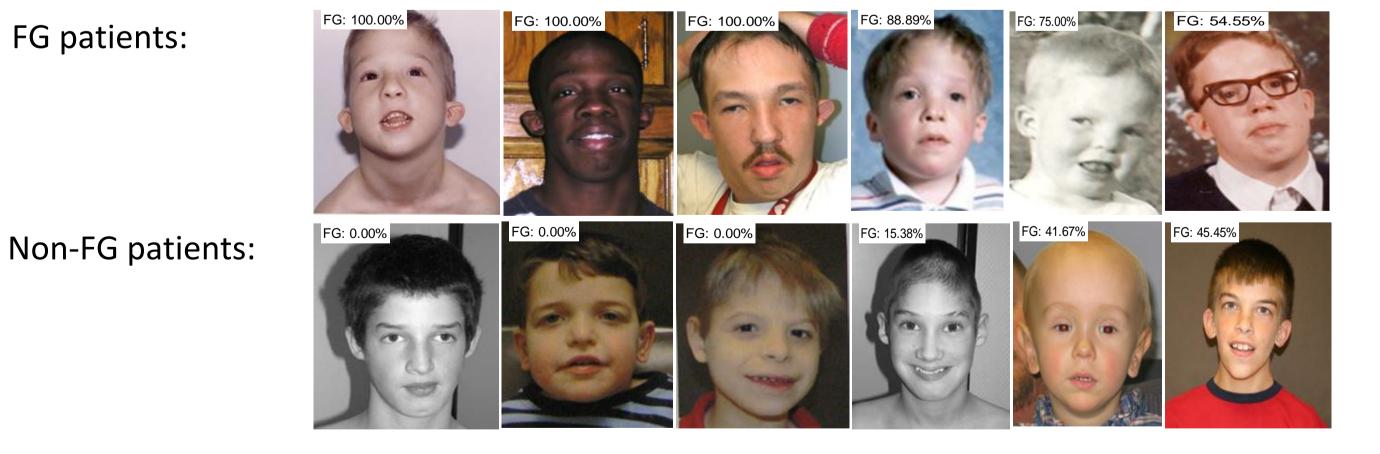


Outer eye corners

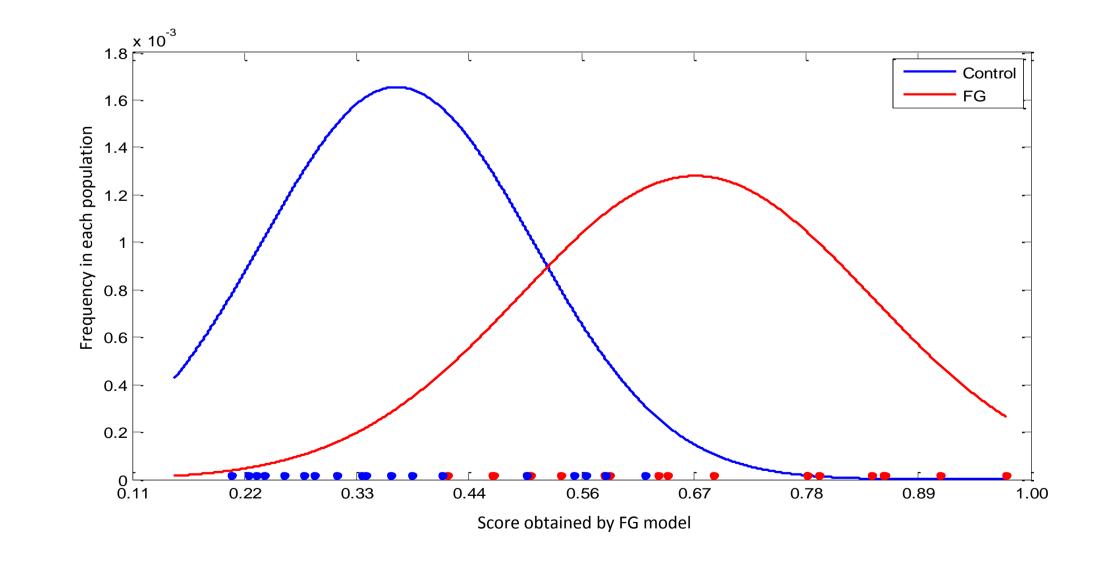
## Results

### The average ratio of the "Hit-Miss" (in %) in the cross validation process

FG patients:



### **Detection scores obtained for FG patients and non-FG control**



The statistical power of the test in correct recognition of FG patients with the p.Arg961Trp mutation was estimated using the conventional recognition metric of average area under the ROC curve and was 90%, which is considered very high.

# This demonstrates that computer-based analysis can be successfully used in supporting experts for the correct recognition of patients with FG.