

# OSAC RESEARCH NEEDS ASSESSMENT FORM



**Title of research need:** Development of infrastructure to compile and share raw electronic data for training and tool development

**Describe the need:**

We have entered the age of “big data” and artificial intelligence. The forensic DNA community has yet to take full advantage of current computational power, methods, or establish an efficient means of sharing data. Unfortunately, the limited availability of large and diverse DNA profile databases currently hinders the implementation of computationally advanced solutions that could profoundly impact DNA mixture interpretation and impedes wide scale assessments of currently used methods of interpretation. The value of “big data” has only recently been realized in the forensic community with the development and assessment of new software tools and statistical models that have already positively impacted the quality of forensic DNA analysis. However, the means to develop these tools and assessment methods are reliant on the availability of large-curated datasets.

Although many research, developmental, and validation projects have compiled large datasets, many are not available due to confidentiality or other related concerns. In addition, available sets may lack the diverse and comprehensive types of DNA profiles (or other related types of data) that would permit robust software development of comparative studies. This research need calls for the development of large and diverse data sets that are accessible to the public and the infrastructure to support such databases. The infrastructure includes: (1) the development of computational resources to support housing of large data sets, (2) the means to properly curate the databases—quality assurance/control methods and tools, documentation and monitoring throughout the “life” of the database, and (3) the development of legal, policy and data sharing framework allowing for the use and updating of databases among academic, governmental and private institutions. The PROVEDIt dataset<sup>21</sup> is a principal example of the type of database and architecture that can be used effectively to realize the benefits of these types of datasets/databases.

**Keyword(s):** Database, validation, NGS, DNA mixture, mixture interpretation, artificial intelligence

**Submitting subcommittee(s):** Human Biology **Date Approved:** 05/16/2025

**Background Information:**

1. Does this research need address a gap(s) in a current or planned standard? (ex.: Field identification system for on scene opioid detection and confirmation)

Yes, ANSI/ASB Standard 020, Standard for Validation Studies of DNA Mixtures, and Development and Verification of a Laboratory’s Mixture Interpretation Protocol

<sup>1</sup> PROVEDIt Initiative (Project Research Openness for Validation with Experimental Data) is a publically available resource of 25,000+ DNA profiles <sup>1</sup>.

2. Are you aware of any ongoing research that may address this research need that has not yet been published (e.g., research presented in conference proceedings, studies that you or a colleague have participated in but have yet to be published)?

Yes, several research projects and proposals as well as mixture analysis reviews including the recent NIST Review on DNA Mixture Interpretation.

3. Key bibliographic references relating to this research need: (ex.: Toll, L., Standifer, K. M., Massotte, D., eds. (2019). Current Topics in Opioid Research. Lausanne: Frontiers Media SA. doi: 10.3389/978-2-88963-180-3)

1. PRESIDENT'S COUNCIL OF ADVISORS ON SCIENCE AND TECHNOLOGY. President's Council of Advisors on Science and Technology. Forensic Science in Criminal Courts: Ensuring Scientific Validity of Feature-Comparison Methods. (2016).
2. Alfonse, L. E., Garrett, A. D., Lun, D. S., Duffy, K. R. & Grgicak, C. M. A large-scale dataset of single and mixed-source short tandem repeat profiles to inform human identification strategies: PROVEDIt. Forensic Sci. Int. Genet. 32, 62–70 (2018).
3. Butler, J. M., Iyer, H., Press, R., Taylor, M.K., Vallone, P.M. and Willis, S. DNA Mixture Interpretation: A NIST Scientific Foundation Review NISTIR 8351-DRAFT. (2021). <https://doi.org/10.6028/NIST.IR.8351-draft>.

4. Review the annual operational/research needs published by the National Institute of Justice (NIJ) at <https://nij.ojp.gov/topics/articles/forensic-science-research-and-development-technology-working-group-operational#latest>? Is your research need identified by NIJ?

No

5. In what ways would the research results improve current laboratory capabilities?

Generally, the development of these databases can impact the ability of researchers to develop new computational tools to better address the analytical and interpretational needs of the forensic DNA analyst. Specific examples, however, cannot be provided, the value in this is in providing the “raw materials” (data) to be able to explore new computational methods. Additional benefits include the ability to use these data sets for training of new analysts and students training to become forensic practitioners.

6. In what ways would the research results improve understanding of the scientific basis for the subcommittee(s)?

New computational tools, specifically using artificial intelligence, are highly dependent on data availability and the quality of the data. If these databases are used for the training and testing of the newly developed computational tools, the subcommittee will have access to these data and thus be able to better understand the response, use and results of these computational tools.

7. In what ways would the research results improve services to the criminal justice system?

Provide the raw data needed to further develop new computational tools that will improve the confidence and quality of results. Additional benefits include the ability to use these data sets for training of new analysts and students training to become forensic practitioners. These databases and methods will permit independent assessments of new and historically relevant computational tools, allowing for a direct comparison of different methods using the same data sets.

8. Status assessment (I, II, III, or IV):

I

	Major gap in current knowledge	Minor gap in current knowledge
No or limited current research is being conducted	I	III
Existing current research is being conducted	II	IV

*This research need has been identified by one or more subcommittees of OSAC and is being provided as an informational resource to the community.*