

## **Bibliography: DNA Mixture Interpretation**

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## **F. Computer Systems for DNA Mixture Interpretation**

International Society for Forensic Genetics, DNA interpretation software resources  
<http://www.isfg.org/Software>

National Institute of Standards and Technology, Information on DNA mixture interpretation  
<http://www.cstl.nist.gov/strbase/mixture.htm>

### *Quantitative data*

ArmedXpert (NicheVision Forensics)  
<http://www.armedxpert.com>

GeneMapper ID-X (Life Technologies/Applied Biosystems)  
<http://idx.appliedbiosystems.com>

GenoProof Mixture (Qualitytype)  
<http://www.qualitytype.de/en/qualitytype/genoproof-mixture>

FSS I-3<sup>®</sup> I-STReam (Promega Corporation)  
<http://www.promega.com/products/pm/genetic-identity/fssi3>

### *Allele dropout*

FST (Office of the Chief Medical Examiner, New York City)  
[http://www.nyc.gov/html/ocme/html/hss/hssservices\\_provided.shtml](http://www.nyc.gov/html/ocme/html/hss/hssservices_provided.shtml)

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*Genotype probability*

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MCMC (James Curran)

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MixSep (Torben Tvedebrink)

Tvedebrink T (2013), MixSep mixture separation R software (Aalborg University, Denmark)

<http://cran.r-project.org/web/packages/mixsep/>

STRmix (Environmental Science & Research, New Zealand)

Taylor D, Bright J-A, Buckleton, J (2013) STRmix fully continuous expert system

<http://strmix.com>

TrueAllele® Casework (Cybergenetics, Pittsburgh, PA)

<http://www.cybgen.com/systems/casework.shtml>

## G. TrueAllele Casework Publications

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