



**International
Standard**

ISO/IEC 29794-4

**Information technology —
Biometric sample quality —**

**Part 4:
Finger image data**

*Technologies de l'information — Qualité d'échantillon
biométrique —*

Partie 4: Données d'image de doigt

**Second edition
2024-09**

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Published in Switzerland

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Foreword

ISO (the International Organization for Standardization) and IEC (the International Electrotechnical Commission) form the specialized system for worldwide standardization. National bodies that are members of ISO or IEC participate in the development of International Standards through technical committees established by the respective organization to deal with particular fields of technical activity. ISO and IEC technical committees collaborate in fields of mutual interest. Other international organizations, governmental and non-governmental, in liaison with ISO and IEC, also take part in the work.

The procedures used to develop this document and those intended for its further maintenance are described in the ISO/IEC Directives, Part 1. In particular, the different approval criteria needed for the different types of document should be noted. This document was drafted in accordance with the editorial rules of the ISO/IEC Directives, Part 2 (see www.iso.org/directives or www.iec.ch/members_experts/refdocs).

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This document was prepared by Joint Technical Committee ISO/IEC JTC 1, *Information technology*, Subcommittee SC 37, *Biometrics*.

This second edition cancels and replaces the first edition (ISO/IEC 29794-4:2017), which has been technically revised.

The main changes are as follows:

- algorithms for normalization of finger image quality components have been added, along with new quality algorithm identifiers for the unique identification of the quality measures defined in this document;
- [Annex A](#) has been technically revised to reflect a new conformance test set.

A list of all parts in the ISO/IEC 29794 series can be found on the ISO and IEC websites.

Any feedback or questions on this document should be directed to the user's national standards body. A complete listing of these bodies can be found at www.iso.org/members.html and www.iec.ch/national-committees.

Introduction

This document specifies finger image quality measures. A reference implementation of the normative measures — NFIQ 2 — is available at Reference [16], which is described in more detail by the developers in Reference [1].

The quality of finger image data is determined by the degree to which the finger image data fulfils specified requirements for the targeted application. Information on quality is therefore useful in many applications. ISO/IEC 19784-1 allocates a quality field and specifies the allowable range for the scores, with a recommendation that the score be divided into four categories with a qualitative interpretation for each category. Finger image quality fields are provided in the finger image data interchange formats standardized in ISO/IEC 19794-4 and ISO/IEC 39794-4. Finger feature data interchange formats standardized in ISO/IEC 19794-2, ISO/IEC 19794-3, ISO/IEC 19794-8 and ISO/IEC 39794-2 provide finger image quality fields for the source image. To facilitate the interpretation and interchange of finger image quality scores, this document specifies how to calculate the finger image quality score of plain finger images with a spatial sampling rate of 196,85 px/cm and a bit depth of 8 bit for the greyscale pixel intensity values scanned from inked fingerprint cards or captured using optical area sensors based on frustrated total internal reflection.

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Information technology — Biometric sample quality —

Part 4: Finger image data

1 Scope

This document establishes:

- terms and definitions for quantifying finger image quality;
- methods used to quantify the quality of finger images; and
- standardized encoding of finger image quality;

for finger images at 196,85 px/cm spatial sampling rate and a bit depth of 8 bit for the greyscale pixel intensity values scanned or captured using optical area sensors in direct contact with friction ridges.

2 Normative references

The following documents are referred to in the text in such a way that some or all of their content constitutes requirements of this document. For dated references, only the edition cited applies. For undated references, the latest edition of the referenced document (including any amendments) applies.

ISO/IEC 2382-37, *Information technology — Vocabulary — Part 37: Biometrics*

ISO/IEC 39794-1, *Information technology — Extensible biometric data interchange formats — Part 1: Framework*

ISO/IEC 29794-1, *Information technology — Biometric sample quality — Part 1: Framework*

3 Terms and definitions

For the purposes of this document, the terms and definitions given in ISO/IEC 2382-37, ISO/IEC 29794-1 and the following apply.

ISO and IEC maintain terminology databases for use in standardization at the following addresses:

- ISO Online browsing platform: available at <https://www.iso.org/obp>
- IEC Electropedia: available at <https://www.electropedia.org/>

3.1

foreground region

set of all pixels of a finger image that form valid finger image patterns

Note 1 to entry: The most evident structural characteristic of a valid finger image is a pattern of interleaved ridges and valleys.

3.2

local region

block of $m \times n$ pixels of the foreground region, where m and n are smaller than or equal to the width and the height of the foreground region respectively

3.3

finger image quality assessment algorithm

algorithm to calculate a quality measure

Note 1 to entry: “Quality assessment algorithm” and “quality algorithm” are synonyms.

3.4

trim

removal of pixels from the top, left, bottom and right sides of a finger image that do not comprise the foreground region

Note 1 to entry: The steps for trimming an image to form the foreground region are defined in [6.1.5.2](#).

4 Abbreviated terms

CBEFF	Common Biometric Exchange File Format
DFT	discrete Fourier transform
DT	determine threshold
FDA	frequency domain analysis
FJFX	FingerJet Fingerprint Feature Extractor, Open Source Edition
LCL	local clarity
NFIQ	NIST Fingerprint Image Quality
OCL	orientation certainty level
OFL	orientation flow
QSND	quality score normalisation dataset
RVU	ridge valley uniformity
TIR	total internal reflection

5 Conformance

A finger image quality assessment algorithm conforms to this document if it conforms to the normative requirements of [Clause 6](#).

A finger image quality block shall conform to this document if its structure and data values conform to the formatting requirements of [Clause 7](#) (finger image quality block) and if its quality values are computed using the methods specified in [6.2](#) and [6.4](#).

A finger image quality assessment implementation conformant to this document may use the biometric organization identifier of ISO/IEC JTC 1/SC 37, which is 257 (101_{Hex}), if it has been tested following the conformance testing methodology in [Clause A.2](#).

Conformance to normative requirements of [Clause 7](#) is achieved by Level 1 and Level 2 conformance as specified in ISO/IEC 39794-1:2019, Annex C. Conformance to normative requirements of [6.2](#) and [6.4](#) is achieved by Level 3 conformance as specified in ISO/IEC 39794-1:2019, Annex C.

The conformance test assertion in [Annex A](#) shall apply.

6 Finger image quality measures

6.1 Overview

6.1.1 General

This clause establishes measures for predicting the utility of a finger image. Image quality measures from a single image are useful to ensure the acquired image is suitable for recognition.

A complete finger image quality analysis shall examine both the local and global structures of the finger image. Fingerprint local structure constitutes the main texture-like pattern of ridges and valleys within a local region while valid global structure puts the ridges and valleys into a smooth flow for the entire fingerprint. The quality of a finger image is determined by both its local and global structures. This clause describes the features and characteristics of finger images at both local and global structures that are to be used as quality components for quantifying finger image quality.

For applying the algorithms as described in 6.2 and 6.3, the finger image shall have a spatial sampling rate of 196,85 pixels per centimetre (500 pixels per inch), a bit depth of 8 bit for the greyscale pixel intensity values, with friction ridges represented by greyscale pixel intensity values lower than those for valleys. The algorithms were developed using images of finger friction ridges in contact with an electronic capture device and inked fingerprints digitized with an electronic scanner. The imaging devices and scanners are considered free from geometric distortion and exhibit greyscale linearity and uniformity.

ISO/IEC 29794-1 requires that quality components be mapped to an integer value between 0 and 100, inclusive.

6.1.2 Methods for mapping to the desired value range

6.1.2.1 Sigmoid function

The mapping of values between 0 and 1 inclusive is accomplished for several quality components with the sigmoid function as shown in [Formula \(1\)](#):

$$\text{sigmoid}(x, x_0, w) = \left(1 + \exp\left(\frac{x_0 - x}{w}\right) \right)^{-1} \quad (1)$$

where

- x is a native quality measure value;
- x_0 is the inflection point at which the function has the value 0,5;
- w is a scaling parameter determining the width of the region in which the function transitions from ε to $1 - \varepsilon$;
- ε is an infinitesimally small positive quantity.

The values computed from the sigmoid function will be mapped to the target value ranges (0 to 100) in subsequent clauses.

6.1.2.2 Known ranges

When the range of values for a given quality measure is known (e.g. from 1 to 250, inclusive), the known range function is used, as shown in [Formula \(2\)](#):

$$\left\lfloor \frac{x - \min(x)}{\max(x) - \min(x) + \varepsilon} 101 \right\rfloor \quad (2)$$

where

- x is a native quality measure value;
- $\lfloor x \rfloor$ is the floor function giving the greatest integer $\leq x$
- ε is an infinitesimally small positive quantity.

6.1.3 Constituent of local quality measures

A finger image is partitioned into local regions such that each local region contains sufficient ridge-valley information, preferably having at least two clear ridges, while not overly constraining high curvature ridges. For images with a spatial sampling rate of 196,85 pixels per centimetre (500 pixels per inch), the ridge separation usually varies between 8 pixels to 12 pixels.^[2] A ridge separation comprises a ridge and a valley. In order to cover two clear ridges, the local region size has to be greater than 24 pixels in both width and height. The size for each local region shall be (32×32) pixels, which is sufficient to cover two clear ridges. Instead of Cartesian coordinate, curvilinear coordinate along the ridge can also be used.

NOTE The size of the local region used during computation of q_{OFL}^{σ} (6.2.16.3), q_{OFL}^{μ} (6.2.16.2), $q_{\text{COH}}^{\text{sum}}$ (6.2.14), and $q_{\text{COH}}^{\text{rel}}$ (6.2.15) in the reference implementation NFIQ 2 prior to version 2.3.0 deviates from size specified in this subclause.

6.1.4 Constituent of global quality measures

A global quality measure shall be computed over the whole finger image after trim to assess the utility of the sample for fingerprint recognition.

6.1.5 Image preprocessing

6.1.5.1 Description

A segmentation process follows where each local region is labelled as background or foreground. There are several segmentation approaches, such as using the average magnitude of the pixel-intensity gradient in each local region.^[2]

This document does not prescribe segmentation methods, but notes that performing segmentation influences several quality components. Constant or near constant areas of the input image shall be removed according to 6.1.5.2 prior to computing quality using the measures specified in 6.2 and 6.3.

See Annex C for the area consideration.

6.1.5.2 Removal of near constant rows and columns in image

Prior to computing quality components, fingerprint images shall be trimmed to remove near constant rows and columns on the margins. Pixel intensities take values $[0, 255]$ for an 8-bit greyscale image. As a first approximation of the region of interest, image columns and rows which are near constant white background are removed. Using the algorithm specified below, a fixed threshold, T_{μ} , is set to the greyscale pixel intensity of 250 to obtain the image without near constant areas.

The algorithm is visualized in Figure 1 and specified as follows:

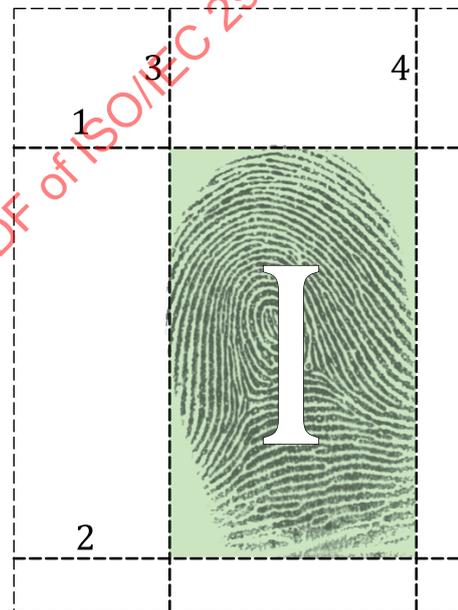
- a) For each row R_i in \hat{I} , starting from the top:
 - 1) compute the row arithmetic mean μ_{row} ;
 - 2) on the first occurrence where $\mu_{\text{row}} \leq T_{\mu}$ set $\text{top} = R_i$;

- 3) on the last occurrence where $\mu_{\text{row}} \leq T_{\mu}$ set $\text{bottom} = R_i$.
- b) For each column C_i in \hat{I} , starting from the left:
 - 1) compute the column arithmetic mean μ_{col} ;
 - 2) on the first occurrence where $\mu_{\text{col}} \leq T_{\mu}$ set $\text{left} = C_i$;
 - 3) on the last occurrence where $\mu_{\text{col}} \leq T_{\mu}$ set $\text{right} = C_i$.
- c) extract the trimmed region of interest, I , as the pixels of \hat{I} encompassed between and including the rows top (a2) and bottom (a3) and the columns left (b2) and right (b3).

where

\hat{I} is the matrix of grey levels corresponding to the pixels of an image;

I is the matrix of grey levels corresponding to the pixels of an image after trim.



a) Image prior to trimming near constant rows and columns on the margins

b) Results of following the steps in [6.1.5.2](#)

Key

- 1 top
- 2 bottom
- 3 left
- 4 right

NOTE 1 In Figure 1 a), the area overlaid in green is a visualisation of \hat{I} .

NOTE 2 In Figure 1 b), the area overlaid in green is a visualisation of I .

NOTE 3 Each subfigure within Figure 1 contains a dashed black border.

Figure 1 — Example of removing near constant white rows and columns from an image

6.1.5.3 Foreground segmentation based on local standard deviation

For quality components that require a foreground mask to indicate regions containing the fingerprint, an algorithm using local standard deviation is adopted.

The algorithm is specified as follows:

- a) Normalize I to zero mean and unit standard deviation to produce I' .
- b) For each local region V in I' :
 - 1) compute the standard deviation of V as σ_v ;
 - 2) mark the corresponding local region in I_{mask} as foreground if $\sigma_v > 0,1$.

6.1.5.4 Computing the dominant ridge flow orientation for a local region from pixel-intensity gradients

The dominant ridge flow orientation is determined by computing the pixel-intensity gradient information and then determining the orientation of the principal variation axis.

The numerical gradient of the local region is determined using finite central difference for all interior pixels in x -direction and y -direction, as shown in [Formulae \(3\)](#) and [\(4\)](#):

$$f_x = \frac{I(x+1, y) - I(x-1, y)}{2} \quad (3)$$

$$f_y = \frac{I(x, y+1) - I(x, y-1)}{2} \quad (4)$$

With f_x and f_y , the dominant ridge flow orientation, $\text{angle}(\mathbf{V})$, is determined analytically using the sine and cosine doubled angle determined from the arithmetic means of the pixel-intensity gradient covariances, as shown in [Formulae \(5\)](#) to [\(12\)](#):

$$a = \overline{f_x^2} \quad (5)$$

$$b = \overline{f_y^2} \quad (6)$$

$$c = \overline{f_x f_y} \quad (7)$$

$$C = \begin{bmatrix} a & c \\ c & b \end{bmatrix} \quad (8)$$

$$d = \sqrt{c^2 + (a-b)^2} \quad (9)$$

$$\sin(\theta) = \frac{c}{d} \quad (10)$$

$$\cos(\theta) = \frac{a-b}{d} \quad (11)$$

$$\text{angle}(\mathbf{V}) = \frac{1}{2} \arctan\left(\frac{\sin(\theta)}{\cos(\theta)}\right) \quad (12)$$

NOTE In Formulae (5), (6) and (7), the use of the overbar indicates the mean of the value.

6.1.6 Image examples

For algorithms operating in a block-wise manner the trimmed input image is subdivided into local regions according to an overlay grid. This is demonstrated in [Figure 2 b\)](#), in which the local region $V(6,1)$ is used as an example in local processing and is marked up using a bold blue line. [Figure 2 c\)](#) shows an enlarged view of $V(6,1)$ and [Figure 2 d\)](#) shows $V(6,1)$ rotated according to its dominant ridge orientation computed using [Formula \(12\)](#).

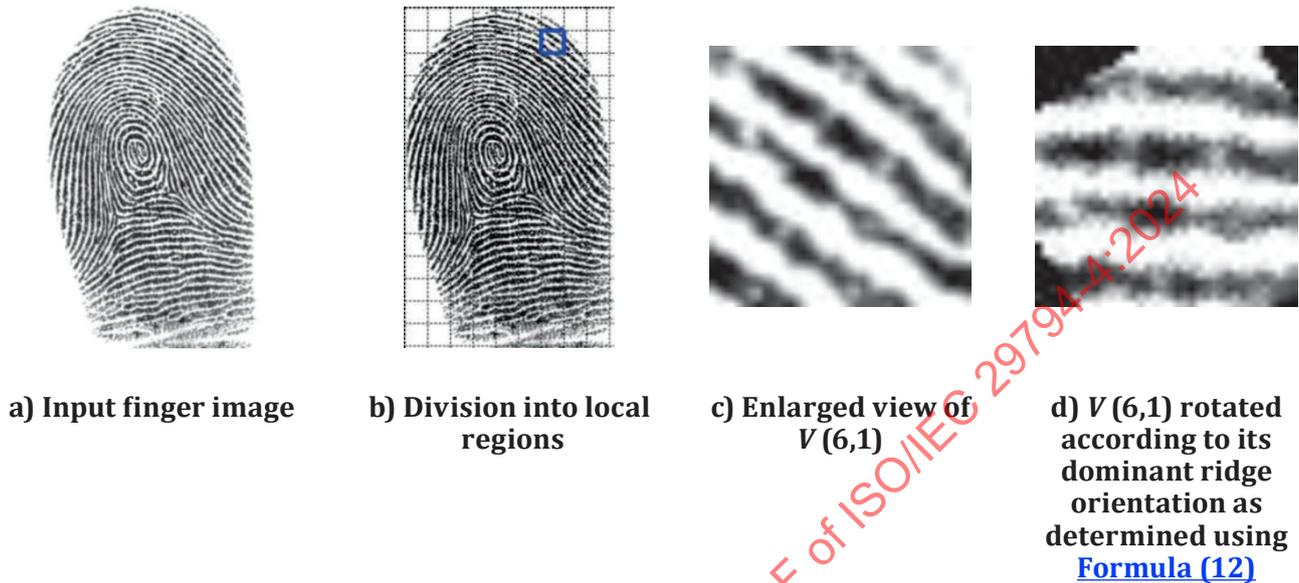


Figure 2 — Example of computing the dominant ridge flow orientation for a local region

6.2 Normative contributive quality components

6.2.1 General

[Subclause 6.2](#) specifies algorithms for computing finger image quality components that contribute to the ISO/IEC 29794-4 quality feature vector and to the computation of the unified quality score.

6.2.2 Orientation certainty level

6.2.2.1 Description

The orientation certainty level (OCL)^[3] of a local region is a measure of the consistency of the orientations of the ridges and valleys contained within this local region. The feature computes local quality and operates in a block-wise manner.

The finger image within a (32×32) pixels local region [as shown in [Figure 2 c\)](#)] generally consists of dark ridge lines separated by white valley lines along the same orientation. The consistent ridge orientation and the appropriate ridge and valley structure are distinguishable local characteristics of the fingerprint local region.

The pixel-intensity gradient (dx, dy) at a pixel describes the direction of the maximum pixel-intensity change and its strength. By performing principal component analysis on the pixel-intensity gradients in a local region, an orthogonal basis for the local region can be formed by finding its eigenvalues and eigenvectors. The resultant first principal component contains the largest variance contributed by the maximum total gradient change in the direction orthogonal to ridge orientation. The direction is given by the first eigenvector and the value of the variance corresponds to the first eigenvalue, λ_{\max} . On the other hand, the resultant second principal component has the minimum change of gradient in the direction of ridge flow which corresponds to the second eigenvalue, λ_{\min} . The ratio between the two eigenvalues thus gives an

indication of the strength of the energy concentrated along the dominant direction with two vectors pointing to the normal and tangential direction of the average ridge flow respectively.

6.2.2.2 Computing the eigenvalues and local orientation certainty

From the covariance matrix C [Formula (8)] the eigenvalues λ_{\min} and λ_{\max} are computed as shown in Formulae (13) and (14):

$$\lambda_{\min} = \frac{a+b - \sqrt{(a-b)^2 + 4c^2}}{2} \quad (13)$$

$$\lambda_{\max} = \frac{a+b + \sqrt{(a-b)^2 + 4c^2}}{2} \quad (14)$$

This yields the local orientation certainty level shown in Formula (15):

$$q_{\text{OCL}}^{\text{local}} = \begin{cases} 1 - \frac{\lambda_{\min}}{\lambda_{\max}}, & \text{if } \lambda_{\max} > 0 \\ 0, & \text{otherwise.} \end{cases} \quad (15)$$

which is a ratio in the interval [0,1] where 1 is highest certainty level and 0 is lowest.

NOTE The orientation certainty level fails to predict recognition performance when some marks or residue exist in the samples that have strong orientation strength, such as those exhibited by latent prints left by the previous user of a capture device.

6.2.2.3 OCL algorithm

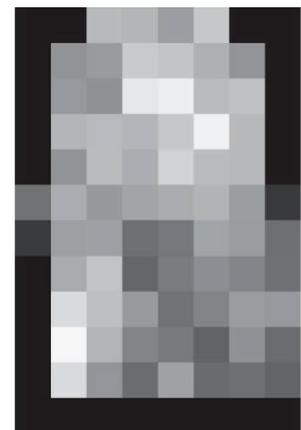
For each local region V in I :

- compute the pixel-intensity gradient of V with the centred differences method [Formulae (3), (4)];
- compute the covariance matrix C [Formula (8)];
- compute the eigenvalues of C to obtain $q_{\text{OCL}}^{\text{local}}$ [Formulae (13), (14), (15)].

Figure 3 visualizes the processing steps.



a) Current local region with the ratio between eigenvalues marked as ellipse



b) Original, untrimmed image and its $q_{\text{OCL}}^{\text{local}}$ values, mapped to values 0-255

Figure 3 — Processing steps of orientation certainty level quality algorithm

6.2.3 Local clarity

6.2.3.1 Description

Good quality fingerprints exhibit clear ridge-valley structure. Thus, the local clarity (LCL),^[4] which is the measure of the ridge-valley structure clarity, is a useful indicator of the quality of a fingerprint. The feature computes local quality and operates in a block-wise manner.

To perform ridge-valley structure analysis, the foreground of the finger image is quantized into local regions of size (32×32) pixels.^[3] Inside each local region, an orientation line, which is perpendicular to the ridge direction, is computed. At the centre of the local region along the ridge direction, a local region of size (32×16) pixels shall be extracted and transformed to a vertically aligned local region.

A linear regression (or least square fitting) is applied to calculate the determine threshold (DT), which is a line positioned at the centre of the local region V used to segment the local region into the ridge and valley regions. Regions with grey levels lower than DT are classified as ridges. Otherwise, they are classified as valleys.

Since good finger images cannot have ridges that are too close or too far apart, the nominal ridge and valley thickness can be used as a measure of the quality of the finger image captured. Similarly, ridges that are unreasonably thick or thin indicate that the finger image has not been captured properly (due to pressing too hard or too softly, for example), or that the image is a residual sample. Thus, the finger image quality can be determined by comparing the ridge and valley thickness to each of their nominal range of values. Any value out of the nominal range can imply a bad quality ridge pattern. To normalize the range of the thickness values, a pre-set maximum thickness is used. The maximum ridge or valley thickness, W_{\max} , for a good finger image is estimated at 20 pixels for a 196,85 pixels per centimetre (500 pixels per inch) scanner spatial sampling rate. The pre-set value of 20 pixels for a 196,85 pixels per centimetre (500 pixels per inch) scanner spatial sampling rate is obtained from the median of the typical ridge separation of $(8 - 12)$ pixels,^[2] and assuming that any ridge separation will not exceed twice the median value. This will ensure that the pre-set value is indeed the maximum to limit the value of the normalized ridge and valley thickness between 0 and 1. The ridge thickness, W_r , and valley thickness, W_v , are then normalized with respect to the maximum thickness, W_{\max} .

With the ridge and valley separated as above, a clarity test can be performed in each segmented rectangular 2D region.

For local regions with good clarity, the pixel-intensity distribution of ridges and the pixel-intensity distribution of the valleys have a very small overlapping area and thus $q_{\text{LCL}}^{\text{local}}$ is high. The following factors affect the size of the total overlapping area:

- a) noise on ridge and valley;
- b) water patches on the image due to wet fingers;
- c) incorrect orientation angle due to the effect of directional noise;
- d) scar across the ridge pattern;
- e) highly curved ridges;
- f) ridge endings, bifurcations, delta and core points;
- g) incipient ridges, sweat pores, and dots.

Factors a) to c) are physical noise found in the image. Factors d) to g) are actual physical characteristics of the fingerprint. See [Annex B](#) for the factors influencing fingerprint image quality.

6.2.3.2 Computing the ridge valley signature of a local region

Given the local region V , the ridge valley signature S is obtained according to [Formula \(16\)](#):

$$S(x) = \frac{\sum_{y=1}^{16} V_{x,y}}{16} \quad (16)$$

where

$V_{x,y}$ is the grey level at point (x, y) ;

x is the index along x -axis.

6.2.3.3 Determining the proportion of misclassified pixels

[Formula \(17\)](#) specifies the calculation of the proportion of pixels misclassified respectively as valley, and [Formula \(18\)](#) specifies the calculation of the proportion of pixels misclassified as ridge.

$$\alpha = \frac{v_B}{v_T} \quad (17)$$

$$\beta = \frac{r_B}{r_T} \quad (18)$$

where

α is the proportion of pixels misclassified as valley;

β is the proportion of pixels misclassified as ridge;

v_B is the number of pixels in the valley region with intensity lower than DT;

v_T is the total number of pixels in the valley region;

r_B is the number of pixels in the ridge region with intensity higher than DT;

r_T is the total number of pixels in the ridge region.

6.2.3.4 Determining the normalized ridge and valley width

The normalized valley width \overline{W}_v and the normalized ridge width \overline{W}_r are determined according to [Formulae \(19\)](#) and [\(20\)](#):

$$\overline{W}_v = \frac{W_v}{\left(\frac{S \times 2,54}{125}\right) W^{\max}} \quad (19)$$

$$\overline{W}_r = \frac{W_r}{\left(\frac{S \times 2,54}{125}\right) W^{\max}} \quad (20)$$

where

S is the spatial sampling rate in pixels per centimetre of the capture device;

W^{\max} is the estimated ridge or valley width for an image with 49,21 pixels per centimetre (125 pixels per inch) spatial sampling rate;

W_v and W_r are the observed valley and ridge widths, respectively.

According to Reference [\[2\]](#), $W^{\max} = 5$ is reasonable for 49,21 pixels per centimetre (125 pixels per inch) spatial sampling rate. By extension, $\left(\frac{S \times 2,54}{125}\right) W^{\max}$ is 20 for a spatial sampling rate of 196,85 pixels per centimetre (500 pixels per inch).

6.2.3.5 Computing the local clarity

The local quality value $q_{\text{LCL}}^{\text{local}}$ is the constrained average value of α and β with a range between 0 and 1, as shown in [Formula \(21\)](#):

$$q_{\text{LCL}}^{\text{local}} = \begin{cases} 1 - \frac{\alpha + \beta}{2}, & \text{if } (W_v^{\text{nmin}} < \overline{W}_v < W_v^{\text{nmax}}), (W_r^{\text{nmin}} < \overline{W}_r < W_r^{\text{nmax}}) \\ 0, & \text{otherwise.} \end{cases} \quad (21)$$

where W_r^{nmin} and W_v^{nmin} are the minimum values for the normalized ridge and valley width, calculated according to [Formulae \(22\)](#) and [\(23\)](#) respectively:

$$W_r^{\text{nmin}} = \frac{3}{\overline{W}_r} \quad (22)$$

$$W_v^{\text{nmin}} = \frac{2}{\overline{W}_v} \quad (23)$$

and where W_r^{nmax} and W_v^{nmax} are the maximum values for the normalized ridge and valley width, calculated according to [Formulae \(24\)](#) and [\(25\)](#) respectively:

$$W_r^{\text{nmax}} = \frac{10}{\overline{W}_r} \quad (24)$$

$$W_v^{\text{nmax}} = \frac{10}{\overline{W}_v} \quad (25)$$

NOTE Particular regions inherent in a fingerprint will negatively affect q_{LCL}^{local} . For example, ridge endings and bifurcations or areas with high curvature such as those commonly found in core and delta points.

6.2.3.6 LCL algorithm

For each local region V in I :

- a) rotate V such that dominant ridge flow is perpendicular to x-axis using nearest neighbour interpolation;
- b) crop rotated V such that no invalid regions are included;
- c) with V obtain the ridge-valley signature S (6.2.3.2);
- d) determine DT using linear regression on S ;
- e) for each element $S(x)$, set threshold $T(x)$ of x being ridge or valley based on DT;
- f) classify columns in V as ridge (1) or valley (0) with $P(x) = \begin{cases} 1, & \text{if } S(x) < T(x); \\ 0, & \text{otherwise.} \end{cases}$;
- g) determine ridge-valley transition vector C from P ;
- h) compute the vector W containing ridge and valley widths from C ;
- i) determine normalized ridge width and valley width \bar{W}_r and \bar{W}_v (6.2.3.4);
- j) determine the proportion of misclassified pixels α and β (6.2.3.3);
- k) compute the local quality value q_{LCL}^{local} (6.2.3.5).

NOTE The reference implementation fills invalid regions with 0-valued pixels.

[Figure 4](#) visualizes the processing steps.

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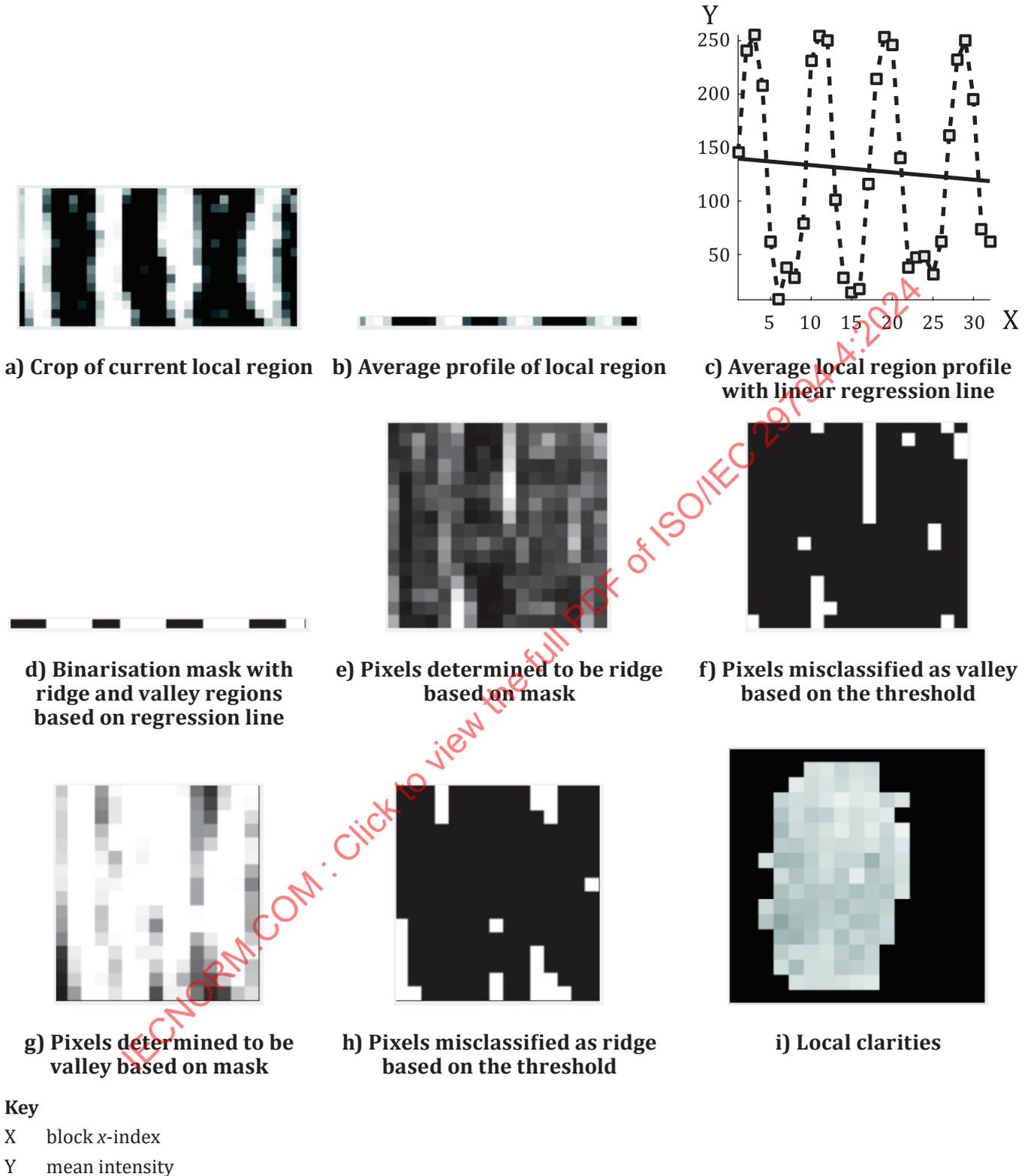


Figure 4 — Processing steps of local clarity algorithm

6.2.4 Frequency domain analysis (FDA)

6.2.4.1 Description

Frequency domain analysis (FDA) computes local quality and operates in a block-wise manner. A one-dimensional signature of the ridge-valley structure is extracted and the discrete Fourier transform (DFT) is computed on the signature to determine the frequency of the sinusoid following the ridge-valley structure.^[6]

The ridge-valley signature of a high quality sample is a periodic signal, which can be approximated either by a square wave or a sinusoidal wave. In the frequency domain, an ideal square wave should exhibit a dominant frequency with sideband frequency components (sinc function). A sinusoidal wave consists of one dominant frequency and minimum components at other non-dominant frequencies.

For each local region, a signature perpendicular to the dominant ridge flow orientation is computed.

The FDA described in 6.2.4 computes the one-dimensional signatures by performing averaging along the ridge flow direction. The averaging process filters out noise along the ridge and valley flow and provides a modelling of a smooth changing signal in a direction perpendicular to ridge flow.

6.2.4.2 Computing the local FDA quality component

The local quality value is computed by using Formula (26):

$$q_{\text{FDA}}^{\text{local}} = \begin{cases} 1, & \text{if } F_{\text{max}} = A_1 \text{ or } F_{\text{max}} = A_{|A|} \\ \frac{A_{F_{\text{max}}} + C(A_{F_{\text{max}-1}} + A_{F_{\text{max}+1}})}{\sum_{F=1}^{|A|/2} A_F}, & \text{otherwise.} \end{cases} \quad (26)$$

where

0,3 is the attenuation parameter C ;

A_x is the amplitude at frequency index x .

The value of $q_{\text{FDA}}^{\text{local}}$ is set to 1 when the maximum frequency, F_{max} , amplitude occurs at index $F_{\text{max}} = A_1$ or $F_{\text{max}} = A_{|A|}$.

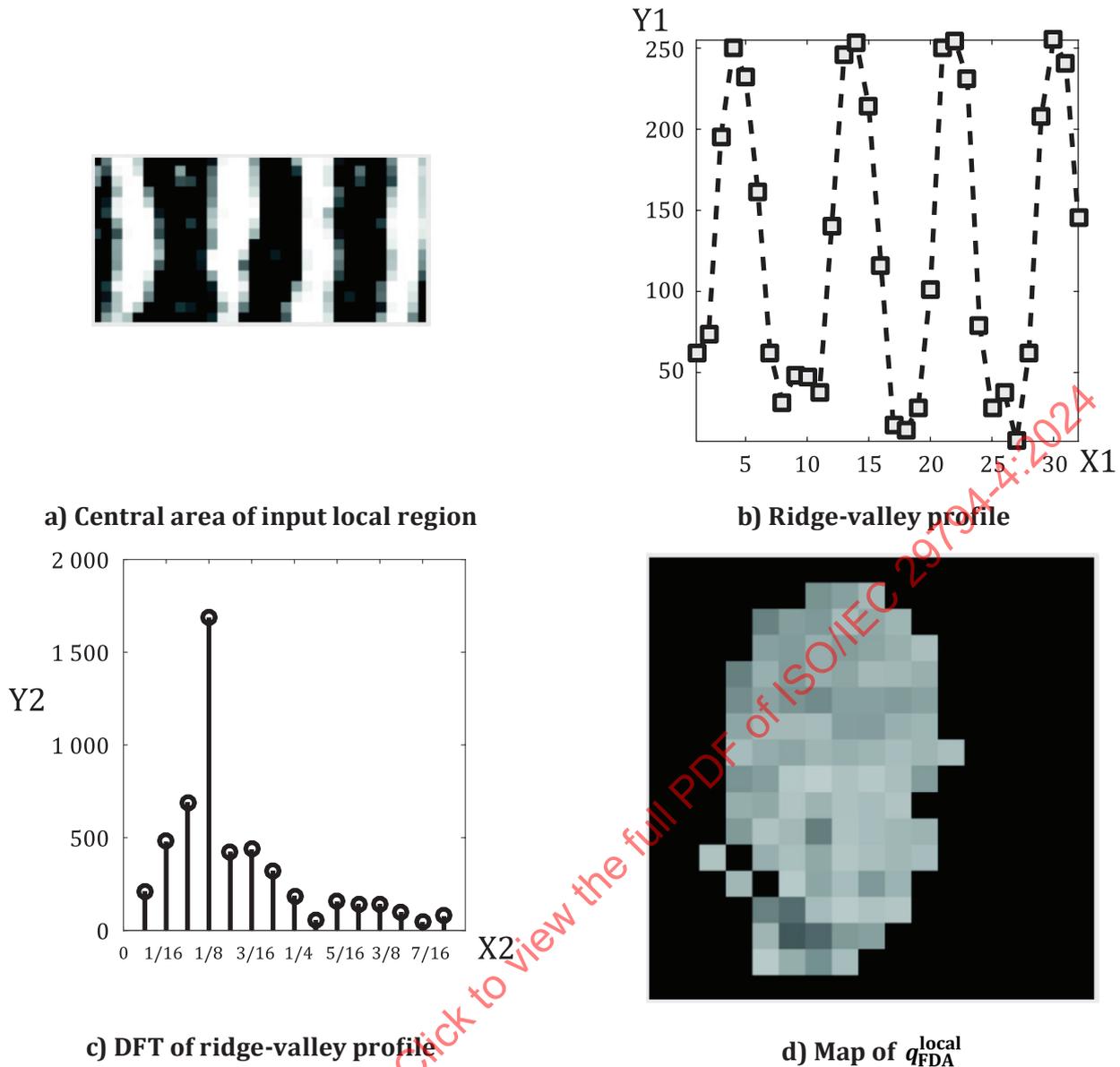
6.2.4.3 FDA algorithm

For each local region V in I :

- pad V with a 2-pixel border;
- rotate V with nearest neighbour interpolation such that dominant ridge flow is perpendicular to the x-axis with nearest neighbour interpolation;
- crop V such that no invalid regions are included;
- with V , obtain the ridge-valley signature S (6.2.3.2);
- compute the DFT of S to obtain the magnitude representation A ;
- discard the DC component (i.e. the first component of A);
- determine F_{max} as the index with the largest magnitude in A ;
- compute $q_{\text{FDA}}^{\text{local}}$ of V using A and F_{max} (6.2.4.2).

NOTE The reference implementation fills invalid regions with 0-valued pixels.

Figure 5 visualizes the processing steps.



- Key**
- X1 mean intensity
 - Y1 block x-index
 - X2 magnitude
 - Y2 cycle/pixel

Figure 5 — Processing steps of FDA quality algorithm

6.2.5 Ridge valley uniformity

6.2.5.1 Feature description

Ridge valley uniformity (RVU) is a measure of the consistency of the ridge and valley widths.^[3] The expectation for finger image with clear ridge and valley separation is that the ratio between ridge and valley widths remains fairly constant throughout the finger image.

The ratio of ridge thickness to valley thickness should be constant and close to 1 throughout the whole image for a good quality finger image. The feature computes local quality and operates in a block-wise manner.

NOTE The ridge valley uniformity depends on the spatial sampling rate.

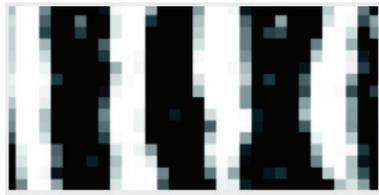
6.2.5.2 RVU algorithm

For each local region V in I :

- a) determine dominant ridge flow orientation $\text{angle}(V)$ of V ;
- b) rotate V such that $\text{angle}(V)$ is perpendicular to x-axis;
- c) crop V such that no invalid regions are included;
- d) with V , obtain the ridge-valley signature S (6.2.3.2);
- e) determine DT using linear regression on S ;
- f) for each $S(x)$, compute threshold $T(x) = x \times \text{DT}(1) + \text{DT}(0)$;
- g) binarize S using T ;
- h) classify ridge and valley in S as $P(x) = \begin{cases} 1, & \text{if } S(x-1) < T(x) \\ 0, & \text{otherwise.} \end{cases}$;
- i) compute ridge-valley transition vector as $C(x) = \begin{cases} 1, & \text{if } P(x-1) \neq P(x) \\ 0, & \text{otherwise.} \end{cases}$;
- j) drop first and last transition from S using C to remove incomplete ridges or valleys and obtain S' ;
- k) compute $q_{\text{RVU}}^{\text{local}}$ as the ratio of the width of ridges against valleys in S' .

[Figure 6](#) visualizes the processing steps.

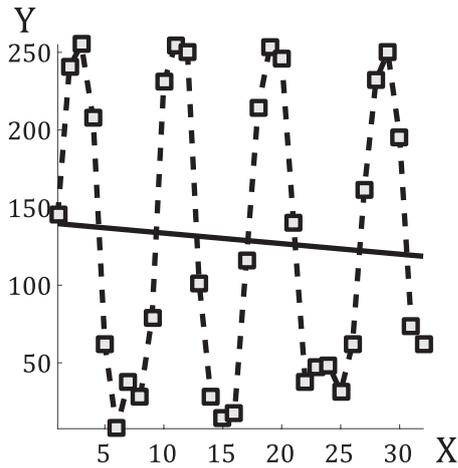
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a) Crop of current local region



b) Average profile of local region



c) Average profile with regression line



d) Local native quality component values displayed as the ratio of the width of ridges against valleys

Key

- X block x-index
- Y mean intensity

Figure 6 — Processing steps of ridge valley uniformity quality algorithm

6.2.6 Orientation flow

6.2.6.1 Description

Orientation flow (OFL) is a measure of ridge flow continuity which is based on the absolute orientation difference between a local region and its 8-neighbourhood of local regions.

Orientation flow is a good indicator to describe the quality of a good fingerprint pattern because, in general, the flow of the ridge direction changes gradually, except in an area with a delta or a core. The feature computes local quality and operates in a block-wise manner.

6.2.6.2 Local region-wise absolute orientation difference

The ridge flow is determined as a measure of the absolute difference between a local region and its neighbouring local regions. The absolute difference $D(i, j)$ for local region $V(i, j)$ is computed using the dominant ridge flow orientations of this local region and of its neighbours, as shown in [Formula \(27\)](#):

$$D(i, j) = \frac{\sum_{m=-1}^1 \sum_{n=-1}^1 \min(|\theta_{V(i,j)} - \theta_{V(i-m,j-n)}|, 360^\circ - |\theta_{V(i,j)} - \theta_{V(i-m,j-n)}|)}{8} \tag{27}$$

[Formula \(27\)](#)^[2] takes the minimum angle differences and 360° minus the angle difference because of the circularity of angles (e.g. the difference between angles of 2° and 358° is 4°).

6.2.6.3 Local orientation flow quality

The local orientation quality q_{OFL}^{local} for the local region orientation difference $D(i, j)$ is calculated according to [Formula \(28\)](#):

$$q_{OFL}^{local} = \begin{cases} \frac{D(i, j) - \theta_{min}}{90^\circ - \theta_{min}}, & \text{if } D(i, j) > \theta_{min} \\ 0, & \text{otherwise.} \end{cases} \tag{28}$$

where $\theta_{min} = 4^\circ$ is the threshold for minimum angle difference to consider and the units of $D(i, j)$ are degrees.

6.2.6.4 OFL algorithm

- a) Determine the dominant ridge flow orientation angle(V) of local region V and I .
- b) For each local region V in I :
 - 1) compute the absolute orientation difference $D(i, j)$ using angle(V) ([6.2.6.2](#));
 - 2) compute the local orientation quality q_{OFL}^{local} ([6.2.6.3](#)).

[Figure 7](#) visualizes the processing steps.

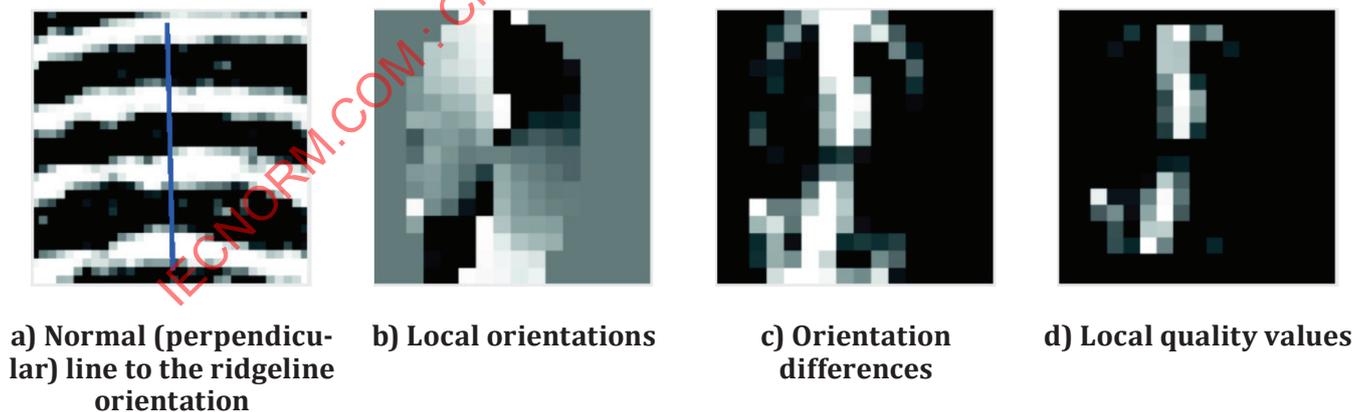


Figure 7 — Processing steps of orientation flow quality algorithm

6.2.7 MU

6.2.7.1 Description

MU is the arithmetic mean of the pixel intensities of all pixels in the input image. The feature computes global quality.

6.2.7.2 MU algorithm

Compute Q_{MU} as the arithmetic mean of pixel intensities in I .

6.2.7.3 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating-point q_{MU} values shall be mapped to a quality component Q_{MU} in the range from 0 to 100 using [Formula \(2\)](#), where x is q_{MU} , $\min(q_{\text{MU}})$ is 0 and $\max(q_{\text{MU}})$ is 255. See [Formula \(29\)](#).

$$Q_{\text{MU}} = \left\lfloor \frac{q_{\text{MU}}}{255 + \varepsilon} 101 \right\rfloor \quad (29)$$

6.2.8 MMB

6.2.8.1 Description

MMB is the arithmetic mean of all local region arithmetic means in the greyscale input image. The feature computes local quality and operates in a block-wise manner.

6.2.8.2 MMB algorithm

- a) For each local region V in I :
 - 1) compute the arithmetic mean of the pixel intensities in V as $q_{\text{MMB}}^{\text{local}}$.
- b) Compute Q_{MMB} as the arithmetic mean of set of $q_{\text{MMB}}^{\text{local}}$.

6.2.8.3 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating-point q_{MMB} values shall be mapped to a quality component Q_{MMB} in the range from 0 to 100 using [Formula \(2\)](#), where x is q_{MMB} , $\min(q_{\text{MMB}})$ is 0 and $\max(q_{\text{MMB}})$ is 255. See [Formula \(30\)](#).

$$Q_{\text{MMB}} = \left\lfloor \frac{q_{\text{MMB}}}{255 + \varepsilon} 101 \right\rfloor \quad (30)$$

6.2.9 Minutiae count in finger image

6.2.9.1 Description

FingerJet Fingerprint Feature Extractor, Open Source Edition (FJFX)^[5] is the minutiae extractor selected for the reference implementation due to licensing conditions (including commercial use cases and redistribution), computational complexity, and platform independence. FJFX provides a count of detected minutiae in the finger image. The minutiae count has a bearing on the mated comparison score. The feature computes global quality.

6.2.9.2 MIN^{CNT} algorithm

$q_{\text{MIN}}^{\text{CNT}}$ is the number of detected minutiae in the finger image as determined by the minutiae extractor.

6.2.9.3 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the $q_{\text{MIN}}^{\text{CNT}}$ values shall be limited to 100 using [Formula \(31\)](#), yielding the quality component $Q_{\text{MIN}}^{\text{CNT}}$:

$$Q_{\text{MIN}}^{\text{CNT}} = \begin{cases} q_{\text{MIN}}^{\text{CNT}} & \text{if } q_{\text{MIN}}^{\text{CNT}} \leq 100 \\ 100 & \text{if } q_{\text{MIN}}^{\text{CNT}} > 100 \end{cases} \quad (31)$$

6.2.10 Minutiae count in centre of mass region

6.2.10.1 Description

This quality component is the minutiae count in a (200 × 200) pixels local region centred on the centre of mass of the detected minutiae as described in [6.2.9](#). The quality component computes local quality at the minutiae locations.

6.2.10.2 MIN^{COM} algorithm

$q_{\text{MIN}}^{\text{COM}}$ is the number of minutiae occurring within a (200 × 200) pixels local region centred at the centre of mass of the locations of all detected minutiae in the finger image as described in [6.2.9](#).

6.2.10.3 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating-point $q_{\text{MIN}}^{\text{COM}}$ values shall be limited to 100 using [Formula \(32\)](#), yielding the quality component $Q_{\text{MIN}}^{\text{COM}}$:

$$Q_{\text{MIN}}^{\text{COM}} = \begin{cases} q_{\text{MIN}}^{\text{COM}} & \text{if } q_{\text{MIN}}^{\text{COM}} \leq 100 \\ 100 & \text{if } q_{\text{MIN}}^{\text{COM}} > 100 \end{cases} \quad (32)$$

6.2.11 Minutiae quality based on local image mean

6.2.11.1 Description

For each minutia location detected as described in [6.2.9](#), a local quality based on image statistics is computed. The reported quality value is aggregated as the count of local qualities which occurs in the specified range. The feature computes local quality at the minutiae locations.

6.2.11.2 MIN^{MU} algorithm

$q_{\text{MIN}}^{\text{MU}}$ is computed by first determining the local quality of each minutia detected as described in [6.2.9](#), using [Formula \(33\)](#):

$$q_{\text{MIN}}^{\text{localMU}} = \frac{\mu(\mathbf{I}) - \mu(\mathbf{V})}{\sigma(\mathbf{I})} \quad (33)$$

where

$\mu(\mathbf{I})$ is the arithmetic mean of the finger image;

$\mu(\mathbf{V})$ is the arithmetic mean of a (32×32) pixels local region centred on the minutia;

$\sigma(\mathbf{I})$ is the standard deviation of the finger image.

$q_{\text{MIN}}^{\text{MU}}$ is finally computed as the percentage of $q_{\text{MIN}}^{\text{localMU}}$ values greater than 0 and less than or equal to 0,5, as shown in [Formula \(34\)](#):

$$q_{\text{MIN}}^{\text{MU}} = \frac{\sum_{x=1}^{q_{\text{MIN}}^{\text{CNT}}} \begin{cases} 1, & 0 < q_{\text{MIN}_x}^{\text{localMU}} \leq 0,5 \\ 0, & \text{otherwise.} \end{cases}}{q_{\text{MIN}}^{\text{CNT}}} \quad (34)$$

6.2.11.3 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating-point $q_{\text{MIN}}^{\text{MU}}$ values shall be mapped to a quality component $Q_{\text{MIN}}^{\text{MU}}$ in the range from 0 to 100 by multiplication with 100, using [Formula \(2\)](#), where x is $q_{\text{MIN}}^{\text{MU}}$, $\min(q_{\text{MIN}}^{\text{MU}})$ is 0 and $\max(q_{\text{MIN}}^{\text{MU}})$ is 255. See [Formula \(35\)](#).

$$Q_{\text{MIN}}^{\text{MU}} = \left\lfloor \frac{q_{\text{MIN}}^{\text{MU}}}{255 + \varepsilon} 101 \right\rfloor \quad (35)$$

6.2.12 Minutiae quality based on local orientation certainty level

6.2.12.1 Description

For each minutia location detected, as described in [6.2.9](#), a local orientation certainty level is computed. The reported quality value is aggregated as the count of local qualities which exceed the specified value. The feature computes local quality at the minutiae locations.

6.2.12.2 MIN^{OCL} algorithm

$q_{\text{MIN}}^{\text{OCL}}$ is computed by first determining the local quality of each minutia detected as described in [6.2.9](#) as shown in [Formula \(36\)](#):

$$q_{\text{MIN}}^{\text{localOCL}} = q_{\text{OCL}}^{\text{local}}(\mathbf{V}) \quad (36)$$

where $q_{\text{OCL}}^{\text{local}}(\mathbf{V})$ is the local orientation certainty level ([6.2.2](#)) for the (32×32) pixels local region \mathbf{V} centred on the minutia.

q_{MIN}^{OCL} is finally computed as the percentage of $q_{MIN}^{localOCL}$ values greater than 0,8, as shown in [Formula \(37\)](#):

$$q_{MIN}^{OCL} = \frac{\sum_{x=1}^{q_{MIN}^{CNT}} \begin{cases} 1, & q_{MIN_x}^{localOCL} > 0,8 \\ 0, & \text{otherwise.} \end{cases}}{q_{MIN}^{CNT}} \quad (37)$$

6.2.12.3 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating-point q_{MIN}^{OCL} values shall be mapped to a quality component Q_{MIN}^{OCL} in the range from 0 to 100 by multiplication with 100, using [Formula \(2\)](#), where x is q_{MIN}^{OCL} , $\min(q_{MIN}^{OCL})$ is 0 and $\max(q_{MIN}^{OCL})$ is 1, as shown in [Formula \(38\)](#):

$$Q_{MIN}^{OCL} = \left\lfloor \frac{q_{MIN}^{OCL}}{1 + \varepsilon} 101 \right\rfloor \quad (38)$$

6.2.13 Region of interest image mean

6.2.13.1 Description

The region of interest for the finger image is the foreground region of the image containing the fingerprint. The mean pixel intensity in this area is computed over the set of (32×32) pixels local regions which have a least one pixel contained in the region of interest. The feature computes global quality.

NOTE The quality component is highly correlated with q_{MU} ([6.2.7](#)) and q_{MMB} ([6.2.8](#)).

6.2.13.2 Determine the region of interest

- a) Erode the finger image I with a 5×5 structuring element to obtain \bar{I} .
- b) Apply normalized Gaussian blur filter (each weight is divided by the sum of all weights) with kernel size 41×41 and standard deviation of 6,5 to \bar{I} to obtain G .
- c) Binarize G to obtain B .

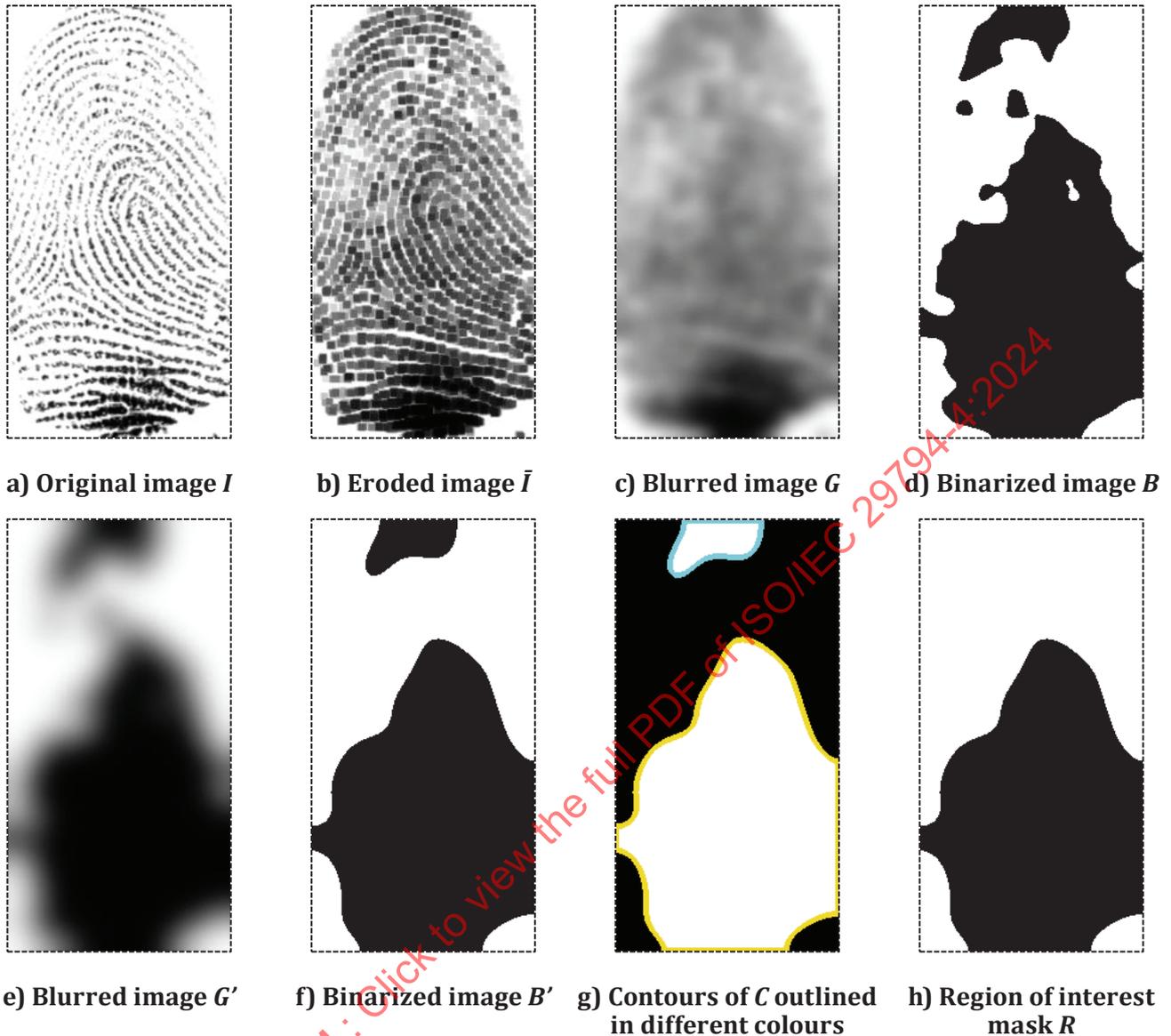
NOTE The reference implementation uses Otsu's method^[7] for binarization.

- d) Apply normalized Gaussian blur filter (each weight is divided by the sum of all weights) with kernel size 91×91 and standard deviation of 14,0 to B to obtain G' .
- e) Binarize G' using Otsu's method to obtain B' .
- f) Determine the contours of inverted B' to obtain C .

NOTE The reference implementation uses Suzuki's method^[8] for determining contours.

- g) Contours in the hierarchy of C that have a parent shall be set to 0 valued pixels.
- h) 0 valued pixel regions that are not the largest 0 valued pixel area shall be set to 1 valued pixels.
- i) The resulting binary mask R contains the region of interest as a region of 0 valued pixels.

[Figure 8](#) visualizes the processing steps.



NOTE Each subfigure contains a dashed black border.

Figure 8 — Processing steps of region of interest algorithm

6.2.13.3 AREA algorithm

- a) Determine the region of interest R (6.2.13.2).
- b) For each 32×32 local region V in I :
 - 1) if V has at least 1 pixel contained in foreground of R , mark the local region as foreground.

NOTE Up to and including in version NFIQ 2.2, V is implemented as a (16×16) pixels region. The values, however, are the same as when local regions of size (32×32) pixels are used.

- c) Compute q_{AREA}^{μ} as the arithmetic mean of the grey levels of all pixels in the set of local regions V marked as foreground.

6.2.13.4 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating-point q_{AREA}^{μ} values shall be mapped to a quality component Q_{AREA}^{μ} in the range from 0 to 100 using [Formula \(2\)](#), where x is q_{AREA}^{μ} , $\min(q_{\text{AREA}}^{\mu})$ is 0 and $\max(q_{\text{AREA}}^{\mu})$ is 255, as shown in [Formula \(39\)](#):

$$Q_{\text{AREA}}^{\mu} = \left\lfloor \frac{q_{\text{AREA}}^{\mu}}{255 + \varepsilon} 101 \right\rfloor \quad (39)$$

6.2.14 Region of interest orientation map coherence sum

6.2.14.1 Description

The orientation map coherence sum quantifies the coherence of the estimated finger image orientation field. The coherence map is computed according to the coherence method specified in Reference [9]. The feature computes local quality and operates in a block-wise manner.

6.2.14.2 Computing the gradient field

The gradient field $\mathbf{g} = (g_x, g_y)^T$ of pixel intensity $I(i, j)$ of I is as shown in [Formulae \(40\)](#) and [\(41\)](#):

$$\begin{aligned} g_x(i, j) &= I(i+1, j) - I(i-1, j) / 2 \quad \text{for } 1 \leq i \leq I_w - 1, 0 \leq j \leq I_h \\ g_x(0, j) &= I(1, j) - I(0, j) \quad \text{for } 0 \leq j \leq I_h \\ g_x(I_w, j) &= I(I_w, j) - I(I_w - 1, j) \quad \text{for } 0 \leq j \leq I_h \end{aligned} \quad (40)$$

and

$$\begin{aligned} g_y(i, j) &= (I(i, j+1) - I(i, j-1)) / 2 \quad \text{for } 0 \leq i \leq I_w - 1, 1 \leq j \leq I_h - 1 \\ g_y(i, 0) &= I(i, 1) - I(i, 0) \quad \text{for } 0 \leq i \leq I_w \\ g_y(i, I_h) &= I(i, I_h) - I(i, I_h - 1) \quad \text{for } 0 \leq i \leq I_w \end{aligned} \quad (41)$$

where I_w and I_h are respectively the width and height of I in pixels.

6.2.14.3 Computing the coherence of a local region

The coherence of a local region V is computed from its pixel-intensity gradient field \mathbf{g}_s as shown in [Formula \(42\)](#):

$$\text{coh}(V) = \frac{|\sum \mathbf{g}_s(i, j)|}{\sum |\mathbf{g}_s(i, j)|} \quad (42)$$

where $|x|$ denotes the Euclidean norm of x and the sums are taken over all pixels in V .

NOTE \mathbf{g}_s is defined in [6.2.14.4](#).

6.2.14.4 COH^{SUM} algorithm

- Compute the pixel-intensity gradient field \mathbf{g} of I ([6.2.14.2](#)).
- Compute the square gradient field as $\mathbf{g}_s = (g_x^2 - g_y^2, 2g_x g_y)^T$.
- Determine the region of interest R ([6.2.13.2](#)).

d) For each (16×16) pixels local region V in I :

- 1) if V has at least 1 pixel contained in foreground of R , compute the coherence of V as $\text{coh}(V)$, otherwise set $\text{coh}(V)=0$.

NOTE Up to and including in version NFIQ 2.2, V is implemented as a (16×16) pixels region.

e) Compute $q_{\text{COH}}^{\text{sum}}$ as the sum of the $\text{coh}(V)$ of all V .

6.2.14.5 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating point $q_{\text{COH}}^{\text{sum}}$ values shall be mapped to a quality component $Q_{\text{COH}}^{\text{sum}}$ in the range from 0 to 100 and rounded to the nearest integer using [Formula \(2\)](#), where x is $q_{\text{COH}}^{\text{sum}}$, $\min(q_{\text{COH}}^{\text{sum}})$ is 0 and $\max(q_{\text{COH}}^{\text{sum}})$ is 3 150. See [Formula \(43\)](#):

$$Q_{\text{COH}}^{\text{sum}} = \left\lfloor \frac{q_{\text{COH}}^{\text{sum}}}{3\,150 + \varepsilon} 101 \right\rfloor \quad (43)$$

NOTE $q_{\text{COH}}^{\text{REL}}$ ([6.2.15.2](#)) is a value between 0,0 and 1,0. The reference implementation's minutiae extractor implementation defined in [6.2.9](#) imposes a maximum image size of $(800 \times 1\,000)$ pixels, which can fit 3 150 (16×16) pixels blocks. Therefore, the maximum value of $q_{\text{COH}}^{\text{sum}}$ in the reference implementation is 3 150. The value of $q_{\text{COH}}^{\text{sum}}$ between 0 and 3 150 can therefore be normalized to derive $Q_{\text{COH}}^{\text{sum}}$.

6.2.15 Region of interest relative orientation map coherence sum

6.2.15.1 Description

The relative orientation map coherence sum is the average of local region orientation coherence as determined by [6.2.14](#). The feature computes local quality and operates in a block-wise manner.

6.2.15.2 COH^{REL} algorithm

a) Compute $q_{\text{COH}}^{\text{sum}}$ and store the number of local regions V which have at least one pixel contained in R as n ([6.2.14](#)).

b) Compute $q_{\text{COH}}^{\text{REL}} = \frac{q_{\text{COH}}^{\text{sum}}}{n}$

6.2.15.3 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating-point $q_{\text{COH}}^{\text{REL}}$ values shall be mapped to a quality component $Q_{\text{COH}}^{\text{REL}}$ in the range from 0 to 100 using [Formula \(2\)](#), x is $q_{\text{COH}}^{\text{REL}}$, $\min(q_{\text{COH}}^{\text{REL}})$ is 0 and $\max(q_{\text{COH}}^{\text{REL}})$ is 1, as shown in [Formula \(44\)](#):

$$Q_{\text{COH}}^{\text{REL}} = \left\lfloor \frac{q_{\text{COH}}^{\text{REL}}}{1 + \varepsilon} 101 \right\rfloor \quad (44)$$

6.2.16 Quality feature vector composition

6.2.16.1 Description

Quality features specified in [6.2.2](#) to [6.2.6](#) provide a map of values for local regions in the finger image. The quality features specified in [6.2.7](#) to [6.2.15](#) provide scalar quality values for the finger image.

The specified features shall be composed such that a fixed length feature vector is obtained for use by a classification system. Thus, the features in 6.2.2 to 6.2.6 are aggregated using arithmetic mean specified in 6.2.16.2, standard deviation specified in 6.2.16.3 and histogram specified in 6.2.16.4 for inclusion in the final feature vector specified in 6.2.16.5.

6.2.16.2 Mean of local quality values

6.2.16.2.1 Basic calculation

The mean quality value over an $N \times M$ matrix of local quality values is computed as shown in Formula (45):

$$\mathbf{G} = \frac{1}{N \times M} \sum_{i=1}^N \sum_{j=1}^M \mathbf{L}(i, j) \quad (45)$$

where

- \mathbf{L} are local quality values $q_{\text{OCL}}^{\text{local}}$ (6.2.2), $q_{\text{LCL}}^{\text{local}}$ (6.2.3), $q_{\text{FDA}}^{\text{local}}$ (6.2.4), $q_{\text{RVU}}^{\text{local}}$ (6.2.5) or $q_{\text{OFL}}^{\text{local}}$ (6.2.6);
- \mathbf{G} are the aggregated mean quality values q_{OCL}^{μ} , q_{LCL}^{μ} , q_{FDA}^{μ} , q_{RVU}^{μ} , or q_{OFL}^{μ} , in respect of \mathbf{L} .

6.2.16.2.2 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating-point aggregated mean quality values produced in 6.2.16.2.1 shall be mapped to a quality component in the range from 0 to 100 and rounded to the nearest integer.

For q_{OCL}^{μ} , q_{LCL}^{μ} , q_{FDA}^{μ} and q_{OFL}^{μ} , refer to Formula (2) from 6.1.2.1 to yield Q_{OCL}^{μ} , Q_{LCL}^{μ} , Q_{FDA}^{μ} and Q_{OFL}^{μ} , respectively. For q_{RVU}^{μ} , Formula (46) shall be used for mapping to yield Q_{RVU}^{μ} :

$$Q_{\text{RVU}}^{\mu} = \left\lfloor 100 \cdot \text{sigmoid} \left(q_{\text{RVU}}^{\mu}, 1, \frac{1}{2} \right) + \frac{1}{2} \right\rfloor \quad (46)$$

6.2.16.3 Standard deviation of local quality values

6.2.16.3.1 Basic calculation

The standard deviation over an $N \times M$ matrix of local quality values is computed as shown in Formula (47):

$$\mathbf{G} = \left(\frac{1}{N \times M - 1} \sum_{i=1}^N \sum_{j=1}^M (\mathbf{L}(i, j) - \mathbf{F})^2 \right)^{\frac{1}{2}} \quad (47)$$

where

- \mathbf{L} are local quality values $q_{\text{OCL}}^{\text{local}}$ (6.2.2), $q_{\text{LCL}}^{\text{local}}$ (6.2.3), $q_{\text{FDA}}^{\text{local}}$ (6.2.4), $q_{\text{RVU}}^{\text{local}}$ (6.2.5), or $q_{\text{OFL}}^{\text{local}}$ (6.2.6);
- \mathbf{G} are the aggregated standard deviation quality values q_{OCL}^{σ} , q_{LCL}^{σ} , q_{FDA}^{σ} , q_{RVU}^{σ} , or q_{OFL}^{σ} , in respect of \mathbf{L} ;
- \mathbf{F} are the aggregated mean quality values q_{OCL}^{μ} , q_{LCL}^{μ} , q_{FDA}^{μ} , q_{RVU}^{μ} , or q_{OFL}^{μ} , in respect of \mathbf{L} .

6.2.16.3.2 Mapping to the desired value range

To fit into a quality block as specified in ISO/IEC 29794-1, the floating-point aggregated standard deviation quality values produced in [6.2.16.3.1](#) shall be mapped to a quality component in the range from 0 to 100 and rounded to the nearest integer.

For q_{OCL}^{σ} , q_{LCL}^{σ} , q_{FDA}^{σ} , and q_{OFL}^{σ} , which range from 0 to 1, [Formula \(2\)](#) shall be used for mapping to yield Q_{OCL}^{σ} , Q_{LCL}^{σ} , Q_{FDA}^{σ} , and Q_{OFL}^{σ} , respectively. For q_{RVU}^{σ} , [Formula \(48\)](#) shall be used for mapping to yield Q_{RVU}^{σ} :

$$Q_{RVU}^{\sigma} = \left\lfloor 100 \cdot \text{sigmoid} \left(q_{RVU}^{\sigma}, 1, \frac{1}{2} \right) + \frac{1}{2} \right\rfloor \quad (48)$$

6.2.16.4 Histogram of local quality

Local quality values from orientation certainty ([6.2.2](#)), local clarity ([6.2.3](#)), frequency domain analysis ([6.2.4](#)), ridge valley uniformity ([6.2.5](#)) and orientation flow ([6.2.6](#)) shall be represented as fixed-length histograms with 10 bins to capture the distribution of local qualities.

The boundaries defining each bin for each of the features are specified as shown in [Formula \(49\)](#):

$$\begin{aligned} B_{FDA} &= \{-\infty; 0,268; 0,304; 0,33; 0,355; 0,38; 0,407; 0,440; 0,5; 1; \infty\} \\ B_{LCL} &= \{-\infty; 0; 0,7; 0,740; 0,77; 0,79; 0,81; 0,83; 0,85; 0,87; \infty\} \\ B_{OCL} &= \{-\infty; 0,337; 0,479; 0,579; 0,655; 0,716; 0,766; 0,81; 0,852; 0,898; \infty\} \\ B_{OFL} &= \{-\infty; 0,01715; 0,035; 0,05570; 0,081; 0,115; 0,17180; 0,25690; 0,47580; 0,748; \infty\} \\ B_{RVU} &= \{-\infty; 0,5; 0,667; 0; 1; 1,25; 1,5; 2; 24; 30; \infty\} \end{aligned} \quad (49)$$

For each of FDA, LCL, OCL, OFL and RVU, a histogram is computed using the specified bin boundaries where the i -th bin in the histogram is given by the interval shown in [Formula \(50\)](#):

$$\begin{aligned} &(B_Q^i, B_Q^{i+1}), \quad \text{for } 1=i \\ &B_Q^i, B_Q^{i+1}), \quad \text{for } 1 < i \leq |B_Q| \end{aligned} \quad (50)$$

where B_Q is B_{FDA} , B_{LCL} , B_{OCL} , B_{OFL} , or B_{RVU} .

The histograms of local qualities are specified according to their bin boundaries as defined in [Formulae \(51\)](#) and [\(52\)](#) where the i -th bin in the histogram contains the cardinality of the multiset that contains values bounded by the histogram boundaries.

$$\begin{aligned}
 q_{\text{FDA}}^i &= \left\{ \left\{ B_{\text{FDA}}^i \leq q_{\text{FDA}}^{\text{local}} < B_{\text{FDA}}^{i+1} \right\} \right\}, \text{ for } 1 \leq i \leq |B_{\text{FDA}}|, \\
 q_{\text{LCL}}^i &= \left\{ \left\{ B_{\text{LCL}}^i \leq q_{\text{LCL}}^{\text{local}} < B_{\text{LCL}}^{i+1} \right\} \right\}, \text{ for } 1 \leq i \leq |B_{\text{LCL}}|, \\
 q_{\text{OCL}}^i &= \left\{ \left\{ B_{\text{OCL}}^i \leq q_{\text{OCL}}^{\text{local}} < B_{\text{OCL}}^{i+1} \right\} \right\}, \text{ for } 1 \leq i \leq |B_{\text{OCL}}|, \\
 q_{\text{OFL}}^i &= \left\{ \left\{ B_{\text{OFL}}^i \leq q_{\text{OFL}}^{\text{local}} < B_{\text{OFL}}^{i+1} \right\} \right\}, \text{ for } 1 \leq i \leq |B_{\text{OFL}}|, \\
 q_{\text{RVU}}^i &= \left\{ \left\{ B_{\text{RVU}}^i \leq q_{\text{RVU}}^{\text{local}} < B_{\text{RVU}}^{i+1} \right\} \right\}, \text{ for } 1 \leq i \leq |B_{\text{RVU}}|.
 \end{aligned} \tag{51}$$

The histogram for a single feature represented by its bins is written as:

$$\begin{aligned}
 q_{\text{FDA}} &= \{q_{\text{FDA}}^i\}, \text{ for } 1 \leq i < |B_{\text{FDA}}|, \\
 q_{\text{LCL}} &= \{q_{\text{LCL}}^i\}, \text{ for } 1 \leq i < |B_{\text{LCL}}|, \\
 q_{\text{OCL}} &= \{q_{\text{OCL}}^i\}, \text{ for } 1 \leq i < |B_{\text{OCL}}|, \\
 q_{\text{OFL}} &= \{q_{\text{OFL}}^i\}, \text{ for } 1 \leq i < |B_{\text{OFL}}|, \\
 q_{\text{RVU}} &= \{q_{\text{RVU}}^i\}, \text{ for } 1 \leq i < |B_{\text{RVU}}|.
 \end{aligned} \tag{52}$$

6.2.16.5 ISO/IEC 29794-4 quality feature vector

The quality feature vector that is input to the unified quality score computation (see 6.4) is specified as shown in [Formula \(53\)](#):

$$\begin{aligned}
 Q_{29794-4} = \{ & q_{\text{OCL}}^{\mu}, q_{\text{LCL}}^{\mu}, q_{\text{FDA}}^{\mu}, q_{\text{RVU}}^{\mu}, q_{\text{OFL}}^{\mu}, \\
 & q_{\text{OCL}}^{\sigma}, q_{\text{LCL}}^{\sigma}, q_{\text{FDA}}^{\sigma}, q_{\text{RVU}}^{\sigma}, q_{\text{OFL}}^{\sigma}, \\
 & q_{\text{OCL}}, q_{\text{FDA}}, q_{\text{LCL}}, q_{\text{RVU}}, q_{\text{OFL}}, \\
 & q_{\text{MU}}^{\text{REL}}, q_{\text{MMB}}, q_{\text{COH}}^{\text{REL}}, q_{\text{COH}}^{\text{sum}}, q_{\text{AREA}}^{\mu}, \\
 & q_{\text{MIN}}^{\text{CNT}}, q_{\text{MIN}}^{\text{COM}}, q_{\text{MIN}}^{\text{MU}}, q_{\text{MIN}}^{\text{OCL}} \}
 \end{aligned} \tag{53}$$

6.3 Non-normative quality measures

6.3.1 General

[Subclause 6.3](#) specifies algorithms for computing further finger image quality components, which, however, are not used in the computation of the unified quality score.

6.3.2 Radial power spectrum

6.3.2.1 Description

The radial power spectrum is a measure of maximal signal power in a defined frequency band of the global radial Fourier spectrum. Ridges can be locally approximated by means of a single sine wave, hence high energy concentration in a narrow frequency band corresponds to consistent ridge structures.

Since the ridges of a finger image can be locally approximated by one sine wave, large values of sine wave energy can represent strong ridges. The robustness of the ridge structure can be used to measure finger image quality. \dot{F} is decided as the maximum Radial Fourier spectrum value within the reasonable Fourier domain. The reasonable Fourier domain refers to the region of neither the highest nor the lowest frequency. The higher the value of \dot{F} , the better the finger image quality.

6.3.2.2 Variables

Name	Default	Description
r_{\min}	0,143 cycles/pixel	Lower bound of frequency band
r_{\max}	0,077 cycles/pixel	Upper bound of frequency band
Δ_r	1 pixel	Sampling step between annular bands in the frequency spectrum
θ	180	Degrees of the spectrum to consider

6.3.2.3 Algorithm

- Compute the magnitude of the 2D-DFT $\dot{F}(u, v)$ of input image, discarding the DC component.
- Transform $\dot{F}(u, v)$ into polar coordinates and normalize to the range of $[0, 1]$.
- Determine the maximum energy (i.e. the maximum magnitude of all frequencies) to compute Q_{POW} (6.3.2.5).

6.3.2.4 Magnitude of frequency bands polar coordinates

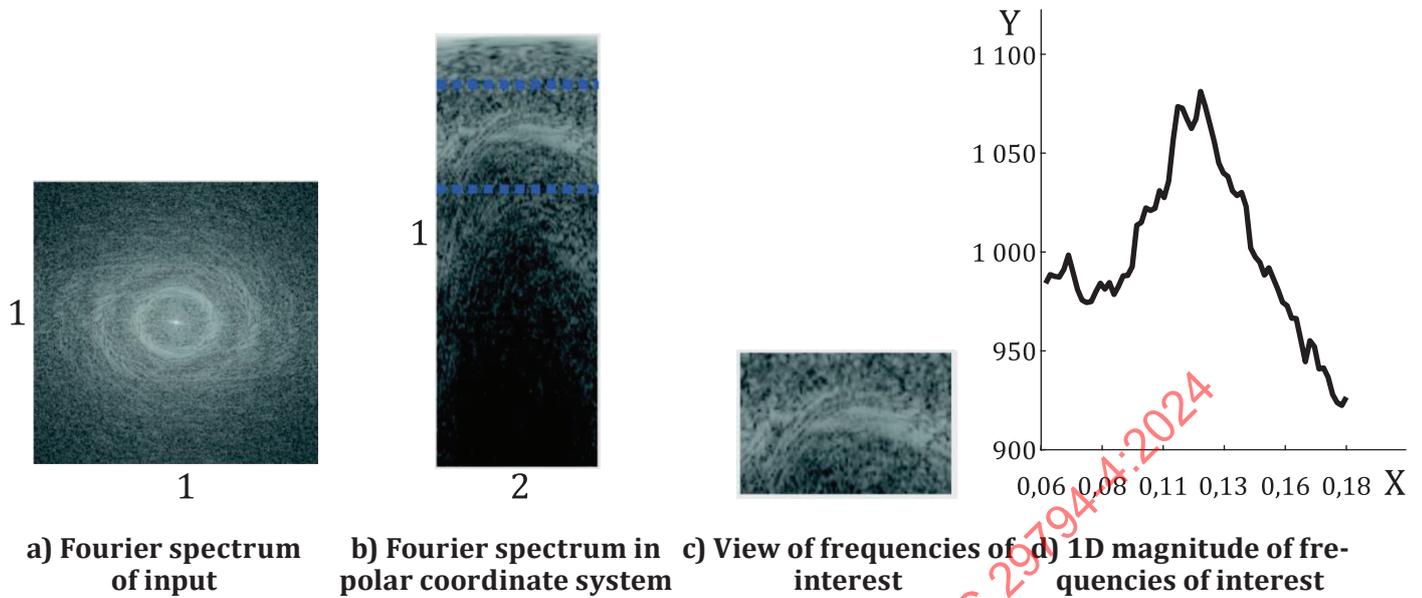
J , the magnitude of the annular band between r and $r + \Delta_r$ in the polar Fourier spectrum $\dot{F}(\alpha, r)$, is computed as shown in [Formula \(54\)](#):

$$J(r) = \frac{\sum_{\alpha=0}^{\pi} \sum_r^{r+\Delta_r} \dot{F}(\alpha, r)}{\sum_{\alpha=0}^{\pi} \sum_{r_{\min}}^{r_{\max}} \dot{F}(\alpha, r)} \quad (54)$$

where

- α is the angle;
- r is the radius.

$\dot{F}(\alpha, r)$ is the Spectrum $f(p, q)$ representation in polar coordinate system (α, r) , see [Figure 9](#).



Key

- X cycles/pixel
- Y magnitude
- 1 cycles/pixel
- 2 radians

Figure 9 — Processing steps of radial power spectrum algorithm

6.3.2.5 Determine quality from energy distribution

The quality feature q_{POW} is found as shown in [Formula \(55\)](#):

$$q_{POW} = \max_{r \in [\Delta r]} |J(r)| \tag{55}$$

NOTE The radial power spectrum depends on the spatial sampling rate. The given defaults assume 196,85 pixels per centimetre (500 pixels per inch).

6.3.3 Gabor filter bank

6.3.3.1 Feature description

The Gabor quality feature operates on a per-pixel basis by calculating the standard deviation of the Gabor filter bank responses.^[10] The size of the filter bank is used to determine a number of filters oriented evenly across the half circle. The strength of the response at a given location corresponds to the agreement between filter orientation and frequency in the location neighbourhood. For areas in the fingerprint image with a regular ridge-valley pattern there will be a high response from one or a few filter orientations. In areas containing background or unclear ridge-valley structure the Gabor response of all orientations will be low and constant.

6.3.3.2 Variables

Name	Default	Description
σ_x	6	2D Gaussian standard deviation in x -direction
σ_y	6	2D Gaussian standard deviation in y -direction
n	4	Size of filter bank (orientations of the Gabor wave)
f	0,1	Gabor filter frequency
θ	—	An orientation of a Gabor filter

6.3.3.3 Algorithm

- Convolve input image with a 2D Gaussian kernel with $\sigma = 1$ and subtract it from the input image I to give \tilde{I} .
- Compute the Gabor response of \tilde{I} for each orientation θ .
- Convolve the magnitude (complex modulus) of each Gabor response with a 2D Gaussian kernel with $\sigma = 4$.
- Compute the standard deviation of the Gabor magnitude response values at each location yielding a map of standard deviations.
- Sum the map of standard deviations and normalize according to number of sample points to produce the final Gabor quality.

Figure 10 visualizes the processing steps.

6.3.3.4 Gabor filter

The general form of the complex 2D Gabor^[11] filter, h_{Cx} , in the spatial domain is given by Formula (56):

$$h_{Cx}(x, y; f, \theta, \sigma_x, \sigma_y) = \exp\left(-\frac{1}{2}\left(\frac{x_\theta^2}{\sigma_x^2} + \frac{y_\theta^2}{\sigma_y^2}\right)\right) \exp(j2\pi fx_\theta) \quad (56)$$

where x_θ and y_θ are calculated using Formulae (57) and (58) respectively:

$$x_\theta = x \sin \theta + y \cos \theta \quad (57)$$

$$y_\theta = x \cos \theta - y \sin \theta \quad (58)$$

and f is the frequency (cycles/pixel) of the sinusoidal plane wave along the orientation θ . The size of the Gaussian smoothing window is determined by σ_x, σ_y .

The filter bank size n is used to compute the differently oriented Gabor filters composing the filter bank. Computing θ given n is done as shown in Formula (59):

$$\theta = \frac{k-1}{n\pi}, k=1, \dots, n \quad (59)$$

NOTE The Gabor quality feature is spatial sampling rate dependent. The given defaults assume 196,85 pixels per centimetre (500 pixels per inch).

6.3.3.5 Computing the Gabor quality from the Gabor filter response

Let G be a matrix with standard deviations of local responses resulting from convolution of I and Gabor filter of orientation n . The Gabor quality q_{GAB} is computed as shown in Formula (60):

$$q_{GAB} = \frac{1}{X \times Y} \sum_x \sum_y G(x, y) \quad (60)$$

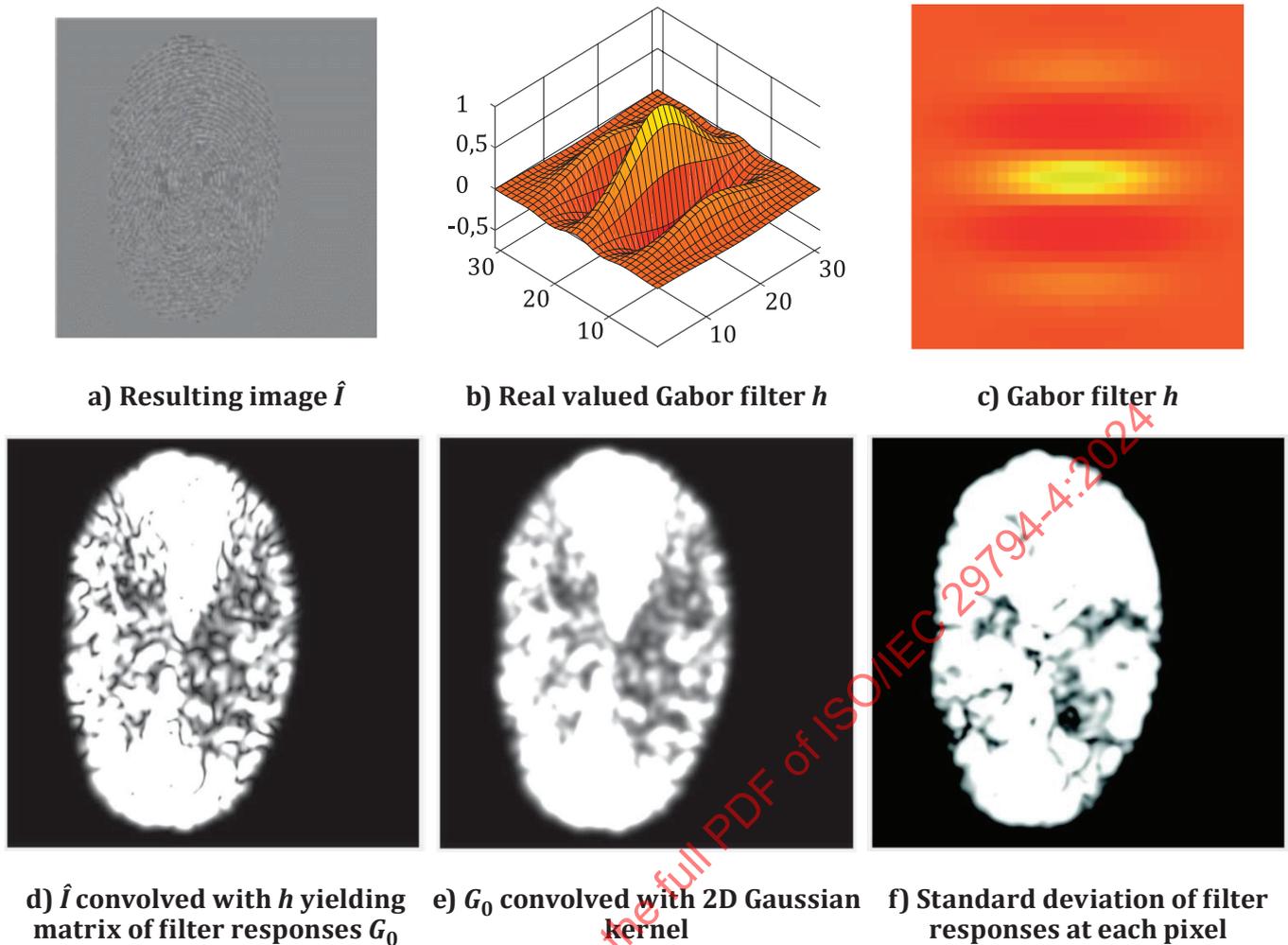


Figure 10 — Processing steps of Gabor algorithm

6.4 Unified quality score

6.4.1 Methodology for combining quality components

In order to obtain a single or unified output from several quality components described in the earlier clauses, it is necessary to combine those values to produce a single scalar quality score as required in the quality field of ISO/IEC 29794-1. Combining quality components shall be done such that the overall quality score is predictive of performance. There are various methods that can be used to combine quality components, e.g. weighted averaging, the use of pattern classifiers and other nonlinear computations.

6.4.2 Training method

Pattern classifiers are mathematical models that can intelligently learn a concept and predict an output when presented with new and even unseen samples. To apply pattern classification to combine the finger image quality analysis measures, it is necessary to train the pattern classifier by providing the values for the quality measures computed for training samples, along with the ground-truth classification [described in a) and b) below] for each of these samples. Once the pattern classifier is well-trained, given the values of the quality measures, it will be able to provide an overall quality score for the finger image.

The feature vector, $Q_{29794-4}$ (6.2.16.5), shall be the input to the pattern classifier. Training the pattern classifier can be performed using a corpus of finger images with pre-assigned quality categories or scores such as the QSND corpus, on the output of one or many quality algorithms. For all the samples in the corpus,

the feature vectors are computed. They are then paired with the quality category or score and fed into the pattern classifier for training.

With the feature vector specified in [6.2.16.5](#), a random forest shall be trained for binary classification, where Class 0 represents images of very low utility and Class 1 represents images of very high utility. The trained random forest outputs class membership along with its probability score. This score is the probability that a given image belongs to class 1 multiplied by 100 and rounded to its closest integer.

The training set shall be chosen such that:

- a) class 1 (or high utility) consists of images with NFIQ 1.0^[12] value of 1 (with activation score > 0,7) and mated similarity score in the 90th percentile for each of the comparison score providers;
- b) class 0 (or low utility) consists of images with NFIQ 1.0 value of 5 (with activation score > 0,9) and mated similarity score smaller than a decision threshold value that corresponds to false match rate of 1 in 10 000, i.e. false reject at false match rate of 0,000 1.

7 Finger image quality block

7.1 Binary encoding

In binary biometric data blocks, quality data should be encoded as specified in ISO/IEC 39794-1.

Quality measures for finger images encoded in the format specified in ISO/IEC 39794-4 should be placed into the quality blocks of that format. CBEFF quality fields should not be used in place of ISO/IEC 39794-4 quality fields but rather as supplementary data. The prescribed use of CBEFF quality fields may be supplied by each CBEFF patron format standard and is beyond the scope of this document. Multiple quality measures calculated by the same algorithm (i.e. same quality algorithm vendor identifier and same quality algorithm identifier) shall not be present in a single biometric data interchange record.

7.2 XML encoding

In XML documents, quality data shall be encoded as specified in ISO/IEC 39794-1.

7.3 Quality algorithm identifiers

The owner of the quality algorithms defined in this document is ISO/IEC JTC 1/SC 37. Its organization identifier is 257 (101_{Hex}). [Table 1](#) lists the quality algorithm identifiers for the quality measures defined in this document, which have been registered in accordance with ISO/IEC 19785-2 to identify the quality measures defined in this document.

The biometric organization identifier of ISO/IEC JTC 1/SC 37 shall be used as the quality algorithm vendor identifier if and only if a reference implementation approved by ISO/IEC JTC 1/SC 37 is used to compute the quality score. An approved reference implementation is posted at Reference [\[16\]](#).

If a unified quality score is calculated and reported in a finger image quality block, the normative quality measures defined in [6.2](#) should be calculated first. The values of the normative quality components defined in [6.2](#) may be reported in the finger image quality block, but this is not mandatory. Calculation of the non-normative quality components defined in [6.3](#) is optional. Their values cannot be reported in an ISO/IEC 29794-1 quality block because they are not mapped to integers in the range 0 to 100.

NOTE 1 The unified quality score summarizes the quality components in [Formula \(53\)](#) in [6.2.16.5](#).

NOTE 2 Revisions to the reference implementation NFIQ 2 have corrected software defects which calculated the unified quality score incorrectly. Tagged revisions to this reference implementation are listed in [Table 1](#).

Table 1 — Quality algorithm identifiers

Quality algorithm identifier Hex	Quality algorithm identifier decimal	Description	Symbol	Governing subclause
21 _{Hex}	33	Unified quality score, reference implementation NFIQ 2 tag v2.0.0	$Q_{\text{NFIQ2 v2.0.0}}$	6.4
22 _{Hex}	34	Mean of local orientation certainty level	Q_{OCL}^{μ}	6.2.16.2.2
23 _{Hex}	35	Standard deviation of local orientation certainty level	Q_{OCL}^{σ}	6.2.16.3.2
24 _{Hex}	36	Mean of local clarity	Q_{LCL}^{μ}	6.2.16.2.2
25 _{Hex}	37	Standard deviation of local clarity	Q_{LCL}^{σ}	6.2.16.3.2
26 _{Hex}	38	Mean of local frequency domain analysis	Q_{FDA}^{μ}	6.2.16.2.2
27 _{Hex}	39	Standard deviation of local frequency domain analysis	Q_{FDA}^{σ}	6.2.16.3.2
28 _{Hex}	40	Mean of local ridge valley uniformity	Q_{RVU}^{μ}	6.2.16.2.2
29 _{Hex}	41	Standard deviation of local ridge valley uniformity	Q_{RVU}^{σ}	6.2.16.3.2
2A _{Hex}	42	Mean of local orientation flow	Q_{OFL}^{μ}	6.2.16.2.2
2B _{Hex}	43	Standard deviation of orientation flow	Q_{OFL}^{σ}	6.2.16.3.2
2C _{Hex}	44	MU	Q_{MU}	6.2.7.3
2D _{Hex}	45	MMB	Q_{MMB}	6.2.8.3
2E _{Hex}	46	Minutiae count	$Q_{\text{MIN}}^{\text{CNT}}$	6.2.9.3
2F _{Hex}	47	Minutiae count in centre of mass	$Q_{\text{MIN}}^{\text{COM}}$	6.2.10.3
30 _{Hex}	48	Minutiae quality based on image mean	$Q_{\text{MIN}}^{\text{MU}}$	6.2.11.3
31 _{Hex}	49	Minutiae quality based on orientation certainty level	$Q_{\text{MIN}}^{\text{OCL}}$	6.2.12.3
32 _{Hex}	50	Region of interest image mean	Q_{AREA}^{μ}	6.2.13.4
33 _{Hex}	51	Region of interest orientation map coherence sum	$Q_{\text{COH}}^{\text{sum}}$	6.2.14.5
34 _{Hex}	52	Region of interest relative orientation map coherence sum	$Q_{\text{COH}}^{\text{REL}}$	6.2.15.3
35 _{Hex}	53	<i>Reserved for future use</i>		
36 _{Hex}	54	<i>Reserved for future use</i>		
37 _{Hex}	55	Unified quality score, reference implementation NFIQ 2 tag v2.1.0	$Q_{\text{NFIQ2 v2.1.0}}$	6.4
38 _{Hex}	56	Unified quality score, reference implementation NFIQ 2 tag v2.2.0	$Q_{\text{NFIQ2 v2.2.0}}$	6.4
39 _{Hex}	57	Unified quality score, reference implementation NFIQ 2 tag v2.3.0	$Q_{\text{NFIQ2 v2.3.0}}$	6.4
3A _{Hex} –40 _{Hex}	58–64	<i>Reserved for future use</i>		

Annex A (normative)

Conformance test assertions

A.1 Overview

In order to achieve the objective of this document, it is necessary to test biometric products to ensure their conformance. Conformant implementations are a necessary prerequisite for achieving interoperability among implementations. Therefore, there is a need for a standardized conformance testing methodology, test assertions and test procedures as applicable to specific modalities addressed by this document. The test assertions cover all normative contributive quality components, so that the conformity results produced by the test suites will reflect the real degree of conformity of the implementations to ISO/IEC 29794-4 finger image quality blocks.

This annex specifies conformance test criteria and a conformance test set to determine whether any implementation claiming conformance to this document conforms to the reference implementation NFIQ 2.

A.2 Conformance test set

To verify conformance, implementations of quality assessment algorithms claiming conformance with this document should run on all images in the conformance test set. In order to conform, the absolute value of the difference between the computed output value and the target unified quality score $Q_{\text{NFIQ2 v2.3.0}}$ in [Table A.1](#) shall not be more than 1 (i.e. 1 % of the maximum value 100). The contents of [Table A.1](#) are available in comma-separated values format from Reference [15]. The values of [Table A.1](#) mapped 0-100 to fit into a quality block as specified in ISO/IEC 29794-1 are shown in [Table A.2](#), which is also available in comma-separated values format from the same location.

The finger images in the conformance test set are selected from databases from the National Institute of Standards and Technology entitled Special Database 300[13] and Special Database 302.[14] The conformance test set can be requested from Reference [15].

NFIQ 2, the source code of which is available from Reference [16], is a reference implementation for this document. The pre-compiled release of version 2.3.0 of this reference implementation produces no differences from the values in [Table A.1](#) and [Table A.2](#).

Table A.1 — Target quality measure values for conformance test set

Identifier	$Q_{NF102.v2.3.0}$	μ_{FDA}	σ_{FDA}	COM η_{MIN}	CNT η_{MIN}	MU η_{MIN}	OC η_{MIN}	μ_{AREA}	μ_{LCL}	σ_{LCL}	η_{MMB}	η_{MU}	μ_{OCL}	σ_{OCL}	μ_{OFL}	σ_{OFL}	REF η_{COH}	SUM η_{COH}	μ_{RVU}	σ_{RVU}	
00001052_plain_500_10	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
00001057_plain_500_03	90	0,463 32	0,199 47	32	55	0,327 27	0,709 09	144,924 44	0,809 47	0,169 21	164,344 8	163,218 3	0,823 69	0,150 01	0,319 42	0,297 81	0,701 99	83,537 43	1,116 62	0,809 59	
00001308_plain_500_08	68	0,484 27	0,200 02	16	44	0,431 81	0,590 90	161,906 96	0,808 12	0,211 51	180,626 8	179,957 4	0,776 32	0,152 70	0,184 06	0,226 64	0,691 81	88,551 78	1,143 47	0,654 04	
00001428_plain_500_07	76	0,557 55	0,260 46	31	88	0,261 36	0,295 45	167,289 85	0,761 09	0,241 69	185,912 0	184,928 9	0,751 32	0,169 34	0,355 75	0,356 01	0,614 64	70,069 32	1,143 28	0,850 57	
00001610_plain_500_10	12	0,641 90	0,331 30	149	189	0,417 98	0,031 74	160,571 34	0,621 38	0,281 23	179,685 7	177,541 5	0,598 00	0,219 88	0,268 39	0,237 78	0,474 84	51,758 05	1,306 79	1,040 58	
00001655_plain_500_04	39	0,469 04	0,260 95	89	147	0,136 05	0,027 21	157,981 14	0,723 24	0,219 17	183,894 9	182,441 8	0,639 19	0,174 25	0,315 25	0,353 09	0,526 51	75,291 37	1,206 79	0,947 92	
00001665_plain_500_08	4	0,588 50	0,336 70	76	192	0,322 91	0,000 00	162,107 97	0,636 12	0,269 49	182,571 8	178,355 9	0,549 24	0,177 12	0,341 58	0,349 34	0,381 28	50,329 59	1,302 55	1,104 72	
00001723_plain_500_05	56	0,563 47	0,294 84	30	57	0,210 52	0,456 14	148,064 01	0,625 86	0,348 42	173,630 7	170,416 4	0,737 52	0,184 16	0,282 37	0,289 60	0,658 78	61,267 12	1,201 38	1,029 67	
00001880_plain_500_10	4	0,679 52	0,344 09	126	139	0,071 94	0,021 58	128,760 82	0,551 92	0,331 58	166,315 7	159,544 8	0,593 72	0,202 79	0,446 70	0,288 57	0,442 04	33,595 33	1,551 33	1,442 46	
00002302_N_500_palm_10	21	0,538 61	0,294 13	79	95	0,431 57	0,000 00	168,144 49	0,636 48	0,244 32	195,682 2	188,512 8	0,490 44	0,176 42	0,354 47	0,277 60	0,350 88	23,860 29	1,097 05	0,757 18	
00002302_N_500_palm_10-core	20	0,385 97	0,188 51	42	42	0,500 00	0,000 00	151,182 40	0,646 67	0,199 94	164,357 2	158,231 2	0,420 74	0,170 77	0,351 83	0,034 62	0,308 08	5,545 44	1,195 46	1,106 64	
00002304_R_500_slap_01	92	0,427 95	0,168 51	36	122	0,336 06	0,393 44	170,452 38	0,699 12	0,295 96	203,153 7	200,842 2	0,723 43	0,206 64	0,339 20	0,350 95	0,659 46	87,709 16	1,087 26	0,545 64	
00002304_R_500_slap_08	31	0,434 30	0,221 52	61	117	0,452 99	0,034 18	172,139 85	0,697 34	0,210 92	197,780 8	195,593 9	0,557 03	0,206 55	0,348 40	0,476 77	0,436 27	30,975 19	1,134 86	1,037 72	
00002310_R_500_slap_05	43	0,411 91	0,240 05	63	102	0,470 58	0,107 84	139,712 62	0,672 34	0,251 73	175,390 5	168,578 9	0,656 03	0,193 54	0,286 93	0,254 65	0,502 90	38,723 81	1,084 45	0,603 60	
00002314_R_500_slap_01	88	0,449 30	0,176 02	31	65	0,538 46	0,061 53	182,084 25	0,751 96	0,222 58	205,687 4	203,327 6	0,618 94	0,156 00	0,388 26	0,377 85	0,486 36	64,686 58	1,132 77	0,680 04	
00002315_R_500_slap_05	3	0,512 71	0,318 45	160	194	0,603 09	0,000 00	165,131 64	0,599 62	0,257 92	190,255 8	183,072 2	0,446 12	0,211 15	0,422 23	0,294 54	0,322 05	25,120 05	1,255 39	0,945 04	
00002315_R_500_slap_06	77	0,426 30	0,191 11	53	167	0,437 12	0,149 70	154,464 74	0,754 26	0,170 61	181,857 0	176,641 8	0,719 14	0,167 05	0,351 18	0,360 34	0,578 95	94,949 16	1,098 45	0,568 24	
00002317_R_500_slap_01	93	0,450 46	0,135 03	31	101	0,732 67	0,267 32	157,765 41	0,804 73	0,143 86	186,808 6	180,155 4	0,699 27	0,164 66	0,365 07	0,339 37	0,552 76	100,602 46	1,080 25	0,583 93	
00002319_R_500_slap_04	57	0,416 80	0,166 56	57	83	0,578 31	0,024 09	190,588 26	0,551 87	0,367 72	203,360 5	202,140 9	0,596 19	0,145 46	0,495 25	0,444 51	0,423 94	34,339 88	1,068 92	0,567 02	
00002319_R_500_slap_09	68	0,425 72	0,165 13	37	92	0,652 17	0,000 00	191,283 63	0,597 28	0,316 63	213,879 1	210,579 0	0,530 51	0,155 19	0,471 02	0,367 95	0,404 71	42,494 51	1,059 33	0,502 13	
00002319_R_500_slap_10	42	0,456 04	0,185 62	25	44	0,727 27	0,022 72	182,774 80	0,746 70	0,214 13	208,544 3	205,425 9	0,627 70	0,143 09	0,131 11	0,173 42	0,497 49	35,819 41	1,098 49	0,423 06	

Table A.1 (continued)

Identifier	Q_{NF1Q2} v2.3.0	q_{FDA}^{μ}	σ_{FDA}	COM q_{MIN}	CNT q_{MIN}	MU q_{MIN}	OC q_{MIN}	q_{AREA}^{μ}	q_{LCL}^{μ}	q_{LCL}^{σ}	q_{MMB}	q_{MU}	q_{OCL}^{μ}	q_{OCL}^{σ}	q_{OFL}^{μ}	q_{OFL}^{σ}	REL q_{COH}	sum q_{COH}	q_{RVU}^{μ}	σ_{RVU}
00002320_R_500_slap_06	47	0.42858	0.24771	19	177	0.29943	0.12429	145,43682	0.71815	0.18211	166,1679	161,9692	0.65248	0.17532	0.28954	0.30765	0.49243	82,23627	1,27433	0.97102
00002320_R_500_slap_08	7	0.45679	0.29690	117	206	0.37378	0.00970	158,41936	0.61032	0.26588	170,9628	171,3416	0.55029	0.18690	0.37024	0.36315	0.40440	44,88846	1,40971	1.30282
00002321_R_500_slap_07	21	0.45436	0.28205	105	170	0.57058	0.03529	189,22152	0.63872	0.24981	207,2029	200,8144	0.52574	0.20088	0.39363	0.31168	0.36980	32,54291	1,33493	1.10649
00002325_R_500_slap_04	22	0.43526	0.25587	121	133	0.23308	0.02255	136,81272	0.67383	0.23128	166,0485	163,9574	0.57759	0.15729	0.39543	0.33461	0.42009	40,32860	1,08247	0.89145
00002325_R_500_slap_08	31	0.45122	0.27264	67	141	0.17730	0.07092	130,35263	0.63210	0.26675	166,0399	160,5075	0.58983	0.19491	0.35736	0.30169	0.44301	45,18792	1,39276	1.32870
00002326_R_500_slap_02	89	0.42614	0.17253	34	120	0.40833	0.07500	172,35130	0.78010	0.14203	196,7841	192,8457	0.70793	0.16366	0.25438	0.26119	0.54189	73,69823	1,17719	0.69288
00002326_R_500_slap_05	1	0.46410	0.30197	146	240	0.42500	0.00833	183,40061	0.57526	0.30177	204,0556	202,2356	0.54135	0.18374	0.39789	0.34513	0.38222	37,07560	1,28363	0.97683
00002330_R_500_slap_04	49	0.40640	0.20211	67	102	0.64705	0.09803	163,81129	0.61877	0.30373	187,0958	182,5866	0.61544	0.17234	0.30186	0.31809	0.46101	43,79608	1,19145	0.65415
00002334_R_500_slap_05	29	0.40060	0.24249	80	89	0.52809	0.12359	144,66230	0.63830	0.27880	171,2261	164,4904	0.73171	0.13436	0.22758	0.24435	0.55158	34,19825	1,22902	0.86013
00002334_R_500_slap_07	83	0.42282	0.17412	46	82	0.63414	0.24390	162,14006	0.68994	0.28622	192,8766	185,9058	0.73360	0.16996	0.36531	0.31007	0.57786	54,31970	1,09665	0.51755
00002340_R_500_slap_08	28	0.39378	0.25225	98	147	0.57823	0.02040	195,31862	0.49002	0.33550	209,2274	208,6055	0.58743	0.17632	0.39812	0.34801	0.42523	35,71937	1,22760	0.82427
00002342_R_500_slap_05	5	0.50947	0.34114	184	236	0.46186	0.00000	162,25277	0.53838	0.31390	187,1672	182,8439	0.57635	0.14674	0.16868	0.19665	0.41020	36,91830	1,47206	1.22852
00002342_R_500_slap_09	16	0.42596	0.29197	79	154	0.38311	0.00649	162,20780	0.64134	0.25634	191,7997	188,8025	0.53286	0.20565	0.17350	0.20598	0.42236	46,88284	1,30787	1.12823
00002349_N_500_palm_05	26	0.44514	0.25764	63	101	0.50495	0.00000	105,33809	0.74323	0.14073	147,7897	141,9276	0.60414	0.17844	0.17262	0.20475	0.45447	40,90226	1,20163	0.82210
00002349_R_500_slap_07	13	0.46734	0.27942	99	152	0.51973	0.01973	133,31449	0.67565	0.21882	164,7814	160,8776	0.57054	0.15122	0.34748	0.31467	0.42779	35,93463	1,26174	1.04322
00002352_R_500_slap_01	85	0.47295	0.17776	24	99	0.76767	0.42424	149,14017	0.78090	0.23245	183,1052	173,9969	0.79222	0.13322	0.32466	0.35580	0.67875	114,70967	1,09700	0.64507
00002352_R_500_slap_10	50	0.39573	0.21589	45	78	0.71794	0.06410	184,67767	0.68934	0.21255	204,8418	201,1275	0.60261	0.15147	0.28007	0.27831	0.45270	42,10146	1,34500	0.93840
00002353_N_500_palm_10	39	0.45946	0.27086	79	134	0.41791	0.42537	146,79700	0.40517	0.39510	187,9935	179,2686	0.75006	0.20417	0.31030	0.31536	0.64351	49,55072	1,19208	0.74133
00002353_R_500_slap_06	48	0.40821	0.18306	60	183	0.30601	0.57923	128,02555	0.50404	0.38199	165,4250	161,4834	0.83112	0.14486	0.25666	0.30383	0.75977	100,29013	1,08693	0.60578
00002353_R_500_slap_10	35	0.44050	0.27035	88	130	0.55384	0.26923	182,83794	0.33650	0.37711	213,1230	208,7437	0.68275	0.16835	0.25766	0.26638	0.52761	41,15399	1,20848	1.09268
00002356_N_500_palm_05	47	0.42250	0.17295	52	89	0.33707	0.02247	130,13473	0.80628	0.05272	161,3879	158,5110	0.70056	0.12726	0.39644	0.36789	0.53416	57,68931	1,39617	1.32811

Table A.1 (continued)

Identifier	$Q_{NF1Q2\ v2.3.0}$	μ_{FDA}	σ_{FDA}	COM η_{MIN}	CNT η_{MIN}	MU η_{MIN}	OC η_{MIN}	μ_{AREA}	μ_{LCL}	σ_{LCL}	η_{MMB}	η_{MU}	μ_{OCL}	σ_{OCL}	μ_{OFL}	σ_{OFL}	REL η_{COH}	sum η_{COH}	μ_{RVU}	σ_{RVU}
00002357_R_500_slap_02	86	0.43232	0.16451	31	53	0.54717	0.30188	143,47705	0.80190	0.13555	166,6973	163,7842	0.75774	0.13024	0.37613	0.26737	0.60605	59,39376	1,08529	0.59116
00002357_R_500_slap_04	60	0.36192	0.13958	63	96	0.82291	0.23958	117,19903	0.74625	0.19523	144,1611	139,4977	0.76321	0.14243	0.29938	0.26794	0.60815	63,85630	1,16083	0.57848
00002357_R_500_slap_08	44	0.41594	0.21712	97	143	0.61538	0.07692	124,57630	0.71119	0.22460	154,4342	144,7081	0.68957	0.15437	0.37146	0.33608	0.53354	55,48875	1,20359	0.83136
00002358_R_500_slap_05	24	0.42752	0.22324	47	68	0.35294	0.00000	133,27844	0.69111	0.22465	173,7606	165,40775	0.62375	0.14380	0.11837	0.14079	0.46792	36,96607	1,06143	0.64883
00002359_J_500_palm_10	44	0.39863	0.17283	58	105	0.37142	0.07619	134,31180	0.73910	0.20659	163,2521	163,2521	0.60742	0.23142	0.28428	0.25485	0.51346	49,80605	1,19037	0.94461
00002361_R_500_slap_03	47	0.39788	0.18445	78	128	0.69531	0.03906	165,05254	0.65028	0.24456	181,3926	176,9338	0.59523	0.18898	0.28979	0.29030	0.43104	48,27695	1,18200	0.79856
00002362_N_500_palm_05	34	0.53332	0.29189	42	75	0.24233	0.18666	131,47815	0.73821	0.19029	159,3834	157,7113	0.68979	0.17344	0.21771	0.20428	0.54859	45,53303	1,24907	1,06423
00002362_R_500_slap_03	34	0.50776	0.28383	97	124	0.41129	0.07032	183,62964	0.59440	0.33616	204,4770	201,1449	0.62411	0.20615	0.24482	0.25712	0.49344	40,95569	1,34713	1,17365
00002362_R_500_slap_05	31	0.46830	0.26942	64	98	0.47959	0.09183	152,93190	0.66449	0.24899	181,3446	172,6756	0.66096	0.14221	0.24242	0.26419	0.49231	37,41570	1,21145	1,18458
00002362_R_500_slap_09	34	0.45367	0.25067	100	132	0.25000	0.06060	135,49762	0.70934	0.20841	166,3893	159,9462	0.65703	0.21099	0.26101	0.28986	0.53504	52,96941	1,18427	0.75563
00002364_R_500_slap_09	28	0.45211	0.26275	80	124	0.34677	0.04838	144,88302	0.62102	0.29609	169,9758	167,7797	0.56654	0.22304	0.59085	0.32526	0.42871	38,15535	1,12277	0.72587
00002367_R_500_slap_03	50	0.37378	0.13817	37	50	0.68000	0.00000	180,97500	0.73733	0.19381	199,4070	194,7487	0.65196	0.10719	0.38793	0.30393	0.48427	32,93098	1,05582	0,60012
00002368_J_500_palm_10	16	0.45438	0.30270	123	159	0.35849	0.03773	150,55885	0.62226	0.27679	171,0265	167,7420	0.64945	0.15930	0.25861	0.25378	0.49983	38,98740	1,26094	0,87108
00002371_J_500_palm_07	83	0.43749	0.16213	43	78	0.51282	0.39743	155,14987	0.78213	0.15719	182,7011	176,7565	0.76053	0.18560	0.29503	0.30470	0.66031	69,99326	1,08556	0,51590
00002373_R_500_slap_06	94	0.43313	0.16094	34	149	0.28187	0.34228	150,17132	0.75775	0.23192	183,9638	179,1577	0.75267	0.20437	0.36612	0.29513	0.69709	99,68445	1,07524	0,69674
00002374_R_500_slap_10	21	0.50232	0.29005	48	70	0.15714	0.00000	121,70483	0.75013	0.18990	168,5035	163,6173	0.61304	0.13328	0.07089	0.06164	0.45125	29,33178	1,30430	1,11433
00002375_J_500_palm_02	82	0.42963	0.13888	24	49	0.42857	0.55102	196,25108	0.75105	0.25144	208,5436	207,6294	0.80142	0.13539	0.23040	0.25069	0.67443	66,76941	1,12118	0,63284
00002376_R_500_slap_03	85	0.42495	0.16956	25	86	0.48837	0.04651	159,91945	0.74508	0.16841	180,0838	176,3743	0.61329	0.16339	0.26603	0.30382	0.44992	62,98890	1,23875	0,90125
00002376_R_500_slap_05	51	0.46156	0.22698	50	98	0.27551	0.04081	129,03228	0.74550	0.18369	166,8466	156,60883	0.62783	0.19227	0.26336	0.28705	0.47022	58,30821	1,21898	0,92994
00002383_J_500_palm_05	45	0.43139	0.20425	40	75	0.52000	0.06666	179,22749	0.75442	0.15101	196,8924	196,0434	0.66242	0.15231	0.19381	0.23485	0.48435	36,32658	1,20580	0,85998
00002383_R_500_slap_06	91	0.41231	0.12735	29	77	0.50649	0.46753	142,26136	0.81800	0.14916	169,1027	162,4066	0.81219	0.11721	0.27202	0.31624	0.69350	111,65397	1,11160	0,57554

Table A.1 (continued)

Identifier	$Q_{NF1Q2\ v2.3.0}$	q_{FDA}^{μ}	σ_{FDA}	COM q_{MIN}	CNT q_{MIN}	MU q_{MIN}	OCL q_{MIN}	q_{AREA}^{μ}	q_{LCL}^{μ}	q_{LCL}^{σ}	q_{MMB}	q_{MU}	q_{OCL}^{μ}	q_{OCL}^{σ}	q_{OFL}^{μ}	q_{OFL}^{σ}	REL q_{COH}	sum q_{COH}	q_{RVU}^{μ}	q_{RVU}^{σ}
00002384_N_500_palm_05	6	0,503 32	0,305 32	105	160	0,306 25	0,000 00	183,724 36	0,604 58	0,244 26	205,594 4	201,857 4	0,427 94	0,175 69	0,376 19	0,274 17	0,281 76	21,695 80	1,291 81	1,131 62
00002384_R_500_slap_04	3	0,526 60	0,337 41	148	255	0,360 78	0,000 00	170,213 54	0,524 71	0,325 36	201,371 0	198,721 1	0,506 08	0,186 96	0,149 19	0,206 62	0,351 61	34,458 25	1,291 40	0,906 75
00002386_R_500_slap_04	0	0,486 39	0,342 60	151	255	0,631 37	0,007 84	159,132 66	0,516 21	0,318 13	180,591 3	176,749 2	0,505 23	0,166 10	0,246 39	0,245 56	0,342 40	35,267 25	1,369 31	1,382 98
00002391_R_500_slap_07	31	0,398 13	0,230 62	120	184	0,581 52	0,000 00	197,651 30	0,674 78	0,223 62	211,736 5	207,948 4	0,558 95	0,135 31	0,425 29	0,317 82	0,416 59	41,659 02	1,189 21	0,857 49
00002392_R_500_slap_04	81	0,434 36	0,187 14	35	90	0,644 44	0,333 33	127,115 64	0,790 53	0,167 50	159,124 4	149,603 6	0,725 56	0,193 96	0,339 71	0,319 50	0,587 15	75,155 48	1,100 83	0,653 12
00002392_R_500_slap_08	78	0,421 84	0,136 53	52	97	0,680 41	0,247 42	149,357 46	0,770 67	0,198 76	169,904 4	165,527 5	0,709 65	0,163 15	0,313 90	0,304 37	0,553 76	67,559 62	1,117 32	0,727 55
00002396_R_500_slap_01	38	0,408 80	0,233 48	96	247	0,582 99	0,024 29	112,242 17	0,662 27	0,245 96	149,509 0	142,072 5	0,614 83	0,190 15	0,271 05	0,311 76	0,455 16	89,666 56	1,284 96	0,866 90
00002396_R_500_slap_08	21	0,371 67	0,233 41	119	244	0,704 91	0,012 29	168,999 54	0,568 18	0,285 84	188,545 3	182,151 5	0,524 85	0,199 29	0,170 27	0,208 53	0,378 91	41,680 05	1,284 60	1,030 64
00002399_J_500_palm_08	55	0,438 52	0,220 03	74	107	0,205 60	0,177 57	179,370 54	0,629 03	0,287 31	204,159 4	198,865 5	0,709 03	0,202 38	0,188 24	0,235 01	0,638 41	46,604 24	1,225 44	0,930 26
00002401_N_500_palm_02	67	0,438 18	0,166 40	19	33	0,393 93	0,303 03	133,583 12	0,805 04	0,109 26	156,644 3	150,503 8	0,735 76	0,142 65	0,388 38	0,338 35	0,557 89	49,652 88	1,173 04	0,686 17
00002401_R_500_slap_06	80	0,402 51	0,161 44	25	95	0,368 42	0,494 73	125,042 43	0,620 91	0,086 68	158,300 9	150,446 7	0,795 37	0,170 71	0,250 24	0,280 85	0,678 81	107,253 46	1,222 29	0,711 48
00002401_R_500_slap_08	75	0,361 41	0,122 03	32	59	0,559 32	0,186 44	158,808 71	0,792 87	0,063 69	174,766 2	171,730 8	0,697 88	0,168 90	0,335 27	0,315 98	0,540 12	51,851 78	1,148 11	0,695 03
00002403_R_500_slap_04	30	0,388 23	0,226 85	87	151	0,450 33	0,000 00	151,456 37	0,663 29	0,233 29	178,386 1	169,681 3	0,610 28	0,104 65	0,181 85	0,228 71	0,448 06	55,112 41	1,275 31	0,935 66
00002405_R_500_slap_01	75	0,452 83	0,116 09	18	62	0,612 90	0,822 58	152,043 28	0,825 09	0,172 05	182,762 4	176,003 6	0,879 61	0,096 74	0,230 11	0,296 14	0,797 28	126,768 94	1,048 18	0,404 72
00002406_R_500_slap_05	4	0,536 36	0,339 87	122	156	0,512 82	0,000 00	134,097 43	0,632 65	0,253 94	163,682 9	158,369 2	0,616 26	0,129 53	0,189 56	0,167 88	0,445 20	30,274 08	1,386 92	1,352 78
00002406_R_500_slap_08	9	0,459 66	0,304 77	156	223	0,506 72	0,008 96	173,851 00	0,593 12	0,283 03	193,388 8	188,586 8	0,563 67	0,154 57	0,218 83	0,211 07	0,396 86	35,320 64	1,321 95	1,213 15
00002408_R_500_slap_02	60	0,380 21	0,181 21	58	94	0,638 29	0,042 55	168,962 25	0,707 43	0,221 31	190,095 3	186,181 7	0,657 14	0,132 69	0,318 39	0,255 14	0,493 27	43,901 38	1,116 58	0,734 06
00002408_R_500_slap_08	46	0,409 21	0,227 82	78	99	0,797 98	0,040 40	173,707 48	0,688 45	0,229 32	190,961 0	185,570 1	0,648 62	0,129 10	0,505 03	0,321 25	0,477 40	44,876 48	1,234 46	0,901 98
00002409_R_500_slap_02	63	0,443 66	0,212 34	37	73	0,369 86	0,013 69	169,465 32	0,739 08	0,181 22	192,998 0	192,584 1	0,628 94	0,141 75	0,274 63	0,265 94	0,478 28	53,089 25	1,259 79	1,111 73
00002409_R_500_slap_09	15	0,463 50	0,298 85	101	204	0,617 64	0,019 60	120,635 36	0,602 17	0,297 18	143,365 3	140,667 5	0,667 74	0,182 26	0,181 74	0,222 32	0,503 17	57,362 04	1,356 99	1,039 35
00002415_R_500_slap_05	31	0,431 00	0,257 77	44	77	0,584 41	0,129 87	122,935 78	0,683 84	0,199 41	156,023 2	150,791 4	0,633 61	0,169 44	0,316 60	0,386 57	0,500 96	37,071 13	1,149 41	0,717 62

Table A.1 (continued)

Identifier	Q_{NF1Q2} v2.3.0	μ_{FDA}	σ_{FDA}	COM η_{MIN}	CNT η_{MIN}	MU η_{MIN}	OC η_{MIN}	μ_{AREA}	μ_{LCL}	σ_{LCL}	η_{MMB}	μ_{MU}	μ_{OCL}	σ_{OCL}	μ_{OFL}	σ_{OFL}	REL η_{COH}	sum η_{COH}	μ_{RVU}	σ_{RVU}
00002415_R_500_slap_09	48	0.407 61	0.179 35	34	56	0.678 57	0.267 85	93,882 52	0.784 32	0.151 87	125,222 0	119,013 8	0.801 07	0.132 15	0.187 77	0.204 86	0.677 99	56,273 33	1,194 66	0.820 60
00002418_R_500_slap_06	72	0.418 27	0.193 67	39	133	0.270 67	0.240 60	116,802 02	0.750 22	0.200 58	152,219 8	144,958 1	0.754 26	0.152 57	0.308 07	0.312 50	0.623 69	107,276 26	1,238 64	0.880 33
00002420_R_500_slap_04	63	0.421 55	0.197 44	51	81	0.419 75	0.185 18	159,025 47	0.678 63	0.269 70	188,739 9	183,469 3	0.658 17	0.205 37	0.272 68	0.243 30	0.496 83	44,715 43	1,196 52	0.747 76
00002420_R_500_slap_09	58	0.405 06	0.211 66	44	85	0.235 29	0.317 64	119,814 25	0.713 32	0.251 45	161,253 9	152,519 34	0.776 34	0.132 59	0.235 45	0.212 76	0.618 83	60,646 19	1,207 82	0.729 29
00002424_R_500_slap_01	94	0.451 53	0.163 60	25	85	0.258 82	0.247 05	165,057 91	0.788 65	0.174 03	192,594 2	189,355 1	0.692 59	0.169 70	0.307 89	0.292 47	0.551 27	85,447 10	1,083 16	0.489 97
00002428_R_500_slap_02	93	0.446 93	0.145 66	31	67	0.417 91	0.507 46	158,139 36	0.821 80	0.135 88	186,372 3	179,148 0	0.771 29	0.184 72	0.286 13	0.270 81	0.662 86	69,600 33	1,039 49	0.541 10
00002433_J_500_palm_08	94	0.439 28	0.143 24	24	62	0.580 64	0.580 64	164,819 34	0.765 80	0.219 99	187,839 0	179,121 6	0.769 93	0.181 30	0.233 20	0.267 43	0.677 84	89,475 83	1,101 19	0.572 88
00002433_J_500_palm_09	86	0.449 22	0.150 16	30	83	0.301 20	0.522 16	144,687 16	0.800 92	0.156 19	175,659 4	168,941 9	0.739 54	0.230 42	0.238 74	0.256 09	0.664 24	96,315 35	1,095 20	0.492 30
00002433_N_500_palm_07	84	0.437 17	0.125 25	24	65	0.553 84	0.646 15	128,568 38	0.811 17	0.162 76	155,676 8	151,752 3	0.826 81	0.130 74	0.232 10	0.251 57	0.702 74	93,465 28	1,083 06	0.441 01
00002434_N_500_palm_04	66	0.441 40	0.205 67	49	77	0.207 79	0.233 76	155,598 72	0.774 70	0.155 93	186,805 5	181,819 4	0.705 60	0.214 22	0.316 79	0.331 31	0.610 25	45,159 18	1,125 60	0.603 98
00002435_N_500_palm_03	82	0.436 20	0.154 19	27	45	0.444 44	0.511 11	179,418 28	0.709 93	0.301 01	199,906 2	193,717 53	0.825 53	0.139 41	0.262 62	0.256 95	0.700 32	60,927 90	1,073 11	0.459 12
00002437_S_500_slap_08	74	0.430 80	0.163 31	21	51	0.196 07	0.274 51	151,040 39	0.756 02	0.230 20	178,135 5	172,826 1	0.776 96	0.097 72	0.209 89	0.258 75	0.645 57	63,911 68	1,119 81	0.604 63
00002439_Q_500_palm_02	79	0.415 66	0.161 13	55	88	0.568 18	0.227 27	179,827 68	0.743 34	0.242 94	196,555 2	193,014 1	0.739 51	0.109 30	0.300 38	0.246 30	0.571 39	59,996 68	1,076 97	0.442 05
00002439_S_500_slap_10	33	0.428 45	0.233 24	77	97	0.432 99	0.000 00	180,749 23	0.667 80	0.227 81	203,817 1	199,744 8	0.577 43	0.156 72	0.303 39	0.233 18	0.429 98	26,229 21	1,273 90	0.954 76
00002440_R_500_slap_03	94	0.429 62	0.140 98	34	86	0.802 32	0.069 76	190,435 99	0.776 10	0.187 43	202,165 6	200,584 4	0.665 53	0.182 05	0.282 64	0.336 28	0.520 18	71,785 85	1,082 83	0.494 21
00002443_Q_500_palm_10	55	0.440 78	0.202 44	28	49	0.408 16	0.326 53	139,353 19	0.787 39	0.157 64	172,063 7	164,200 0	0.740 94	0.189 95	0.158 57	0.192 96	0.612 11	43,460 11	1,089 14	0.483 93
00002445_J_500_palm_02	81	0.478 20	0.221 78	35	65	0.292 30	0.323 07	136,475 83	0.791 50	0.161 50	170,631 2	160,012 7	0.762 36	0.172 67	0.317 60	0.294 36	0.657 33	69,677 61	1,153 15	0.584 84
00002445_S_500_slap_05	48	0.463 44	0.263 30	62	90	0.455 55	0.100 00	90,216 01	0.775 89	0.130 85	124,642 6	121,375 8	0.737 20	0.144 77	0.284 72	0.286 25	0.575 55	59,857 64	1,242 90	0.859 94
00002447_R_500_slap_01	38	0.385 97	0.223 74	94	255	0.654 90	0.007 84	189,521 87	0.602 53	0.304 60	209,229 5	205,250 91	0.600 91	0.153 94	0.344 07	0.320 76	0.445 51	73,064 16	1,267 93	0.904 37
00002447_R_500_slap_06	56	0.388 97	0.195 29	89	229	0.790 39	0.017 46	174,964 77	0.708 83	0.222 35	194,283 6	190,370 6	0.622 49	0.171 00	0.324 41	0.322 83	0.467 01	79,859 65	1,167 87	0.871 59
00002454_N_500_palm_02	78	0.454 44	0.190 16	29	69	0.420 29	0.304 34	146,107 13	0.782 25	0.167 86	172,307 4	168,318 1	0.742 03	0.143 55	0.277 19	0.251 76	0.608 64	70,602 48	1,227 08	1.051 87

Table A.1 (continued)

Identifier	Q_{NF1Q2} v2.3.0	μ_{FDA}	σ_{FDA}	COM η_{MIN}	CNT η_{MIN}	MU η_{MIN}	OC η_{MIN}	μ_{AREA}	μ_{LCL}	σ_{LCL}	η_{MMB}	η_{MU}	μ_{OCL}	σ_{OCL}	μ_{OFL}	σ_{OFL}	REL η_{COH}	sum η_{COH}	μ_{RVU}	σ_{RVU}
00002455_R_500_slap_01	89	0.442 20	0.178 15	21	106	0.754 71	0.018 86	186,410 70	0.745 55	0.189 57	203,278 6	199,010 1	0.639 59	0.133 30	0.254 00	0.305 73	0.478 73	85,215 12	1,123 91	0.599 40
00002458_J_500_palm_02	61	0.389 66	0.153 64	73	122	0.491 80	0.000 00	180,321 71	0.679 01	0.265 89	196,343 2	195,890 8	0.658 82	0.173 80	0.349 37	0.285 92	0.497 57	48,264 53	1,068 08	0.506 15
00002458_N_500_palm_09	19	0.466 97	0.272 71	149	216	0.444 44	0.101 85	157,781 98	0.495 44	0.345 76	181,481 3	176,448 3	0.553 60	0.243 35	0.351 30	0.345 13	0.411 62	42,808 72	1,232 90	0.887 90
00002458_S_500_slap_02	25	0.474 94	0.278 20	144	147	0.687 07	0.095 23	179,955 85	0.578 76	0.329 50	192,105 0	192,290 6	0.663 77	0.194 95	0.465 92	0.319 43	0.507 26	42,610 17	1,193 94	1,084 11
00002462_N_500_palm_02	33	0.486 83	0.243 58	28	65	0.523 07	0.046 15	169,284 43	0.746 00	0.142 51	198,309 9	192,148 6	0.560 41	0.228 86	0.237 00	0.234 96	0.456 20	41,971 24	1,366 95	1,115 15
00002462_R_500_slap_01	89	0.424 81	0.156 82	29	114	0.342 10	0.157 89	160,314 16	0.786 88	0.170 10	182,371 6	177,982 3	0.720 55	0.157 35	0.267 15	0.296 58	0.563 31	100,833 10	1,144 24	0.725 26
00002462_R_500_slap_03	17	0.457 30	0.251 35	65	111	0.360 36	0.027 02	147,501 95	0.703 51	0.199 04	167,209 1	165,045 7	0.549 52	0.186 88	0.418 38	0.415 20	0.376 23	38,375 62	1,411 53	1,143 52
00002463_R_500_slap_06	69	0.428 00	0.162 99	41	90	0.744 44	0.000 00	167,166 86	0.766 36	0.106 87	188,558 0	185,720 7	0.609 06	0.091 20	0.280 48	0.295 79	0.443 78	71,006 20	1,151 73	0.642 66
00002466_N_500_palm_04	77	0.463 24	0.189 65	29	63	0.190 47	0.238 09	134,530 12	0.787 41	0.174 59	172,132 5	167,328 0	0.708 81	0.199 04	0.429 54	0.386 42	0.575 94	65,657 39	1,054 88	0.460 80
00002475_J_500_palm_04	23	0.410 00	0.238 25	78	105	0.304 76	0.000 00	194,398 86	0.620 45	0.261 44	222,065 2	219,182 4	0.456 28	0.154 84	0.487 13	0.390 49	0.329 00	18,095 49	1,272 31	1,084 15
00002475_J_500_palm_10	1	0.529 61	0.313 09	67	101	0.346 53	0.000 00	190,169 86	0.636 31	0.225 91	215,546 3	209,455 5	0.421 45	0.170 73	0.334 60	0.235 34	0.323 49	22,320 88	1,365 77	1,097 75
00002475_S_500_slap_01	10	0.438 13	0.283 18	86	255	0.486 27	0.015 68	150,615 84	0.590 91	0.297 18	178,803 6	171,129 6	0.564 81	0.146 55	0.283 94	0.288 16	0.400 93	59,738 90	1,329 90	1,209 44
00002475_S_500_slap_04	20	0.381 27	0.213 56	76	186	0.322 58	0.000 00	133,442 38	0.687 56	0.189 07	162,899 8	154,048 0	0.495 06	0.155 26	0.503 48	0.320 32	0.329 10	33,897 78	1,415 60	1,336 38
00002478_N_500_palm_05	3	0.586 18	0.329 85	90	144	0.381 94	0.055 55	140,624 84	0.627 88	0.243 68	183,770 1	178,336 5	0.536 97	0.251 09	0.369 08	0.337 99	0.394 74	33,948 04	1,423 46	1,412 54
00002478_Q_500_palm_03	31	0.452 01	0.244 56	93	156	0.179 48	0.083 33	151,369 19	0.586 47	0.312 47	173,962 7	169,096 9	0.605 20	0.234 65	0.339 87	0.346 92	0.474 44	48,867 84	1,188 27	0,922 44
00002484_Q_500_palm_05	20	0.423 90	0.245 42	58	66	0.484 84	0.075 75	161,833 16	0.647 95	0.249 93	188,810 1	181,393 6	0.638 62	0.155 26	0.255 02	0.265 16	0.478 46	25,837 16	1,425 75	1,231 47
00002489_J_500_palm_02	84	0.452 32	0.132 07	33	56	0.553 57	0.678 57	144,438 03	0.832 60	0.129 65	172,341 9	168,705 7	0.803 12	0.166 12	0.291 08	0.269 75	0.724 06	82,543 41	1,036 70	0,331 18
00002494_J_500_palm_04	40	0.438 41	0.202 71	50	75	0.400 00	0.013 33	170,754 12	0.679 11	0.214 74	199,548 4	196,474 2	0.490 06	0.184 19	0.266 58	0.184 43	0.399 29	24,357 03	1,203 93	0,813 94
00002494_J_500_palm_10	13	0.525 58	0.298 40	78	100	0.440 00	0.080 00	141,302 49	0.699 39	0.184 28	181,080 9	169,765 30	0.597 30	0.200 65	0.140 36	0.099 81	0.458 84	31,201 23	1,224 47	1,000 43
00002494_S_500_slap_03	75	0.381 61	0.135 54	34	56	0.410 71	0.142 85	170,496 41	0.763 61	0.129 55	194,590 3	187,838 3	0.681 54	0.155 94	0.324 99	0.293 93	0.532 25	50,564 14	1,159 99	0,588 91
00002495_J_500_palm_03	86	0.478 20	0.185 19	24	59	0.644 06	0.254 23	182,556 75	0.801 11	0.191 12	203,431 1	199,554 1	0.701 42	0.183 21	0.364 73	0.352 67	0.585 68	78,482 36	1,135 12	1,026 44

Table A.1 (continued)

Identifier	$Q_{NF1Q2\ v2.3.0}$	μ_{FDA}	σ_{FDA}	COM η_{MIN}	CNT η_{MIN}	MU η_{MIN}	OC η_{MIN}	μ_{AREA}	μ_{LCL}	σ_{LCL}	η_{MMB}	η_{MU}	μ_{OCL}	σ_{OCL}	μ_{OFL}	σ_{OFL}	REL η_{COH}	sum η_{COH}	μ_{RVU}	σ_{RVU}
00002495_S_500_slap_08	83	0,46179	0,16234	24	52	0,75000	0,23076	182,77830	0,76287	0,23545	201,5593	198,2629	0,65630	0,20598	0,30970	0,32627	0,53312	59,17648	1,13182	0,64084
00002497_N_500_palm_08	16	0,49304	0,34219	47	50	0,50000	0,00000	223,29629	0,49523	0,33384	241,2249	241,2835	0,37130	0,23069	0,14697	0,07717	0,35444	12,05122	1,25546	0,82725
00002497_N_500_palm_09	0	0,56000	0,34077	169	219	0,47488	0,00000	195,70155	0,54530	0,29471	227,0198	224,5644	0,41638	0,20429	0,42953	0,23779	0,35540	17,77020	1,36079	1,37437
00002502_Q_500_palm_05	29	0,48832	0,26915	54	61	0,77049	0,06557	113,60927	0,73135	0,12532	149,4669	139,1367	0,59346	0,19895	0,37408	0,27058	0,45290	33,51516	1,43200	1,43022
00002504_N_500_palm_03	83	0,44128	0,16722	42	113	0,38938	0,31858	153,28360	0,69560	0,29662	185,8852	183,6091	0,73863	0,19557	0,32904	0,31113	0,65130	74,90033	1,17871	0,67056
00002504_S_500_slap_06	93	0,44950	0,16859	36	87	0,58620	0,24137	155,71579	0,77363	0,16211	176,2892	175,0218	0,68387	0,18236	0,34230	0,30628	0,54632	89,59673	1,15977	0,87416
00002505_N_500_palm_10	12	0,57333	0,31513	30	44	0,43781	0,18181	147,60927	0,66681	0,25460	187,7016	179,7347	0,59771	0,20185	0,15279	0,24003	0,46019	34,05472	1,27926	1,07343
00002514_Q_500_palm_03	72	0,43407	0,15839	29	40	0,45000	0,30000	170,45351	0,79458	0,16188	198,3571	193,2994	0,73227	0,13514	0,30440	0,29443	0,59187	37,88015	1,12803	0,47172
00002516_S_500_slap_09	48	0,44698	0,25081	70	123	0,63414	0,06504	142,75625	0,75421	0,15906	168,4901	162,8628	0,66369	0,17167	0,20186	0,21585	0,50026	49,52576	1,28680	0,96569
00002521_Q_500_palm_10	13	0,45471	0,27447	107	244	0,62295	0,01229	150,56760	0,60043	0,27207	171,3819	166,0930	0,53849	0,17251	0,22846	0,21640	0,38513	45,83083	1,30934	1,23989
00002523_J_500_palm_03	70	0,47002	0,21026	36	54	0,61111	0,01851	206,48474	0,74794	0,15394	222,1027	218,3158	0,59699	0,19561	0,32496	0,27386	0,44682	33,06466	1,14953	0,75956
00002523_J_500_palm_05	8	0,46941	0,29825	103	123	0,47967	0,00000	189,25872	0,59874	0,27243	212,3879	208,0666	0,47163	0,19327	0,30969	0,24281	0,37379	19,81125	1,45088	1,34989
00002523_S_500_slap_04	48	0,41938	0,21120	78	113	0,70796	0,02654	180,19840	0,74398	0,14263	205,0209	196,0977	0,62041	0,14974	0,38895	0,27383	0,47453	40,33516	1,21783	0,85807
00002523_S_500_slap_05	19	0,43000	0,25078	122	145	0,72413	0,00689	191,23440	0,63127	0,25967	210,2600	206,0959	0,60029	0,18497	0,19461	0,21160	0,43883	28,96334	1,37342	1,42120
00002528_N_500_palm_05	59	0,43420	0,22570	40	74	0,45945	0,31081	127,23412	0,74614	0,19186	161,3593	153,1599	0,73221	0,18568	0,29203	0,29057	0,62165	66,51666	1,18349	0,71708
00002531_Q_500_palm_09	35	0,47681	0,27513	70	94	0,38297	0,13829	130,16226	0,73514	0,21821	154,8473	151,9704	0,71183	0,18619	0,33895	0,30691	0,56829	46,03169	1,23192	1,05229
00002533_N_500_palm_10	27	0,45695	0,26658	136	202	0,61386	0,03465	96,38731	0,66134	0,25380	137,9797	132,9817	0,63563	0,20287	0,35011	0,28141	0,49877	47,38394	1,25607	0,99186
00002533_Q_500_palm_09	50	0,41828	0,21713	97	164	0,61585	0,16463	92,49101	0,72409	0,20861	125,6309	125,0877	0,71744	0,22106	0,21350	0,25429	0,59223	63,96182	1,18954	0,88829
00002533_S_500_slap_08	58	0,45642	0,24179	52	58	0,44827	0,24137	132,45085	0,68872	0,28096	166,9048	156,8935	0,80731	0,13636	0,43772	0,28369	0,67054	46,93791	1,07420	0,57807
00002534_N_500_palm_05	54	0,41306	0,20576	34	43	0,34883	0,23255	132,04778	0,73331	0,19653	175,3678	160,7380	0,68702	0,13300	0,33668	0,28362	0,54979	37,93554	1,14345	0,59991
00002544_S_500_slap_03	88	0,43976	0,13352	30	75	0,52000	0,25333	156,18909	0,79820	0,13941	177,4049	173,2623	0,68493	0,16064	0,22237	0,25829	0,54666	75,43962	1,10302	0,58680

Table A.1 (continued)

Identifier	$Q_{NF1Q2\ v2.3.0}$	μ_{FDA}	σ_{FDA}	COM η_{MIN}	CNT η_{MIN}	MU η_{MIN}	OC η_{MIN}	μ_{AREA}	μ_{LCL}	σ_{LCL}	η_{MMB}	η_{MU}	μ_{OCL}	σ_{OCL}	μ_{OFL}	σ_{OFL}	REL η_{COH}	sum η_{COH}	μ_{RVU}	σ_{RVU}
00002544_S_500_slap_08	84	0.47254	0.18357	22	88	0.56818	0.12500	170,52992	0.77708	0.19978	190,2335	187,5830	0.68955	0.15903	0.18896	0.26012	0.57346	74,55018	1,09530	0.60753
00002547_N_500_palm_08	70	0.37934	0.13075	38	65	0.55384	0.23076	124,87903	0.77959	0.16425	152,4587	143,1632	0.72009	0.12495	0.22919	0.21060	0.55219	55,21912	1,13037	0.76290
00002547_Q_500_palm_03	73	0.42658	0.16652	40	70	0.31428	0.11428	134,74216	0.78236	0.13208	167,7124	157,4012	0.68057	0.13148	0.34712	0.28663	0.51704	52,22101	1,20749	0.70155
00002547_S_500_slap_04	65	0.41348	0.21313	35	62	0.19354	0.09677	130,82492	0.76775	0.14353	177,3228	172,8140	0.67690	0.12271	0.30771	0.27392	0.51121	44,47534	1,20816	0.82118
00002551_S_500_slap_10	48	0.39181	0.22087	66	74	0.25675	0.08108	184,40687	0.42431	0.35958	219,5983	213,2676	0.56934	0.21516	0.26655	0.17263	0.53508	22,47368	1,26672	1.06271
00002554_J_500_palm_08	17	0.46878	0.29590	73	103	0.27184	0.00000	198,09211	0.60331	0.27749	221,9887	218,3980	0.51186	0.21527	0.35822	0.37210	0.43181	32,81759	1,26896	0.91266
00002554_S_500_slap_07	26	0.40902	0.24116	158	204	0.54902	0.00000	210,64146	0.50927	0.32081	223,5411	220,3435	0.48048	0.15237	0.25549	0.27643	0.33765	29,03813	1,29317	1.01313
00002555_S_500_slap_08	64	0.41557	0.19315	40	68	0.35294	0.50000	107,03741	0.70929	0.26055	151,7984	139,7428	0.77789	0.17528	0.28939	0.29736	0.67833	65,11969	1,16213	0.60821
00002559_S_500_slap_05	10	0.45995	0.29612	129	231	0.41125	0.03896	182,56494	0.56406	0.30149	207,4267	203,2530	0.58349	0.19600	0.28432	0.30747	0.42529	43,38022	1,28146	1.05481
00002559_S_500_slap_07	62	0.40788	0.19510	36	100	0.33000	0.03000	178,41424	0.63498	0.28616	203,9667	202,6570	0.64456	0.11331	0.18796	0.21518	0.49874	50,87235	1,14076	0.80673
00002559_S_500_slap_09	62	0.39213	0.18444	38	87	0.43678	0.22988	152,23717	0.70520	0.22131	180,2472	177,3790	0.70971	0.16567	0.19011	0.26240	0.56766	70,38997	1,10564	0.66585
00002560_N_500_palm_10	18	0.43104	0.21133	24	26	0.15384	0.00000	162,27037	0.70682	0.21601	197,2127	189,3814	0.59805	0.19336	0.23370	0.24262	0.47607	26,65994	1,13710	0.49857
00002564_S_500_slap_09	4	0.48967	0.33058	176	255	0.37647	0.00392	171,14308	0.58826	0.25711	193,0487	188,5095	0.42176	0.17920	0.45676	0.33311	0.29116	24,16699	1,24740	1.10907
00002568_J_500_palm_04	22	0.43491	0.25618	122	172	0.31976	0.02325	128,24252	0.59836	0.29801	155,3699	147,3058	0.55774	0.23756	0.45936	0.34551	0.44205	38,45851	1,14334	0.90132
00002568_J_500_palm_10	31	0.44769	0.26836	137	166	0.56626	0.03012	170,06861	0.49140	0.36361	196,8166	188,4401	0.63857	0.19944	0.22857	0.27924	0.46635	33,57744	1,22597	0.97313
00002569_N_500_palm_08	20	0.41530	0.25248	104	191	0.63350	0.00000	190,51134	0.65981	0.23524	203,2511	199,4955	0.55328	0.17294	0.50864	0.39608	0.40074	42,07846	1,32108	1.09103
00002569_N_500_palm_09	9	0.44078	0.29086	144	255	0.42352	0.00000	186,02354	0.62558	0.26101	204,1864	201,6569	0.51196	0.14938	0.59620	0.36095	0.34376	38,50217	1,22607	0.96544
00002569_S_500_slap_01	74	0.39807	0.14906	39	90	0.74444	0.14444	168,37022	0.77158	0.17410	185,6544	183,2469	0.74075	0.10657	0.37806	0.35930	0.59032	93,27056	1,11080	0.64570
00002569_S_500_slap_03	37	0.39387	0.19255	82	151	0.72185	0.00000	161,95617	0.74900	0.16475	181,2401	178,6969	0.60118	0.17351	0.42437	0.34962	0.44267	52,67780	1,22905	0.99402
00002569_S_500_slap_08	16	0.43641	0.28213	88	208	0.62019	0.01442	192,37669	0.64949	0.24495	204,3258	200,5118	0.61774	0.16838	0.36469	0.32946	0.44232	49,54055	1,30073	1.06970
00002571_S_500_slap_07	88	0.41769	0.11847	25	52	0.59615	0.36538	200,54226	0.79014	0.18981	216,1900	213,7541	0.74792	0.15067	0.26125	0.26553	0.64516	67,09679	1,09638	0.44948

Table A.1 (continued)

Identifier	$Q_{NF1Q2\ v2.3.0}$	μ_{FDA}	σ_{FDA}	COM η_{MIN}	CNT η_{MIN}	MU η_{MIN}	OC η_{MIN}	μ_{AREA}	μ_{LCL}	σ_{LCL}	η_{MMB}	η_{MU}	μ_{OCL}	σ_{OCL}	μ_{OFL}	σ_{OFL}	REL η_{COH}	sum η_{COH}	μ_{RVU}	σ_{RVU}
00002571_S_500_slap_10	81	0.44684	0.17485	33	51	0.35294	0.49019	172,01864	0.76370	0.25861	202,6237	198,9730	0.78116	0.15428	0.23521	0.23792	0.70947	64,56231	1,13656	0.41446
00002573_Q_500_palm_03	67	0.43867	0.17580	40	80	0.42500	0.36250	128,83008	0.78010	0.13989	158,5815	152,3574	0.75213	0.17158	0.39277	0.36161	0.62024	64,50494	1,13598	0.72320
00002573_Q_500_palm_04	58	0.42713	0.20052	74	127	0.51181	0.22834	138,90406	0.69712	0.23880	171,2604	164,2434	0.71148	0.18628	0.29442	0.32025	0.57310	57,31045	1,11872	0.71935
00002574_S_500_slap_06	86	0.44317	0.17627	30	115	0.32173	0.45217	135,76049	0.77714	0.20731	171,8981	163,8972	0.77558	0.21283	0.28113	0.31557	0.67914	101,87137	1,10695	0.63350
00002577_N_500_palm_09	40	0.48944	0.29130	54	109	0.19266	0.11009	120,84003	0.67922	0.24344	174,1594	165,8749	0.61506	0.21526	0.29786	0.36608	0.57458	49,41453	1,21319	1,18301
00002578_N_500_palm_04	62	0.42789	0.20143	45	97	0.67010	0.06185	171,82829	0.69210	0.23302	195,2009	188,7145	0.59501	0.14805	0.35336	0.30239	0.43305	42,87232	1,08956	0.65659
00002579_J_500_palm_05	34	0.46258	0.26197	114	154	0.46103	0.03896	141,96700	0.68997	0.22445	176,8952	171,7177	0.64837	0.17418	0.25750	0.22275	0.53288	51,15679	1,18042	0.88077
00002579_J_500_palm_08	89	0.45753	0.17757	29	64	0.62500	0.43750	160,28641	0.80818	0.14978	175,9763	173,4564	0.80667	0.14804	0.34351	0.32975	0.68948	84,80684	1,17648	0.53533
00002581_Q_500_palm_10	20	0.47133	0.29289	157	197	0.81218	0.00000	179,66015	0.57297	0.31766	205,3902	202,0176	0.58795	0.16727	0.36738	0.30155	0.43010	35,69879	1,16970	0.75640
00002581_S_500_slap_08	66	0.42456	0.19126	42	70	0.62857	0.07142	176,44046	0.66088	0.27127	193,8795	187,6766	0.69112	0.12001	0.39499	0.31428	0.53326	55,45952	1,14624	0.63709
00002584_J_500_palm_09	3	0.47823	0.31357	146	233	0.72103	0.00000	236,67901	0.52740	0.32138	239,6515	239,4468	0.46393	0.17866	0.36870	0.32745	0.32161	39,23643	1,33424	1,25081
00002586_S_500_slap_01	90	0.45363	0.14630	37	92	0.59782	0.19565	158,86986	0.79497	0.14274	182,5607	179,0008	0.63250	0.19130	0.33267	0.30805	0.46983	81,28152	1,13860	0.62004
00002588_J_500_palm_07	51	0.41865	0.21625	53	71	0.57746	0.08450	184,90571	0.69043	0.22987	204,9403	199,2879	0.65415	0.17980	0.21530	0.18674	0.50259	32,66848	0,96804	0,47638
00002591_S_500_slap_01	79	0.39019	0.11693	35	70	0.64285	0.78571	110,58360	0.82573	0.10415	148,2867	135,8804	0.86273	0.11029	0.40975	0.30470	0.72341	81,74625	1,15230	0,51247
00002593_Q_500_palm_05	31	0.38818	0.19085	110	155	0.69677	0.00000	195,61318	0.61553	0.25758	210,7440	209,4408	0.51555	0.12336	0.36297	0.29028	0.35487	28,03545	1,30114	1,08132
00002610_S_500_slap_06	94	0.42272	0.13842	33	150	0.55333	0.10666	172,13219	0.76066	0.20110	194,9162	188,4745	0.62293	0.22400	0.26563	0.29162	0.48116	89,97789	1,12261	0,60480
00002616_Q_500_palm_07	13	0.42679	0.25780	102	154	0.23376	0.00000	169,41641	0.64058	0.23765	193,2178	191,0845	0.49509	0.15269	0.26546	0.27219	0.34289	26,74578	1,24222	0,91013
00002624_Q_500_palm_07	75	0.42460	0.12855	35	63	0.58730	0.04761	188,28525	0.75310	0.20858	203,0196	201,4754	0.66121	0.14603	0.20599	0.21680	0.49525	47,04949	1,12214	0,43117
00002626_S_500_slap_04	65	0.46099	0.16248	21	39	0.43589	0.15384	174,36840	0.81275	0.13535	195,4373	193,2091	0.71856	0.15645	0.30865	0.28488	0.54761	55,85637	1,15270	0,77588
00002633_S_500_slap_09	51	0.43482	0.22173	56	110	0.50000	0.30909	131,30952	0.62501	0.32346	160,0805	152,3477	0.75974	0.14247	0.22561	0.22396	0.61354	62,58128	1,16198	0,71967
00002633_S_500_slap_10	44	0.40513	0.24755	94	135	0.51111	0.04444	206,83605	0.53372	0.31415	220,2791	218,5820	0.52103	0.21522	0.24816	0.22963	0.40602	28,42206	1,22276	1,22256

Table A.1 (continued)

Identifier	$Q_{NF1Q2\ v2.3.0}$	q_{FDA}^{μ}	σ_{FDA}	q_{FDA}^{σ}	q_{COM}^{σ}	q_{CNT}^{σ}	q_{MU}^{σ}	q_{OCL}^{σ}	q_{AREA}^{μ}	q_{LCL}^{μ}	q_{LCL}^{σ}	q_{MMB}	q_{MU}	q_{OCL}^{μ}	q_{OCL}^{σ}	q_{OFL}^{μ}	q_{OFL}^{σ}	q_{REL}^{σ}	q_{COH}^{σ}	q_{RVU}^{μ}	q_{RVU}^{σ}
00002636_S_500_slap_07	67	0,395 16	0,141 23	36	70	0,314 28	0,100 00	184,937 70	0,691 79	0,262 60	208,819 4	203,163 2	0,607 90	0,227 28	0,497 70	0,376 65	0,513 76	42,129 00	1,125 01	0,668 82	
00002636_S_500_slap_09	56	0,418 21	0,246 52	53	101	0,574 25	0,069 30	204,927 51	0,473 30	0,357 57	220,789 0	218,458 0	0,611 40	0,206 21	0,276 60	0,275 47	0,510 34	42,358 18	1,223 30	0,945 06	
00002640_Q_500_palm_04	9	0,460 36	0,298 68	108	164	0,121 95	0,000 00	142,969 40	0,507 57	0,305 71	186,548 6	180,449 3	0,413 07	0,170 59	0,422 34	0,255 26	0,295 81	18,340 54	1,312 27	0,960 76	
00002641_J_500_palm_04	47	0,480 96	0,206 25	70	98	0,367 34	0,112 24	137,501 12	0,747 25	0,198 05	167,677 5	163,093 5	0,668 55	0,193 47	0,383 50	0,312 17	0,543 90	45,143 93	1,112 00	0,563 03	
00002641_S_500_slap_03	71	0,436 24	0,174 90	43	57	0,578 94	0,526 31	148,084 20	0,720 50	0,272 23	175,318 2	170,668 2	0,854 34	0,089 58	0,311 57	0,274 25	0,748 80	54,662 49	1,113 71	0,722 77	
00002644_S_500_slap_01	80	0,438 84	0,173 34	28	74	0,364 86	0,662 16	101,390 70	0,823 68	0,131 02	145,951 8	136,926 2	0,846 92	0,124 56	0,300 48	0,300 57	0,726 12	121,262 94	1,161 63	0,534 15	
00002645_S_500_slap_01	96	0,440 92	0,142 95	38	88	0,659 09	0,215 90	164,439 03	0,789 40	0,155 27	187,067 0	183,370 6	0,665 81	0,201 00	0,366 92	0,325 31	0,512 82	74,873 01	1,037 30	0,482 47	
00002645_S_500_slap_10	25	0,460 83	0,253 19	67	78	0,192 30	0,054 10	162,512 41	0,635 81	0,273 09	193,386 0	188,109 0	0,573 17	0,191 27	0,360 73	0,244 14	0,447 52	27,299 06	1,317 04	1,240 72	
00002649_Q_500_palm_09	35	0,473 26	0,274 34	67	154	0,603 89	0,129 87	116,601 16	0,556 41	0,342 60	142,316 3	136,462 0	0,665 39	0,265 40	0,308 14	0,375 02	0,536 37	71,338 43	1,190 68	0,784 05	
00002650_S_500_slap_07	65	0,399 06	0,133 41	36	73	0,301 37	0,246 57	183,810 48	0,738 36	0,247 32	207,789 7	202,246 9	0,687 99	0,252 72	0,430 97	0,348 33	0,629 18	59,772 18	1,117 13	0,632 73	
00002650_S_500_slap_08	64	0,391 98	0,145 37	74	141	0,297 87	0,127 66	195,690 12	0,673 97	0,305 53	212,492 5	208,893 5	0,668 69	0,231 96	0,331 65	0,284 99	0,582 68	62,929 52	1,183 14	0,871 25	
blank_fd249	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
solid_0_500x500	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
solid_0_800x1000	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
solid_0_800x1001	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
solid_0_801x1000	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
solid_250_500x500	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
solid_251_500x500	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
solid_255_500x500	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA

Key

- NA value not computed (not applicable)
- $Q_{NF1Q2\ v2.3.0}$ unified quality score
- q_{FDA}^{μ} mean of local frequency domain analysis
- q_{FDA}^{σ} standard deviation of local frequency domain analysis
- q_{COM}^{σ} minutiae count in centre of mass
- q_{MMB} MMB
- q_{MU} MU
- q_{OCL}^{μ} mean of local orientation certainty level
- q_{OCL}^{σ} standard deviation of local orientation certainty level
- q_{OFL}^{μ} mean of local orientation flow

Table A.1 (continued)

Identifier	$Q_{NF1Q2\ v2.3.0}$	q_{FDA}^{μ}	σ_{FDA}	q_{MIN}^{COM}	q_{MIN}^{CNT}	q_{MIN}^{MU}	q_{MIN}^{OCL}	q_{AREA}^{μ}	q_{LCL}^{μ}	q_{LCL}^{σ}	q_{MMB}	q_{MU}	q_{OCL}^{μ}	q_{OCL}^{σ}	q_{OFL}^{μ}	q_{OFL}^{σ}	q_{COH}^{REL}	q_{COH}^{sum}	q_{RVU}^{μ}	q_{RVU}^{σ}	
q_{MIN}^{CNT}	minutiae count									q_{OFL}^{σ}											
q_{MIN}^{MU}	minutiae quality based in image mean									q_{COH}^{sum}											
q_{MIN}^{OCL}	minutiae quality based on orientation certainty level									q_{COH}^{REL}											
q_{AREA}^{μ}	region of interest image mean									q_{RVU}^{μ}											
q_{LCL}^{μ}	mean of local clarity									q_{RVU}^{σ}											
q_{LCL}^{σ}	standard deviation of local clarity																				

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Table A.2 — Target quality component values mapped 0-100 for conformance test set

Identifier	Q_{FDA}^H	Q_{FDA}^S	Q_{COM}^{MIN}	Q_{CNT}^{MIN}	Q_{MU}^{MIN}	Q_{OCL}^{MIN}	Q_{AREA}^H	Q_{LCL}^H	Q_{LCL}^S	Q_{MMB}	Q_{MU}	Q_{OCL}^H	Q_{OCL}^S	Q_{OFL}^H	Q_{OFL}^S	Q_{COH}^{REL}	Q_{COH}^{SUM}	Q_{RVU}^H	Q_{RVU}^S
00001052_plain_500_10	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA
00001057_plain_500_03	46	20	32	55	33	71	57	81	17	65	64	83	15	17	30	70	2	56	41
00001308_plain_500_08	48	20	16	44	43	59	64	81	21	71	71	78	15	11	22	69	2	57	33
00001428_plain_500_07	56	26	31	88	26	29	66	76	24	73	73	75	17	19	35	62	2	57	43
00001610_plain_500_10	64	33	100	100	42	3	63	62	28	71	70	60	22	15	24	47	1	65	52
00001655_plain_500_04	47	26	89	100	13	2	62	73	22	72	72	64	17	17	35	53	2	60	47
00001665_plain_500_08	59	34	76	100	32	0	64	64	27	72	70	55	17	18	35	38	1	65	55
00001723_plain_500_05	56	29	30	57	21	46	58	63	35	68	67	74	18	15	29	66	1	60	51
00001880_plain_500_10	68	34	100	100	7	2	50	55	33	65	63	59	20	23	29	44	1	75	71
00002302_N_500_palm_10	54	29	79	95	50	0	66	64	24	77	74	49	17	19	28	35	0	55	38
00002302_N_500_palm_10-core	38	19	42	42	43	0	59	65	20	65	62	42	17	19	3	31	0	60	55
00002304_R_500_slap_01	43	17	36	100	33	39	67	70	29	80	79	73	20	18	35	66	2	54	29
00002304_R_500_slap_08	43	22	61	100	45	3	68	70	21	78	77	56	20	19	48	44	0	57	52
00002310_R_500_slap_05	41	24	63	100	47	10	55	67	25	69	66	66	19	16	25	50	1	54	31
00002314_R_500_slap_01	45	17	31	65	54	6	72	75	22	81	80	62	15	20	38	49	2	57	35
00002315_R_500_slap_05	51	32	100	100	60	0	65	60	26	75	72	45	21	22	29	32	0	62	47
00002315_R_500_slap_06	43	19	53	100	44	15	61	76	17	72	69	72	16	19	36	58	3	55	30
00002317_R_500_slap_01	45	13	31	100	73	26	62	81	14	73	71	70	16	19	34	55	3	54	30
00002319_R_500_slap_04	42	16	57	83	58	2	75	55	37	80	80	60	14	26	44	42	1	53	30
00002319_R_500_slap_09	42	16	37	92	65	0	75	60	31	84	83	53	15	24	37	40	1	53	27
00002319_R_500_slap_10	46	18	25	44	73	2	72	75	21	82	81	63	14	8	17	50	1	55	24
00002320_R_500_slap_06	43	25	19	100	30	12	57	72	18	65	64	65	17	16	31	49	2	63	49
00002320_R_500_slap_08	46	29	100	100	37	0	62	61	26	67	67	55	18	20	36	40	1	69	65
00002321_R_500_slap_07	45	28	100	100	57	3	74	64	25	82	79	53	20	21	31	37	1	66	55

Table A.2 (continued)

Identifier	Q_{FDA}^{μ}	Q_{FDA}^{σ}	Q_{COM}^{μ}	Q_{COM}^{σ}	Q_{CNT}^{μ}	Q_{CNT}^{σ}	Q_{OCL}^{μ}	Q_{OCL}^{σ}	Q_{AREA}^{μ}	Q_{LCL}^{μ}	Q_{LCL}^{σ}	Q_{MMB}	Q_{MU}	Q_{OCL}^{μ}	Q_{OCL}^{σ}	Q_{OFL}^{μ}	Q_{OFL}^{σ}	Q_{COH}^{μ}	Q_{COH}^{σ}	Q_{COH}^{μ}	Q_{COH}^{σ}	Q_{RVU}^{μ}	Q_{RVU}^{σ}
00002325_R_500_slap_04	43	25	72	100	23	2	54	68	23	65	64	58	15	21	33	42	1	54	45				
00002325_R_500_slap_08	45	27	67	100	17	7	51	63	26	65	63	59	19	19	30	44	1	69	66				
00002326_R_500_slap_02	43	17	34	100	41	7	68	78	14	77	76	71	16	14	26	54	2	59	35				
00002326_R_500_slap_05	46	30	100	100	42	0	72	58	30	80	80	54	18	21	34	38	1	64	49				
00002330_R_500_slap_04	41	20	67	100	65	9	64	62	30	74	72	62	17	16	32	46	1	59	33				
00002334_R_500_slap_05	40	24	80	89	53	12	57	64	28	67	65	73	13	13	24	55	1	61	43				
00002334_R_500_slap_07	42	17	46	82	64	24	64	69	28	76	73	74	17	19	31	58	1	55	28				
00002340_R_500_slap_08	39	25	98	100	58	2	47	49	33	82	82	59	17	21	35	42	1	61	41				
00002342_R_500_slap_05	51	34	100	100	46	0	64	54	31	74	72	58	14	10	19	41	1	72	61				
00002342_R_500_slap_09	43	29	79	100	38	0	64	64	25	75	74	53	20	10	20	42	1	65	56				
00002349_N_500_palm_05	44	26	63	100	50	0	41	75	14	58	56	61	18	10	20	45	1	60	41				
00002349_R_500_slap_07	47	28	99	100	52	1	52	68	22	65	63	57	15	19	31	43	1	63	52				
00002352_R_500_slap_01	47	17	24	99	77	42	59	78	23	72	68	80	13	17	35	68	3	55	33				
00002352_R_500_slap_10	39	21	45	78	72	6	73	69	21	81	79	60	15	15	28	45	1	67	47				
00002353_N_500_palm_10	46	27	79	100	42	42	58	40	39	74	71	75	20	17	31	64	1	59	37				
00002353_R_500_slap_06	41	18	60	100	30	58	50	50	38	65	63	83	14	14	30	76	3	54	31				
00002353_R_500_slap_10	44	27	88	100	55	27	72	33	38	84	82	68	17	14	26	53	1	60	55				
00002356_N_500_palm_05	42	17	52	89	34	2	51	81	5	63	62	70	12	21	37	53	1	69	66				
00002357_R_500_slap_02	43	16	31	53	55	30	56	80	13	66	64	76	13	20	27	61	1	54	31				
00002357_R_500_slap_04	36	14	63	96	83	24	46	75	19	57	55	77	14	16	27	61	2	58	30				