Forensic Investigative Genetic Genealogy (FIGG)

Dr. Claire L. Glynn

Professor of Forensic Science Founding Director - FIGG Program Executive Director - Lee Institute

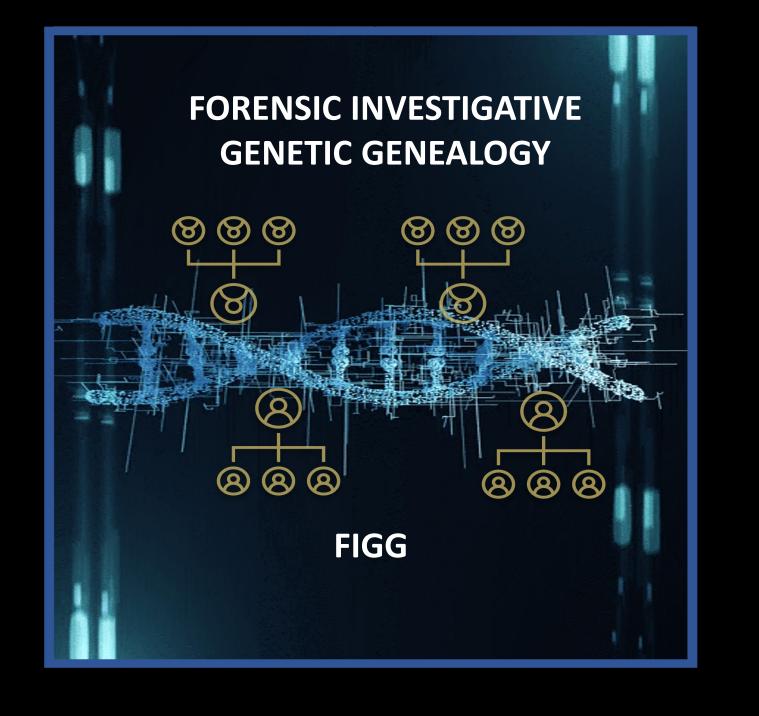


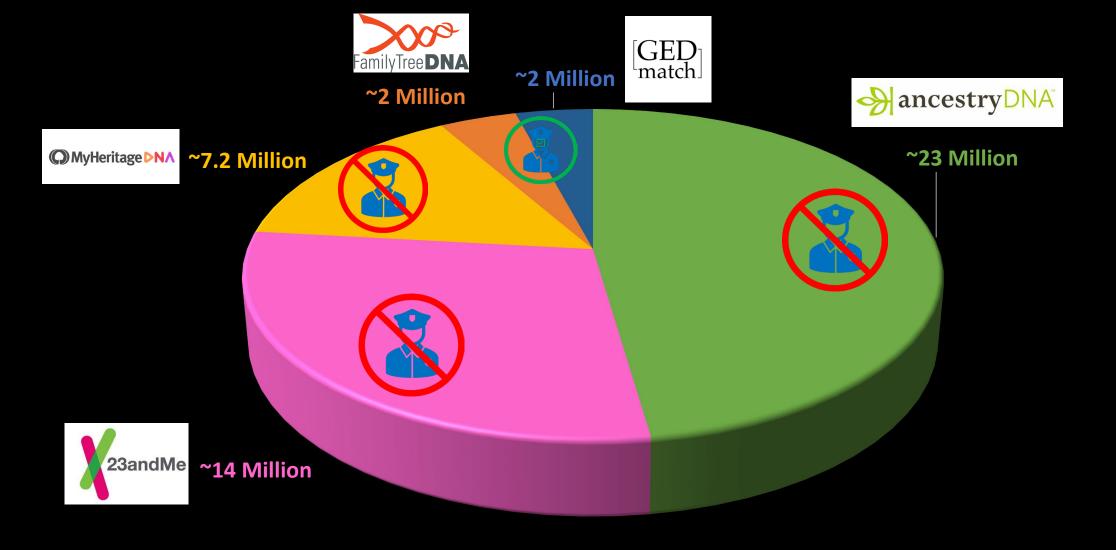
HENRY C. LEE COLLEGE OF CRIMINAL JUSTICE AND FORENSIC SCIENCES

Department of Forensic Science







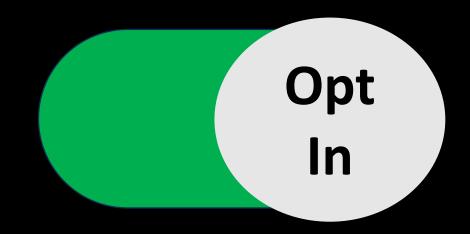


Consumer DNA: >48 MILLION PEOPLE FIGG Databases: <4 MILLION PEOPLE





All users can:

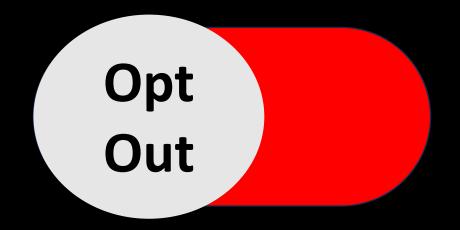


for Law Enforcement Comparisons





All users can:



for Law Enforcement Comparisons

1.

FIGG Case File



)

Genetic Genealogy

SNP Sequencing

Case Assessment

3

5

Family Trees
Review/Report

4.

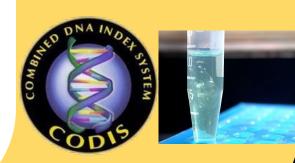
Case Assessment



FIGG Checklist:

- ✓ Case Criteria/Type meets:
 - ☑ DOJ Interim Guidelines
 - ✓ Individual State Laws
 - ✓ Database Terms of Service/Use
- ✓ Forensic STR Profile Developed:
 - ✓ Uploaded to CODIS
 - **✓** State-level Familial Search (if permitted)
- ✓ Sufficient DNA Remaining (Quantity & Quality)
- ✓ All other Investigative Leads Exhausted





SNP Sequencing



Forensic DNA Analysis

Short Tandem Repeats (STRs)

CODIS 20 Core (US) / DNA17/23+ (Europe)

Person 2: ACTGTCGATAGATAGATAGATAGATAGATAGATAGATAACTGTC

SNP Sequencing

Single Nucleotide Polymorphisms (SNPs)

600,000 – >1,000,000 markers

Person 1 : GCTGTATGAGTAGAAGATCGAT

Person 2 : GCTGTATGACGAGAGATCGAT

SNP Sequencing







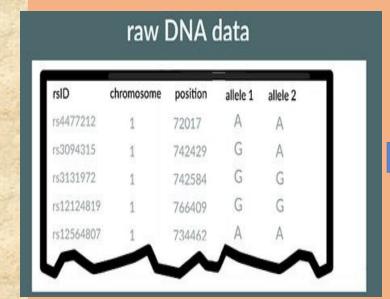


SNP Microarray (~600,000 SNPs)

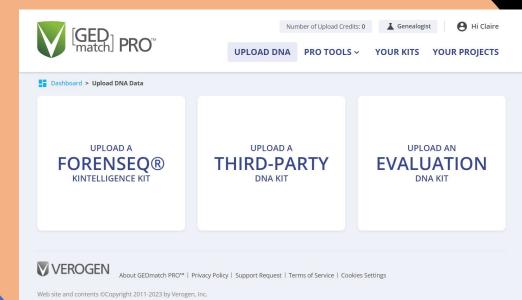
Targeted SNP Kit – Kintelligence (~10,000 SNPs)

Whole Genome
Sequencing
(~1 million SNPs)





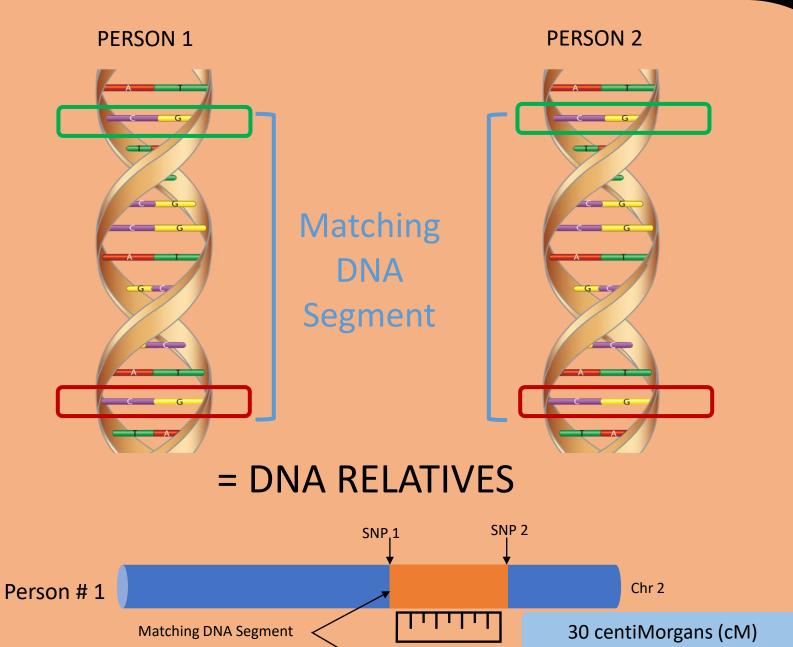
Upload DNA Data to FIGG Approved Databases





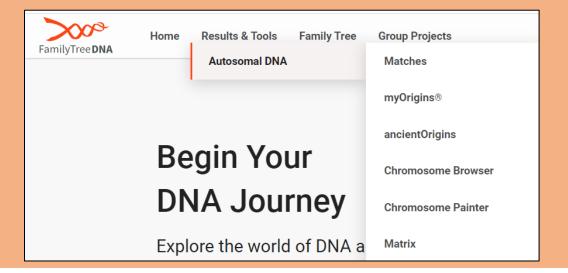


Person # 2



Chr 2







Number of Upload Credits: 0

▲ Genealogist

Hi Claire

UPLOAD DNA PRO TOOLS ~ **YOUR KITS**

YOUR PROJECTS

GEDmatch PRO™ Dashboard

This is your GEDmatch PRO™ dashboard where you can quickly access your latest uploaded kits and projects you are a working on.

Your Kits

Upload New Kit +

One-to-One Kinship DNA Comparison

One-to-Many Segment-Based

One-to-Many Kinship

AutoKinship

Admixture (Eurogenes/K13)

Are Your Parents Related?

Kit Diagnostic Utility

Kit Evaluation

Multiple Kit Analysis

One-to-One Autosomal DNA Comparison

One-to-One Q Matching

Segment Search

Triangulation

User Lookup

All Pro Tools

Α

Admixture (Eurogenes/K13)

Are Your Parents Related?

AutoKinship

Kit Diagnostic Utility

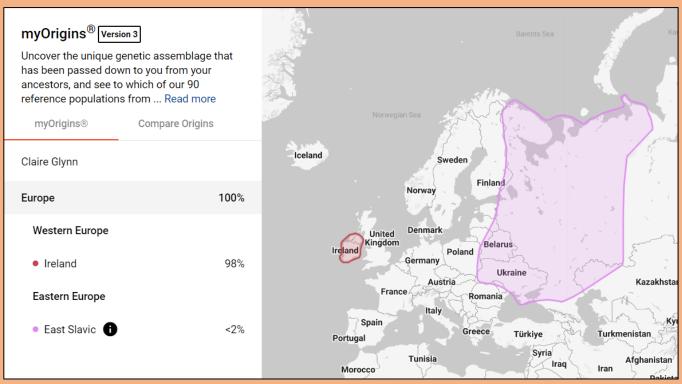
Kit Evaluation

М

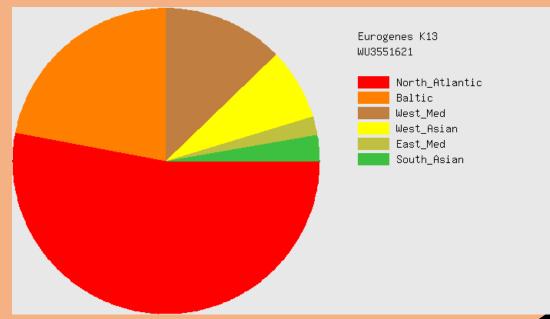
Multiple Kit Analysis

One-to-Many Kinship





Biogeographic Ancestry Estimation



Shared DNA

Genetic Genealogy

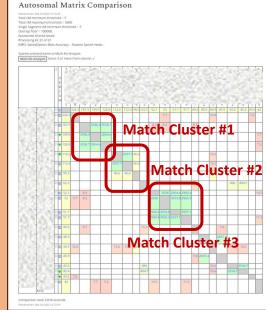


| Visualization Options | | | | Asosomal | | | | Haplogroup X-DNA | | | | | | | |
|-----------------------|--------|----------------------|------------------------|-------------------------|-----------------|-------|----------|------------------|-------|------|-----|------------|-----------|-----------------------|-----------|
| 0 | Kit ↑ | Name (* => alias) | Email ↑ | GED ① ↑ WikiTree | Age (days) ↑ | Sex ↑ | Total cM | Largest ↑ | Gen ↑ | Mt ↑ | Y ↑ | Total cM ↑ | Largest ↑ | Source ↑ | Overlap ↑ |
| 0 | 12345 | Cillian Murphy | peakyB@gmail.com | | 2594 | М | 115.3 | 29.9 Q | 3.48 | | | 0 | 0 | Migration - F2 - A | 187790 |
| 0 | 54321 | Ryan Gosling | ken@aol.com | GED | 645 | М | 75.3 | 40.3 Q | 3.79 | | | 0 | 0 | MyHeritage | 78376 |
| 0 | 23451 | Al Pacino | thegodfather@gmail.com | | 3082 | М | 74.9 | 23.4 Q | 3.79 | | | 0 | 0 | Migration - F2 - F | 197171 |
| | 34512 | Emma Stone | estone@outlook.com | | 645 | F | 74.8 | 39.7 Q | 3.79 | | | 0 | 0 | MyHeritage | 79805 |
| 0 | 45123 | Emily Blunt | eblunt@gmail.com | | 2344 | F | 74.7 | 21.6 Q | 3.79 | | | 0 | 0 | Migration - F2 - A | 187542 |
| 0 | 543216 | Robert Downy Jr | rdj@outlook.com | | 1907 | М | 65.8 | 47.7 Q | 3.88 | | | 0 | 0 | AncestryDNA V2 | 213355 |
| 0 | 13452 | Christopher Nolan | cnolan@gmail.com | | 966 | М | 54.2 | 37.9 Q | 4.02 | | I | 0 | 0 | FTDNA | 79615 |

*Kit #, Names, and Emails altered for Privacy

Shared DNA Matches i.e., Genetic Relatives





Relationship Probabilities for 115 total shared cM Relationship Type Probability 2C1R Group 32.3% 2C Group 9.1% 2C, 1C2R, Half-1C1R, Half-2G-Aunt/Uncle/Niece **3C1R Group** 8.9% 3C1R, Half-3C, Half-2C2R, 2C3R **4C Group** 3.7% 4C, 3C2R, Half-3C1R, Half-2C3R 4C1R Group 0.5% 4C1R, Half-4C, Half-3C2R, 3C3R 5C Group 0.2% 5C, 4C2R, Half-4C1R, Half-3C3R Relationship prediction algorithm developed by Brit Nicholson. Read the methodology here.

CLOSE



Verify identities of DNA matches:

- John Smith: johnsmith84@gmail.com
- Some use aliases

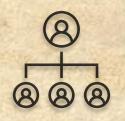


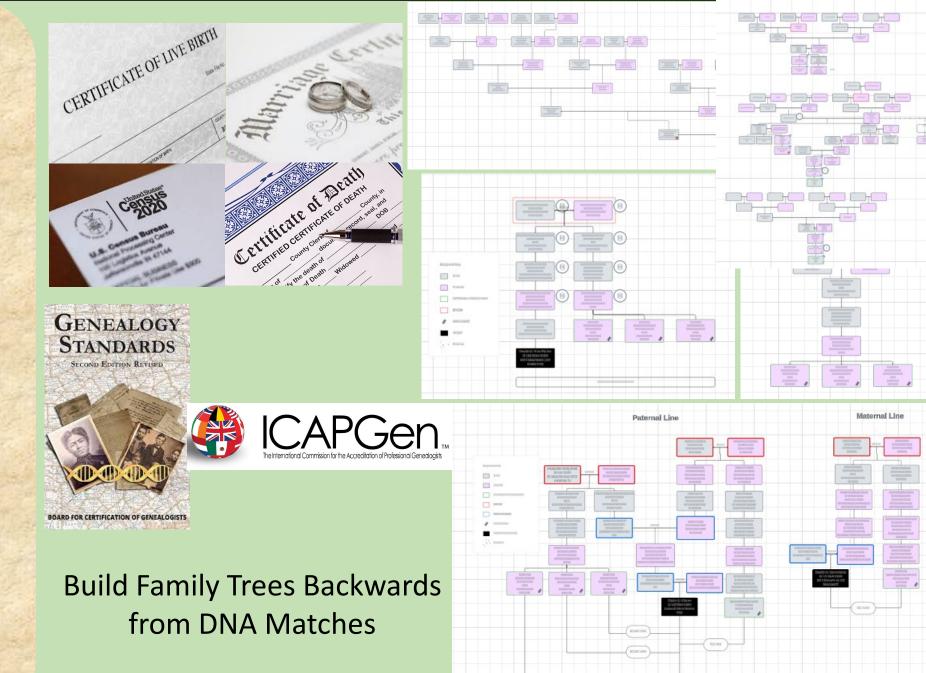


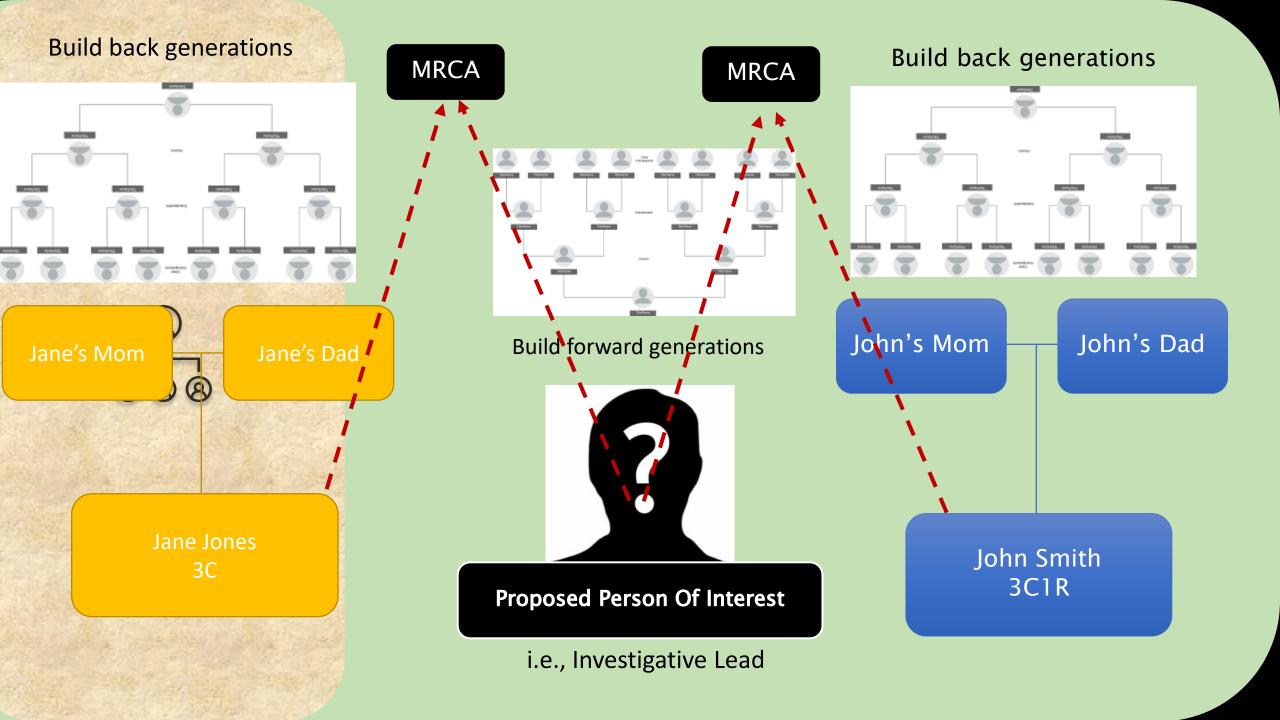




Family Trees



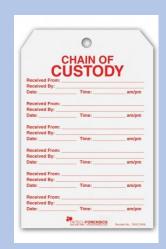




Review/ Report



- Every step of a FIGG investigation should be documented.
 - Every genetic genealogy tool performed
 - ✓ Genealogical records used to build family trees
- ✓ Final report detailing the FIGG analysis should be produced.
- ✓ Any other competent FIGG analyst should be able to interpret the findings and reach the same conclusion.
- ✓ Administrative and Technical Review









Quality Assurance Standards Documents

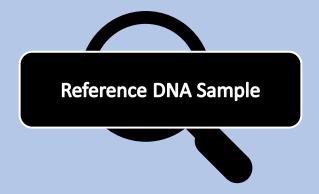




Review/ Report



Investigative Lead provided to Law Enforcement Agency



Violent Crime Suspect:

Overt or Covert DNA Collection

Unidentified Human Remains:

Closest Living Relative DNA Collection

STR Profile Comparison

| Locus | Forensic Unknown | Reference Known |
|-------|------------------|-----------------|
| D3 | 15, 16 | 15, 16 |
| vWA | 14, 16 | 14, 16 |
| D16 | 9, 10 | 9, 10 |
| D2S1 | 14, 15 | 14, 15 |
| D8 | 12, 13 | 12, 13 |
| D21 | 28, 31 | 28, 31 |
| D18 | 12, 15 | 12, 15 |
| D10 | 14.15 | 14.15 |

MATCH − INCLUDED ✓

| rua | 24, 20 | 24, 20 |
|-----|--------|--------|
| D1 | 13, 16 | 13, 16 |

NO MATCH − EXCLUDED *****

| D12 | 18, 19 | 18, 19 | | | |
|--------|----------|----------|--|--|--|
| D22 | 11, 16 | 11, 16 | | | |
| SE33 | 17, 25.2 | 17, 25.2 | | | |
| D7 | 7, 12 | 7, 12 | | | |
| CSF1PO | 11, 12 | 11, 12 | | | |
| D13 | 11 | 11 | | | |
| TP0X | 8 | 8 | | | |
| D5 | 11 | 11 | | | |
| Amelo | XY | XY | | | |
| DYS391 | 11 | 11 | | | |

COLD CASES 2018 - 2024 Case 003-0907196

cases have benefited from Forensic Investigative Genetic Genealogy in the United States

FIGG: 2018 - 2024

Building the plane while flying it



Law Forensic Enforcement Scientists Agencies **FIGG** Legal **FIGG Experts Professionals**

National Technology Validation and Implementation Collaborative

Forensic Investigative Genetic Genealogy (FIGG) Working Group



Forensic Science International: Synergy
Volume 7, 2023, 100446



National Technology Validation and Implementation Collaborative (NTVIC): Guidelines for establishing Forensic Investigative Genetic Genealogy (FIGG) programs

Ray A. Wickenheiser (2) 🖾 , Jennifer Naugle, Brian Hoey, Rylene Nowlin, Swathi A. Kumar, Mark A. Kubinski, Claire Glynn, Raymond Valerio, Lance Allen, Stephanie Stoiloff, Jennifer Kochanski, Christi Guerrini, Anne Marie Schubert



Increase # of people in FIGG Databases:

Increase # of cases solved









FIGG Potential:

Why wait until a case is cold?

- **☑** Prevent Serial Offenders
- **☑** Deter Potential Offenders
- **☑** Prevent Wrongful Convictions
- **☑** Clear All Backlogs/Cold Cases

1934



Method Validation

Legislation









2024



Protecting:

Maintain Public Trust:
✓ Adhere to policies, laws, database terr

✓ Adhere to policies, laws, database terms of use, and best practices

Follow Forensic & Criminal Justice Standards:

✓ Education & Training ✓ Documentation✓ Chain of Custody ✓ Oversight

Public Safety Individual Privacy

Increase Awareness:

✓ Increase #'s in Databases = Increase #'s of Cases

Through the Responsible & Ethical use of FIGG



FORENSIC INVESTIGATIVE GENETIC GENEALOGY

UNIVERSITY OF NEW HAVEN

Scan Here to Learn about our Graduate Certificate program in FIGG





University of New Haven

HENRY C. LEE COLLEGE OF CRIMINAL JUSTICE AND FORENSIC SCIENCES

Department of Forensic Science

Connect with me on Linked in



Claire Glynn

Professor - Forensic Expert - Forensic Investigative Genetic Genealogy (FIGG)



