

# Computational diagnostic methods on 2D photographs: A review of the literature

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#### 25 ABSTRACT

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27 Here we provide a literature review of all the methods reported to date for analyzing 2D 28 pictures for diagnostic purposes. Pubmed was used to screen the MEDLINE database 29 using MeSH (Medical Subject Heading) terms and keyworks. The different recognition 30 steps and the main results were reported. All human studies involving 2D facial 31 photographs used to diagnose one or several conditions in healthy populations or in 32 patients were included. We included 1515 articles and 27 publications were finally 33 retained. 67% of the articles aimed at diagnosing one particular syndrome versus 34 healthy controls and 33% aimed at performing multi-class syndrome recognition. Data 35 volume varied from 15 to 17,106 patient pictures. Manual or automatic landmarks were 36 one of the most commonly used tools in order to extract morphological information 37 from images, in 22/27 (81%) publications. Geometrical features were extracted from 38 landmarks based on Procrustes superimposition in 4/27 (15%). Textural features were 39 extracted in 19/27 (70%) publications. Features were then classified using machine 40 learning methods in 89% of publications, while deep learning methods were used in 41 11%. Facial recognition tools were generally successful in identifying rare conditions in 42 dysmorphic patients, with comparable or higher recognition accuracy than clinical 43 experts.

44 Keywords: literature review; machine learning; deep learning; photograph;
45 dysmorphology; diagnosis

46 **Declarations of interest:** none.

#### 48 INTRODUCTION

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50 Shape analysis on 2D pictures can involve computations based on pre-defined rules; in 51 machine learning approaches, these rules are replaced by the ability of the model to 52 learn from examples (1). Applications of machine learning are increasing in healthcare 53 (1), in a variety of fields such as radiology (2), dermatology (3) and surgery (4). Inputs 54 or features are converted into outputs or labels. Deep learning is defined as a subtype of 55 machine learning in which artificial neural networks are used. These neural networks can encode relationships between features and labels that are not accessible to the 56 57 human brain, because of their complexity or because they do not match the way our 58 brain processes data (1).

59 Examples of relevant inputs in healthcare are X-rays, texts, pathology slices and clinical 60 photographs. Among these inputs, clinical photographs are particularly important in the 61 diagnosis of craniofacial malformations. In fact, 30-40% of the 7000 rare diseases have 62 craniofacial anomalies (5). Many among these 2-3000 syndromes have minor facial 63 features or are very uncommon: their screening requires the eye of an expert in 64 dysmorphology. Due to this specific clinical issue, there have been a recent increase in 65 the number of scientific publications dedicated to the diagnosis of rare conditions using 66 computational methods on facial photographs (6).

Here we provide a literature review of all the methods used for computational diagnosis
on facial photographs. This overview will be of use for physicians and scientists
interested in syndrome recognition, and more generally, in facial recognition.

70

72 **METHODS** 

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We performed a review of the literature according to the framework described by Arksey and O'Malley (7). We aimed at reporting all articles describing the computerized identification of a disease and/or a syndrome based on 2D facial photographs. PubMed was used to screen the MEDLINE database using MeSH (Medical Subject Heading) terms along with keywords from titles and abstracts (see request Table 1).

80 All human studies involving 2D facial photographs used to diagnose one or several 81 conditions in healthy or patient populations were included. Only studies with diagnosis 82 confirmation by clinical experts and/or molecular investigations were retained. Articles 83 on prenatal diagnosis, extra-facial diseases and dermatological diseases of the face (e.g. 84 facial acnea or naevi) were excluded. Studies assessing treatment results, such as 85 craniofacial surgery results, were excluded, as well as studies dedicated to the detection 86 of pain or other emotions. We also excluded studies based on the diagnosis of patients 87 using a pre-existing algorithm, such as *Face2Gene* (8).

88 One reviewer (QH) screened all the titles and/or abstracts. When exclusion criteria were 89 not clear after reading title and abstract, the full text was assessed. The relevant 90 publications were then charted with the following items: year of publication, first 91 author, country of affiliation, study design or objective, disease of interest, origin of 92 data and number of patients. We reported the pre-processing of the pictures, *i.e.* every 93 treatment the authors applied to the photograph before extracting features or using 94 classifiers. Concerning feature extraction, we reported the use of: (1) landmarks, with 95 manual and/or automatic placement, (2) geometric features, *i.e.* distances, angles or

96 other morphometric data and (3) texture features, based on the spatial arrangement of 97 intensities on the image. We listed the classifiers and the validation methods of machine 98 learning designs. We considered that authors had used machine learning methods when 99 predictions were made on a test set with a classifier or a model trained on a training set.

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101 **RESULTS** 

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With the pre-defined request, we listed 1515 results on PubMed. After applying all the exclusion criteria on reading title, abstract and full text when necessary, we included 27 articles. These publications were fully read and charted (Table 2). The region of affiliation of the first author was North America – USA only in 37%, Europe – Germany, UK and Croatia in 33%, and Asia – China, India and Turkey in 30%.

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#### 9 1. Study designs and conditions

110 We defined two types of studies depending of the number of diseases to be diagnosed.

The first and most common study design – in 18/27 (67%) publications – intended to diagnose one condition in the general population versus healthy controls using facial photographs. Facial pictures of the following conditions were tested versus healthy controls: Down syndrome (10–13), Cornelia de Lange syndrome (16), Rubinstein-Taybi syndrome (16), 22q11.2 deletion syndrome (17), Laron syndrome (18), Williams-Beuren syndrome (19), acromegaly (20–24), Cushing disease (25), Turner syndrome (26), Chronic Fatigue Syndrome (27) and schizophrenia (28).

118 The second most prevalent study design – in 9/27 (33%) publications – consisted in 119 using multi-syndrome classification, *i.e.* in differentiating syndromes from each other and from the general population (8,29–38). For example, Ferry et al. (29) aimed at
diagnosing 8 syndromes – Angelman, Apert, Cornelia de Lange, Down, fragile X,
progeria, Treacher-Collins and Williams-Beuren syndromes – in the general population.
Gurovich et al. (8) aimed at diagnosing 216 different syndromes in a population of
affected patients.

- 125 The conditions reported in the included studies were:
- 126 genetic syndromes: Down (29), Angelman (8), Prader-Willi (34), Cornelia de Lange
- 127 (8), X fragile (34), Apert (29), Progeria (29), Treacher-Collins (29), Williams-
- 128 Beuren (32), Sotos (30), Cri-du-chat (34), Smith-Lemli-Opitz (34), 22q11.2 deletion
- 129 (34), Turner (8), Noonan (8), Mucopolysaccharidosis (34), Rubinstein-Taybi (8),
- 130 Hurler (36), Wolf-Hirschhorn (36) and 183 others in (8),
- 131 endocrine conditions with a facial phenotype (e.g. acromegaly and Cushing
  132 syndrome) (23)
- 133 psychiatric conditions: schizophrenia (28) and chronic fatigue syndrome (27).
- 134

#### 135 **2.** Data description

136 The photographic data had diverse origins: (1) pictures taken during clinics 137 (9,10,12,15,17-25,27,30,32-36,39-41), (2) pictures from articles books or 138 (14,16,17,29,38), (3) pictures from pre-existing cohorts (8,37,42,43) and (4) pictures 139 publicly available on the internet (11,29). The data volume varied from 15 pictures of 140 affected patients (9) to 17,106 for Gurovich et al. (8). The average number of affected 141 (non-control) patients in the studies was 844 (+/- 3272); more precisely, 114 (+/- 138) 142 affected individuals were included in average in order to detect one syndrome versus the 143 general population and 2085 (+/- 5309) affected individuals in order perform multisyndrome classification. There was one photograph per patient in all publications.
Authors used age-, sex-, (15,19,22,25,40) and ethnicity-matched controls (12) in 6/27
(22%) studies.

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- 148

### 3. Image pre-processing

Pre-processing corresponded to the preparation of pictures before extracting data and applying classifiers. It is a useful step in machine learning: images are grouped (face, profile), resized and transformed (rotation, cropping, lighting) to optimize analysis.

An automated facial detection system was used in several studies to sort frontal views from lateral views and then crop the pictures. The most common algorithm in this field was the *Haar cascade* tool developed by Viola and Jones (1). One author used manual cropping, scaling and rotation to maximize the classifier performances (9). This approach was not adapted to large datasets. Manual or automated cropping was useful for little standardized databases, for instance when several individuals were present on pictures (*e.g.* the child and his mother) (6).

159

160 *4. Image processing* 

All authors applied two exclusive image processing approaches: (1) use of geometric and/or textural features in order to describe phenotypes or (2) deep learning process in order to detect a syndrome without describing phenotypes.

164

#### 165 Landmarks positioning

166 Landmarks are defined by reproductible points of interest on an image. They are one of167 the most common tools used to extract phenotypic data, as in 22/27 (81%) publications.

In 7/27 (26%) publications, the authors placed landmarks manually and automatic
landmarking was used in 16/27 (59%) publications.

In 4/27 (15%) publications (9,11,24,41), the authors did not use landmarks to analyze
faces and phenotypes and preferred assessing the overall shape and textures.

172

#### 173 Frontalization and facial alignment.

Frontalization is a process modifying a frontal image in order to improve the orientation of the frontal view. Kong et al. (21) used alignment, based on Procrustes superimposition (47,48), in order to compute frontal pictures from photos with capture angles varying from -45° to +45° around the standard coronal plane. The final frontal face was produced using the corresponding symmetric sides, which raised numerous issues in conditions with facial asymmetry.

180

#### 181 Geometric features extraction

Geometric features were extracted from landmarks and used for disease detection. The main advantage of using landmarks was to be more flexible with pose changes and different angles of view. One straightforward approach was to measure Euclidean distances and angles from landmarks (22,23,25,26,35,36).

Another approach to geometrical information extraction, used in 4/27 (15%) publications, was Procrustes analysis (10,16,29,49,50). Procrustes distance was defined as the deformation necessary to fit a dysmorphic face on a control face via an iterative least-squares process. Each of the faces were centered and scaled using a centroid (51) to minimize sum of squared differences between the landmarks. Douglas and Mutsvangwa (52) concluded that Procrustes-based approaches were the best methodsfor shape analysis in terms of prediction accuracy.

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#### 194 *Texture features extraction*

195 A texture extraction system was used in 18/27 (67%) publications, in association with 196 geometric features extraction (8,11,12,14,22,23,25–27,32,37,41,48) or independently 197 (9,11,41). The most common methods were Gabor Wavelet Transformation (GWT) 198 used in 12/27 (44%) (54,55) and/or a Local Binary Patterns (LBP) in 7/27 (26%) 199 (56). These methods provide information about the surface located publications 200 between the landmarks. For example, zygomatic hypoplasia in Treacher Collins 201 syndrome would be better detected by analyzing the shadows on a photo than by 202 measuring angles between landmarks.

203

The information on geometric / texture features created extensive data matrices, with
the need to reduce dimensionality before incorporation into classifiers. For this purpose,
Principal Component Analysis (PCA) was used in 17/27 (70%) publications.

Gurovich et al. (8) did not use any geometric or textural features extraction step for prediction making after facial alignment based on different landmarks. Their approach is referred to as deep learning because it corresponds to a *black box* approach diagnosing dysmorphic features without going through a phenotype description process. When using deep learning, authors cannot easily explain what contributed to the diagnosis of a particular syndrome.

213

214 5. Classification

215 Machine learning was defined by using a training set to make predictions on a 216 validation or test set. Based on this, 3/27 (11%) publications were not using machine 217 learning methods.

In these studies, diverse classifiers were used. The simplest classifiers were Generalized Linear Models (GLM) such as logistic regression and linear discriminant analysis, as used in 4/27 (15%) publications (16,21). Their output was a linear combination of input facial data.

SVM – used in 13/27 (48%) publications – was a classification method aiming at
separating two data sets with distance maximization based on an optimal separating
hyperplane (57). Bounds between datasets and the optimal separating hyperplane were
referred to as 'support vectors' (68).

The k-nearest neighbors (kNN) rule – used in 11/27 (41%) publications – classified each unlabeled example by the majority label of its k-nearest neighbors in the training set (59). It stored instances of previously earned images. Classification was obtained from a majority vote of the nearest neighbors (21). The kNN classifier was particularly adapted for multiclass detection.

Random forests (RT) – used in 2/27 (7%) publications – were another technique (21)
consisting in growing many classification trees. The forest chose the classification tree
having the best prediction. In addition, a subset of features was randomly selected from
the optional features in order to grow the tree at each node. Random forests were
particularly adapted for large number of weak classifiers.

Deep learning – Convolutional Neural Networks (CNN) or Artificial Neural Networks
(AAN) – was used in 3/27 (11%) publications. Kong et al. (21) also used CNN with
convolutional layers, pooling layers and fully-connected layers. Gurovich et al. (8) used

Deep Convolutional Neural Networks (DCNN) (60) for each of the cropped facial areas,
which were then combined by averaging to form a model for a multiclass problem.

Kong et al. (21) compared the performances of these machine learning techniques (linear functions, SVM, kNN, RF and CNN): better classification abilities were found in terms of accuracy for SVM with detected features and for CNN without detected features.

245

#### 246 *6. Clinical results*

247 Saraydemir et al. (9) and Zhao et al. (10) found an overall accuracy of respectively 97% 248 and 97.9% with 15 and 24 patients for diagnosing Down syndrome in a healthy 249 population. Both these performances were higher than clinical experts. Kruszka and his 250 team found an accuracy of 95% for detecting 22q11.2 deletion syndrome in 156 patients 251 vs. the general population (17), 94% for Down syndrome (12) in 129 patients, 90% for 252 Williams-Beuren syndrome in 286 patients (19), 94% for Cornelia de Lange in 246 253 patients (15) and 89% for Noonan syndrome in 161 patients (40). Kong et al. (21) used 254 the F1-score in order to evaluate the performances of their classifier in detecting 255 acromegaly in 527 patients. They obtained a F1-score of 92% with landmarks and SVM 256 and 91% without landmarks and with deep learning methods. Accuracies of 86% and 257 82% were obtained to detect acromegaly among a healthy population in 24 (20) and 57 258 patients (22).

Ferry et al. (29) used the *Clustering Improvement Factor* (CIF) after implementing kNN
on 8 syndromes and 1363 patients. Treacher-Collins syndrome was then clustered 23.5fold better than by chance alone, Down syndrome 11.6-fold better, and WilliamsBeuren 9.4-fold-better. The overall accuracy for detecting one syndrome vs. the 7 others

263 was 76% and the overall accuracy for detecting syndromes vs. a healthy population was 264 95%. Another method for displaying results in a multi-syndrome classification design 265 was to provide a Top-5 or Top-10 accuracy, i.e. to evaluate the probability of a 266 condition to be ranked as the 5 or 10 most probable syndromes by the classifier. 267 Gurovich et al. (8), in 17,106 patients with 216 syndromes, found a Top-10 accuracy of 268 91%, a Top-5 accuracy of 85% and a Top-1 accuracy of 61%. Kuru et al. (38) found 269 53% accuracy on a multi-class syndrome detection of 15 conditions with 92 patients. 270 Boehringer et al. (32) had an overall accuracy of 21% for multi-syndrome detection on 271 14 conditions in 202 patients, which is very low compared to the performances of the 272 clinical experts.

273

#### 274 **DISCUSSION**

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276 Machine learning approaches used to diagnose clinical conditions on 2D photographs 277 generally followed similar frameworks (Figure 1): either features were extracted from 278 photographs to diagnose a disease or the syndrome was detected directly without feature 279 extraction by deep learning. In case of features extraction, the steps were: image pre-280 processing, image processing with or without landmark placement, extraction of 281 geometrical and/or textural features, reduction of the dimensions of the input, and 282 classification and validation using training and test sets. Procrustes analysis was a 283 reliable technique for the extraction of morphometric parameters (52). Texture 284 extraction techniques seemed of interest in adding information to the geometric features 285 (27).

Deep learning methods are a promising tool with much space for performance
improvement (8). The number of publications using deep learning was however limited
because this approach requires large data volumes – 17,106 patients in Gurovich et al.
(8) – in order to train the model. The challenge in the future years will be to build large
photographic databases and train neural networks in order to improve predictions.

291 Comparing publications and diagnosis performances of machine learning tools was 292 difficult due to the variety of assessment parameters: Boehringer (32) used accuracy, 293 Zhao (10) precision and recall, Dudding-Byth (37) top-10, top-5 and top-1 accuracies, 294 and Ferry (29) the Clustering Improvement Factor (CIF). Evidence-based comparisons 295 of machine learning in facial recognition would greatly benefit from more homogenous 296 result assessment approaches.

297

#### 298 CONCLUSION

Facial recognition approaches are efficient in identifying rare conditions using 2D photographs. Machine-learning algorithms improve recognition accuracy. Deep learning techniques such as CNNs are promising, but larger datasets are needed. Based on the current results of available algorithms, we can predict that collaborative studies with massive shared datasets could shortly lead to the development of algorithms with almost perfect diagnostic abilities. 305 **REFERENCES** 

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#### 477 FIGURES AND TABLE CAPTIONS

478

479 *Table 1.* Pubmed request.

(diagnosis[Title] OR detection[Title] OR dysmorphic[Title/Abstract] OR dysmorphology[Title/Abstract] OR disease[Title/Abstract] OR syndrome[Title/Abstract])

**AND** (*face*[*Title*/Abstract] *OR faces*[*Title*/Abstract] *OR facial*[*Title*/Abstract])

AND ("face-classification"[Title/Abstract] OR "facial analysis technology"[Title/Abstract] OR "artificial intelligence"[MeSH Terms] OR "biometric identification"[Mesh] OR "Medical informatics"[MeSH Terms] OR "Image Processing, Computer-Assisted"[MeSH Terms])

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481

482 *Table 2.* General characteristics for each of the studies, and method of classification.

483 Publications not using machine learning (n = 3) were excluded. SVM = Support Vector

484 Machine. kNN = k-Nearest Neighbors. RF = Random forests. CNN = Convolutional

485 Neural Network. AAN = Artificial Neural Networks.

			Linear				CNN /
Multisyndrome	Data volume	Number of syndromes	functions	SVM	kNN	RF	ANN
Boehringer, 2006	147	10		$\checkmark$	$\checkmark$		
Vollmar, 2007	200	14		$\checkmark$	$\checkmark$		
Boehringer, 2011	202	14		$\checkmark$	$\checkmark$		
Ferry, 2014	1363	8			$\checkmark$		
Kuru, 2014	92	15			$\checkmark$		
Shu, 2015	13	100		$\checkmark$	$\checkmark$		
Özdemir, 2018	124	5			$\checkmark$		$\checkmark$
Gurovich, 2019	17,106	216					$\checkmark$
Monosyndrome		Condition / syndrome					
Schaefer, 1994	49	Laron	$\checkmark$				

L	24	A		/			
Learned-Miller, 2006	24	Acromegaly		$\checkmark$			
Miller, 2011	24	Acromegaly		$\checkmark$			
Saraydemir, 2011	15	Down		$\checkmark$	$\checkmark$		
Schneider, 2011	57	Acromegaly	$\checkmark$				
Buretic-Tomljanovic, 2012	58	Schizophrenia	$\checkmark$				
Kosilek, 2013	20	Cushing	$\checkmark$				
Zhao, 2014	100	Down		$\checkmark$	$\checkmark$	$\checkmark$	
Chen, 2015	294	Chronic fatigue syndrome			$\checkmark$		
Kong, 2017	527	Acromegaly		$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$
Kruszka, 2017	156	22q11.2		$\checkmark$			
Kruszka, 2017	129	Down		$\checkmark$			
Kruszka, 2017	161	Cornelia de Lange		$\checkmark$			
Chen, 2018	32	Turner		$\checkmark$			
Kruszka, 2018	286	Williams-Beuren		$\checkmark$			
Dowsett, 2019	246	Cornelia de Lange		$\checkmark$			
407							