



# **NFIQ 2.0 – Features for fingerprint quality determination**

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# Outline

Introduction

NFIQ 2.0 Quality features

Quality features

Two ground-truth classes

Quality feature example - frequency domain analysis

Actionable feedback

Speeding up NFIQ 2.0

NFIQ 2.0 and WSQ compression

Alignment with international standard

Contact & further information



- ▶ Starting point for features
  - ▶ NFIQ 1.0
  - ▶ ISO/IEC TR 29794-4:2010
  - ▶ Literature



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- ▶ Implementation of prototype features
- ▶ Hundreds of variations of features; parameter configurations and variations in algorithm steps



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- ▶ Hundreds of variations of features; parameter configurations and variations in algorithm steps
- ▶ Iterative development to arrive at NFIQ 2.0 feature vector
- ▶ Prioritize predictive power and speed of computation
- ▶ Workshops central to development of features



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- ▶ Selecting features
  - ▶ Speed of computation
  - ▶ Contribution to predictive performance





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$$\mathbf{Q}_{\text{NFIQ 2.0}} = \left( Q_{\text{FDA}}^{\mu}, Q_{\text{LCS}}^{\mu}, Q_{\text{OCL}}^{\mu}, Q_{\text{OFL}}^{\mu}, Q_{\text{RVU}}^{\mu}, \right. \\ Q_{\text{FDA}}^{\sigma}, Q_{\text{LCS}}^{\sigma}, Q_{\text{OCL}}^{\sigma}, Q_{\text{OFL}}^{\sigma}, Q_{\text{RVU}}^{\sigma}, \\ \mathbf{Q}_{\text{FDA}}, \mathbf{Q}_{\text{LCS}}, \mathbf{Q}_{\text{OCL}}, \mathbf{Q}_{\text{OFL}}, \mathbf{Q}_{\text{RVU}}, \\ Q_{\text{MU}}, Q_{\text{MMB}}, Q_{\text{COH}}^{\text{rel}}, Q_{\text{COH}}^{\text{sum}}, Q_{\text{AREA}}^{\mu}, \\ \left. Q_{\text{MIN}}^{\text{cnt}}, Q_{\text{MIN}}^{\text{com}}, Q_{\text{MIN}}^{\text{mu}}, Q_{\text{MIN}}^{\text{ocl}} \right).$$



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- ▶ Mean and standard deviation of local features
- ▶ Histogram of local features (boundaries determined from CDF)



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- ▶ Mean and standard deviation of local features
- ▶ Histogram of local features (boundaries determined from CDF)
- ▶ Classifier
  - ▶ Random Forest trained for binary classification
  - ▶ Input: 69 dimensional feature vector
  - ▶ Output: probability of input being Class 1 (high utility) quantized [1, 100]



Name	Capture mode	Type	Number of subjects	Fingers	Number of comparisons per finger	Used for
AZLA	Scanned ink	Operational	240,000	Index and Thumb	120,000 mated. 120,000 non-mated	training + testing
POEBVA	Live scan	Operational	180,000	Index	120,000 mated. 120,000 non-mated	training + testing
VISITIDF	Live scan	Operational	220,000	Index and Thumb	95,000 mated. 120,000 non-mated	training + testing
DHS2	Live scan	Operational	180,000	Index	120,000 mated. 120,000 non-mated	training + testing
IQMI	Scanned ink	Operational	250,000	10 fingers	250,000 mated. 250,000 non-mated	testing
BAK	Live scan	Operational	342,000 images	10 fingers	—	testing
BAK + SD 29	Scanned ink	Public	209	10 fingers	1912 mated. 35,791 non-mated	testing
FVC 2000 DB1	Live scan	Public	110	8 fingers	—	compliance testing
FVC 2000 DB3	Live scan	Public	110	8 fingers	—	compliance testing
FVC 2002 DB1	Live scan	Public	110	8 fingers	—	compliance testing

► Data from operational sources (Optical sensors)



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- ▶ Data from operational sources (Optical sensors)
- ▶ Training set 6629 images (3295 in Class 0 and 3334 in Class 1)
- ▶ Validation set 99797 randomly selected images
- ▶ External validation on BKA data and FBI data



- ▶ Criteria for two classes of samples in training





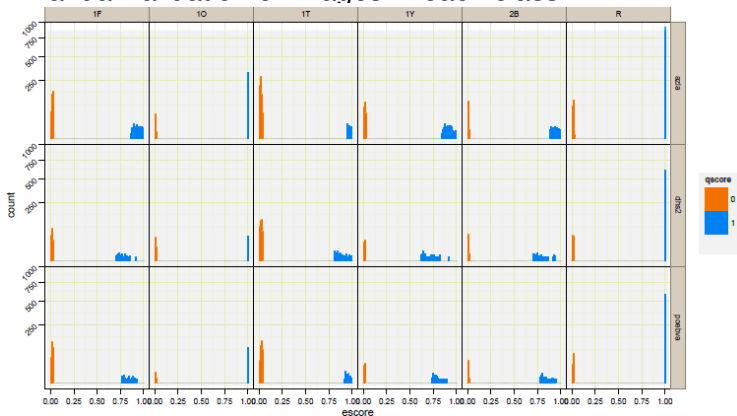
- ▶ Criteria for two classes of samples in training
  - 1 NFIQ=1 ( $S_{act} > 0.7$ ) and  $S_{gen}$  in 90th percentile
  - 0 NFIQ=5 ( $S_{act} > 0.9$ ) and  $S_{gen} < t$  at  $FMR = 10^{-4}$



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## Feature importance ranking

	Name	MeanDreaseGini
$Q_{FDA}^{\sigma}$	Frequency Domain Analysis_Standard Deviation	140.760
$Q_{MIN}^{com}$	FingerJet FX OSE COM Minutiae Count	92.089
$Q_{MIN}^{ocl}$	FingerJet FX OSE OCL MinutiaeQuality	83.027
$Q_{RVU}^{\mu}$	Ridge Valley Uniformity_Mean	69.517
$Q_{FDA}^{\mu}$	Frequency Domain Analysis_Mean	62.229
$Q_{MIN}^{cnt}$	FingerJet FX OSE Total Minutiae Count	57.565
$Q_{RVU}^{\sigma}$	Ridge Valley Uniformity_Standard Deviation	50.946
$Q_{LCS}^7$	Local Clarity Score_Bin_7	50.688
$Q_{LCS}^8$	Local Clarity Score_Bin_8	50.100
$Q_{FDA}^9$	Frequency Domain Analysis_Bin_9	47.844
$Q_{COH}^{sum}$	ROI Orientation Map Coherence Sum	38.104
$Q_{OFL}^2$	Orientation Flow_Bin_2	37.172
$Q_{LCS}^{\mu}$	Local Clarity Score_Mean	36.483
$Q_{RVU}^5$	Ridge Valley Uniformity_Bin_5	35.617
$Q_{RVU}^3$	Ridge Valley Uniformity_Bin_3	35.139
$Q_{AREA}^{\mu}$	ROI Area Mean	34.932
$Q_{OFL}^1$	Orientation Flow_Bin_1	33.751
$Q_{OFL}^0$	Orientation Flow_Bin_0	33.513
$Q_{MU}$	MU	32.914



## ► $Q_{FDA}$ local determination of ridge-valley signature

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### Algorithm 3: fda algorithm

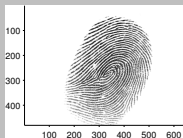
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Input: Fingerprint image  $I$

Output: fda quality score  $Q_{FDA}$

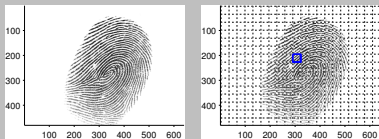
```
1 for each block  $V$  in  $I$  do
2   pad  $V$  with 2 pixel around border
3   rotate  $V$  with nearest neighbour interpolation such that dominant ridge flow is perpendicular to x-axis
4   crop  $V$  such that no invalid regions are included
5   with  $V$  obtain the ridge-valley signature  $T$  (eq. (11))
6   compute the dft of  $T$  to obtain the magnitude representation  $A$ 
7   discard the first component of  $A$ 
8   determine  $F_{max}$  as the index of the largest magnitude in  $A$ 
9   compute  $Q_{FDA}^{local}$  of  $V$  using  $A$  and  $F_{max}$  (eq. (12))
10 end
```

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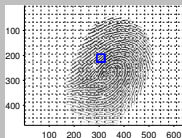
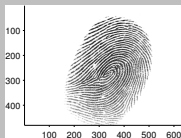


# Quality feature example - frequency domain analysis





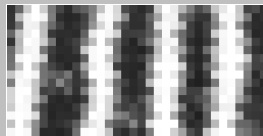
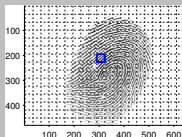
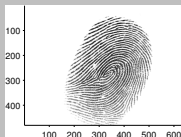
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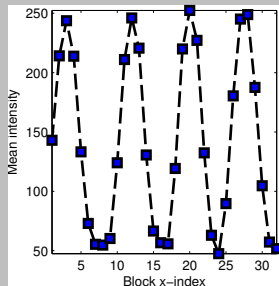
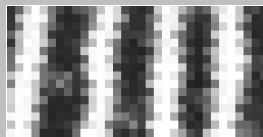
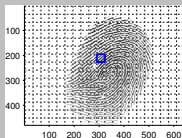
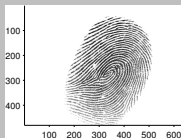






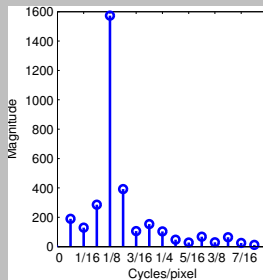
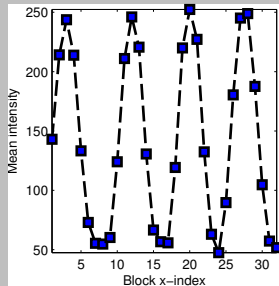
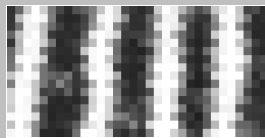
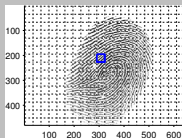
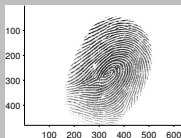
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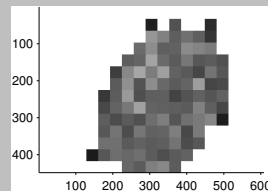
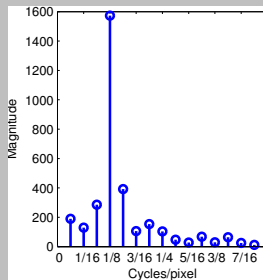
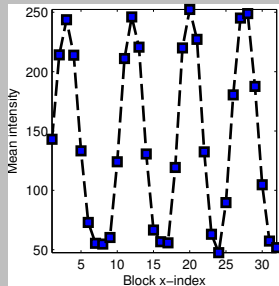
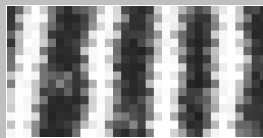
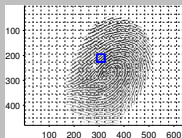
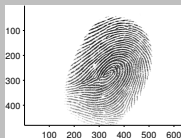


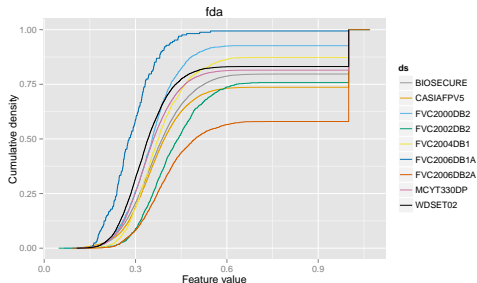
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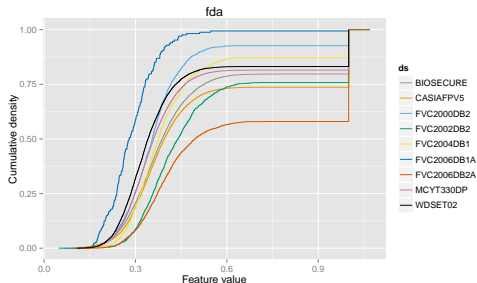




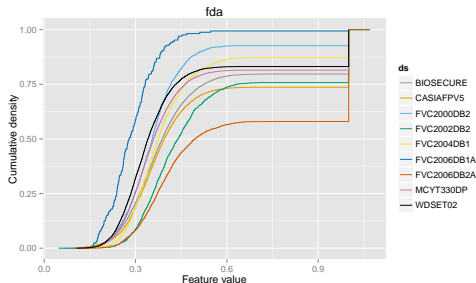
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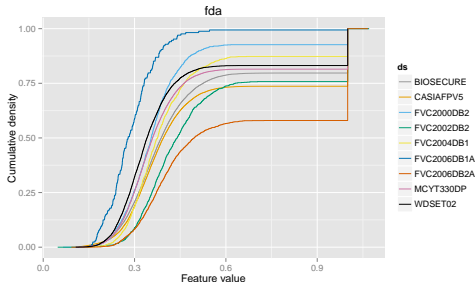


$$B_{\text{FDA}} = \{ -\infty, 0.26800, 0.30400, 0.33000, 0.35500, \\ 0.38000, 0.40700, 0.44000, 0.50000, 1.00000, \infty \} .$$



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- ▶ Local quality values  $\Rightarrow$  fixed length feature vector
- ▶ Mean, std.dev., 10 bin histogram  $\Rightarrow$  12-dimension feature vector



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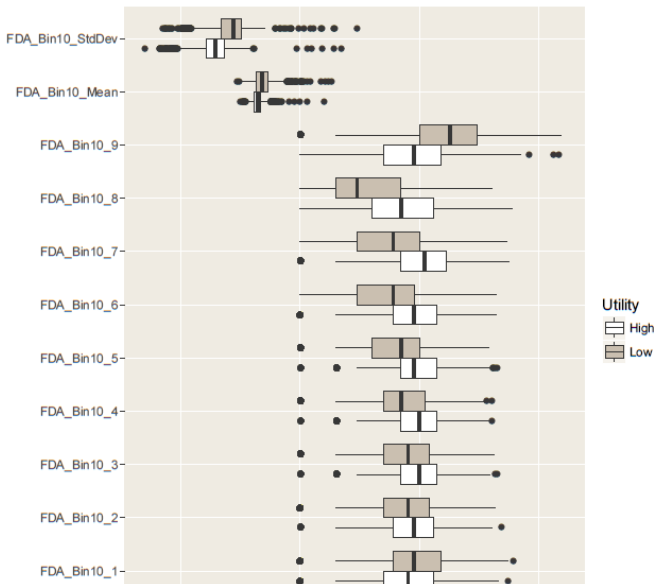
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$$Q_{\text{FDA}} : \{ Q_{\text{FDA}}^{\mu}, Q_{\text{FDA}}^{\sigma}, Q_{\text{FDA}}^1, \dots, Q_{\text{FDA}}^{10} \}$$





# Quality feature example - frequency domain analysis



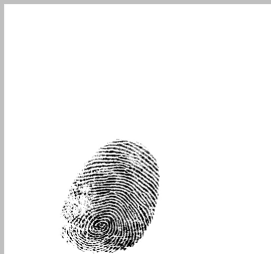


- ▶ Request  $\Rightarrow$  near frame rate quality assessment (10 Hz)





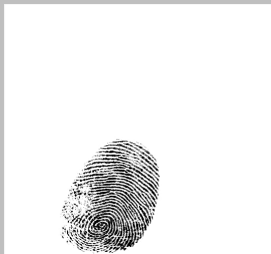
- ▶ Request  $\Rightarrow$  near frame rate quality assessment (10 Hz)
- ▶ Slap sensors provide large finger images



800  $\times$  750 pixel sensor output reproduced at 25% scale



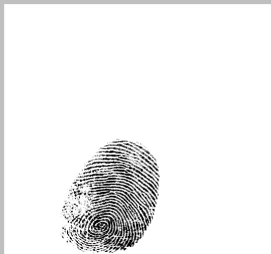
- ▶ Request  $\Rightarrow$  near frame rate quality assessment (10 Hz)
- ▶ Slap sensors provide large finger images
- ▶ Removal of near constant area
- ▶ No processing of background area blocks



800  $\times$  750 pixel sensor output reproduced at 25% scale



- ▶ Request  $\Rightarrow$  near frame rate quality assessment (10 Hz)
- ▶ Slap sensors provide large finger images
- ▶ Removal of near constant area
- ▶ No processing of background area blocks
- ▶ Avoid removing low quality fingerprint areas



800  $\times$  750 pixel sensor output reproduced at 25% scale



$330 \times 286 = 94380$   
(15.7%)



$$330 \times 286 = 94380 \\ (15.7\%)$$



$$330 \times 286 - (10 \times (32 \times 32)) = 84140 \\ (13.9\%)$$



- ▶ Demand for actionable feedback from quality algorithm
- ▶ More than a quality score - helps to answer the why
- ▶ Provide information  $\Rightarrow$  improve quality at recapture





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  - ▶ empty image ( $\mu > 250$ )
  - ▶ uniform image pixel intensity ( $\sigma = 1.0$ )
  - ▶ no or few minutiae detected ( $N_{min} < 5$ )
  - ▶ small foreground area ( $N_{fgnd} < 50000$ )



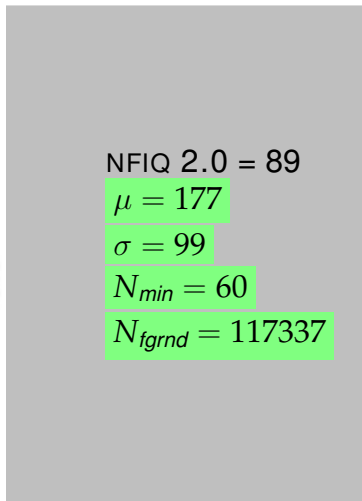
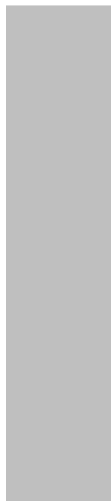


empty image  
few minutiae

$$\mu > 250$$
$$N_{min} < 5$$

uniform image intensity  
small foreground

$$\sigma = 1.0$$
$$N_{fgrnd} < 50000$$



NFIQ 2.0 = 89

$$\mu = 177$$

$$\sigma = 99$$

$$N_{min} = 60$$

$$N_{fgrnd} = 117337$$



empty image  
few minutiae

$$\mu > 250$$
$$N_{min} < 5$$

uniform image intensity  
small foreground

$$\sigma = 1.0$$
$$N_{fgnd} < 50000$$



NFIQ 2.0 = 21

$$\mu = 220$$

$$\sigma = 64$$

$$N_{min} = 40$$

$$\uparrow N_{fgnd} = 36887$$



empty image  
few minutiae

$$\mu > 250$$
$$N_{min} < 5$$

uniform image intensity  
small foreground

$$\sigma = 1.0$$
$$N_{fgnd} < 50000$$



$$\text{NFIQ 2.0} = 1$$

$$\mu = 196$$

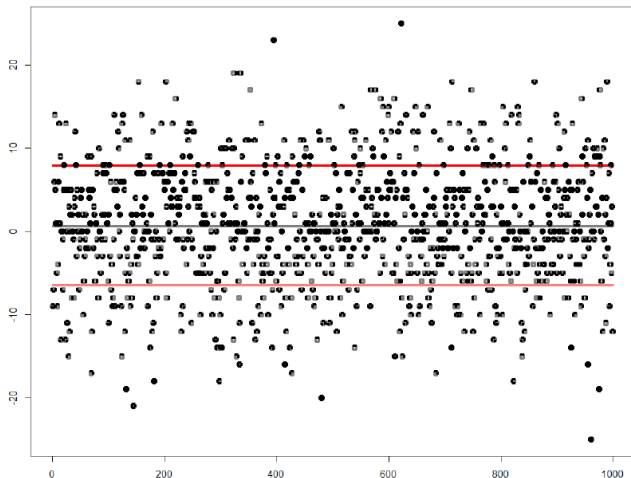
$$\sigma = 79$$

$$\uparrow N_{min} = 0$$

$$\uparrow N_{fgnd} = 16262$$



- Deviation between uncompressed and WSQ compressed (factor 8). 1000 images, MCYT 330 DP.

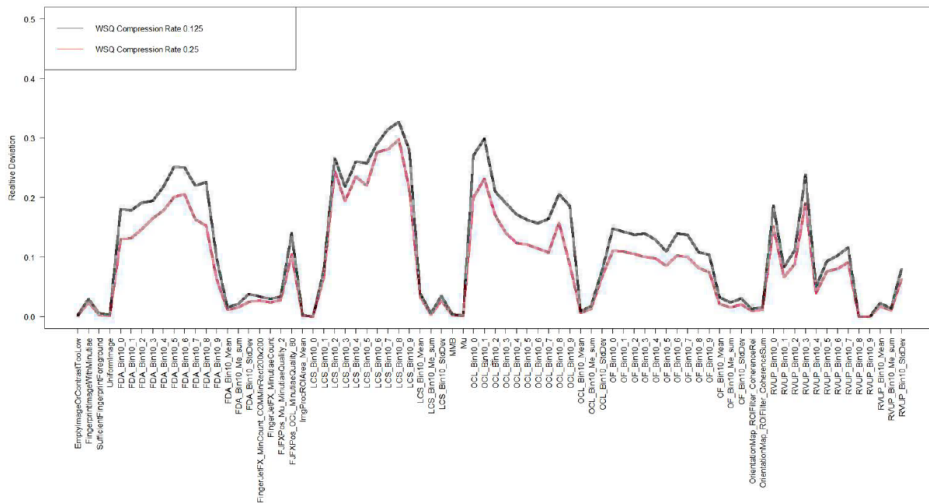






- Fingerprint boundary artifact at WSQ compression (factor 8). Gamma adjusted.







- ▶ Standardization of features a priority throughout NFIQ 2.0 development



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- ▶ 29794-4 – biometric sample quality – finger image data
  - ▶ current status is 3rd Committee Draft
  - ▶ progression to Draft International Standard in May 2016
  - ▶ projected release as International Standard in 2017



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  - ▶ progression to Draft International Standard in May 2016
  - ▶ projected release as International Standard in 2017
- ▶ NFIQ 2.0 effectively a reference implementation of 29794-4 at this point
  - ▶ Open source, publicly available



# Thanks for your attention

Martin A. Olsen

Contact: `martin.olsen@{cased.de; ntnu.no}`

**NFIQ 2.0** `nist.gov/itl/iad/ig/development_nfiq_2.cfm`

**Prototype quality features** `share.nbl.nislabs.no/public`