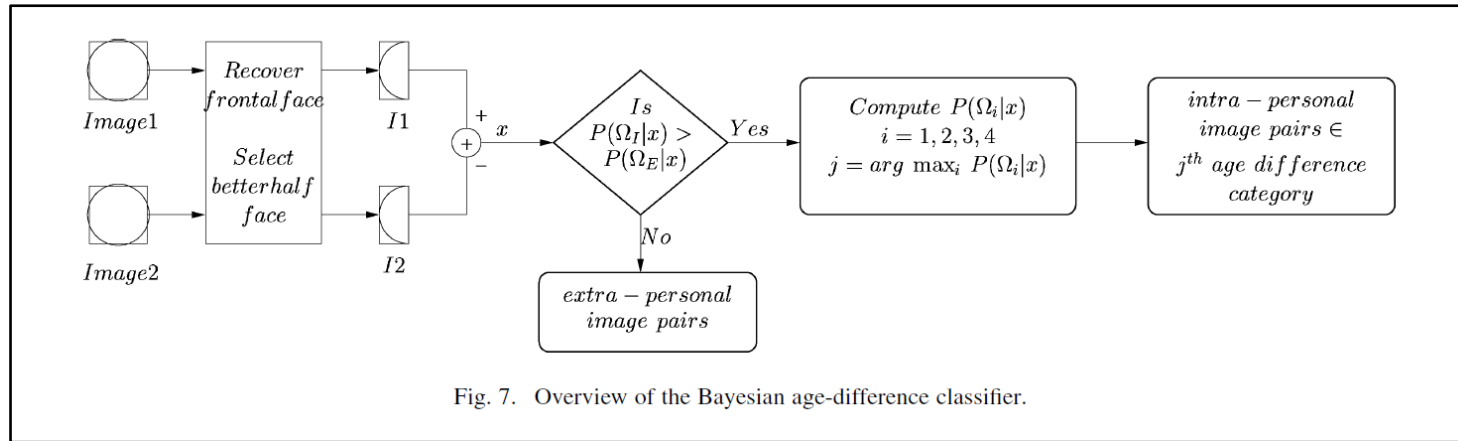


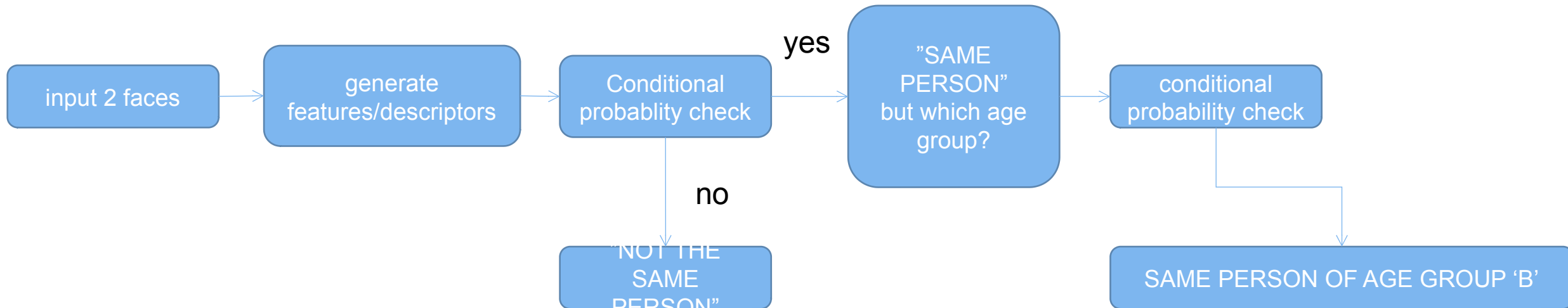
Identification of time-invariant  
unique features of Objects

- A. The following research in biometrics revolves around temporally influenced features and algorithms.
1. Facial verification across age.
  2. Facial verification across age using discriminative methods.
  3. Latent Fingerprint matching using descriptor based hough transform
  4. Fingerprint Spoof Detection: Temporal Analysis of Image Sequence
  5. Iris recognition performance in children: a longitudinal study

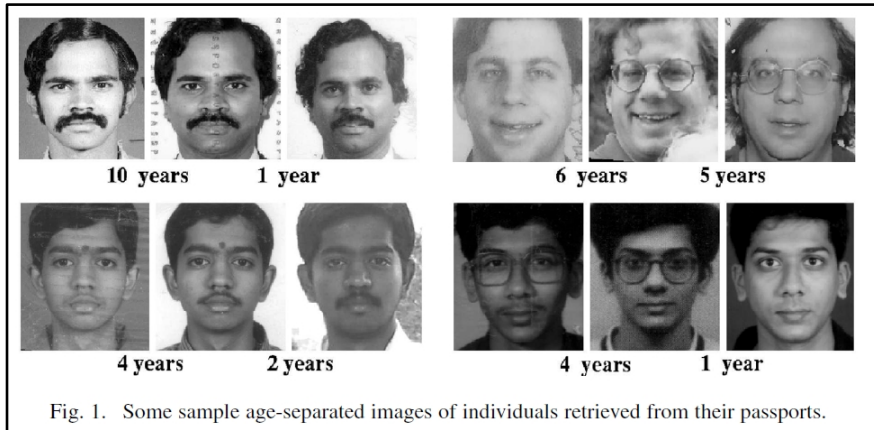
# 1. Facial verification across age



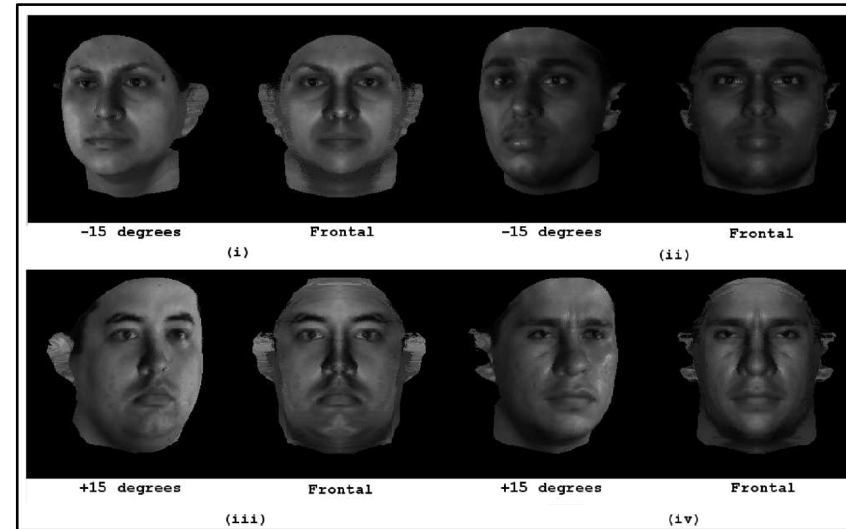
Making the features time invariant by removing artifacts that accumulate with time, like illumination, intensity. facial hair etc.,



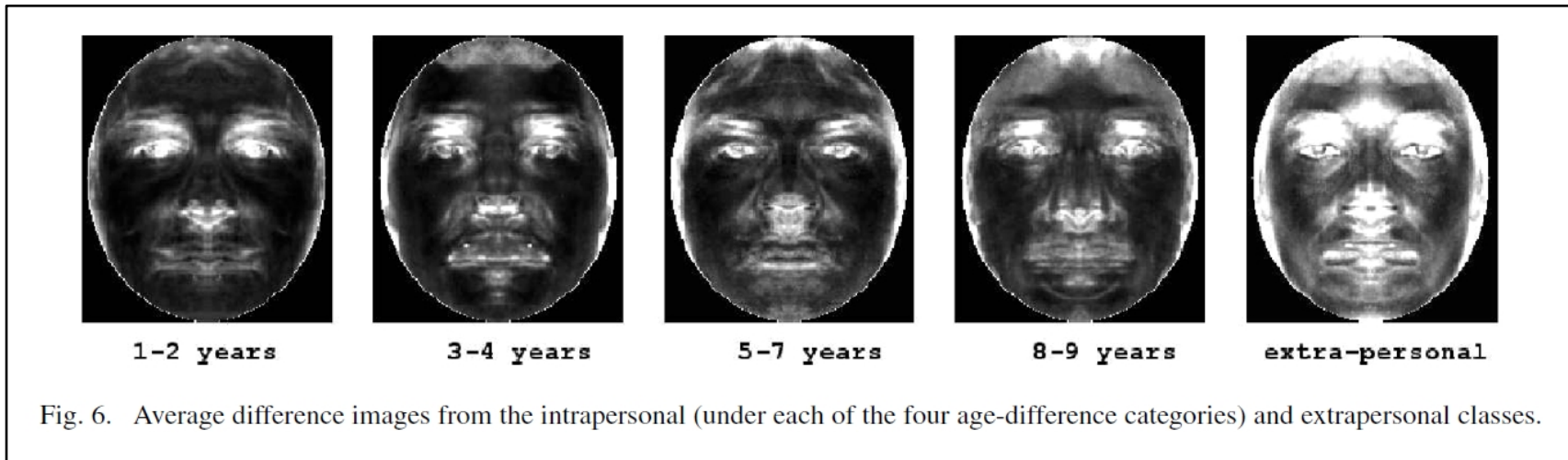
# 1. Facial verification across age



The time separated image samples



Face posture normalisation by unwrapping algorithms. Removing time induced artifacts as much as possible.



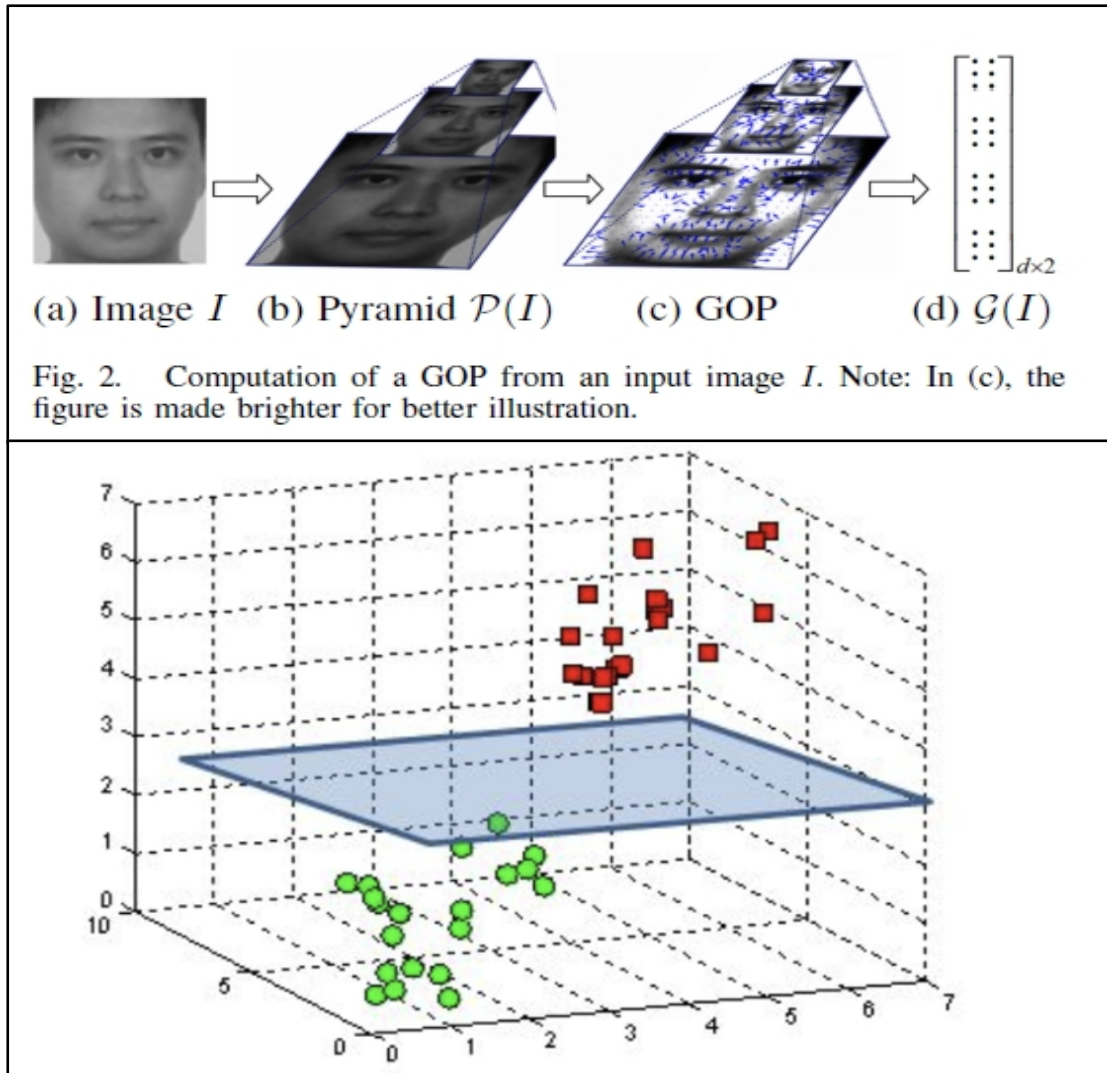
first 4 images:  
Intra personal  
difference image

last image:  
Extra personal  
difference image

# 1. Facial verification across age

- Useful ideas from this paper:
  - Identifying and Removing time induced artifacts can make the features time-invariant.
  - Using Bayesian framework as a classifier can yield good results **only when the features changes predictably with time.**
  - for example: we know that “facial wrinkles” are features in older images only.
  - Also the error rate increases with increase in Age gap.

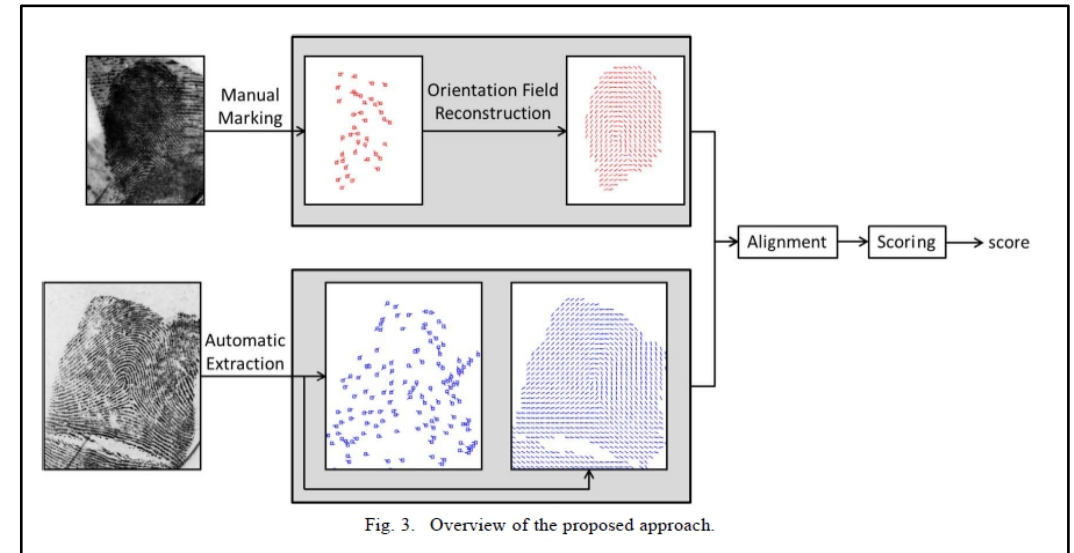
## 2. Facial verification across age using discriminative methods



- The paper proposes GOPs (Gradient Orientation Pyramid) as features for classifying OLD vs NEW images.
- The GOP features of input image is classified as “same person” or “different person” by using the classifier called SVM (Support Vector Machine).

### 3. Latent Fingerprint matching using descriptor based hough transform

- Latent fingerprints are fingerprints that are formed by sweat and oil on the skin.
- Such prints are not visible with naked eye and are collected carefully.
- The idea is to extract Miniteaue descriptors and use Hough transforms for alignment.



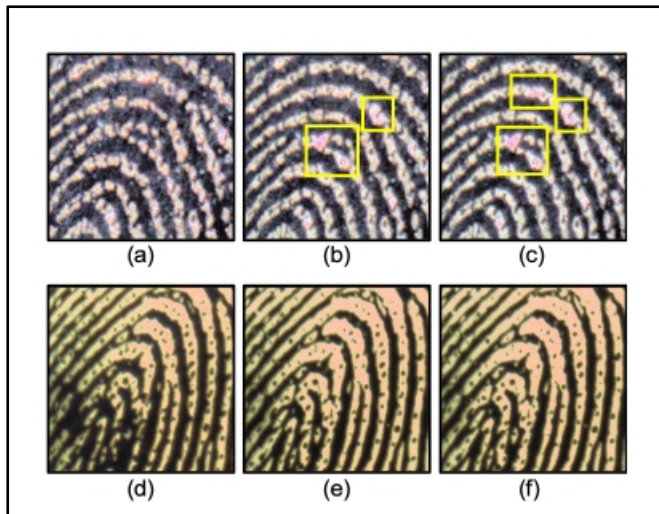
### 3. Latent Fingerprint matching using descriptor based hough transform

- Useful ideas from this paper
  - Latent fingerprints are usually contaminated by lot of external influence. The randomness in this influence can represent the randomness of the world and time.
  - The identification of features like 'Orientation field' at minutia points, even if only few are identified, it will help in matching with a template.
  - The usage of descriptor based Hough transforms to extract the orientation information of the lines on fingerprint is uniquely effective for fingerprints.
  - 2d representation of the fingerprints is 'set of lines'.

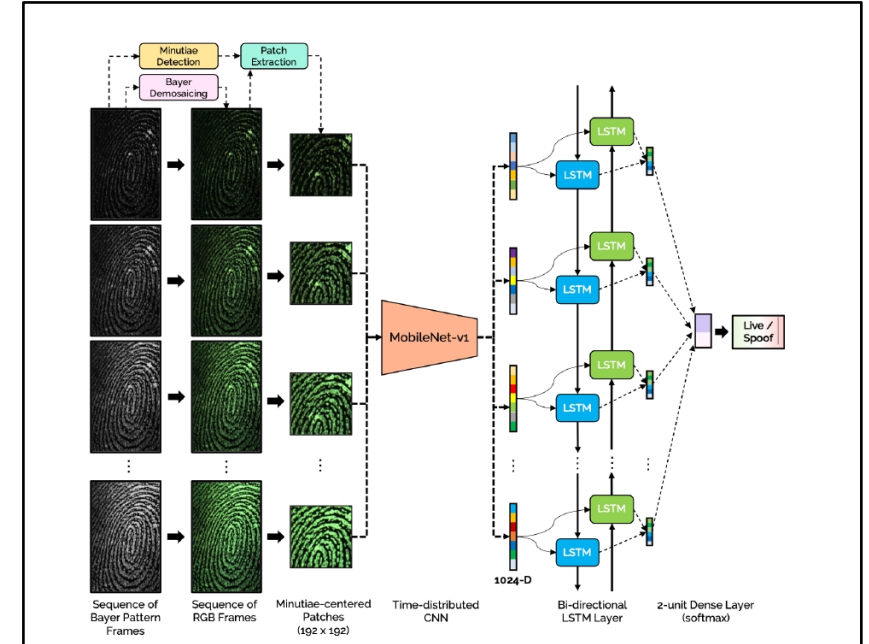


# 4. Fingerprint Spoof Detection: Temporal Analysis of Image Sequence

- Fingerprint spoof: Trying to fool the system by scanning fingerprint impressions made from any thing, but not human.



The preprocessing filters highlight the intricate skin imperfections as feature and compare with previous detections (LSTM).



The architecture of CNN-LSTM model used in this paper.

# 5. Iris recognition performance in children: a longitudinal study

- The study was performed on Children between ages 4-11 for 3 years every 6 months.
- Pupil data collected is combined with other data like
  - Time difference
  - Enrollment Age
  - Dilation
  - Delta dilation
- The matching score is a function of both dilation data and time difference.

$$\begin{aligned} MS \sim & \beta_0 + \beta_1 TD + \beta_2 EA + \beta_3 PD + \beta_4 \Delta D \\ & + \beta_5 EA^2 + \beta_6 \Delta PD^2 + \beta_7 \Delta D^2 + b_{0i} \\ & + b_{1i} TD + b_{3i} PD + b_{4i} \Delta D + b_{6i} PD^2 \end{aligned}$$

Time difference as a parameter is highlighted

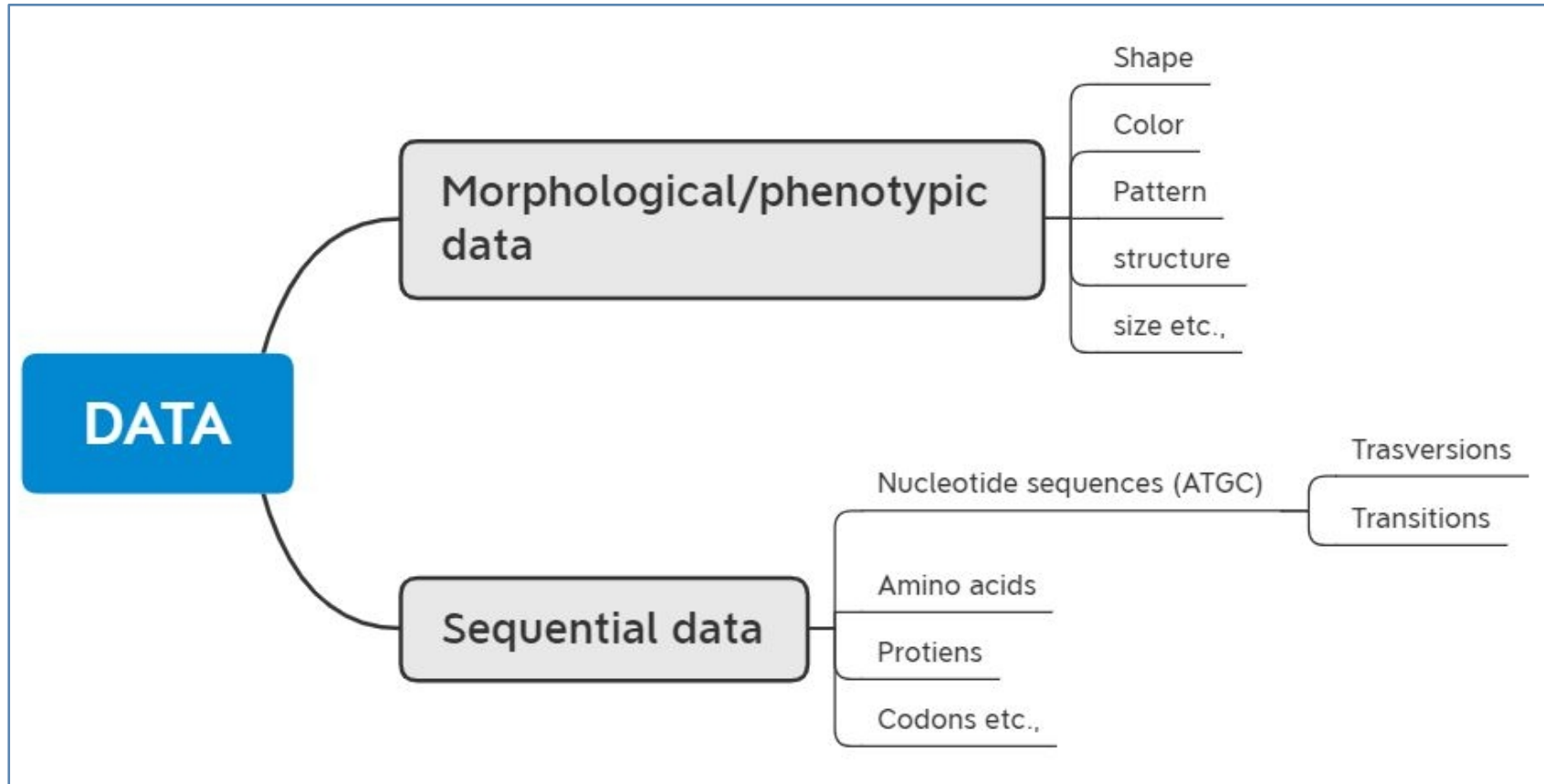
# 5. Iris recognition performance in children: a longitudinal study

- Useful ideas from this paper:
  - The tiny variations/changes in the pupil dilation measure will contribute 29 to 45 times more than that of due to **Time difference**.
  - The feature that is being studied in this paper varies very little with time. Dilation is computed as ratio of pupil radius and iris radius.

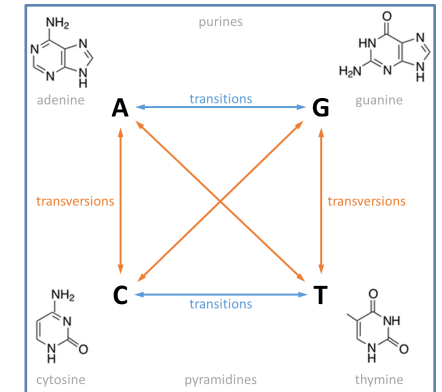
$\Delta D > \text{Subject} > \text{Dilation of the probe image} > \text{Time Difference}$

*Interestingly, we can infer from the model that dilation difference between two comparing images (in our case from 2 different TFs) adds in the most variability to the match score than any other significant factor. Aging effect contributes to the least variability in MS.*

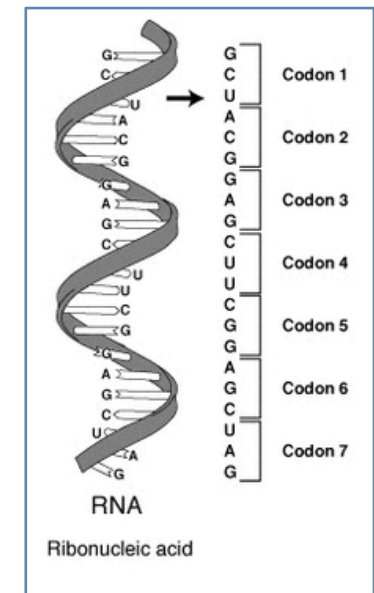
# B. Phylogenetics



Various formats of DATA in phylogenetics

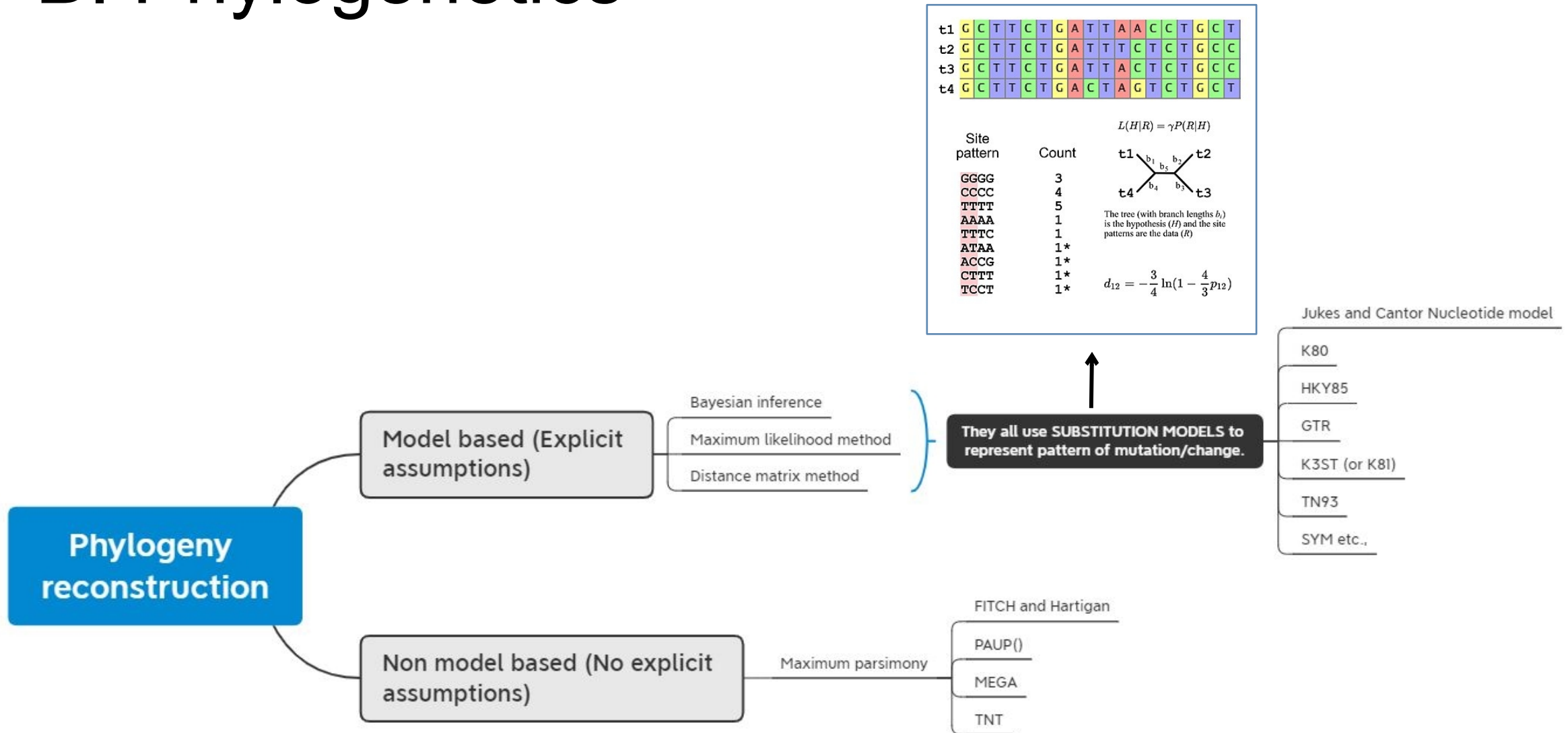


Transition and Transversion



Nucleotide and CODON sequences

## B. Phylogenetics



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  - <https://arxiv.org/abs/2011.02698>
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  - [https://www.researchgate.net/publication/224356996\\_A\\_Non-generative\\_Approach\\_for\\_Face\\_Recognition\\_Across\\_Aging](https://www.researchgate.net/publication/224356996_A_Non-generative_Approach_for_Face_Recognition_Across_Aging)
- Face Verification across Age Progression using discriminative method
  - <https://ieeexplore.ieee.org/document/5353681>
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  - <https://ieeexplore.ieee.org/document/1709980>
- Fingerprint Spoof Detection\_Temporal Analysis of Image Sequence 2020
  - <https://ieeexplore.ieee.org/document/9304921>
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  - <https://arxiv.org/abs/2101.06346>
- Latent Fingerprint Matching Using Descriptor-Based Hough Transforms
  - <https://ieeexplore.ieee.org/document/6117483>
- Phylogenetics\_beyond\_biology
  - [https://www.researchgate.net/publication/325898381\\_Phylogenetics\\_beyond\\_biology](https://www.researchgate.net/publication/325898381_Phylogenetics_beyond_biology)
- Spatio-Temporal Non-Rigid Registration of 3D Point Clouds of Plants\_chebrolu2020icra
  - <http://www.ipb.uni-bonn.de/pdfs/chebrolu2020icra.pdf>
  - [https://youtu.be/uGkep\\_aelBc](https://youtu.be/uGkep_aelBc)

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  - chapter 4. genetic distances and nucleotide substitution models
  - chapter 5. Phylogenetic inference based on Distance methods
  - chapter 6. Phylogenetic inference using Maximum likelihood methods
  - chapter 7. Bayesian Phylogenetic analysis using MrBayes
  - chapter 8. Phylogenetic inference based on Parsimony and Other methods
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