





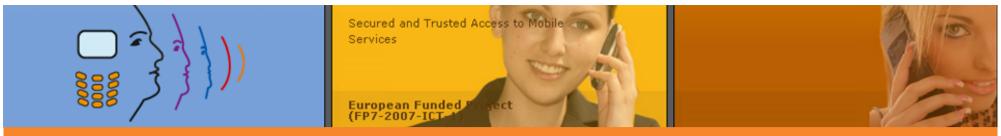
D3.3 Advanced Unimodal System

UMAN

MOBIO Review Meeting, Sep.16-17, 2009

EyePmedia – 1020 Renens





D3.3 UMAN Advanced Unimodal System

- Recap
 - Shape
 - Appearance
 - Matching
- Guided candidate selection
- Hierarchical implementation
- Results

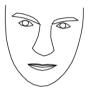






Recap: Shape

- Photograph lots of people
- Label anatomical features on the face
 - Eyes
 - Nose
 - Mouth
- Compute mean shape and modes of deformation











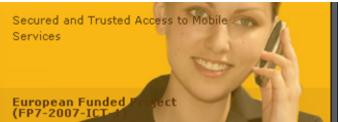












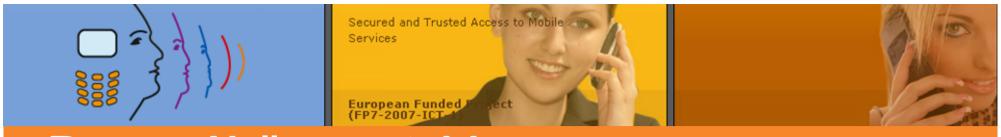


Recap: Appearance

- Photograph lots of people
- Resample image region around each labelled feature
- Compute mean of samples to give template
 - May pre-process image first
 - E.g. compute gradients

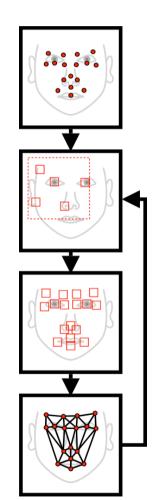




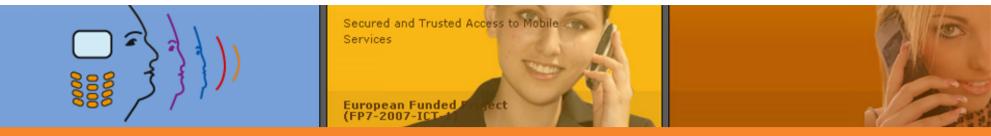


Recap: Naïve matching

- Guess where features are
 - Mean shape wrt detected face
- Search a nearby region for patches that match template
- Pick best match for each feature
- Project selected candidates onto nearest 'valid' face
- Repeat to convergence

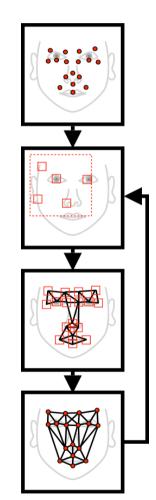






Guided candidate selection

- Best match for each feature selected independently
 - Ignores spatial constraints
 - E.g. both eyes could be assigned to same image patch
- Can be remedied using a Markov Random Field prior
 - Approximates joint distribution
 - Efficient graph solvers (e.g. dynamic programming)

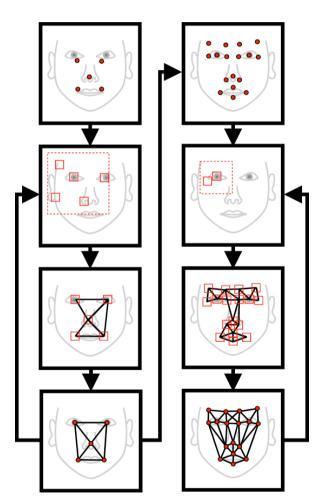




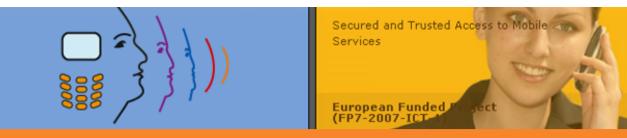


Hierarchical implementation

- Initial uncertainty in scale and orientation is high
 - Need large search regions
 - False matches to undistinctive features are 'distracting'
- Start with most salient features only
 - Eyes, nose, mouth

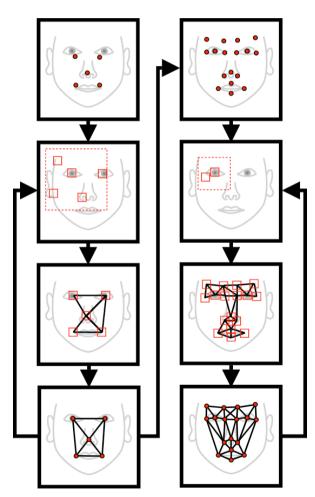


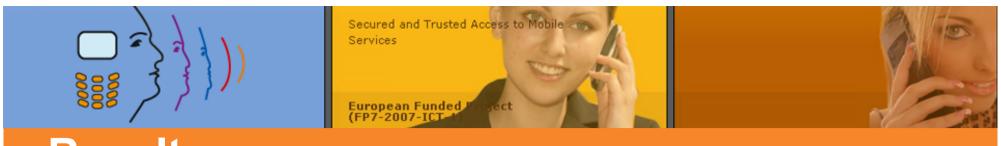




Hierarchical implementation

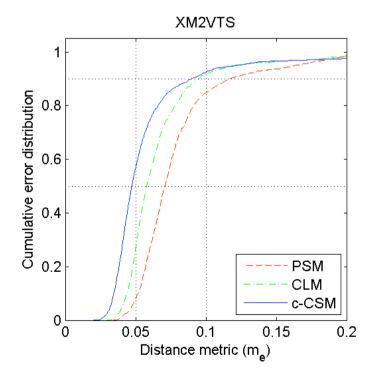
- Uncertainty in scale and orientation gets smaller
- Can make better prediction of other feature positions
- Smaller search regions
 - Fewer false matches
 - More efficient

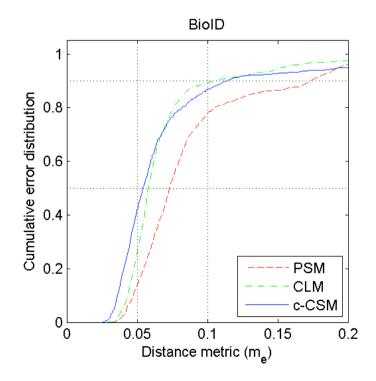




Results

- Tested on standard benchmarks
 - XM2VTS, BioID







Discussion

- Results compare favourably with previous methods
 - Clearly outperforms baseline on XM2VTS
 - Outperforms baseline on BioID in ~70% of cases
- Recently presented at BMVC (Sep 2009)



Ongoing research

- Improved image representations
 - More accurate detections (i.e. close to true solution)
 - Fewer false matches
- Optimized Markov Random Field structure
 - Maximize accuracy for a given complexity
 - Minimize complexity for a given accuracy
- Exploiting sequence dynamics for efficient tracking of facial features



Thank you for your attention