T-Model Fingerprint Calculator v. 9.9 Legend

RIDGE FEATURE VALUES

Ridge Feature Shape	<u>Value</u>	Ridge Feature Position	<u>Value</u>
Continuous Ridge Unit (.45mm x .45mm)*	1.15	0-2 Intervening Ridges To Nearest Level II Neighbor	1
Pore*	5		
Ending Ridge Unit In Funnel	10	3 Intervening Ridges To Nearest Level II Neighbor	4
Ending Ridge Unit Not In Funnel	14.25		
Ending/Bifurcating Ridge Unit In Funnel	14.375	4 Intervening Ridges To Nearest Level II Neighbor	10
Bifurcating Ridge Unit In Funnel	18.75		
Ending/Bifurcating Ridge Unit Not In Funnel	20.5	5 Intervening Ridges To Nearest Level II Neighbor	62.5
Bifurcating Ridge Unit Not In Funnel	26.75		
Dot (Nearest Level2 Neighbor in Same Furrow>1mm)	40	6 Intervening Ridges To Nearest Level II Neighbor	976
2 Dots In Furrow <1mm apart. Value Per Dot:	10		
3 Dots In Furrow <1mm apart. Value Per Dot:	6	7 Intervening Ridges To Nearest Level II Neighbor	38,125
4 Dots In Furrow <1mm apart. Value Per Dot:	4.5		
5 Dots In Furrow <1mm apart. Value Per Dot:	4	8 Intervening Ridges To Nearest Level II Neighbor	3,723,144
6 Dots In Furrow <1mm apart. Value Per Dot:	3.75		
Core Area (1mm x 1mm)	209	9 Intervening Ridges To Nearest Level II Neighbor	908,970,832
Delta Area (Y Shape) (1mm x1mm)	190		
Delta Area (Non-Y Shape) (1mm x 1mm)	570	10 Intervening Ridges to Nearest Level II Neighbor**	554,791,767,578
* Values for CRU and Pore tentative pending close-mat	ch testing.	** Value omitted from calculator due to extreme rarity.	

RIDGE FEATURE REDUCTION FACTORS

Ridge Feature Clarity and Reliability (Analysis)						<u>ysis)</u>	Ridge Feature Quality of Agreement (Comparison)			
	DISTORTION LEVEL					Reduction		idge Type and Path Agrees	Spatial Relationship To Nearest Neighbor Agrees	Reduction Factor
<u>Grade</u>	<u>None</u>	<u>Low</u>	<u>Moderate</u>	<u>High</u>	Very High	Factor	Grade	Agrees	Nearest Neighbor Agrees	Pactor
A	Yes	No	No	No	No	1	Α	Yes	Yes	1
В	No	Yes	No	No	No	0.75	С	No	Yes	0.5
С	No	No	Yes	No	No	0.5	F	Yes/No	No	No Value
D	No	No	No	Yes	No		A - Excellent	_		
F	No	No	No	No	Yes		C - Satisfactory F - Unsatisfactory			

GUIDELINES

No Distortion Ridge feature appears visually clear and reliable. Ridge feature appears visually unclear or unreliable. Moderate Distortion Ridge feature appears visually unclear and unreliable.

High Distortion Ridge feature appears obstructed, however the orientation and relative position are reliably predictable.

Very High Distortion Ridge feature appears too distorted to analyze.

Note: 1/P denotes neutralization of ridge feature value, e.g. value equals 1.

GUIDELINES

Ridge Feature Type: Ending Ridge, Bifurcation, etc. Example #1: Ridge feature in latent print is an ending ridge and the ridge feature in the exemplar print is an ending ridge. The ridge feature types agree. Example #2: Ridge feature in the latent print is an ending ridge and the ridge feature in the exemplar print is a bifurcation. The ridge feature types do not agree.

Ridge Path: Ridge path, ie., an ending ridge unit slants to right, left, or not, or the ridge angle of separation, i.e., the angle of separation in a bifurcation, is large or small.

Spatial Relationship To Nearest Neighbor: Difference in distal relationship is less than 20% and difference in angle of rotation relationship is less than 10 degrees.

T-MODEL FORMULAE

T-Value (Total Discriminating Value)

T-Value = value 1 x value 2 x value 3 x value n...

where.

value 1 = value for ridge feature no. 1 (shape x position x clarity x agreement) value 2 = value for ridge feature no. 2 (shape x position x clarity x agreement) value 3 = value for ridge feature no. 3 (shape x position x clarity x agreement) value n = value for ridge feature no. n (shape x position x clarity x agreement)

FINGERPRINT PARTS

FINGERPRINT MATCH PROBABILITY

Fingerprint Match Probability (FMP) = 1/T-Value

If FMP < 1/Relevant Population (e.g., Number of People x 10 Fingers x Fingerprint Parts), then "Match"

Same as,

If T-Value > Relevant Population, then "Match"

(T) ^ (P) = 10 ^ 120 P = Fingerprint Parts

T = T-Value

Note: 10 ^ 120 = T-Value for Average Latent (Flat) Fingerprint

ESTIMATED NUMBER OF FINGERPRINT LOOK-ALIKES

L = RP / T

L = Estimated Number of Look-alikes (conservative, upper-bound number) RP = Relevant Population (e.g., Number of People x 10 Fingers x Fingerprint Parts) T = T-Value

T-Model Fingerprint Calculator v. 9.9