

### 20th International Forensic Science Managers Symposium

INTERPOL Headquarters, Lyon, France 10 November 2022



# Recent Advances in Forensic Biology and Forensic DNA Typing 2019-2022

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**Points of view are mine** and do not necessarily represent the official position or policies of the National Institute of Standards and Technology. Certain commercial entities are identified in order to specify experimental procedures as completely as possible. **In no case does such identification imply a recommendation or endorsement** by the National Institute of Standards and Technology, nor does it imply that any of the entities identified are necessarily the best available for the purpose.

## **Overview of Papers Reviewed**

**20<sup>th</sup> International Forensic Science Managers Symposium** 

Years Examined

2019, 2020, 2021, 2022 (through October 2022)

**Number of Papers** 

636 + 137 = 773

Number of Journals

**96** (FSI Genetics = 240)

**Topics Covered** 

15+

17<sup>th</sup> INTERPOL Review on DNA (2010-2013) examined **114 articles** (Jolicoeur 2013) 18<sup>th</sup> INTERPOL Review on DNA (2013-2016) examined **75 articles** (Laurent & Pene 2016) 19<sup>th</sup> INTERPOL Review on DNA (2016-2019) examined **235 articles** (Butler & Willis 2019)

The 2016-2019
INTERPOL DNA Review

Interpol review of forensic biology and forensic DNA typing 2016-2019

John M. Butler<sup>®</sup>, She<mark>i</mark>la Willis

National Institute of Standards and Technology, USA

(2020) 2: 352-367

## Discussed 235 references from 35 journals



19th INTERPOL International Forensic Science
Managers Symposium
Lyon, France

7-10 October 2019 Review Papers Review of forensic biology and DNA publications from 2016 to 2019

- Category selection and article selection:
  - 1. Core Loci Expansion
  - 2. Rapid Analysis of STR Markers
  - Investigative Genetic Genealogy
  - 4. Next-Generation Sequencing
  - 5. DNA Mixture Interpretation and Probabilistic Genotyping Software
  - 6. DNA Transfer and Activity Level Evaluations
  - 7. Forensic Biology and Body Fluid Identification
  - 8. DNA Phenotyping
  - 9. Privacy and Ethical Issues
  - 10. Guidance Documents (SWGDAM, OSAC, ASB, ENFSI, UK Regulator)
  - 11. Contamination Avoidance and DNA Success Rates
  - 12. Recent Special Issues and Review Articles of Note

https://www.interpol.int/content/download/14458/file/Interpol%20Review%20Papers%202019.pdf

## **Search Strategy**

Seeking to learn from Ménard, H., et al. (2021) Research trends in forensic science: a scientometric approach to analyze the content of the INTERPOL reviews. *WIREs Forensic Sci.* 2: e1147. doi: 10.1002/wfs2.1447

- Scopus and Web of Science searched "forensic DNA" from March 7, 2022, combined (4087)
- Removed duplicates and those not associated with forensic DNA based on title (1891)
  - Sorted by 26 MVP topics and added 6 other topic areas
  - Removed Chinese-only articles (e.g., Fa Yi Xue Za Zhi) (44)
  - Removed German-only articles (e.g., Rechtsmedizin) (9)
  - Resulted in **1779 articles** (68 pages of 8pt font single-spaced separated in 32 categories)
- Added FSI Reports, which were not in Scopus (40 from manual search of 256 total on 3/29/22)
- Added WIREs Forensic Science, which were not in Scopus (28 from manual search 3/31/22)
- Added other articles as identified in specific searches
- 3/31/2022: references from which to start writing the review (1860)
  - Continuing to add additional references as they become available and located with supplemental searches
- 6/13/2022: performed another Scopus search "forensic DNA" from 2019 to 2022 (3,059)
- 8/26/2022: performed another Scopus search "forensic DNA" from 2019 to 2022 (3,188)
- 11/1/2022: performed another Scopus search "forensic DNA" from 2019 to 2022 (3,361)

## **Topics Covered: Forensic Biology and DNA Typing**

| 4 |      | 4   |    |   |    | 4.5 |        |   |
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|   | <br> | H   | ı  | u | u  | GΠ  | U      | ı |

Hyperlinks to documents in tables and 50 footnotes to relevant websites

- 1. Books, Special Issues, and Review Articles of Note
- 2. Guidance Documents (SWGDAM, OSAC, ASB, NIFS, ENFSI, UK Forensic Science Regulator)
- 2. Advancements in Current Practices (Practitioner Focused)
  - 1. Rapid DNA Analysis
  - 2. Use of DNA Databases (Familial Searching, Investigative Genetic Genealogy, Privacy and Ethical Issues, SAKs)
  - 3. Forensic Biology and Body Fluid Identification
  - 4. DNA Collection and Extraction
  - 5. DNA Typing
  - 6. DNA Interpretation at the Source or Sub-Source Level (Probabilistic Genotyping Software)
  - 7. DNA Interpretation at the Activity Level (DNA Transfer)
- 3. Emerging Technologies, Research Studies, and Other Topics (Researcher Focused)
  - 1. Next-Generation Sequencing
  - 2. DNA Phenotyping (Ancestry, Appearance, Age Predictions) + Supplemental File (N=30+51+56)
  - 3. Lineage Markers (Y-chromosome, mtDNA, X-chromosome)
  - 4. New Markers and Approaches (Microhaplotypes, InDels, Proteomics, Human Microbiome)
  - 5. Kinship Analysis, Human Identification, and Disaster Victim Identification
  - 6. Non-Human DNA Testing and Wildlife Forensics

# articles

70

39

23

83

34

32

35

50

45

82

27 **+137** 

67

69

30

26

## Sources: Top 30 Countries (# documents published)

## 1. United States (717)

- 2. China (639)
- 3. India (256)
- 4. Australia (254)
- 5. United Kingdom (226)
- 6. Italy (185)
- 7. Germany (175)
- 8. Spain (136)
- 9. Netherlands (132)
- 10. Brazil (97)

- 11. Japan (94)
- 12. Switzerland (92)
- 13. Portugal (71)
- 14. South Korea (69)
- 15. Austria (66)
- 16. New Zealand (64)
- 17. Canada (60)
- 18. Poland (60)
- 19. France (57)
- 20. Malaysia (54)

- 21. Denmark (52)
- 22. Pakistan (49)
- 23. Russian Federation (47)
- 24. Sweden (47)
- 25. Saudi Arabia (47)
- 26. Thailand (44)
- 27. Norway (43)
- 28. South Africa (43)
- 29. Belgium (38)
- 30. Indonesia (38)

Based on Scopus search "forensic DNA" and "2019 to 2022" (1 Nov 2022; 3,361 document results)

## Sources: Top 30 Journals (# documents published)

| 1. | FSI | Genetics | (466) |  |
|----|-----|----------|-------|--|
|----|-----|----------|-------|--|

- 2. Int J Legal Med (288)
- 3. Forensic Sci Int (201)
- 4. FSI Genetics Sup (198)
- 5. J Forensic Sci (116)
- 6. Legal Med (88)
- 7. Sci Justice (69)
- 8. Genes (65)
- 9. Australian J Forensic Sci (64)
- 10. Scientific Reports (56)

- 11. Electrophoresis (55)
- 12. J Forensic Med (51)
- 13. Annals Human Biol (49)
- 14. Forensic Sci Tech (43)
- 15. Front Genet (38)
- 16. Indian J Forensic Med Tox (33)
- 17. Forensic Sci Res (32)
- 18. J Forensic Legal Med (29)
- 19. FSI Reports (25)
- 20. Forensic Sci Med Path (25)

- 21. PLoS One (25)
- 22. Egyptian J Forensic Sci (23)
- 23. FSI Synergy (22)
- 24. Mol Genet Gen Med (17)
- 25. Mol Biol Reports (17)
- ) 26. Rechtsmedizin (17)
  - 27. Forensic Genet Res Prog (15)
  - 28. Med Sci Law (15)
  - 29. Russian J Genetics (14)
  - 30. Gene (13)

Based on Scopus search "forensic DNA" and "2019 to 2022" (1 Nov 2022; 3,361 document results)

## in FSI Genetics only

## 1. Duncan Taylor (29) – Australia

- 2. Walther Parson (27) Austria
- 3. Jo-Anne Bright (21) New Zealand
- 4. Bruce Budowle (18) USA
- 5. Chris Phillips (18) Spain
- 6. Manfred Kayser (17) Netherlands
- 7. Yiping Hou (16) China (Chengdu)
- 8. Wojciech Branicki (12) Poland
- 8. Leonor Gusmão (12) Brazil
- 8. Adrian Linacre (12) Australia
- 8. Titia Sijen (12) Netherlands
- 8. Catarina Xavier (12) Austria

## In all indexed journals

## 1. Bruce Budowle (49) – USA

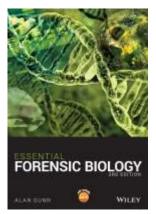
- 2. Walther Parson (47) Austria
- 3. Pankaj Shrivastava (43) India (Sagar)
- 4. Chengtao Li (39) China (Shanghai)
- 5. Duncan Taylor (38) Australia
- 6. Adrian Linacre (36) Australia
- 7. Jo-Anne Bright (35) New Zealand
- 8. Chao Liu (33) China (Guangzhou)
- 9. Bofeng Zhu (33) China (Guangzhou)
- 10. Guanglin He (32) China (Xiamen)

## **Topics Covered: Forensic Biology and DNA Typing**

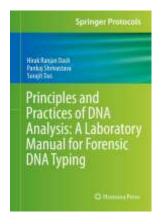
#### 1. Introduction

- 1. Books, Special Issues, and Review Articles of Note
- 2. Guidance Documents (SWGDAM, OSAC, ASB, NIFS, ENFSI, UK Forensic Science Regulator)
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  - 5. Kinship Analysis, Human Identification, and Disaster Victim Identification
  - 6. Non-Human DNA Testing and Wildlife Forensics

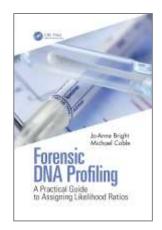
## Recent Books on Forensic Biology and Forensic DNA Typing (2019-2022)



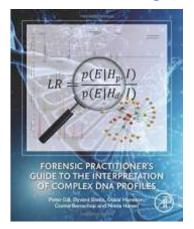
Essential Forensic Biology, Third Edition (2019, Wiley)



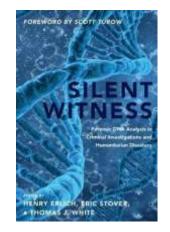
Principles and Practices of DNA Analysis: A Laboratory Manual for Forensic DNA Typing (2020, Humana Press)



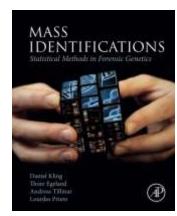
Forensic DNA Profiling: A Practical Guide to Assigning Likelihood Ratios (2020, CRC Press)



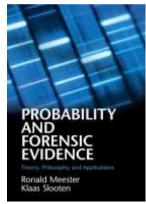
Forensic Practitioner's Guide to the Interpretation of Complex DNA Profiles (2020, Elsevier)



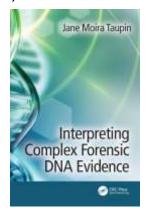
Silent Witness: Forensic DNA
Evidence in Criminal Investigations
and Humanitarian Disasters
(2020, Oxford University Press)



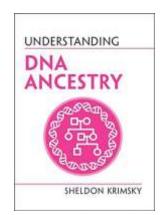
Mass Identifications: Statistical Methods in Forensic Genetics (2021, Elsevier)



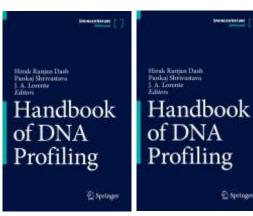
Probability and Forensic Evidence: Theory, Philosophy, and Applications (2021, Cambridge University Press)



Interpreting
Complex Forensic
DNA Evidence
(2021, CRC Press)

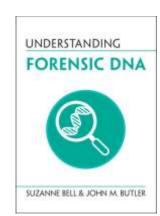


Understanding DNA Ancestry (2022, Cambridge University Press)



Handbook of DNA Profiling, 2 Volumes (2022, Springer)

1206 pages with 54 chapters from 115 contributors representing 17 countries



Understanding Forensic DNA (2022, Cambridge University Press)



## FSI Genetics Special Issues

(both guest edited by Manfred Kayser)

#### **Trends and Perspectives in Forensic Genetics (2018/2019)**

- 1. <u>Introduction to special issue</u>
- 2. DNA transfer in forensic science: A review
- 3. Evaluation of forensic genetics findings given activity level propositions: A review
- 4. Probabilistic genotyping software: An overview
- 5. Match probabilities for Y-chromosomal profiles: A paradigm shift
- 6. <u>Microhaplotypes in forensic genetics</u>
- 7. From next generation sequencing to now generation sequencing in forensics
- 8. Next generation database search algorithm for forensic mitogenome analyses
- 9. <u>Forensic human identification with targeted microbiome markers using nearest neighbor classification</u>
- Estimating the postmortem interval using microbes: Knowledge gaps and a path to technology adoption
- 11. Large scale DNA identification: The ICMP experience
- 12. Recent progress, methods and perspectives in forensic epigenetics

## Forensic Genetics: Unde venisti et quo vadis? (2021/2022)

- 1. Editorial/Prelude (written by ISFG Prize Winners)
- Massively parallel sequencing is unlocking the potential of environmental trace evidence
- The germlines of male monozygotic (MZ) twins: Very similar, but not identical
- Investigative genetic genealogy: Current methods, knowledge and practice
- Forensic transcriptome analysis using massively parallel sequencing
- 6. <u>Interpreting NUMTs in forensic genetics: Seeing the forest for the trees</u>
- 7. <u>Capture enrichment and massively parallel sequencing for human identification</u>
- 8. Forensic proteomics
- Progress in forensic bone DNA analysis: Lessons learned from ancient DNA
- 10. <u>Integrating the human microbiome in the forensic toolkit:</u> Current bottlenecks and future solutions
- 11. Forensic DNA Phenotyping (ancestry, appearance, age)
- 12. Microfluidic Technology in Forensic DNA Analysis

https://www.sciencedirect.com/journal/forensic-science-international-genetics/special-issue/10TSDS4360H

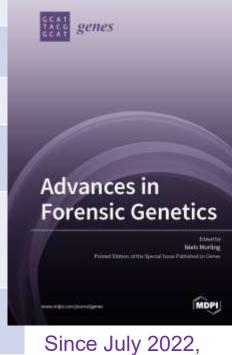
https://www.sciencedirect.com/journal/forensic-science-international-genetics/special-issue/10D6PT650B2

## Recent or Forthcoming Special Virtual Issues Related to Forensic Genetics from the Online Journal genes

As of 10/31/2022

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| Special Issue Title (publication dates)   | Editor(s)  | # Articles |
|---|--|------------|
| "Forensic Genetics and Genomics" (2020-2021)  | Emiliano Giardina & Michele Ragazzo                | 12         |
| "Forensic Mitochondrial Genomics" (2020-2021)   | Mitch Holland & Charla Marshall                    | 11         |
| "Advances in Forensic Genetics" (2021-2022)   | Niels Morling                                      | 25         |
| "State-of-the-Art in Forensic Genetics" (2022)  | Chiara Turchi                                      | 10         |
| "Trends in Population Genetics and Identification—<br>Impact on Anthropology" (2022)                                | Antonio Amorim, Veronica<br>Gomes, & Luisa Azevedo | 7          |
| "Identification of Human Remains for Forensic and Humanitarian Purposes: From Molecular to Physical Methods" (2023) | Elena Pilli & Cristina Cattaneo                    | 1          |
| "Improved Methods in Forensic and DNA Analysis" (2023)  | Marie Allen  | 3          |
| "Forensic DNA Mixture Interpretation and Probabilistic Genotyping" (2023)   | Michael Coble                                      |            |
| "Advances in Forensic Molecular Genetics" (2023)  | Erin Hanson & Claire Glynn                         |            |



available as a <u>518</u> page PDF file or a \$130 printed book



## **Special Issue** (Niels Morling, editor): Advances in Forensic Genetics

an Open Access Journal by MDPI

https://www.mdpi.com/journal/genes/special issues/Advances Forensic Genetics

## 25 articles published (2021-2022), topics covered include:

- PGS Review: EuroForMix, DNAStatistX, STRmix
- OpenArray for forensic phenotyping
- Skin pigmentation and genetic ancestry
- Eye color prediction
- Ancestry informative markers (VISAGE)
- Single cell analysis for forensic phenotyping
- Animal forensic genetics
- Predicting visible traits in dogs (CaDNAP)
- Single cell analysis for mixture interpretation
- New STR panel for cross-species bird DNA
- Ancient DNA methods improve Korean/WW2 IDs

- DNA transfer review and recent progress
- Bayesian Networks for DNA transfer questions
- SNP markers for investigative genetic genealogy (FORCE panel)
- DNA sampling in burglary investigations
- Body fluid ID and tissues
- Microbiome analysis
- Software options for forensic sequencing
- ChrY and mtDNA statistics/assessment
- Ethical decision-making as lived practice
- Aged rootless hair shafts in Romanov relics

### **Guidance Documents – United States**

With hyperlinks to each document

### **SWGDAM** (see <a href="https://www.swgdam.org/publications">https://www.swgdam.org/publications</a>)

- 1. SWGDAM (Apr 2019) Mitochondrial DNA Analysis Revisions Related to NGS
- 2. SWGDAM (Apr 2019) Addendum to Interpretation Guidelines to Address NGS
- 3. SWGDAM (Feb 2020) Overview of Investigative Genetic Genealogy
- 4. SWGDAM (July 2020) Report on Y-Screening of Sexual Assault Evidence Kits (SAEKs)
- 5. SWGDAM (July 2020) <u>Training Guidelines</u>
- 6. SWGDAM (Jan 2022) YHRD Updates for U.S. Laboratories
- 7. SWGDAM (March 2022) <u>Interpretation Guidelines for Y-Chromosome STR Typing by Forensic DNA Laboratories</u>
- 8. SWGDAM (March 2022) <u>Supplemental Information for the SWGDAM Interpretation Guidelines for Y-Chromosome STR Typing by Forensic DNA Laboratories</u>

## FBI (drafted by SWGDAM)

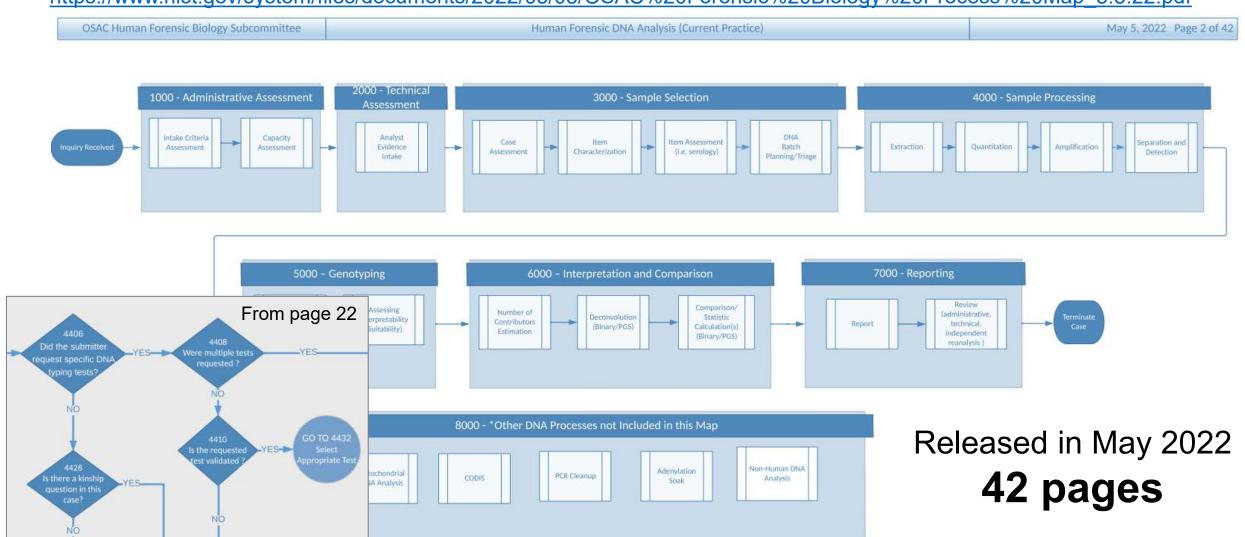
- 1. FBI Quality Assurance Standards for Forensic DNA Testing Laboratories (effective July 1, 2020)
- 2. FBI Quality Assurance Standards for DNA
  Databasing Laboratories (effective July 1, 2020)
- 3. FBI Quality Assurance Standards Audit for Forensic DNA Testing Laboratories
- 4. FBI Quality Assurance Standards Audit for DNA Databasing Laboratories
- 5. <u>Guidance Document for the FBI Quality Assurance</u>
  Standards for Forensic DNA Testing and DNA
  Databasing

FBI – A Guide to All Things Rapid DNA (13 pages) <a href="https://www.fbi.gov/file-repository/rapid-dna-guide-january-2022.pdf/view">https://www.fbi.gov/file-repository/rapid-dna-guide-january-2022.pdf/view</a>

### **Guidance Documents – United States**

DNA Process Map produced by OSAC Human Forensic Biology Subcommittee (with SWGDAM input)

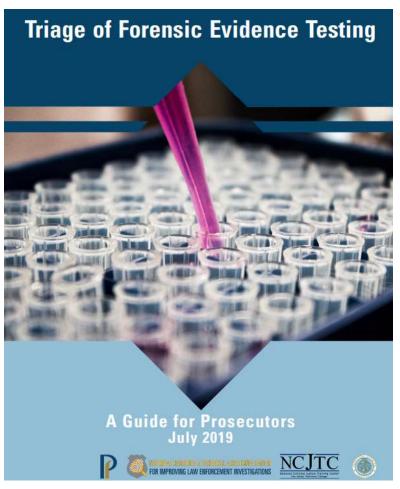
https://www.nist.gov/system/files/documents/2022/05/05/OSAC%20Forensic%20Biology%20Process%20Map 5.5.22.pdf



### **Guidance Documents – United States**

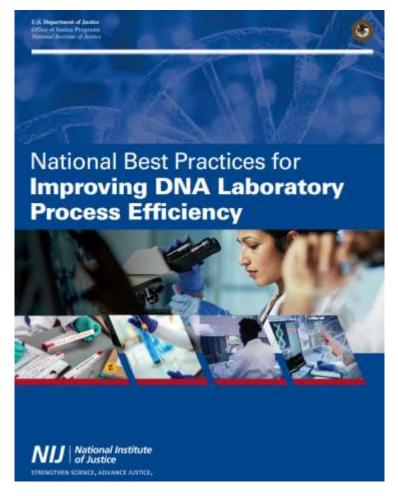
how best to maximize the resources of a public forensic laboratory, with a focus on the prosecutor's perspective

- Bureau of Justice Assistance (BJA) Triage of Forensic Evidence Testing: A Guide for Prosecutors (July 2019)
- National Institute of Justice (NIJ) National Best Practices for Improving DNA **Laboratory Process** Efficiency (May 2022)



https://bja.ojp.gov/library/publications/triage-forensicevidence-testing-guide-prosecutors (49 pages)

provides forensic DNA laboratories with a roadmap for managing expected increases in case submissions due to stakeholder demand



https://www.ojp.gov/pdffiles1/nij/304051.pdf (106 pages)

With 67 recommendations

## **Guidance Documents – UK**

### **Forensic Science Regulator**

- 1. FSR-C-100, Issue 7 Codes of Practice and Conduct (2021)
- FSR-C-108, Issue 2 DNA Analysis: Codes of Practice and Conduct (2020)
- 3. FSR-C-116, Issue 1 Sexual Assault Examination: Requirements for the Assessment, Collection and Recording of Forensic Science Related Evidence (2020)
- 4. FSR-C-118, Issue 1 Development of Evaluative Opinions (2021)
- 5. <u>FSR-G-201</u>, <u>Issue 2 Validation</u> (2020)
- 6. <u>FSR-G-202</u>, <u>Issue 2</u>, <u>The Interpretation of DNA Evidence</u> (<u>Including Low-Template DNA</u>) (2020)
- FSR-P-300, Issue 2 Validation Use of Casework Material (2020)
- 8. FSR-P-302, Issue 2 DNA Contamination Detection: The Management and Use of Staff Elimination DNA Databases (2020)
- FSR-G-206, Issue 2 The Control and Avoidance of Contamination in Scene Examination involving DNA Evidence Recovery (2020)
- FSR-G-207, Issue 2 The Control and Avoidance of Contamination in Forensic Medical Examinations (2020)

### **Forensic Science Regulator**

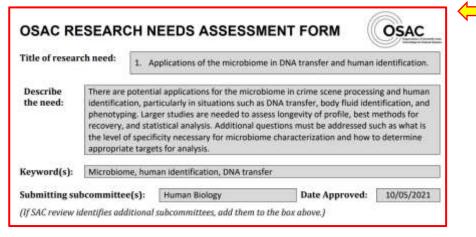
- FSR-G-208, Issue 2 The Control and Avoidance of Contamination in Laboratory Activities involving DNA Evidence Recovery Analysis (2020)
- 12. FSR-G-212, Issue 1 Guidance for the Assessment, Collection and Recording of Forensic Science Related Evidence in Sexual Assault Examinations (2020)
- 13. FSR-G-213, Issue 2 Allele Frequency Databases and Reporting Guidance for the DNA (Short Tandem Repeat) Profiling (2020)
- 14. <u>FSR-G-217</u>, <u>Issue 2 Cognitive Bias Effects Relevant to Forensic Science Examinations</u> (2020)
- 15. FSR-G-222, Issue 3 DNA Mixture Interpretation (2020)
- 16. <u>FSR-G-223</u>, <u>Issue 2 Software Validation for DNA Mixture</u> Interpretation (2020)
- 17. FSR-G-224, Issue 1 Proficiency Testing Guidance for DNA Mixture Analysis and Interpretation (2020)
- 18. <u>FSR-G-227</u>, <u>Issue 1 Y-STR Profiling</u> (2021)
- 19. FSR-G-228, Issue 1 -- DNA Relationship Testing using Autosomal Short Tandem Repeats (2021)
- 20. FSR-G-229, Issue 1 Methods Employing Rapid DNA Devices (2021)

## Information Tables in This INTERPOL DNA Review

- **Table 1**: Top ten journals with forensic DNA articles published from 2019 to 2022 based on a Scopus.com search on June 13, 2022
- **Table 2**: Guidance documents (70 total) related to forensic DNA published from 2019 to 2022 (with hyperlinks to each document)
- **Table 3**: Research and development needs in forensic biology as identified by the OSAC Human Forensic Biology Subcommittee (with hyperlinks to each document)
- **Table 4**: Summary of 20 rapid DNA instrument validation and evaluation studies published from 2019 to 2022
- **Table 5**: STR kits assessed with 24 published validation studies from 2019 to 2022

## **OSAC Research & Development Needs**

#### Human Forensic Biology





- 1. Applications of the Microbiome in DNA Transfer and Human Identification
- 2. Assessing DNA Background and Transfer Scenarios in Forensic Casework
- 3. <u>Best Practices to Minimize Potential Biases in the Generation and Interpretation of</u> DNA Profiles
- 4. Best Practices for Reporting Likelihood Ratios or Other Probabilistic Results in Court
- 5. <u>Characterization, Development and Validation of Methods in Single Cell Isolation and</u> Analysis
- 6. Characterization, Optimization and Comparison of DNA Sequencing Methods
- 7. Characterizing the Presence and Prevalence of Cell-Free DNA
- Development of Infrastructure to Compile and Share Raw Electronic Data for Training and Tool Development
- 9. <u>Efficiency, Throughput and Speed Improvements in Rapid DNA Instrumentation Through the Development of Direct PCR Methods</u>
- 10. Efficient Collection of DNA at the Scene and from Evidence Items
- 11. <u>Establishing the Value and Designing a Process for Including Flanking Region SNPs in Massive Parallel Sequencing Based on STRP Casework</u>
- 12. Improving the Recovery of Male DNA from Sexual Assault Kits
- 13. Methods in Forensic Genealogy
- 14. Non-PCR Based Methods for DNA Amplification and/or Detection
- 15. Optimization of DNA Extraction for Low Level Samples
- 16. <u>Software Solutions for Low Template and High Order DNA Mixture Interpretation in Sequence and Fragment-Based Methods</u>
- 17. Software Solutions for Y-STR Mixture Deconvolution
- 18. Solutions in Phenotyping and Ancestry Analyses

https://www.nist.gov/organization-scientific-area-committees-forensic-science/osac-research-and-development-needs

## Rapid DNA Studies (N=20) Published from 2019 to 2022

| Publication             | Instrument                               | Cartridge/Kit            | Test Performed and Success Rates Reported  |
|-------------------------|--|--------------------------|--|
| Amick &<br>Swiger 2019  | RapidHIT ID                              | ACE and EXT              | Performed SWGDAM internal validation studies including known and database-type samples, reproducibility, precision, sensitivity, stochastic effects, mixtures, contamination assessment, and concordance studies   |
| Carney et al.<br>2019   | ANDE 6C                                  | A-Chip                   | Conducted SWGDAM developmental validation (across 6 labs, 2045 swabs, 13 instruments): species specificity, limit of detection, stability, inhibitors, reproducibility, reference material, mixtures, precision, concordance, signal strength, peak height ratio, stutter, non-template addition, resolution, and contamination assessment; <b>first-pass success rate (1338 samples with 20 CODIS core loci) = 92%</b> ; successfully interpreted >2000 samples with over 99.99% concordant alleles; data package led to receiving NDIS approval in June 2018 |
| Shackleton et al. 2019a | RapidHIT ID                              | NGM SElect<br>Express    | Described development studies that included process optimization, sensitivity, repeatability, contamination checks, inhibition, swab age, concordance, and overall performance; success rate (124 samples) = 84.5% gave a full profile   |
| Shackleton et al. 2019b | RapidHIT 200                             | NGM SElect<br>Express    | Performed some protocol adjustments that enhanced slightly the sensitivity with mock crime scene samples (dilutions of blood and cell line DNA)  |
| Yang et al.<br>2019     | MiDAS                                    | PowerPlex ESI<br>16 Plus | Described protocols for analysis of reference samples with a fully automated integrated microfluidic system (MiDAS), which is not commercially available   |
| Romsos et al.<br>2020   | ANDE 6C,<br>RapidHIT ID,<br>RapidHIT 200 | A-Chip, ACE              | Reported results from the July 2018 rapid DNA maturity assessment with multiple instruments organized by NIST; the average success rate for obtaining the 20 CODIS core loci was 85% (n=240)   |

## STR Kits with Validation Studies (N=24) Published from 2019 to 2022

| Publication            | STR Kit/Primer Set                                    | Comments   |
|------------------------|---|--|
| Al Janaahi et al. 2019 | VeriFiler Plus  | Validation studies (sensitivity, peak height ratio, precision, reproducibility, thresholds, mixtures, concordance) |
| Alsafiah et al. 2019   | SureID 23comp Human Identification                    | Validation studies (following ENFSI and SWGDAM guidelines); has 17 non-CODIS STRs                                  |
| Bai et al. 2019        | DNATyper25  | Validation studies (following SWGDAM and China National Standard); has 20 non-CODIS STRs                           |
| Cho et al. 2021        | Investigator 24plex QS, PowerPlex Fusion, GlobalFiler | Examined 189 casework samples and compared performance across the three kits                                       |
| Fan et al. 2021        | STRtyper-32G  | Developmental validation studies (SWGDAM); has 10 non-CODIS STRs   |
| Green et al. 2021      | VeriFiler Plus  | Developmental validation studies (SWGDAM); concordance checked with Huaxia Platinum kits                           |
| Hakim et al. 2020      | Investigator 24plex GO!                               | Validation studies; concordance with GlobalFiler   |
| Harrel et al. 2021     | Investigator 24plex QS and GO!                        | Assessment of sample quality metrics in both kits  |
| Jiang et al. 2021a     | STRscan-17LC kit                                      | Validation studies (SWGDAM)  |
| Jiang et al. 2021b     | Novel 8-dye STR multiplex                             | Validation studies (SWGDAM); 18 STRs plus AMEL; detection with GA118-24B Genetic Analyzer                          |
| Lenz et al. 2020       | VersaPlex 27PY system                                 | Developmental validation studies (SWGDAM); includes D6S1043  |
| Li et al. 2021         | SureID S6 system                                      | Validation studies (SWGDAM); concordance with Huaxia Platinum kit; uses lyophilized reagents                       |
| Liu et al. 2019        | 19 autosomal and 27 Y-STRs                            | Validation studies (Chinese National Standard); 47 loci (Fusion 6C, GlobalFiler, Yfiler Plus) with 6-dyes          |
| Qu et al. 2019         | Microreader 20A ID system                             | Developmental validation studies (SWGDAM)  |
| Qu et al. 2021         | Novel 6-dye, 31-plex                                  | Developmental validation studies (SWGDAM and Chinese National Standard); 29 STRs, AMEL, Y-InDel                    |
| Wang et al. 2020a      | 21plex with DYS391 and ABO                            | Describes a 21plex with 18 autosomal STRs, ABO blood group locus, DYS391, and AMEL                                 |
| Wang et al. 2020b      | Investigator 26plex QS kit                            | Validation studies (SWGDAM); concordance with AGCU Expressmarker 22 kit  |
| Xie et al. 2020        | AGCU Expressmarker 16+22Y                             | Developmental validation studies (SWGDAM)  |
| Xie et al. 2022        | Novel 26plex  | Validation studies (SWGDAM); multiple STRs on chromosomes 13, 18, 21, and X for prenatal diagnosis                 |
| Yin et al. 2021        | Microreader 28A ID System                             | Developmental validation (SWGDAM); concordance with AGCU Expressmarker 22 kit                                      |
| Zhang et al. 2020      | SiFaSTR 21plex_NCII                                   | Developmental validation (SWGDAM); describes 18 new non-CODIS STR loci   |
| Zhang et al. 2021      | AGCU Expressmarker 30 Kit                             | Developmental validation (SWGDAM); includes 6 non-CODIS STR loci; concordance with AGCU Expressmarker 22 kit       |
| Zheng et al. 2019      | SiFaSTR 23-plex panel                                 | Developmental validation (SWGDAM and Chinese National Standard)  |
| Zhong et al. 2019      | Huaxia Platinum PCR kit                               | Developmental validation (SWGDAM and Chinese National Standards)   |

## **Recent Major Conferences** on Forensic DNA (2019-2022)



American Academy of Forensic Sciences (AAFS)

2020

(Anaheim, CA)

2019

(Baltimore, MD)

**2021 AAFS** 

MEETING

2021

(virtual)

European Academy of Forensic Science (EAFS)

2022

(Seattle, WA)

International Symposium on Human Identification





347 extended abstracts published

2021

2020

O Promega | Powered for 32 years TERNATIONAL SYMPOSIUM

ON HUMAN IDENTIFICATION ORLANDO, FL | SEPT. 13-17, 2021



Washington, D.C., USA Aug/Sept 2022

International Society

for Forensic Genetics

Prague, Czech Republic

Sept **2019** 

WASHINGTON, DC INTERNATIONAL SOCIETY FOR FORENSIC GENETICS

extended abstracts published

~126



ISFG Proceedings published in Forensic Science International: Genetics Supplement Series







## MVPs (Most Valuable Publications) of Forensic DNA AAFS 2021 MVPs Workshop



## MVPs of Forensic DNA:

Examining the Most Valuable Publications in the Field

John M. Butler, PhD

Co-Chair

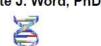
Robin W. Cotton, PhD

BOSTON
UNIVERSITY

Mechthild K. Prinz, PhD



Charlotte J. Word, PhD



- 4-hour virtual workshop with pre-recorded videos from each presenter
- 116-page handout (<u>available on STRBase</u>)
  - 240 slides and a 33-page reference list
- 480 references in 26 categories + 17 books
  - Built on lists from SWGDAM July 2020 <u>Training Guidelines</u> and OSAC October 2020 <u>Informative Literature for Forensic Biology and DNA</u>
  - Valuable input from co-presenters is gratefully acknowledged: Robin Cotton, Mecki Prinz, and Charlotte Word
- Discussed principles, MVP trends (article citations, journals used), and value of the #1 article in each category
  - Covered 17 of 26 categories in 10 modules
  - Also discussed training standards and value of a knowledge base for forensic DNA analysts

### 2022 Revised (and Reduced) MVPs = 85 references

https://strbase.nist.gov/pub\_pres/AAFS2022-W2-NIST-Forensic-DNA-Activities-FINAL.pdf (pp. 77-84)

## FSI Genetics Publications between ISFG 2019 and ISFG 2022



18 volumes (18 issues)

439 articles





**2019** (v38) (v39) (v40) (v41) (v42) (v43)

**2020** (v44) (v45) (v46) (v47) (v48) (v49)

**2021** (v50) (v51) (v52) (v53) (v54) (v55)

**2022** (v56) (v57) (v58) (v59) (v60) (v61)

439 Articles Published in FSI Genetics between ISFG 2019 and ISFG 2022 (vol. 43 to vol. 60)

- Audree, S. E., Hipp, M. J., Kennerly, S. R., & Weir, B. S. (2020). Analyzing population structure for forensis STR markers in next generation sequencing data. Forensis: Science International-Greenics, 49, 102364. doi:10.1016/j.fnigen.2020.102364
- Achesh, N., van Weert, A., Birkl, M., van Leeuwen, T. G., Anders, M. C. G., & van Durn, A. (2021). The compatibility of immunolabellary with STR profiling. Foreists: Science International-Genetics, 52, 102485. doi: 10.1016/j.itigen.2021.102485
- Adolfason, E., Qvick, A., Green, H., Kling, D., Gumarsson, C., Jonesson, J., & Green, A. (2021). Technical in-depth compension of two massive parallel DNA-segrencing methods for formalin-fixed paraffin-embedded ussus from victims of radden earline, death. Forensic Science International-Genetics, 33, 102522. doi:10.1016/j.fnjgen.2021.102522
- Agostini, V., Baito, P., Chiri, E., Liauerllo, P., Gratile, G., Principanai, P., ... Piccianii, A. (2020). Ocular swabs on exhanned bodies: An adremative to the collection of "cleasion" stone samples in foreasic genetics. Foreasic Science International-Genetics. 44, 102206. doi:10.1016/j.fnigen.2019.102206
- Agudo, M. M., Annes, H., Roseth, A., Albert, M., Gill, P., & Bloka, O. (2022). A comprehensive characterization of MPS-STR statter artefacts. Forentic Science International-Genetics, 69, 102728. doi:10.1016/j.fsigen.2022.102728

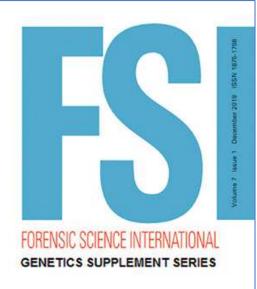
https://strbase.nist.gov/pub\_pres/ FSI-Genetics-439articles-2019to2022-alphabetical-byauthor.pdf



#### Supplement

The 28th Congress of the International Society for Forensic Genetics Prague

Guest Editors: Mechthild Prinz, John M. Butler and Jiri Drabek





## **ISFG 2019 Proceedings**

- Published in December 2019
- FSI Genetics Supplement Series, Volume 7
- 914 pages freely available online
- https://www.fsigeneticssup.com/current
- 347 articles + 1 editorial + 1 corrigendum

The ISFG 2022 Proceedings (volume 8 of FSI Genetics Supplement Series) should be published in December 2022 (currently 126 articles will be included)

## **Topics Covered: Forensic Biology and DNA Typing**

- 1. Introduction
  - 1. Books, Special Issues, and Review Articles of Note
  - 2. Guidance Documents (SWGDAM, OSAC, ASB, NIFS, ENFSI, UK Forensic Science Regulator)
- 2. Advancements in Current Practices (Practitioner Focused)
  - 1. Rapid DNA Analysis
  - 2. Use of DNA Databases (Investigative Genetic Genealogy, Privacy and Ethical Issues, SAKs)
  - 3. Forensic Biology and Body Fluid Identification
  - 4. DNA Collection and Extraction
  - 5. DNA Typing
  - 6. DNA Interpretation at the Source or Sub-Source Level (Probabilistic Genotyping Software)
  - 7. DNA Interpretation at the Activity Level (DNA Transfer)
- 3. Emerging Technologies, Research Studies, and Other Topics (Researcher Focused)
  - 1. Next-Generation Sequencing
  - 2. DNA Phenotyping (Ancestry, Appearance, Age Predictions)
  - 3. Lineage Markers (Y-chromosome, mtDNA, X-chromosome)
  - 4. New Markers and Approaches (Microhaplotypes, InDels, Proteomics, Human Microbiome)
  - 5. Kinship Analysis, Human Identification, and Disaster Victim Identification
  - 6. Non-Human DNA Testing and Wildlife Forensics

## **Investigative Genetic Genealogy Review Article**

Forensic Science International: Genetics 52 (2021) 102474

Contents lists available at ScienceDirect

#### Forensic Science International: Genetics

journal homepage: www.elsevier.com/locate/fsigen



#### Review article

Investigative genetic genealogy: Current methods, knowledge and practice



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- Department of Forensic Sciences, Oslo University Hospital, Oslo, Norway
- <sup>6</sup> Forensic Genetics Unit, Institute of Forensic Sciences, University of Santiago de Compostela, Santiago de Compostela, Spain
- d Research Department of Genetics, Evolution and Environment, University College London, Gower Street, London WC1E 6BT, United Kingdom
- <sup>e</sup> Department of Biomedical and Clinical Sciences, Faculty of Medicine and Health Sciences, Linköping University, Linköping, Sweden

#### ARTICLE INFO

Keywords:
Genetic genealogy
SNP microarrays
Whole-genome-sequencing
Familial searching
Identity by descent
Forensic DNA analysis
Crime investigation

#### ABSTRACT

Investigative genetic genealogy (IGG) has emerged as a new, rapidly growing field of forensic science. We describe the process whereby dense SNP data, commonly comprising more than half a million markers, are employed to infer distant relationships. By distant we refer to degrees of relatedness exceeding that of first cousins. We review how methods of relationship matching and SNP analysis on an enlarged scale are used in a forensic setting to identify a suspect in a criminal investigation or a missing person. There is currently a strong need in forensic genetics not only to understand the underlying models to infer relatedness but also to fully explore the DNA technologies and data used in IGG. This review brings together many of the topics and examines their effectiveness and operational limits, while suggesting future directions for their forensic validation. We further investigated the methods used by the major direct-to-consumer (DTC) genetic ancestry testing companies as well as submitting a questionnaire where providers of forensic genealogy summarized their operation/services. Although most of the DTC market, and genetic genealogy in general, has undisclosed, proprietary algorithms we review the current knowledge where information has been discussed and published more openly.

### Highlights

- Comprehensive review of investigative genetic genealogy from a forensic perspective
- Background outlined for the DNA methodology and long-range familial searching process
- Survey of current direct-to-consumer testing companies connected to investigative genetic genealogy
- Overview of DNA technologies focusing on high-density SNP genotyping

147 references cited with 7 supplemental files

## **Summary of Recent Advances**

### Aiding Investigations

- Phenotyping and Ancestry Testing (VISAGE and beyond)
- Investigative Genetic Genealogy (GEDmatch and growing commercial support)

### Improving Methods

- DNA recovery, extraction, quantitation, amplification chemistry, new kits
- Process mapping, standards and guidance documents

## Speeding and Strengthening Analysis

- Rapid DNA
- Massively Parallel Sequencing

## Innovating Interpretation

- Probabilistic genotyping software for DNA mixtures
- Activity level evaluations using DNA transfer studies

These advances are reported in the scientific literature and summarized in this INTERPOL review so that we can, as Isaac Newton famously stated, "stand on the shoulders of giants" to see further



## Forensics@NIST 2022 (happening this week)

https://www.nist.gov/agenda/forensicsnist-2022

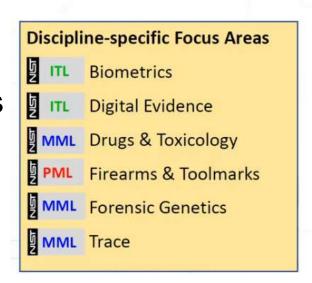
(Free) Virtual Meeting>1625 registrants from 70 countries55 presentations + 5 workshopsRecordings to be available in two weeks

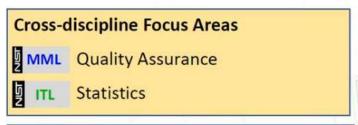
## **NIST Forensic Science Efforts**

- 1. Conduct impactful research
- 2. Facilitate **standards** development
- 3. Assess foundational knowledge

DNA Mixture Interpretation
Digital Evidence
Bitemark Analysis
Firearm Examination

### Research Focus Areas





#### **Future Planning Priorities**

- Computational Forensic Science
- Forensic Science Data
- Training Officers of the Court



#### NIST Center of Excellence











## Thank you for your attention!

John Butler john.butler@nist.gov



https://www.nist.gov/topics/forensic-science

#### **MVPs of Forensic DNA**

2021 (480): <a href="https://strbase.nist.gov/pub\_pres/AAFS2021-W19-Handouts.pdf">https://strbase.nist.gov/pub\_pres/AAFS2021-W19-Handouts.pdf</a> (pp. 3-35)

2022 ( 85): https://strbase.nist.gov/pub\_pres/AAFS2022-W2-NIST-Forensic-DNA-Activities-FINAL.pdf (pp. 77-84)



**Points of view are mine** and do not necessarily represent the official position or policies of the National Institute of Standards and Technology. Certain commercial entities are identified in order to specify experimental procedures as completely as possible. **In no case does such identification imply a recommendation or endorsement** by the National Institute of Standards and Technology, nor does it imply that any of the entities identified are necessarily the best available for the purpose.

| Informative Forensic DNA Reviews and Research Studies (A-to-Z) |  |    | # Articles   |  |
|--|--|----|--------------|--|
| Category<br>Group  | Topic(s) Covered   |    | 85<br>(2022) |  |
| Α  | Plain Language Guides to Forensic DNA Analysis               | 4  | 2            |  |
| В  | Serology and Body Fluid Identification                       | 24 | 3            |  |
| C  | Collection and Storage of Biological Material                | 25 | 2            |  |
| D  | DNA Extraction/Purification, Differential Extraction         | 18 | 2            |  |
| E  | DNA Quantitation, Degraded DNA                               | 10 | 2            |  |
| F  | PCR Amplification, Inhibition, and Artifacts                 | 13 | 3            |  |
| G  | Capillary Electrophoresis Separation and Detection           | 12 | 2            |  |
| Н  | Assessing Sample Suitability & Complexity, Low-Template      | 7  | 2            |  |
| I  | Estimating the Number of Contributors                        | 12 | 4            |  |
| J  | Data Interpretation, Mixture Deconvolution, Interlab Studies | 12 | 4            |  |
| K  | Interpretation: Binary Approaches (CPI, RMP, LR)             | 11 | 5            |  |
| L  | Interpretation: Probabilistic Genotyping Software            | 44 | 4            |  |
| M  | Report Writing and Technical Review                          | 8  | 4            |  |

| Informative Forensic DNA Reviews and Research Studies (A-to-Z) |   |    | # Articles       |  |
|--|---|----|------------------|--|
| <b>Category Group</b>  | Topic(s) Covered  |    | <b>85</b> (2022) |  |
| N  | Court Testimony, Communication, Juror Comprehension         | 22 | 5                |  |
| 0  | Autosomal STR Markers and Kits                              | 29 | 2                |  |
| Р  | Mitochondrial DNA Testing                                   | 11 | 3                |  |
| Q  | Y-Chromosome and X-Chromosome Testing                       | 17 | 4                |  |
| R  | DNA Databases and Investigative Genetic Genealogy           | 14 | 3                |  |
| S  | Statistical Analysis  | 11 | 2                |  |
| Т  | Population Genetics   | 11 | 2                |  |
| U  | DNA Phenotyping (Ancestry, Appearance, Age)                 | 24 | 2                |  |
| V  | New Technologies (Rapid DNA, Massively Parallel Sequencing) | 35 | 5                |  |
| W  | DNA Transfer and Activity Level Reporting                   | 57 | 8                |  |
| X  | Non-Human DNA Testing                                       | 15 | 2                |  |
| Y  | Method Validation, Quality Control, and Human Factors       | 23 | 5                |  |
| Z  | General Forensic Science Topics                             | 11 | 3                |  |

## Testing the Current 26 MVP "A-to-Z" Categories with the 2019-2022 INTERPOL Review

## Starting with 4,087 articles

Scopus & Web of Science "forensic DNA" searches January 2019 to March 2022 (with some additions)

> Removed duplicates and sorted

into 26 categories

MVP "A to Z"

+ 6 additional ones

1,884 articles May 16, 2022

version

| MVP | # articles |
|-----|------------|
| Α   | 9          |
| В   | 56         |
| С   | 116        |
| D   | 100        |
| E   | 27         |
| F   | 38         |
| G   | 5          |
| Н   | 3          |
| - 1 | 10         |
| J   | 20         |
| K   | 6          |
| L   | 63         |
| M   | 2          |

| MVP | # articles |
|-----|------------|
| N   | 18         |
| 0   | 49         |
| Р   | 95         |
| Q   | 117+25     |
| R   | 77         |
| S   | 54         |
| Т   | 147        |
| U   | 172        |
| V   | 105+32     |
| W   | 57         |
| X   | 126        |
| Y   | 22         |
| Z   | 18         |

| Additional Categories  |
|------------------------|
| Human Remains ID (DV   |
| 92                     |
| Microbial & Viral DNA  |
| 59                     |
| Microhaplotypes/InDels |
| 53                     |
| Proteomics             |
| 15                     |
| Sexual Assault Policy  |
| 33                     |
| Other Applications     |
| 48                     |
| (+12 unsorted)         |

## DOJ Uniform Language for Testimony and Reports (ULTRs)

<u>Approved ULTR for the Forensic DNA Discipline –</u>
<u>Autosomal DNA with Probabilistic</u>
<u>Genotyping</u> (effective 3.18.19)

<u>Approved ULTR for the Forensic DNA Discipline – Mitochondrial DNA</u> (effective 3.18.19)

<u>Approved ULTR for the Forensic DNA Discipline – Y-STR DNA</u> (effective 3.18.19)

https://www.justice.gov/olp/uniform-language-testimony-and-reports



Needs Assessment of Forensic Laboratories and Medical Examiner/Coroner Offices



December 2019, 200 pages https://www.justice.gov/olp/page/file/1228306/download

#### UNITED STATES DEPARTMENT OF JUSTICE INTERIM POLICY FORENSIC GENETIC GENEALOGICAL DNA ANALYSIS AND SEARCHING

#### Purpose and Scope<sup>1</sup>

The purpose of this interim policy is to promote the reasoned exercise of investigative, scientific, and prosecutorial discretion in cases that involve forensic genetic genealogical DNA analysis and searching ('FGGS').2 It provides guidance to Department agencies when formulating a thoughtful and collaborative approach to important interdisciplinary decisions in cases that utilize this investigative technique. Collaboration between investigators, laboratory personnel, and prosecutors is important because the decision to pursue FGGS may affect privacy interests, the consumption of forensic samples, and law enforcement's ability to solve violent

#### UNITED STATES DEPARTMENT OF JUSTICE INTERIM POLICY FORENSIC GENETIC GENEALOGICAL DNA ANALYSIS AND SEARCHING

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> Approved: 09.02.2019 Effective: 11.01.2019

EDITORIAL

#### Responsible genetic genealogy

practices...

in forensic

**DNA** analysis

ne scientific development of forensic genetic ge- | or sex crime) for which a CODIS search resulted in no nealogy (FGG), which couples genetic analysis with investigation of publicly available genealogy information, has successfully transformed law enforcement investigations by solving more than 50 cases over the last 18 months in the United States. However, use of FGG by law enforcement has preceded widespread development of best practices to protect the genetic privacy of private citizens who have voluntarily submitted samples to genealogy databases. Absent best practices, use of FGG could lead to compromised cases, diminished use, or the loss of this new investigative tool. Public support for FGG could be jeopardized and confidence ods (SWGDAM; swgdam.org), which recommends

in forensic DNA analysis could be undermined. As the custodian of a national law enforcement DNA database (CODIS), the U.S. Federal Bureau of Investigation (FBI) is looked to by many in the law enforcement and forensic DNA communities for guidance, and its efforts often influence the global community. The emergence of FGG suggests that further discussions on privacy, genomics, and the use of genealogy by law enforcement would be beneficial. Accordingly, the FBI seeks to engage the scientific and bioethics communities in such a dialogue.

Use of FGG involves databases and family trees composed of genetic data of private citizens who are not under suspicion

these databases, potential perpetrators may be uncovered by identifying their close or distant relatives, and then building family trees that can extend over many generations and may include hundreds to thousands of relatives. To date, this approach is only used if crime scene DNA has not matched genetic profiles in the CO-DIS database of known offenders and arrestees. A consensus has emerged that there is no legal prohibition on such use. The question is how it should be done.

Under a recently released interim policy from the U.S. Department of Justice (DOJ), effective this November, federal investigative agencies may develop internal policies and procedures and can utilize FGG if the case involves an unsolved violent crime (homicide

matches, and for which reasonable investigative leads have been pursued. The DOJ Interim Policy is the first substantial attempt to address "how genetic genealogy should be done." The interim guidance restricts investigative agencies to using only public databases or direct-to-consumer genetic genealogy services that provide clear notice to users and the public that law enforcement may access their sites for investigative or unidentified human remains identification purposes.\*

The forensic DNA community is also working on guidance to address the "how to" question. In April, the Scientific Working Group on DNA Analysis Meth-

standards to the FBI for CODIS and issues guidance for the forensic DNA community, formed an interim committee on FGG composed of genealogists, bioethicists, academicians, law enforcement, and forensic scientists, as well as representatives of the European Network of Forensic Science Institutes and the International Society of Forensic Genetics (the author is co-chair of this committee). This group held an FGG technical symposium for SWGDAM membership in July and recently submitted recommendations to SWGDAM leadership that included establishing an FGG Working Group.

With the FBI and other agencies now

moving to develop internal policies and for any crimes. When searching crime scene DNA in | procedures under the DOJ interim policy, the FBI has committed to leading the process of receiving stakeholder input by hosting a symposium on Genetic Privacy and Law Enforcement in 2020. In addition to symposium presentations, a comprehensive discussion of the interim FGG policy should also consider comments solicited from the scientific community on FGG privacy and ethical implications, metrics required by the interim policy, transparency, and SWGDAM guidance and recommendations. To initiate this review, the scientific community and other interested parties are encouraged to provide the FBI with comments at forensicgenealogy@fbi.gov.



is chief biometric scientist at the U.S. Federal Bureau of Investigation (FBI) Laboratory Division in Quantico, VA, USA tfcallaghan2@fbi.gov

-Thomas F. Callaghan

"U.S. Department of Justice, Interim Policy on Forensic Genetic Genealogical DNA Analysis

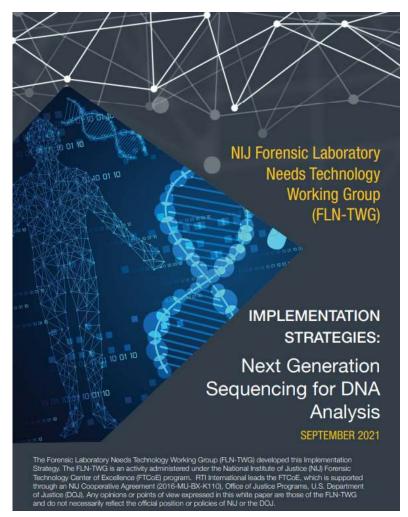
10.1126/science.aaz6578

SCIENCE sciencemag.org

II OCTUBER 2010 + VOL 366 ISSUE 6462 155

#### **NIJ FLN-TWG**

(Forensic Laboratory Needs Technology Working Group)







### **Human Factors Sourcebook**

 https://forensiccoe.org/human\_factors\_forensic\_science\_ sourcebook/

Publication of six articles in *Forensic Science International: Synergy*, March 2022

- 1) Overview of special issue: Human factors in forensic science practice sourcebook
- 2) The need for research-based tools for personnel selection and assessment in the forensic sciences
- 3) The benefits of errors during training
- 4) Challenges to reasoning in forensic science decisions
- 5) Stressors in forensic organizations: Risks and solutions
- 6) <u>Describing communication during a forensic investigation using the Pebbles on a</u>
  Scale metaphor

## Organization of Scientific Area Committees for Forensic Science (OSAC)

- OSAC 2020-N-0007, Best Practice Recommendations for the Management and Use of Quality Assurance DNA Elimination Databases in Forensic DNA Analysis (added April 6, 2021, to OSAC website and sent to <u>ASB</u> for further development and publication).
- OSAC 2020-S-0004, Standard for Interpreting, Comparing and Reporting DNA Test Results Associated with Failed Controls and Contamination Events (added June 1, 2021, to OSAC website and sent to ASB for further development and publication).

#### **Academy Standards Board (ASB)**

- ANSI/ASB Standard 020, Standard for Validation Studies of DNA Mixtures, and Development and Verification of a Laboratory's Mixture Interpretation Protocol, First Edition, 2018 (added to OSAC Registry May 12, 2020).
- ANSI/ASB Standard 040, Standard for Forensic DNA Interpretation and Comparison Protocols, First Edition, 2019 (added to OSAC Registry May 12, 2020).
- ANSI/ASB Standard 022, Standard for Forensic DNA Analysis Training Programs, First Edition, 2019 (added to OSAC Registry September 1, 2020).
- 4. ANSI/ASB Standard 018, Standard for Validation of Probabilistic Genotyping Systems, First Edition, 2020 (added to OSAC Registry May 4, 2021).
- ANSI/ASB Standard 023, Standard for Training in Forensic DNA Isolation and Purification Methods, First Edition, 2020 (added to OSAC Registry August 3, 2021).
- ANSI/ASB Standard 110, Standards for Training in Forensic Serological Methods, First Edition, 2020 (added to OSAC Registry August 3, 2021).
- 7. ANSI/ASB Standard 115, Standard for Training in Forensic Short Tandem Repeat Typing Methods using Amplification, DNA Separation, and Allele Detection, First Edition, 2020 (added to OSAC Registry August 3, 2021).
- 8. ANSI/ASB Standard 116, Standard for Training in Forensic DNA Quantification Methods, First Edition, 2020 (added to OSAC Registry August 3, 2021).

## **Guidance Documents – Australia and Europe**

## **Australian National Institute of Forensic Science (NIFS)**

- Case Record Review in Forensic Biology (September 2019)
- 2. Empirical Study Design in Forensic Science A
  Guideline to Forensic Fundamentals
  (September 2019)
- 3. <u>Transitioning Technology from the Laboratory</u> to the Field Process and Considerations for the Forensic Sciences (December 2019)

## **European Network of Forensic Science Institutes (ENFSI) DNA Working Group**

- DNA Database Management Review and Recommendations (April 2019)
- Guideline for the Training of Staff in Forensic DNA Laboratories (March 2022)

https://www.anzpaa.org.au/forensic-science/our-work/products/publications

## **ISFG DNA Commissions**

## https://www.isfg.org/Publications/DNA+Commission

Provide recommendations and considerations to enable interconnectivity and advance the quality of forensic DNA evidence

- Autosomal STRs and allele nomenclature
  - Bär et al. 1994 allelic ladders & partial repeats (e.g., 9.3)
  - Bär et al. 1997 motif choice & repeat nomenclature
  - Parson et al. 2016 8 considerations with sequence data
- mtDNA
  - Carracedo et al. 2000 guidelines on QC, nomenclature, heteroplasmy, and interpretation
  - Parson et al. 2014 16 recommendations on sequencing, quality control, interpretation, and databases
- Y-STRs
  - Gill et al. 2001 locus & allele nomenclature, allelic ladders
  - Gusmão et la. 2006 repeat nomenclature, new loci
  - Roewer et al. 2020 statistics and report information
- X-STRs
  - <u>Tillmar et al. 2017</u> 10 recommendations on use of X-STRs in kinship analyses, linkage, and statistical calculations

- DNA mixture interpretation and assessing evidence
  - Gill et al. 2006 nine recommendations on mixture interpretation (e.g., LR vs. CPI)
  - Gill et al. 2012 allele drop-in and drop-out using probabilistic methods
  - Gill et al. 2018 formulation of propositions; investigator and evaluator roles
  - Gill et al. 2020 activity level propositions
- Other topics
  - DNA Polymorphisms: Brinkmann et al. 1989, 1992, Bär et al. 1992
  - Disaster Victim Identification: Prinz et al. 2007 12 recommendations
  - Biostatistics in Paternity Testing: Gjertson et al. 2007
  - Non-human DNA: <u>Linacre et al. 2011</u> 13 recommendations
  - STRidER: Bodner et al. 2016 quality control of autosomal STR allele
  - Software Validation: Coble et al. 2016 16 recommendations and expectations

#### **Current (Active) DNA Commissions**

- 1. STR nomenclature with DNA sequencing information (building on STRAND efforts): *K. Gettings*
- 2. Phenotyping (building on VISAGE efforts): *M. Kayser*