Accurate Personal Identification using Finger Vein and Finger Knuckle Biometric Images



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Multimodal Systems

Bimodal Systems

- Simultaneous Imaging, Single Shot
- Finger Imaging \rightarrow Fingerprint and Fingervein
- Finger Imaging \rightarrow Fingerprint and Finger Knuckle
- Hand Imaging → Palmprint, Finger Geometry and Hand Geometry
- Face Imaging → Face and Periocular, Iris and Periocular, ...
 Obscured or Changed

Finger Vein Biometric

Key Advantages

- Orientation Large, Robust and Hidden Biometric Feature
- Vascular Structure \rightarrow Unique and Private Identifier
- Identical Twins → Different Vein Structure
- Not Intrusive
- Not Easily Damaged, Obscured or Changed
- Highly Stable and Repeatable
- Extremely Difficult to Fake



Finger Vein Imaging







Imaging Hardware



Imaging and Illumination (810nm)



M. Kono, H. Ueki, and S. Umemura, "A new method for the identification of individuals by using of vein pattern matching of a finger," *Proc. 5th Symp. Pattern Measurement*, pp. 9–12 (*in Japanese*), Yamaguchi, Japan, 2000.

M. Kono, H. Ueki, and S.-i. Umemura, "Near-infrared finger vein patterns for personal identification," *Applied Optics,* vol. 41, no. 35, pp. 7429-7436, December, 2002

Preprocessing

- Matched Image → Registration
- Orientation Alignment using Finger/Images Shape

Normalized Cross-Correlation Coefficient

- Matching Finger Vein Images (aligned ROI)
 - Similarity Score \rightarrow Cross-Correlation Coefficient
 - y_i = IFFT2 [♣FFT2(p) *FFT2(q)], , j = 1...N
 - ♣ → complex conjugate; * → element-by-element multiplication
 - Normalized Cross Correlation $\rightarrow C = \max[Y_{i,i}]^{1/2}$
- Experimental Results
 - Database \rightarrow 678 volunteers, 2 images/person
 - Genuine \rightarrow 678, Impostors \rightarrow 229, 503 (678×677×1/2)
 - "All 678 individuals were perfectly identified"

Limitations

- Proprietary database → Lack of reproducibility
- Only 2 images/person → Reliable? Commercial Interests?



Repeated Line Tracking (2004)

Line Tracking

Improved Imaging, System





- Small No of Repetitions \rightarrow *Insufficient* feature extraction
- Large No of Repetitions \rightarrow *High* computational cost
- At least \rightarrow 3000 (lower limit)

Tracking Results

Number of times a pixel has been tracked



Infrared image (left) and value distribution in the tracking space (right)

Tracking Results

Comparisons



Manually Labelled, RLT Method, and using Matched Filter

26	8	-18	-32	-18	8	26
26	8	-18	-32	-18	8	26
26	8	-18	-32	-18	8	26
26	8	-18	-32	-18	8	26
26	8	-18	-32	-18	8	26
26	8	-18	-32	-18	8	26
26	8	-18	-32	-18	8	26

Tracking Results

Comparisons





Bright Sample: Repeated Line Tracking and using Matched Filter



Dark Sample: Repeated Line Tracking and using Matched Filter

Matching Binarized Images

- Downsampling, Translation and Matching \rightarrow Highest Score
- Mismatch Ratio (Normalized by vein pixels in two images)
- Database → 678 Volunteers (Same)
- EER $\rightarrow 0.145\%$



Limitations

- No comparison with earlier (Hitachi) work
- Proprietary database → Lack of reproducibility
- Only 2 images/person \rightarrow Least reliable, Commercial

Local Maximum Curvature (2007)

Multiple Profiles





(a) Vertical direction (d1).





(c) Oblique direction (1) (d3).

(d) Oblique direction (2) (d4)

- Computing Curvature
 - **Discrete Lines**
 - Binarization \rightarrow Otsu's Method
 - Same Dataset (678 Subjects)





N. Miura, A. Nagasaka, and T. Miyatake, "Extraction of finger-vein patterns using maximum curvature points" in image profiles," ICICE Transactions, August 2007.

Finger Vein Imaging



25mm

Webcam

FingerVein Images Dataset

Region of Interest Segmentation

Pre-Processing



> Sample Example

Acquired image to segmented ROI











Region of Interest Segmentation

Pre-Processing



Sample Example (Poor Quality)

Acquired image to segmented ROI



- Mask → Estimation of Orientation (centroid & moments)
- Rotational Alignment of ROI

Region of Interest Enhancement

Pre-Processing



Image Enhancement

- Method → HistEq (Img Avg Background Illumination)
- Sample Results



Feature Extraction

Gabor Filter and Morphological Processing

- Set of Filters \rightarrow Extract Vein Structure
- $f(x,y) = \max_{\forall n=1,2,\dots\Omega} \{ \hat{h}_{\theta_n}(x,y) \star v(x,y) \}$







Feature Extraction

Morphological Operations and Feature Encoding

- Morphological Operations \rightarrow Enhance clarity of vein patterns

$$z(x,y) = f(x,y) - (f(x,y) \ominus b) \circledast b$$

• SE \rightarrow *b*, Grey scale erosion/dilation, top-hat operation





Feature Encoding

•
$$R(x,y) = \begin{cases} 255 & if \ z(x,y) > 0 \\ 0 & if \ z(x,y) \le 0 \end{cases}$$

Generating Match Score

Finger Vein Match Score

- Robust \rightarrow Accommodate translational and rotational variations
- Binarized feature map R and $T \rightarrow$ Match score

$$S_{v}(R,T,M_{R},M_{T}) = \min_{\forall i \in [0,2w], \forall j \in [0,2h]} \left(\frac{\sum_{x=1}^{m} \sum_{y=1}^{n} \odot \left(\widehat{R}(x+i,y+j), T(x,y), M_{R}(x+i,y+j), M_{T}(x,y) \right)}{\sum_{x=1}^{m} \sum_{y=1}^{n} M_{R}(x,y) \cap M_{T}(x,y)} \right)$$

• Masks $\rightarrow M_R$, M_T , Automatically generated

$$M = \left\{ (x, y) \middle| \forall (x, y) \in I, I(x, y) \neq I_{bg} \right\}$$

Sample Results



Sample results from different feature extraction methods: (a) *enhanced* finger vein image, (b) output from matched filter, (c) output from repeated line tracking, (d) output from maximum curvature, (e) output from Gabor filters, and (f) output from morphological operations on (e)

HK PolyU Fingervein Database

- World's First Publicly/Freely Accessible Database
- Two Session Database, 6264 Images
- First Session \rightarrow 156 Subjects, Second Session \rightarrow 105 Subjects
- Six Images → Each from Index and Middle Fingers

Two Session Experiments (Protocol A)

Three Sets \rightarrow Individual Fingers and Combination

- Genuine Scores \rightarrow 630 (105 \times 6)
- Imposter Scores \rightarrow 65,520 (105 \times 104 \times 6)
- Combination → Index and Middle Finger. 210 Class
- Genuine Scores \rightarrow 1260 (210 \times 6)
- Imposter Scores \rightarrow 263,340 (210 \times 209 \times 6)

Two Session Experiments (Protocol A)

Comparative Results \rightarrow Individual Fingers and Combination

Approach	Index Finger	Middle Finger	Index and Middle Finger
Even Gabor with Morphological	7.14%	12.39%	9.31%
Repeated line tracking [5]	15.28%	18.59%	16.70%
Maximum curvature [6]	15.41%	18.06%	16.61%
Matchedfilter	8.60%	11.87%	10.00%
Even Gabor	6.50%	10.12%	8.10%
Table 2: Performance from fir	iger vein matchi	ng with various ap	proaches <u>with</u> mask
Approach	Index Finger	Middle Finger	Index and Middle Finger
Even Gabor with Morphological	3.33%	6.99%	4.91%
Repeated line tracking [5]	15.60%	18.18%	16.43%
Maximum curvature [6]	10.96%	11.08%	10.99%
Matched filter	4.84%	7.81%	5.31%
Even Gabor	3.82%	7.08%	4.61%

Table 1: Performance from finger vein matching with various approaches without mask

➤ Two Session Experiments (Protocol A) Comparative Results→ Individual Fingers and Combination



A. Kumar and Y. Zhou, "Human identification using finger images," IEEE Trans. Image Processing, vol. 21, pp. 2228-2244, April 2012

Single Session Experiments (Protocol B, Larger Subjects) Comparative Results Individual Fingers and Combination

	Table 4. Fertormanee from miger vent materning with various approaches without mask							
Approach	Index Finger	Middle Finger	Index and Middle Finger					
Even Gabor with Morphological	1.16%	2.24%	1.71%					
Repeated line tracking [5]	5.57%	7.77%	6.38%					
Maximum curvature [6]	2.59%	3.73%	3.27%					
Matchedfilter	1.70%	1.75%	1.71%					
Even Gabor	0.89 %	1.71%	1.22%					

Table 4: Performance from finger vein matching with various approaches without mask

Table 5: Performance from finger vein matching with various approaches with mask

Approach	Index Finger	Middle Finger	Index and Middle Finger		
Even Gabor with Morphological	0.43%	0.96%	0.65%		
Repeated line tracking [5]	6.54%	9.95%	8.25%		
Maximum curvature [6]	2.20%	3.13%	2.65%		
Matchedfilter	1.88%	2.10%	1.89%		
Even Gabor	0.54%	1.16%	0.80%		

Two Session Experiments (Protocol A, using CNN)

Lightened CNN Architecture



C. Xie and A. Kumar, "Finger Vein Identification using Convolutional Neural Networks," *Technical Report No. COMP-K-25*, The Hong Kong Polytechnic University, Dec. 2016.

Light CNN Architecture

- Light CNN introduced in [A]
- Maxout \rightarrow less parameters
- MFM (Max Feature Map)



Туре	Filter Size /Stride	Output Size	#Params
Conv1	$5 \times 5/1, 2$	$128 \times 128 \times 96$	2.4K
MFM1	-	$128\times128\times48$	-
Pool1	$2 \times 2/2$	$64 \times 64 \times 48$	-
Conv2a	$1 \times 1/1$	$64 \times 64 \times 96$	4.6K
MFM2a	-	$64 \times 64 \times 48$	-
Conv2	$3 \times 3/1, 1$	$64 \times 64 \times 192$	165K
MFM2	-	$64 \times 64 \times 96$	-
Pool2	$2 \times 2/2$	$32 \times 32 \times 96$	-
Conv3a	$1 \times 1/1$	$32 \times 32 \times 192$	18K
MFM3a	-	$32 \times 32 \times 96$	-
Conv3	$3 \times 3/1, 1$	$32 \times 32 \times 384$	331K
MFM3	-	$32 \times 32 \times 192$	-
Pool3	$2 \times 2/2$	$16 \times 16 \times 192$	-
Conv4a	$1 \times 1/1$	$16 \times 16 \times 384$	73K
MFM4a	-	$16 \times 16 \times 192$	-
Conv4	$3 \times 3/1, 1$	$16 \times 16 \times 256$	442K
MFM4	-	$16\times 16\times 128$	-
Conv5a	$1 \times 1/1$	$16 \times 16 \times 256$	32K
MFM5a	-	$16 \times 16 \times 128$	-
Conv5	$3 \times 3/1, 1$	$16 \times 16 \times 256$	294K
MFM5	-	$16 \times 16 \times 128$	-
Pool4	$2 \times 2/2$	$8 \times 8 \times 128$	-
fc1	-	512	4,194K
MFM_fc1	-	256	-
Total	-	-	5,556K

[A] X. Wu et al., "A Light CNN for Deep Face Representation with Noisy Labels," https://arxiv.org/abs/1511.02683 Nov. 2016

C. Xie and A. Kumar, "Finger Vein Identification using Convolutional Neural Networks," *Technical Report No. COMP-K-25*, The Hong Kong Polytechnic University, Dec. 2016.

Experimental Results using Light CNN

EER of 13.27% (Independent Second Session Test Data)



[A] X. Wu et al., "A Light CNN for Deep Face Representation with Noisy Labels," https://arxiv.org/abs/1511.02683. Nov. 2016

DCNN (VGG) with cross entropy loss

> Architecture



K. Simonyan and A. Zisserman, "Very deep convolutional networks for large-scale image recognition," Proc. ICLR, 2015.

Results → Convolutional Neural Network



Two Session Experiments (Comparative Results using Public Database)

Key Conclusions

- Generally SDH delivers superior performance (better ROC and also notable improvement in EER)
- DCNN with cross entropy loss has similar effect on SDH, but cannot combined with SDH directly
- Log scale and the modified TFS structure can improve performance (evident from ROC but less noticeable for EER)
- Triplet loss has similar effect as TFS
- State of art (TIP2012) \rightarrow GAR of over 0.6 @FAR of 1e-05 (slide 24)

In summary, achieved accuracy fails to match those from using the method detailed in TIP 2012 reference

(more details available in the following reference)

C. Xie and A. Kumar, "Finger Vein Identification using Convolutional Neural Networks," *Technical Report No. COMP-K-25*, The Hong Kong Polytechnic University, Dec. 2016.

Synthesizing Finger Vein Images

Summary of Public Databases

Database	Ref.	Size	Sessions	Public
Hong Kong	[KZ12]	6264 images from 156 subjects, 2 fingers per	2	Yes
Polytechnic University		subject		
SDUMLA-HMT	[YLS11]	3816 images, 6 fingers per subjects	1	Yes
University of Twente	[TV13]	1440 images of 60 subjects	2	No
FV-USM	[MASR14]	5904 images of 123 subjects, 492 different	2	Yes
		finger classes		
CFVD	[ZLL+13]	1345 images of 13 subjects, 130 different	2	No
		fingers		

Which is Real? Which is Synthesized?



F. Hillerström and A. Kumar, "On generation and analysis of synthetic finger-vein images for biometrics identification," *Technical Report No. COMP-K-17*, June 2014, <u>http://www.comp.polyu.edu.hk/~csajaykr/COMP-K-17.pdf</u>

Finger Knuckle Identification

Motivation

- Limitations of Traditional Biometrics
- Multimodal Biometrics, Identification At-A-Distance
- Anatomy of Hands \rightarrow Uniqueness of Knuckle, Correlation with DNA



■ Forensic Identification → Only Piece of Evidence from Suspects



Online Finger Knuckle Identification

KnuckleCodes (BTAS 2009)

Automated Segmentation \rightarrow Efficient ROI Matching using KnuckleCodes



A. Kumar and Y. Zhou, "Human identification using knucklecodes," *Proc. 3rd Intl. Conf. Biometrics, Theory and Applications*, BTAS'09, pp. 147-152, Washington DC, USA, Sep. 2009



Localized Radon Transform



Select the direction which results in minimum (maximum) magnitude

Match Score Generation

Matching KnuckleCodes

- Partially Matching Knuckles \rightarrow Translation and Rotation of Fingers
- Matching Score for two Z-bit KnuckleCodes

$$S(\mathbf{R}, \mathbf{T}) = \min_{\forall i \in [0, 2w], \forall j \in [0, 2h]} \left(\sum_{x=1}^{m} \sum_{y=1}^{n} \phi\left(\widehat{\mathbf{R}}(x+i, y+j), \mathbf{T}(x, y)\right) \right)$$
$$w = \operatorname{floor}\left(\frac{m}{3}\right), \mathbf{h} = \operatorname{floor}\left(\frac{n}{3}\right)$$
$$\widehat{\mathbf{R}}(x, y) = \begin{cases} \mathbf{R}(x-w, y-h) & x \in [w+1, w+m], \ y \in [h+1, \ h+n] \\ -1 & \text{otherwise} \end{cases}$$
$$\phi(J_b, K_b) = \begin{cases} 0 & \text{if } J_b = K_b \ \forall b \\ 0 & \text{otherwise} \end{cases} \quad b = 1, 2, -7 \end{cases}$$

• Size of KnuckleCodes \rightarrow One fourth of knuckle image size ($X_p = 2$)

Experiments

- 158 Subjects, 5 Images per Subject, Age group \rightarrow 16-55 year
- Unconstrained (peg-free) imaging
- Five-fold Cross-Validation, Average of Results
- Genuine Scores \rightarrow 790 (158 \times 5)
- Imposter Scores \rightarrow 124030 (158 \times 157 \times 5)
- Comparative Performance using (even) Gabor filters
 - $f = 1/(2\sqrt{2})$, 12 filters, 15×15 mask size



KnuckleCodes generated for knuckle image in (a) using LRT in (b), and using even Gabor filters in (c)

- Results
 - Comparative Receiver Operating Characteristics



Results

Performance Analysis

	Equal Error Rate					
X_p	1		2	3		4
1	13	15	14	13 15		14
Mean (%)	1.15	1.15	1.08	2.78	2.53	6.96
Std deviation (%)	1.57	1.57	1.08	0.96	1.48	0.9

•	Equal Error Rate						
D (Intervals in $0-\pi$)		6	8	10	12	14	16
KnucklaCodas (Padan)	Mean (%)	2.03	1.08	1.29	1.14	1.27	1.29
KnuckieCodes (Radoll)	Std deviation (%)	1.37	1.08	1.59	1.37	1.60	1.24
ViewaltlaCadas (Cabar)	Mean (%)	4.18	11.14	5.82	2.66	3.29	7.59
KnuckieCodes (Gabol)	Std deviation (%)	2.31	1.88	1.04	1.81	2.26	2.24

KnuckleCodes generated for knuckle image in (a) using LRT in (b), and using even Gabor filters in (c)

Results

Cumulative Match Characteristics



Minor Finger Knuckle



Second Minor Finger Knuckle?

A. Kumar, "Importance of being unique from finger dorsal patterns: Exploring minor finger knuckle patterns in verifying human identities," *IEEE Trans. Information Forensics & Security*, vol. 9, pp. 1288-1298, August 2014.

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Collaborators

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- Zhihuan Xu
- Bichai Wang
- Cihui Xie
- Ch. Ravikanth

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