

Response Evaluation In Neurofibromatosis Schwannomatosis  
INTERNATIONAL COLLABORATION

- If sharing any data or information from these slides generated by the REiNS International Collaboration, please acknowledge the authors, group chairs, and specific working group.
- If using any information presented with a citation, please reference the primary source.



# *Biomarker Group Updates*

## *June 17, 2022*

---

**Leadership:** Chetan Bettagowda, MD, PhD and  
Oliver Hanemann, MD

**Presenting:** R. Taylor Sundby, MD



Response Evaluation In Neurofibromatosis Schwannomatosis  
INTERNATIONAL COLLABORATION

## ***REiNS Biomarker Group Members:***

- Chetan Bettegowda
- Oliver Hanemann
- Edina Komlodi-Pasztor
- Gareth Evans
- Meena Upadhyaya
- Steven Rhodes
- Robert Soto
- Taylor Sundby
- Jaishri Blakeley
- Aerang Kim
- Onno Faber
- Vito Grasso
- Herb Sarnoff
- Krizelle Alcantara



# Current Projects

1. Centralized resource with NF1 biobanks and biorepositories
2. Manuscript with proposed common data language/model for NF1 biomarker research



# NF1 Biorepository Resource

- List of international biobanks/biorepositories with NF1/2 and SWN samples
  - Tissue types, contact information
- List of NF Biomarker publications

- 
- Currently limited to NF1, will expand to NF2, SWN
  - Currently .csv, will publish to REiNS website

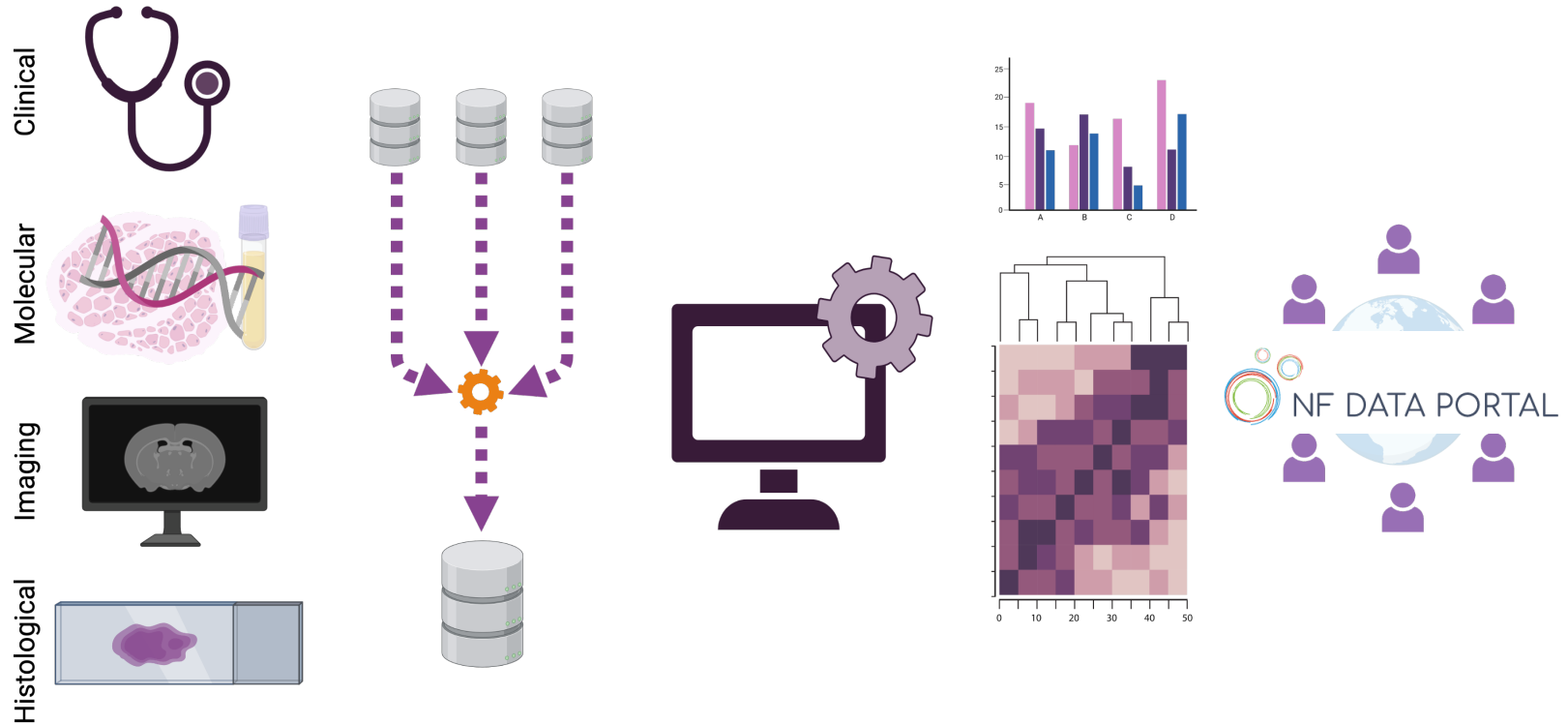


# Manuscript:

*Establishment of a common data language (CDL) and common data model (CDM) for neurofibromatosis biomarker research*



# Multi-institutional collaborations required to study rare diseases



Multi-site  
Data  
Acquisition

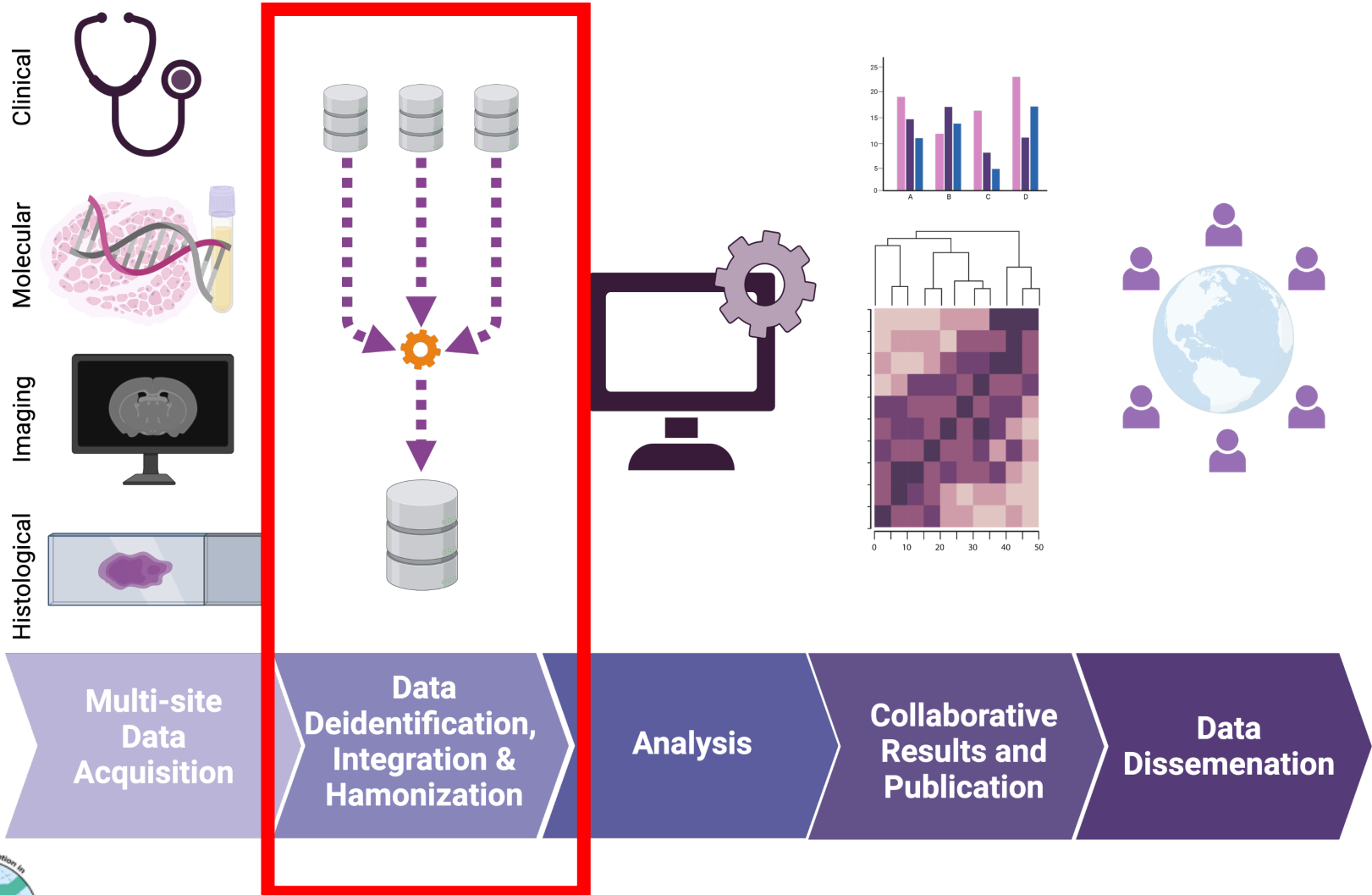
Data  
Deidentification,  
Integration &  
Hamonization

Analysis

Collaborative  
Results and  
Publication

Data  
Dissemination

# Multi-institutional collaborations required to study rare diseases





Challenge: Inter- (and intra-) institutional  
variation in sample collection and annotations  
complicate data harmonization



# Current status and recommendations for biomarkers and biobanking in neurofibromatosis

C. Oliver Hanemann, Jaishri O. Blakeley, Fabio P. Nunes, Kent Robertson, Anat Stemmer-Rachamimov, Victor Mautner, Andreas Kurtz, Michael Ferguson, Brigitte C. Widemann, D. Gareth Evans, Rosalie Ferner, Steven L. Carroll, Bruce Korf, Pierre Wolkenstein, Pamela Knight, Scott R. Plotkin, For the REINS International Collaboration

First published August 15, 2016, DOI: <https://doi.org/10.1212/WNL.0000000000002932>

## RECOMMENDATIONS FOR SAMPLE COLLECTION AND METHODOLOGY

The REINS biomarker group discussed potential barriers to biomarker research for NF. These barriers include the low prevalence of NF1, NF2, and schwannomatosis, which makes coordinated studies technically difficult and expensive; the extreme variability of these conditions requires expert clinical researchers to accurately phenotype patients. Based on this discussion, the group endorsed the following goals to advance the study of biomarkers within the NF community:

1. Build a prospective biorepository of curated samples. The aim would be to collect longitudinal samples from each patient to facilitate the development of early detection and prognostic markers.
2. Standardize tissue collection at participating institutions. The aim would be to collect all tissues using an identical protocol that meets standards set forth by the American Association for Cancer Research–Food and Drug Administration–National Cancer Institute Cancer Biomarker Collaborative<sup>4</sup> and would be linked via a shared, anonymized registry (on a Web site). Participating sites would share common consent, SOPs, quality control (see supplemental data on the *Neurology*® Web site at [Neurology.org](http://Neurology.org)), minimal clinical dataset, and database.
3. Annotate samples with an agreed minimal clinical dataset. The goal is to link anonymously the phenotypic data to deidentified samples in the biorepository. A proposed minimal clinical dataset developed at a consensus meeting in October 2014 is shown in **table 2**. The group anticipates that modifications of this dataset may be required in the future to optimize the utility of biomarker research.
4. Incorporate the decentralized biorepository into existing biorepositories that are used for diagnostic purposes. The biomaterial could thus be used for both diagnosis and research and is not necessarily restricted to an upfront definition of the amount of surplus tissue. Patient care takes preference when allocating the amount of samples used for biomarker investigations.
5. Provide open access to deposited biomarkers to facilitate research. The aim would be to have samples and data open to all qualified researchers with approval of an institutional review board. A biorepository council would govern database requests.
6. An operational and an executive committee will govern requests and audit implementation of SOPs and quality control measures.
7. Incorporate biomarker collection into clinical studies. When feasible, sample collection should be incorporated into prospective clinical trials and natural history studies to help develop pharmacodynamic and predictive biomarkers.
8. Incorporate biomarker collection into routine clinical visits. Patients receiving routine care should be invited to participate in prospective sample collection during routine clinic visits.

## RECOMMENDATIONS FOR BIOMARKERS BASED ON EXISTING DATA

1. Validate individual biomarkers as well as cocktails/signatures of biomarkers.
2. For ongoing and planned NF trials, studies of drug metabolism/pharmacodynamic biomarkers should be drug-specific.
3. For malignant tumors such as MPNST, explore and validate cDNA, cRNA, and circulating tumor cells as biomarkers.
4. Validate use of extracellular vesicles (exosomes) based on encouraging preliminary data as biomarkers of cancer.<sup>22</sup>
5. Using candidate approach, focus on the clinically relevant questions, i.e., total tumor load, presence of plexiform neurofibroma, evidence of malignant transformation, and taking into account the statistical significance in published studies. We recommend validating the following biomarker candidates in patients with NF1: BIRC5/TK1/TOP2A immunohistochemistry, ADM, interferon- $\gamma$ , IGFBP-1, and sAXL.
6. Complement candidate biomarker approach with systematic unbiased approach. Encourage well-powered studies using systematic unbiased approaches, including genomics (DNA, RNA, miRNA next-generation sequencing), metabolomics, and proteomics; that is, further screening with metabolomics, proteomics, expression arrays, and miRNA.



# Common Data Model/Language Recommendations Manuscript

1

Review:

- \* **Variations in practice**
- \* **Examples of CDM** (OMOP, mCODE)

2

**Provider Survey**

3

Recommend:

- \* **Updated minimal clinical annotations**
- \* **Updated minimal sample collection**
- \* **NF and SWM specific CDM/L**

# Part 2: Provider Survey

- Survey of NF1 Providers\* covering:
  - Awareness of 2016 guidelines
  - Variation in current practice
    - Standardized clinical data forms?

*\*Proposed audience: CTF members (discussed with Salvo and Patrice)*



**Table 2 Recommended minimal clinical dataset**

Demographics						
Date:						
Sex	<input type="checkbox"/> male	<input type="checkbox"/> female				
Birth	<input type="checkbox"/> month	<input type="checkbox"/> year				
Diagnosis of NF1, NF2, or schwannomatosis	<input type="checkbox"/> infancy	<input type="checkbox"/> childhood	<input type="checkbox"/> adolescence	<input type="checkbox"/> adulthood	<input type="checkbox"/> unknown	
Inheritance	<input type="checkbox"/> parent affected	<input type="checkbox"/> parent not affected	<input type="checkbox"/> unknown			
Mosaicism	<input type="checkbox"/> patient is mosaic	<input type="checkbox"/> patient not mosaic	<input type="checkbox"/> unknown			
Germline mutation	<input type="checkbox"/> not tested	<input type="checkbox"/> tested + unknown	<input type="checkbox"/> determined: (specify __)			
Clinical status						
Status	<input type="checkbox"/> alive	<input type="checkbox"/> deceased				
WHO performance status	<input type="checkbox"/> 0	<input type="checkbox"/> 1	<input type="checkbox"/> 2	<input type="checkbox"/> 3	<input type="checkbox"/> 4	
Pain	<input type="checkbox"/> not a problem	<input type="checkbox"/> occasional	<input type="checkbox"/> disabling			
Treatment directed at tumor	<input type="checkbox"/> no specific therapy	<input type="checkbox"/> chemotherapy	<input type="checkbox"/> surgery	<input type="checkbox"/> radiation	<input type="checkbox"/> targeted therapy (specify)	<input type="checkbox"/> clinical trial (specify)
NF1						
≥6 Café-au-lait macules	<input type="checkbox"/> absent	<input type="checkbox"/> present	<input type="checkbox"/> unknown			
Skin fold freckling	<input type="checkbox"/> absent	<input type="checkbox"/> present	<input type="checkbox"/> unknown			
Iris Lisch nodules	<input type="checkbox"/> absent	<input type="checkbox"/> present	<input type="checkbox"/> unknown			
Dermal neurofibromas	<input type="checkbox"/> absent	<input type="checkbox"/> scattered	<input type="checkbox"/> dense	<input type="checkbox"/> unknown		
Subcutaneous nodular neurofibromas	<input type="checkbox"/> absent	<input type="checkbox"/> scattered	<input type="checkbox"/> dense	<input type="checkbox"/> unknown		
Diffuse dermal neurofibromas	<input type="checkbox"/> absent	<input type="checkbox"/> scattered	<input type="checkbox"/> dense	<input type="checkbox"/> unknown		
Spinal neurofibromas	<input type="checkbox"/> not imaged	<input type="checkbox"/> absent	<input type="checkbox"/> 1-3 levels	<input type="checkbox"/> all levels (cervical, thoracic, lumbar, sacral)		<input type="checkbox"/> unknown
Plexiform neurofibromas	<input type="checkbox"/> present	<input type="checkbox"/> absent by MRI	<input type="checkbox"/> absent clinically, but no MRI	<input type="checkbox"/> unknown		
Optic glioma	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present, asymptomatic	<input type="checkbox"/> present, symptomatic, not treated		<input type="checkbox"/> present, symptomatic, treated
Heart defect	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present (specify)			
Vascular disease	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present (specify)			
Puberty onset	<input type="checkbox"/> prepubertal	<input type="checkbox"/> precocious	<input type="checkbox"/> normal	<input type="checkbox"/> late	<input type="checkbox"/> unknown	
Stature	<input type="checkbox"/> <5th centile	<input type="checkbox"/> 5th-95th centile	<input type="checkbox"/> >95th centile	<input type="checkbox"/> unknown		
Peripheral neuropathy	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present			
Aqueductal stenosis	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present			
Long bone dysplasia	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present			
Sphenoid dysplasia	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present			
Scoliosis	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present			
Intellectual disability	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present			

Continued



**Table 2** Continued

<b>NF1</b>					
Learning disability	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
Attention deficit disorder	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
Pheochromocytoma	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
Glomus tumor	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
MPNST	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
Glioma (not optic glioma)	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
GIST	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
Leukemia	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
Breast cancer	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
Other tumors	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present (specify ____)		
<b>NF2</b>					
Vestibular schwannoma	<input type="checkbox"/> MRI not done	<input type="checkbox"/> absent by MRI	<input type="checkbox"/> unilateral	<input type="checkbox"/> bilateral	<input type="checkbox"/> unknown
Meningioma	<input type="checkbox"/> MRI not done	<input type="checkbox"/> absent by MRI	<input type="checkbox"/> single	<input type="checkbox"/> multiple	<input type="checkbox"/> unknown
Glioma/ependymoma	<input type="checkbox"/> MRI not done	<input type="checkbox"/> absent by MRI	<input type="checkbox"/> present	<input type="checkbox"/> unknown	
Spinal schwannoma	<input type="checkbox"/> MRI not done	<input type="checkbox"/> absent by MRI	<input type="checkbox"/> single	<input type="checkbox"/> multiple	<input type="checkbox"/> unknown
Dermal schwannoma	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
Nonvestibular cranial schwannoma	<input type="checkbox"/> MRI not done	<input type="checkbox"/> absent by MRI	<input type="checkbox"/> present (specify)	<input type="checkbox"/> unknown	
Lenticular opacity	<input type="checkbox"/> unknown	<input type="checkbox"/> absent	<input type="checkbox"/> present		
<b>Schwannomatosis</b>					
Schwannomas (nonvestibular)	<input type="checkbox"/> only by imaging evidence	<input type="checkbox"/> 1 pathologically confirmed	<input type="checkbox"/> 2 or more, at least 1 pathologically confirmed	<input type="checkbox"/> unknown	
Number of schwannomas	<input type="checkbox"/> single	<input type="checkbox"/> scattered	<input type="checkbox"/> dense	<input type="checkbox"/> unknown	
Vestibular schwannomas	<input type="checkbox"/> not imaged	<input type="checkbox"/> absent by imaging	<input type="checkbox"/> unilateral	<input type="checkbox"/> bilateral	<input type="checkbox"/> unknown
Meningiomas	<input type="checkbox"/> not imaged	<input type="checkbox"/> absent by imaging	<input type="checkbox"/> single	<input type="checkbox"/> multiple	<input type="checkbox"/> unknown
Other schwannomatosis-related tumors (please specify)	<input type="checkbox"/> not investigated	<input type="checkbox"/> absent	<input type="checkbox"/> present	<input type="checkbox"/> unknown	

Abbreviations: GIST = gastrointestinal stromal tumors; MPNST = malignant peripheral nerve sheath tumors; NF1 = neurofibromatosis 1; NF2 = neurofibromatosis 2.



# Part 2: Provider Survey

- Survey of NF Providers\* covering:
  - Awareness of 2016 guidelines
  - Current practice
    - Standardized clinical data forms?
    - Sample types collected
    - Sample frequency
    - Imaging type collected
    - Imaging Frequency
  - General attitude regarding data sharing
  - General attitudes regarding role of biomarkers in NF



# Common Data Model/Language Recommendations Manuscript

1

Review:

- \* **Variations in practice**
- \* **Examples of CDM** (OMOP, mCODE)

2

**Provider Survey**

3

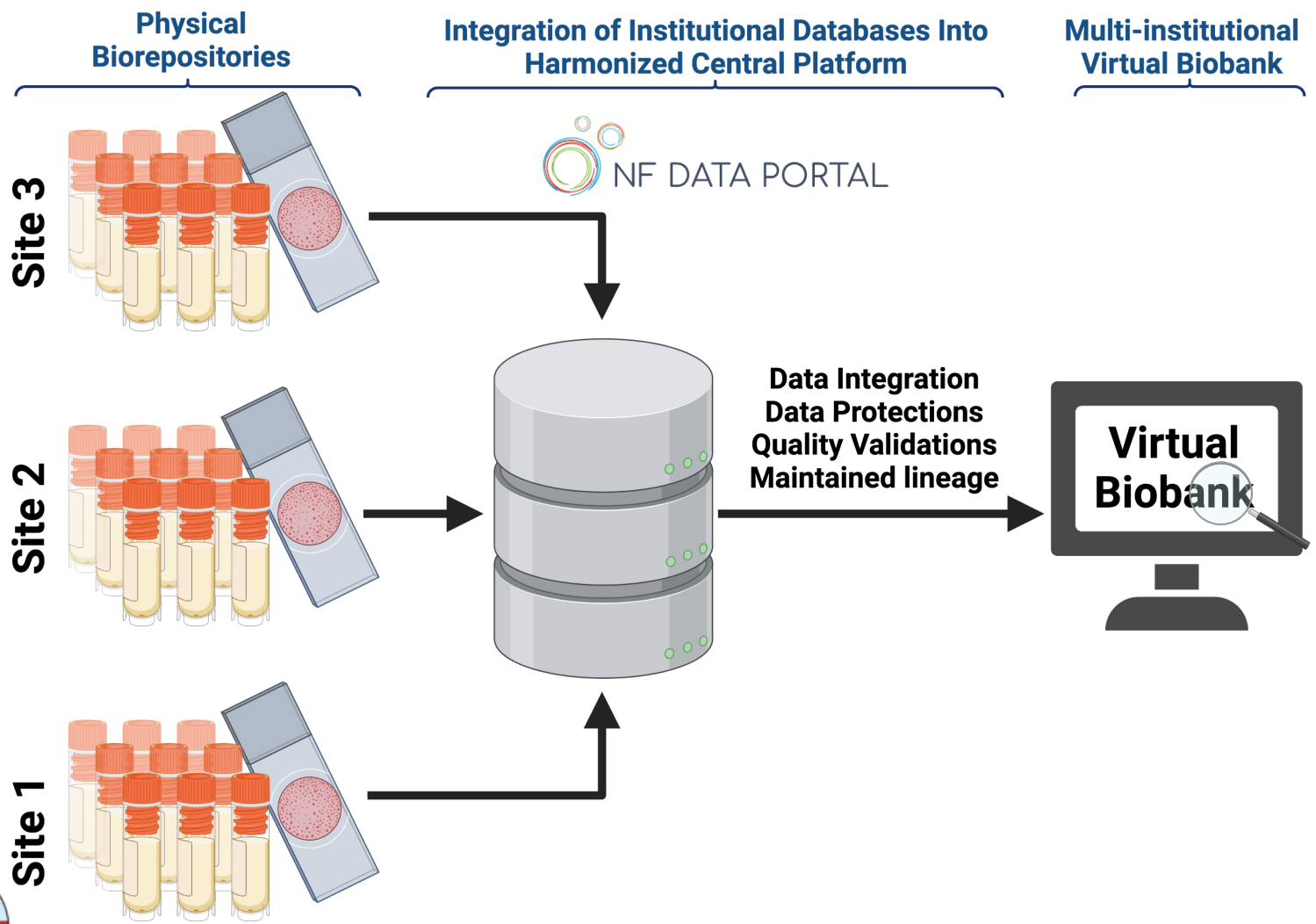
Recommend:

- \* **Updated minimal clinical annotations**
- \* **Updated minimal sample collection**
- \* **NF and SWM specific CDM/L**





# Future Directions: Establishment of a virtual biobank



# Thank you!

Contact: [Taylor.Sundby@NIH.gov](mailto:Taylor.Sundby@NIH.gov)