# Use of search keys to solve more cases with expanded DNA profiles

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### Disclaimer

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### Outline

- Current use of databases
- Direct and indirect matching
- Search keys
- Case flow
- Business case
- Recommendations

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### Current Practice

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#### **Current Practice**

- DNA databases are very effective at producing **investigative leads**
- 206 Accredited Crime Labs in United States using CODIS (NDIS) via STRs
- Forensic Sample: Crime Scene sample from the putative perpetrator
- Direct matching: comparison of the forensic profile to known samples (individually or via a DNA database) where the entire profile is in common for the same source

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#### **CODIS** and Direct Comparison

- As of October 2021, the NDIS contained over 14,836,490 offender profiles, 4,513,955 arrestee profiles, which equals 19,350,445 known profiles available for comparison.
- There were also 1,144,255 forensic profiles in NDIS.
- Those forensic profiles recovered from biological materials deposited at crime scenes have produced 587,773 hits, which equates to roughly a 51.37% hit rate.
- This also means that roughly **556,482** forensic profiles have not been hit upon.
- The unsolved crimes represent a massive opportunity to solve crime and prevent future crimes through investigative leads.

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#### Opportunity

- A larger database means more hits; more investigative leads
  - Directly (physically)
  - Indirectly (scientifically)
- Strategies:
  - Direct Matching
    - Arrestees
    - Lawfully owed DNA samples
    - Unidentified Human Remains (UHRs) Direct and Indirect
  - Indirect Matching
    - Partial matching
    - Familial Searching
    - FIGG
    - EDIM



## Indirect Matching

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#### Indirect matching

- DNA profiles partially match (share common areas) due to biological relatedness
- Biological relatedness conveys known inheritance patterns
- Kinship analysis permits evaluation of indirect match to postulate the level of biological relatedness
- Infer Relationship immediate family, extended family, paternal egal Powered for 33 years and maternal family lines



#### Indirect matching techniques

- Partial matching (discover non-identical profiles sharing significant DNA similarities) – passive search
- Familial Searching active search of state CODIS for immediate family members
- Forensic Investigative Genetic Genealogy (FIGG) active search of genealogy database to find potential kin, then use genealogical researching to develop investigative leads

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Indirect matching increases the size of the database by including biological relatives

#### Descriptions

- Partial matching and familial searching use STRs in common and the relative rarity of STRs to develop a Likelihood Ratio (LR)
- FIGG uses SNPs. SNPs map the genome and are used by algorithms to compare fragments (similarities measured in centimorgans).
  - cM (centimorgans) is the measuring stick (1 cM is roughly 1 million base pairs)
  - shared fragments of DNA where those fragments have the same SNPs
  - shared number and size of fragments in common provides a measure of relatedness

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Forensic Investigative Genetic Genealogy (FIGG) – use of SNPs to search genealogy database to find potential kin sharing portions of DNA (centimorgans), then use genealogical researching to develop investigative leads

## EDIM and Search Keys

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#### Proposed Expanded DNA Indirect Matching (EDIM)(1)

- What is a Search Key?
- Search keys are components (aspects/types) of DNA profiles that are shared by related individuals, but are not unique to that individual, but rather tell something about their family through inheritance
- Use of a search key can effectively search samples for potential relatives
  - Y-STR inherited paternally
  - X Chromosome inherited from maternal family line in males, from both sides in females
  - mtDNA inherited maternally

#### Forensic Y-STR as a search key

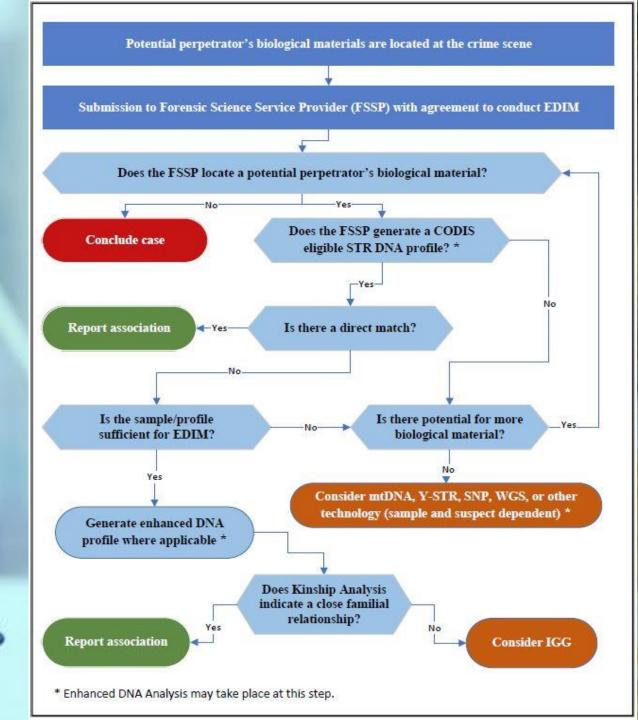
- Y-STRs are inherited paternally
  - Using the Y-STR as an example, expanding the DNA profile beyond the CODIS core loci would include a Y-STR profile
  - Y-STR used to search for candidate male biologically related individuals
  - Once a candidate is located, kinship analysis can be conducted to evaluate the level of relatedness

#### Proposed Expanded DNA Indirect Matching (EDIM)(1)

- 1. Develop an expanded DNA profile (Y-STR, X Chromosome, mtDNA, SNP, WGS)
- 2. Search cases against each other using search keys
- 3. Conduct kinship analysis between cases with the same search key
- 4. Report cases with close relatives
  - a. Cases with previous matches or leads add value to the indirectly matched cases (biological relatives)
- 5. Consider SNPs for IGG for more distant relatives

#### Proposed Expanded DNA Indirect Matching (EDIM) workflow (1)

- Several entry points for NGS (see \*)
- 1. At original DNA analysis
- 2. After CODIS search is unsuccessful
- 3. When other typing methods are not successful
- Is this repetition the best process?
- Our goal is to maximize the evidence, therefore expanded profiles create the most matching opportunities



# Bump up the value

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#### Unidentified Human Remains (UHRs)

- Indirect and direct matching opportunity which qualifies for CODIS
- Direct
  - High risk lifestyle (victim and perpetrator)
  - Opportunity to close open cases
  - Indirect matching
    - Help ID UHRs through EDIM and FIGG
    - ID in turn feeds direct matching
  - Provides closure

Help solve case, particularly if UHR is homicide victim

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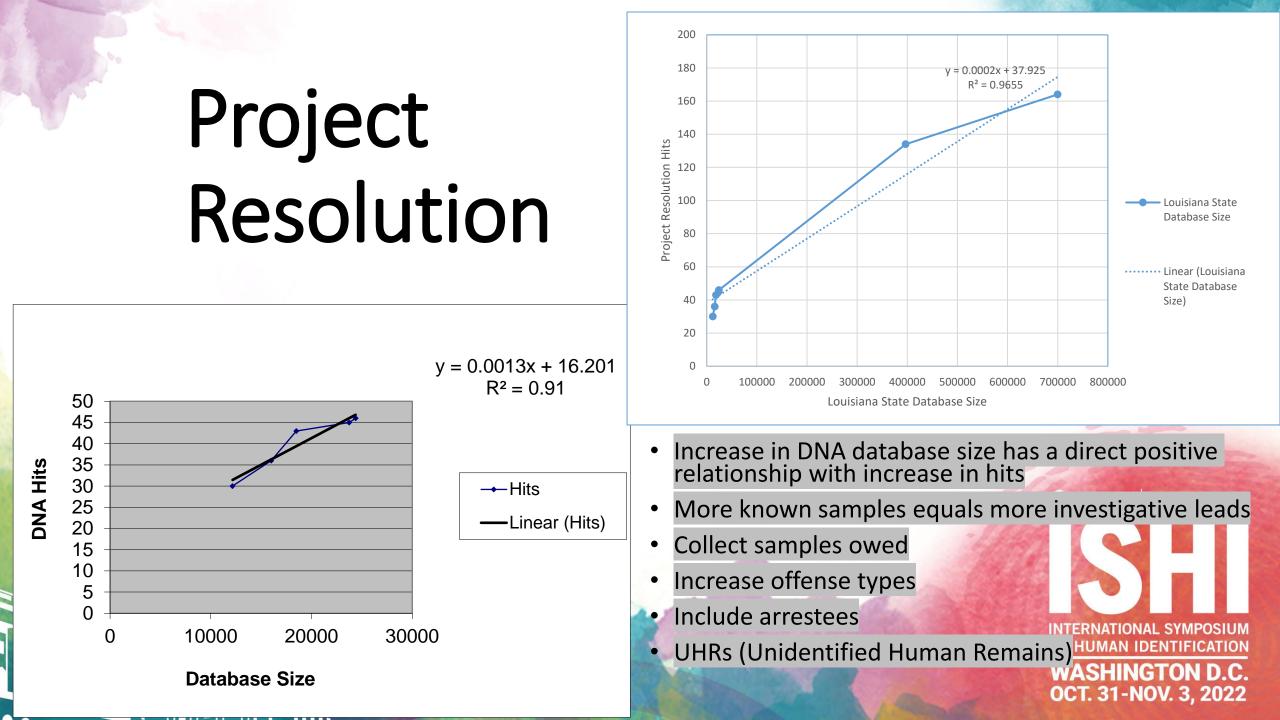
#### Universal DNA Database (for illustration only)

- Cost of \$16.5 Billion (331.4 Million Americans at \$50/sample)
- 42% plus remaining hit rate potential compared to cold case projects (Palm Beach, Acadiana and Detroit)
- 139,380 sexual assaults in US annually
- \$435,419 per sexual assault
- 26.22 preventable sexual assaults per hit
- Opportunity cost is \$66.8 Trillion
- ROI is \$4,050 per \$1 spent



**Project Resolution** – Conduct DNA analysis on 605 case cuttings, 285 CODIS profiles resulted, with 164 hits

	Cost of Sexual Assault	CODIS Hits	Project Resolution Cost	Recidivism Factor	Return on Investment	
Conservative Model	\$111,238	164	\$286 <i>,</i> 000	7	\$446.51	ed for 33 years
Aggressive Model	\$435 <i>,</i> 419	164	\$286 <i>,</i> 000	26.22	\$6546.63	
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Cost per sexual assault	\$435,419	
Number of preventable sexual	26.22	
assaults per hit		
Number of hits	164	
Size of Louisiana SDIS	699,618	
Louisiana SDIS offenders per hit	4,266	
Cost benefit of Project Resolution	\$1,872,336,534	
Analysis cost per database sample	\$50	
Cost of Louisiana SDIS offender	\$33,980,900	
sample analysis		
Cost benefit per \$1 spent	\$53.52	
Return on investment percentage	5,352	

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#### Indirect matching – frequency of relatives

- Depends on biological relatives to provide leads
- Frequency of siblings in the New York State DNA database has been estimated
- In 2017 there were 536 sets of identical twins, with the database size at approximately 700,000 individuals
- Rate of identical twins is 1 in 250 live births
- 536 X 250 = 134,000 sibling pairs
- 134,000/700,000 X 100 = 19.14% (1), so 1 in 5 individuals will have a sibling (low estimate as parent-child relationships are additional)

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#### Business Case for NGS/EDIM

- The estimated number of sexual assault hits annually is 49,964 (7)
- EDIM has potential to increase the hit rate by 19.14%
- This will solve an additional 9,563 sexual assaults annually
- The cost of a single sexual assault is \$435,519 and each solved sexual assault prevents 26.22 sexual assaults (3)
  - Estimated cost of crime savings is \$109.18 Billion
  - Potential to prevent 250,741 sexual assaults

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#### Considerations and Recommendations

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#### **Crime Scene DNA Bioethics**

- Autonomy right to privacy and protection from unwarranted search and seizure
- Discarded material no presumed right to privacy: autonomy is vacated
- Proportionality weighing the benefits and risks of competing options to maximize the overall good while minimizing the overall downsides
- Recidivism damage to society caused by a small number of individuals committing repeat and escalating crimes on new victims
- Objectivity scientific evidence is unbiased, neutral, can be retested, challenged in court, debated (investigative lead)
- Forensic profile is developed pre-suspect and STR profile from lead is confirmed with STRs within an accredited crime lab system for court

#### Why Now?

- We forget that these criminals are out there now committing new crimes and we have their DNA at the crime scene in existing crimes
- NGS/EDIM is not just for cold cases; this is for any case we do not get a CODIS hit upon
- Our mission is to maximize the value of evidence

The huge ROI demonstrates the value of DNA investigative leads and databases



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#### Recommendations

- Increase DNA Database size for direct matching
  - Expand qualifying offenses
  - Include arrestees
  - Maximize UHRS
  - Collect all lawfully owed DNA samples
- Increase the DNA database by utilizing indirect matching
  - NGS/EDIM is not just for cold cases; this is for any case we do ega Powered for 33 years not get a CODIS hit upon

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 Case for an expanded DNA profile on all forensic (crime scene) samples

#### Current

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## Thank you!

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