

# NIDDK DMS Webinar Series

## Metadata and Data Standards

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# Introduction



- Ken Young
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  - Assistant Professor
- Health Informatics Institute
- University of South Florida – Morsani College of Medicine
- TEDDY Data Coordinating Center (DCC) – Tampa, FL

# TEDDY Study

- The Environmental Determinants of Diabetes in the Young (TEDDY) study.
- Primary objective(s) of this multi-center, multi-national, epidemiological study:
  - Identification of infectious agents, dietary factors, or other environmental exposures that are associated with increased risk of autoimmunity and T1DM.
  - Factors affecting specific phenotypic manifestations such as early age of onset or rate of progression, or with protection from the development of T1DM will also be identified.
- The TEDDY study investigates:
  - Genetic and genetic-environmental interactions, including gestational infection or other gestational events.
  - Childhood infections or other environmental factors after birth in relation to the development of prediabetes autoimmunity and T1D.

# TEDDY Study Centers



# TEDDY Study Centers

- Clinical Centers:
  - Colorado Barbara Davis Center, Univ. CO, Denver, CO
  - Finland University of Turku, Turku, Finland
  - Georgia/Florida Augusta University, Augusta, GA
  - Germany Diabetes Research Institute, Munich, Germany
  - Sweden Lund University, Malmö, Sweden
  - Washington Pacific Northwest Diabetes Research Institute, Seattle, WA
- Data Coordinating Center (DCC):
  - University of South Florida Health Informatics Institute, Tampa, FL

# TEDDY Data Assets

- A variety of different data types are collected as a part of the TEDDY study, including clinical metadata and laboratory test result data across various 'omics analytes.
- The TEDDY DCC manages, curates, integrates, and provisions these data assets for analysis by TEDDY and approved external investigators.

# TEDDY Clinical Metadata

- Case-Control Indicators
- Demographics
- Diet
- Family History
- Genotypes
- Household Exposures
- Medical History
- Medications
- Physical Activity
- Pre and Perinatal Exposures
- Psychosocial Stressors
- Test Results

# TEDDY 'Omics Analytes

- Dietary Biomarkers
- Epigenetics
- Exome
- Gene Expression
- Inflammatory Biomarkers
- Lipidomics
- Metabolomics
- Microbiome and Metagenomics
- Proteomics
- RNA Sequencing
- ImmunoChip SNPs
- Urinary Biomarkers
- Whole Genome Sequencing

# TEDDY Data Standards

- The TEDDY study has implemented the following biomedical ontologies to increase interoperability:

Type of Data	Standard/Ontology
Adverse Events/Reactions	<a href="#">CTCAE v5.0</a> (Common Terminology Criteria for Adverse Events)
Diagnoses	<a href="#">WHO ICD-10</a> (International Statistical Classification of Diseases and Related Health Problems) <a href="#">UMLS SNOMED CT</a> (Systemized Nomenclature of Medicine Clinical Terms)
Medications	<a href="#">UMLS RxNorm</a>



# TEDDY Data Standards

- TEDDY has also created unique “TEDDY codes” to capture other clinical data in a standardized way.
- Collecting codes in place of free-text fields improves data accuracy and provides consistency among reported values by restricting abbreviations, synonyms, and misspellings.

**4. Allergies** Does the child have any new allergies? <sup>^</sup>

No  Yes

The child is allergic to:	Code	When did the allergy start? (Age in Months)	If the allergy has stopped, when did it stop (Age in months)?	What symptoms does the child have? Code	Recommended by a health care provider?
	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/> <input type="text"/> <input type="text"/>	<input type="radio"/> No <input type="radio"/> Yes

# TEDDY Data Dictionaries

- To ensure data are findable and reusable, each clinical data set TEDDY shares is accompanied by a data dictionary, which contains the variable name, type, length, and label.
- Data dictionaries can be provided in multiple formats including:
  - RTF
  - CSV
  - DOCX
  - XLSX

**TEDDY**  
*The Environmental Determinants of Diabetes in the Young (TEDDY) study*  
*adverse\_event\_table Form Data Dictionary*  
*File Name: ADVERSE\_EVENT\_TABLE*  
*Number of Observations: 268*  
*Number of Variables: 27*

Variable Name	Variable Type	Variable Length	Variable Label
AERPORTTYPE	Char	9	Initial or Follow-up
REFERGENETICCOUNSELING	Char	7	Was this subject referred for genetic counseling: Yes, No, Unknown
REFERPOSTPARTUMDEPRESSIONCOUNS	Char	7	Was this subject referred for postpartum depression counseling: Yes, No, Unknown
AECATEGORY	Char	31	Allergy/Immunology, Blood/Bone Marrow, Cardiac Arrhythmia, Coagulation, Death not associated with event, Dermatology/Skin, Endocrine, Infection, Musculoskeletal/Soft Tissue, Neurology, Pain, Pulmonary/Upper Respiratory, Syndrome, Vascular
AECAUSALITYBYREPORTER	Char	22	Causality by reporter: Definitely not related, Definitely related, Possibly related, Probably not related, Probably related
AEEEXPECTED	Char	3	Was the adverse event expected
AEPATIENTOUTCOME	Char	35	Fatal, Intervention for AE continues, Not recovered/Not resolved, Recovered/Resolved with sequelae, Recovered/resolved without sequelae, Recovering/resolving, Unknown
AEREASONFORFOLLOWUP	Char	28	Correction of initial report
AESERIOUS	Char	3	Was the adverse event serious

# TEDDY Data Dictionaries

- Dictionaries for omics data vary by data type.
- Examples of dictionaries include:
  - Manifest files (*Generated by Instrument Manufacturer*)
  - Annotation files (*Generated by Lab/TEDDY*)

```
1 Illumina Inc. GenomeStudio version 1.8.0
2 Normalization = none
3 Array Content = HumanHT-12_V4_0_R2_15002873_B.bgx.xml
4 Error Model = none
5 DateTime = 2/20/2013 11:07 AM
6 Local Settings = en-US
7
8 TargetID      ProbeID MIN_Signal-9234985006_A AVG_Signal-9234985006_A MAX_Signal-9234985006_A NARRAYS-923498
9 7A5 6450255 98.5 98.5 98.5 1 NaN 24.646 27 0.66234 106.0 106.0 106.0 1 NaN 35.381 16
10 A1BG 2570615 106.1 106.1 106.1 1 NaN 19.769 16 0.85844 127.9 127.9 127.9 1 NaN 40.544
11 A1BG 6370619 89.0 89.0 89.0 1 NaN 24.256 19 0.31169 106.4 106.4 106.4 1 NaN 26.084
12 A1CF 2600039 91.4 91.4 91.4 1 NaN 19.230 25 0.40000 90.6 90.6 90.6 1 NaN 28.613
13 A1CF 2650615 90.1 90.1 90.1 1 NaN 23.593 24 0.34805 98.9 98.9 98.9 1 NaN 25.446
14 A1CF 5340672 84.3 84.3 84.3 1 NaN 22.901 21 0.16883 94.3 94.3 94.3 1 NaN 23.168
15 A26C3 2000519 102.4 102.4 102.4 1 NaN 28.242 19 0.77273 97.0 97.0 97.0 1 NaN 28.426
16 A26C3 3870044 68.3 68.3 68.3 1 NaN 13.575 21 0.00390 76.4 76.4 76.4 1 NaN 25.610
17 A26C3 7050209 99.1 99.1 99.1 1 NaN 29.676 24 0.68182 115.1 115.1 115.1 1 NaN 44.332
18 A2BP1 1580181 101.9 101.9 101.9 1 NaN 24.926 19 0.76104 109.0 109.0 109.0 1 NaN 35.068
19 A2BP1 5220554 90.9 90.9 90.9 1 NaN 20.848 22 0.38312 84.1 84.1 84.1 1 NaN 22.901
20 A2BP1 5390438 83.8 83.8 83.8 1 NaN 11.051 12 0.15065 95.1 95.1 95.1 1 NaN 22.473
21 A2BP1 6420681 86.5 86.5 86.5 1 NaN 19.291 22 0.23636 94.8 94.8 94.8 1 NaN 28.595
22 A2LD1 4760377 111.3 111.3 111.3 1 NaN 39.396 25 0.91818 86.1 86.1 86.1 1 NaN 28.009
23 A2M 2370438 82.4 82.4 82.4 1 NaN 25.456 20 0.12987 71.4 71.4 71.4 1 NaN 23.945 26
```

Illumina HumanHT-12 v4.0 Manifest File

# TEDDY Data Dictionaries

- Dictionaries for omics data vary by data type.
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  - Manifest files (*Generated by Instrument Manufacturer*)
  - Annotation files (*Generated by Lab/TEDDY*)

Lipid	Retention Index (RI)	Mass to Charge Ratio (MZ)	InChIKey
1_CUDA iSTD [M-H]-	45	339.2653863	HPTJABJPZMULFH-UHFFFAOYSA-N
1_Ceramide (d18:1/17:0) iSTD [M+Cl]-	363	586.4960498	ICWGMOFDULMCFL-QKSCFGQVSA-N
1_Ceramide (d18:1/17:0) iSTD [M+FA-H]-	363	596.5260169	ICWGMOFDULMCFL-QKSCFGQVSA-N
1_FA iSTD (16:0)-d3 [M-H]-	189	258.2521035	IPCSVZSSVZVIGE-FIBGUPNXSA-N
1_LPC (17:0) iSTD [M+FA-H]-	108.6	554.3459919	SRRQPVVYXBTRQK-XMMPXPASA-N
1_LPE (17:1) iSTD [M-H]-	77.4	464.2774753	LNJNONCNASQZOB-HEDKFQSOSA-N
1_MAG (17:0/0:0/0:0) iSTD [M+FA-H]-	183.6	389.29085	SVUQHVRAGMNPWL-UHFFFAOYSA-N
1_PC (12:0/13:0) iSTD [M+FA-H]-	214.2	680.447887	FCTBVSCBBWKZML-WJOKGBTCSA-N
1_PE (17:0/17:0) iSTD [M-H]-	380.4	718.538132	YSFFAUPDXKTJMR-DIPNUNPCSA-N
1_PG (17:0/17:0) iSTD [M-H]-	336.6	749.5339987	ZBVHXVKEMAIWQQ-QPPIDDCLSA-N
1_SM (d18:1/17:0) iSTD [M+FA-H]-	309.6	761.5789623	YMQZQHIESOAPQH-JXGHDCMNSA-N
Ceramide (d32:1) [M+Cl]-	301.2	544.4506	ZKRPGPZHULJKJ-JHRQRACZSA-N
Ceramide (d32:1) [M+FA-H]-	301.8	554.47895	ZKRPGPZHULJKJ-JHRQRACZSA-N
Ceramide (d33:1) [M+Cl]-	322.2	558.4656	QBFXCLDNTKBAPQ-STSAHMJASA-N
Ceramide (d33:1) [M+FA-H]-	321.6	568.49475	QBFXCLDNTKBAPQ-STSAHMJASA-N
Ceramide (d34:0) [M+Cl]-	355.8	574.4973	GCGTXOVNFGTPQ-JHOUSYSJSA-N
Ceramide (d34:0) [M+FA-H]-	356.4	584.52605	GCGTXOVNFGTPQ-JHOUSYSJSA-N
Ceramide (d34:1) [M+Cl]-	342.6	572.4816	YDNKGFDDKKRUKPY-TURZORIXSA-N

# TEDDY Release Notes

- Direct to Investigator Data Releases also receive release notes describing the data freeze date, population, datasets provided, and any relevant notes for the investigator

**v1** **Release\_Notes.txt**  
TEDDY-MP037 · Updated Mar 6, 2023 by Michael Toth

MP37 - Islet Autoantibody Levels from Seroconversion to Time of Type 1 Diabetes Diagnosis

Project: TEDDY  
Release Name: TEDDY-MP037  
Freeze Date: July 31,2022  
Population: TEDDY subjects who are positive for any autoantibody (GADA MIAA IA2A or ZntA8)

Release 1: 2015-03-06 (Released on LabKey)  
- mp37\_gad\_longitudinal.csv  
- mp37\_ia2a\_longitudinal.csv  
- mp37\_miaa\_longitudinal.csv

Release 2: 2023-01-17  
- Child\_Life\_Experiences\_Category\_Placement\_Mar\_25\_2020.docx  
- MP37\_Data Dictionary Additional\_vars\_masked.docx  
- MP37\_KVehik\_NIDDK\_Data Dictionary\_31May2012\_1.rtf  
- MP37\_KVehik\_NIDDK\_Data Dictionary\_31May2012\_2.rtf

# TEDDY Omics Metadata

- TEDDY omics metadata is also shared with data repositories, such as dbGaP and Metabolomics Workbench.

**SRX7944152: Human nasal 16S rRNA: Sample M6820224**

1 ILLUMINA (Illumina MiSeq) run: 1,232 spots, 616,000 bases, 488,711b downloads

**Design:** Total genomic DNA was extracted using the PowerMag Microbiome DNA isolation kit. The V4 region of the 16S rRNA gene was amplified by PCR using modified 515F and R806 primers that additionally contained adapters for MiSeq sequencing and a single 12-mer index barcode (R). Individual libraries constructed from each sample were pooled and sequenced on the Illumina MiSeq platform using the 2x250 bp paired-end read protocol.

**Submitted by:** NIDDK-phs001442

**Study:** The Environmental Determinants of Diabetes in the Young (TEDDY) Parent Study

[PRJNA416160](#) • [SRP235574](#) • [All experiments](#) • [All runs](#)

[show Abstract](#)

**Sample:** DNA/RNA sample from Nasal of a human male participant in the dbGaP study "The Environmental Determinants of Diabetes in the Young Study (TEDDY)"

[SAMN13458218](#) • [SRS5799084](#) • [All experiments](#) • [All runs](#)

**Organism:** [Homo sapiens](#)

**Library:**

**Name:** 221141

**Instrument:** Illumina MiSeq

**Strategy:** AMPLICON

**Source:** METAGENOMIC

**Selection:** PCR

**Layout:** PAIRED

# TEDDY Annotated Forms

- TEDDY eCRFs annotated with the data set name at the top and variable name by each field have been shared with the NIDDK Central Repository as a searchable PDF.
- Annotated forms allow investigators to find data of interest, see how it was collected, and identify related variables.

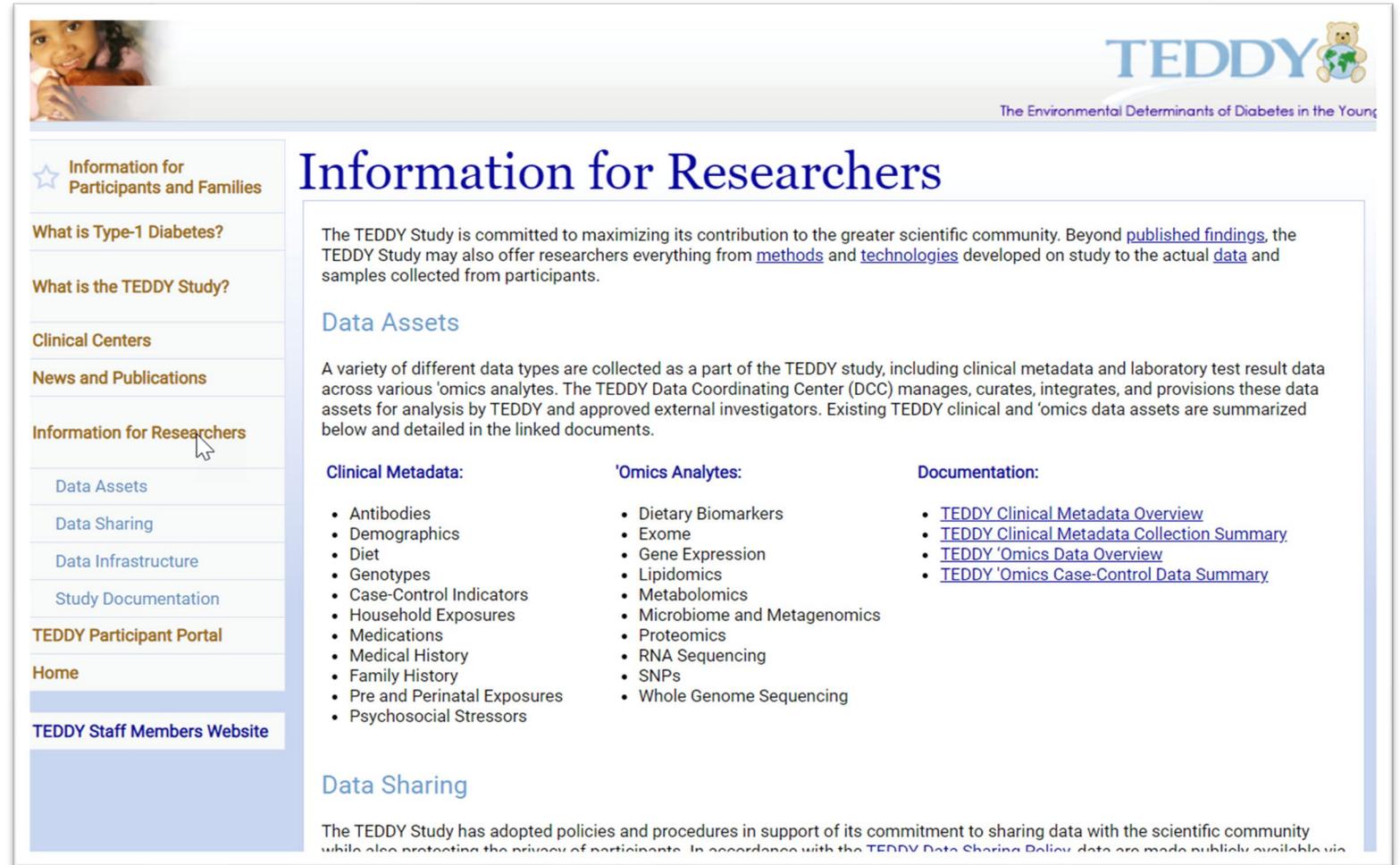
**7a. Acute Illnesses** - Has the child been ill since the last visit? Record all chronic illnesses/conditions on the next page.

No  Yes 2582 ChildIllSinceLastVisit

Date Illness first appeared	ICD-10 Code: ONLY code <u>Symptoms</u> here (ALWAYS is equal to or higher than 38°C or 101°F)	Fever? (temperature higher than 38°C or 101°F)	Diagnosis: ICD-10 Code
<input type="text" value="2583"/> <input type="text" value="2584"/>	<span style="color: red;">Illness Symptoms Codes1_1</span> <input type="text" value="2586"/> <input type="text" value="2587"/> <input type="text" value="2588"/>	<input type="radio"/> No 2739 <input type="radio"/> Yes, Measured <input type="radio"/> Yes, Not Measured	<input type="text" value="2741"/> <span style="color: red;">IllnessDiagnosisCode1_1</span> <input type="radio"/> Diagnosed by parent <input type="radio"/> Diagnosed by health care provider 2740
<input type="text" value="2585"/> <span style="color: red;">DateIllnessAppearedAge1_1</span>	<input type="checkbox"/> No Symptoms 4181 <span style="color: red;">IllnessNoSymptoms1_1</span>	<span style="color: red;">IllnessFever1_1</span>	<span style="color: red;">IllnessDiagnosis1_1</span>

# TEDDY Data Documentation

- On the [TEDDY public website](#), investigators can find documents detailing TEDDY data collection procedures (e.g., protocol, MOO, NCC Design), data availability, and data sharing policy.
- Data documentation available on the public website and across repositories has been developed to make TEDDY data more FAIR.



**Information for Researchers**

The TEDDY Study is committed to maximizing its contribution to the greater scientific community. Beyond [published findings](#), the TEDDY Study may also offer researchers everything from [methods](#) and [technologies](#) developed on study to the actual [data](#) and samples collected from participants.

### Data Assets

A variety of different data types are collected as a part of the TEDDY study, including clinical metadata and laboratory test result data across various 'omics analytes. The TEDDY Data Coordinating Center (DCC) manages, curates, integrates, and provisions these data assets for analysis by TEDDY and approved external investigators. Existing TEDDY clinical and 'omics data assets are summarized below and detailed in the linked documents.

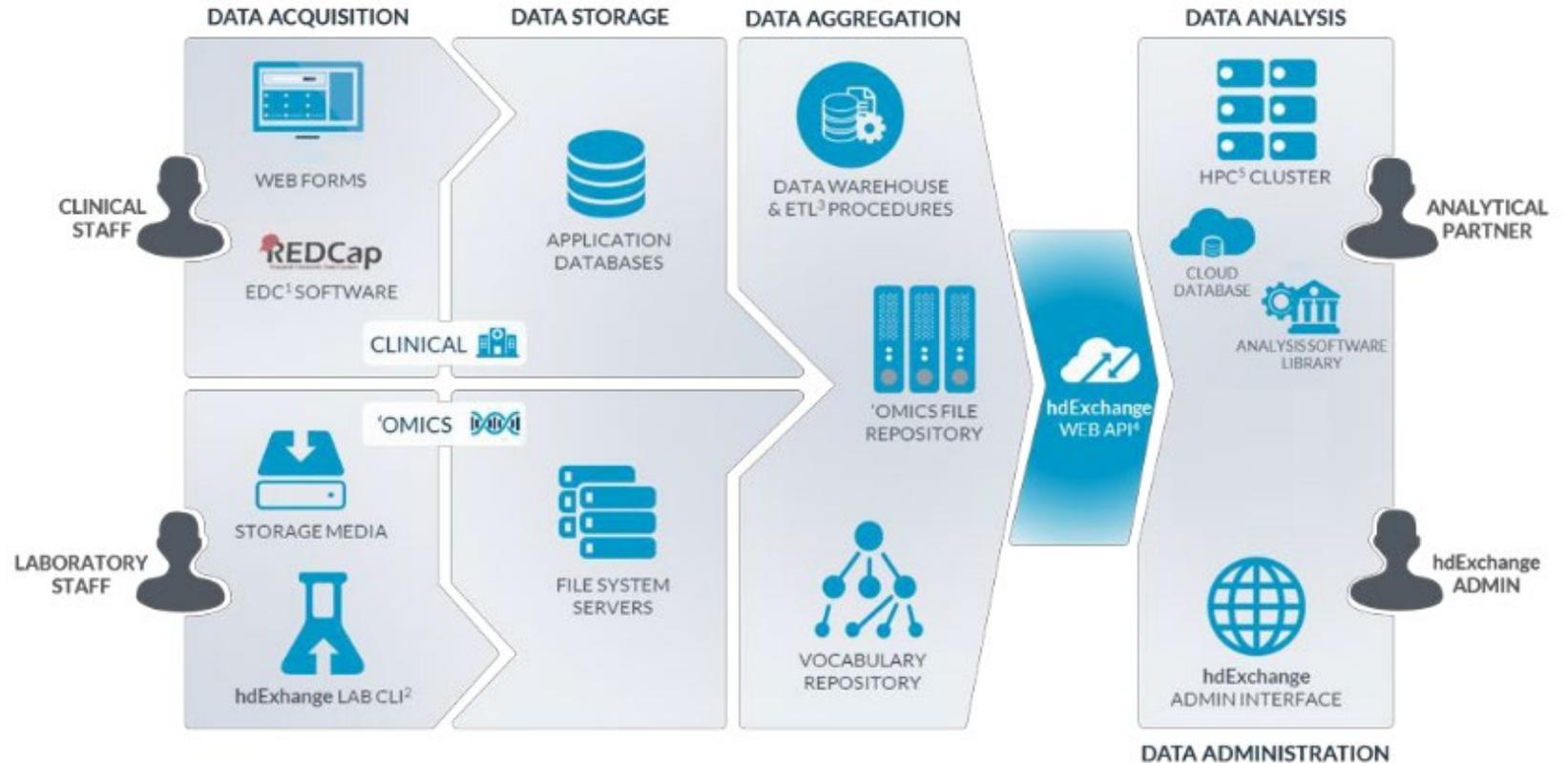
<b>Clinical Metadata:</b>	<b>'Omics Analytes:</b>	<b>Documentation:</b>
<ul style="list-style-type: none"><li>• Antibodies</li><li>• Demographics</li><li>• Diet</li><li>• Genotypes</li><li>• Case-Control Indicators</li><li>• Household Exposures</li><li>• Medications</li><li>• Medical History</li><li>• Family History</li><li>• Pre and Perinatal Exposures</li><li>• Psychosocial Stressors</li></ul>	<ul style="list-style-type: none"><li>• Dietary Biomarkers</li><li>• Exome</li><li>• Gene Expression</li><li>• Lipidomics</li><li>• Metabolomics</li><li>• Microbiome and Metagenomics</li><li>• Proteomics</li><li>• RNA Sequencing</li><li>• SNPs</li><li>• Whole Genome Sequencing</li></ul>	<ul style="list-style-type: none"><li>• <a href="#">TEDDY Clinical Metadata Overview</a></li><li>• <a href="#">TEDDY Clinical Metadata Collection Summary</a></li><li>• <a href="#">TEDDY 'Omics Data Overview</a></li><li>• <a href="#">TEDDY 'Omics Case-Control Data Summary</a></li></ul>

### Data Sharing

The TEDDY Study has adopted policies and procedures in support of its commitment to sharing data with the scientific community while also protecting the privacy of participants. In accordance with the TEDDY Data Sharing Policy, data are made publicly available via

# HII Data Infrastructure

- As TEDDY DCC, HII developed infrastructure to support large scale data and analysis.



# TEDDY Data Sharing

- The TEDDY Study has adopted policies and procedures in support of its commitment to sharing data with the scientific community while also protecting the privacy of participants.
- Data releases have been submitted at different time points and to various repositories, depending on NIH requirements and the nature of the data.
- Each submission is treated as an independent release, possessing uniquely masked subject and sample identifiers.
- Researchers may desire to combine data across these releases for analysis but are unable to do so as a result of the independently masked identifiers. The NIDDK repository can provide repository data release identifier mapping materials to satisfy this demand once the investigators have received approval to access the data.

# Acknowledgements

- Health Informatics Institute at the University of South Florida (USF)
  - Dr. Jeffrey Krischer
  - Dena Tewey
  - Chris Shaffer
- TEDDY Study Group:
  - NIH, TEDDY DCC, clinical center investigators, clinical center staff, and TEDDY study participants.
- NIDDK

