



**FIND**   
Diagnosis for all

Photo: FIND/Katie G.Nelson

# REGISTRATION GUIDE

## FIND VIRTUAL BIOBANK DIRECTORY

CONTACT US: [vbd@finddx.org](mailto:vbd@finddx.org) VBD WEBSITE: <https://vbd.finddx.org> FIND WEBSITE: <http://finddx.org>

# INTRODUCTION

The Foundation for Innovative New Diagnostics (FIND) Virtual Biobank Directory (VBD) aims to give visibility to specimen collections hosted by other organizations or networks worldwide. This is to encourage and support commercial and academic researchers in the development and evaluation of new and existing diagnostic tests for infectious diseases in low-middle-income countries.

## HOW TO REGISTER:

This guide has been designed to assist you in how to register for the VBD. We have included both an overview of what to expect during the process and more detailed information for the different stages in the process. If you need more assistance, don't hesitate to contact us at [vbd@finddx.org](mailto:vbd@finddx.org)

## OVERVIEW:

1

### CHOOSE TO REGISTER AS:

A new biobank OR a new network of resources

*(1) What's the difference?*

2

### FILL IN THE PRELIMINARY INFORMATION:

Biobank/Network name, Admin name

Admin email

*You will receive a confirmation email*

3

### CONFIRM REGISTRATION:

Use this link in the confirmation email to complete your registration and create the profile of your biobank/network

4

### EDIT BIOBANK OR NETWORK GENERAL DETAILS:

Description

Funders

Contact email

Collections and the sample set details

Address details

Capabilities for prospective collections

Institution

Network acceptance

Services offered

Annual stats

The admins

*(1) A **biobank** is any infrastructure that holds or can collect and distribute human samples and data (e.g. biobanks, bioresources, biorepositories, cohorts and clinical trials). A **network** is a group of biobanks that have come together with some common objective or agreed standard. If you are unsure about whether to register as a biobank or network, [please contact us.](#)*

# MANDATORY FIELDS

Some fields in the registration process are mandatory. You may want to collect this information before beginning the registration process:

## BIOBANK DETAILS

- Name
- Description
- Contact email
- Address
- Institution

## COLLECTION

- Disease status
- Year started
- Access condition
- Collection status
- Collection point
- Consent restriction

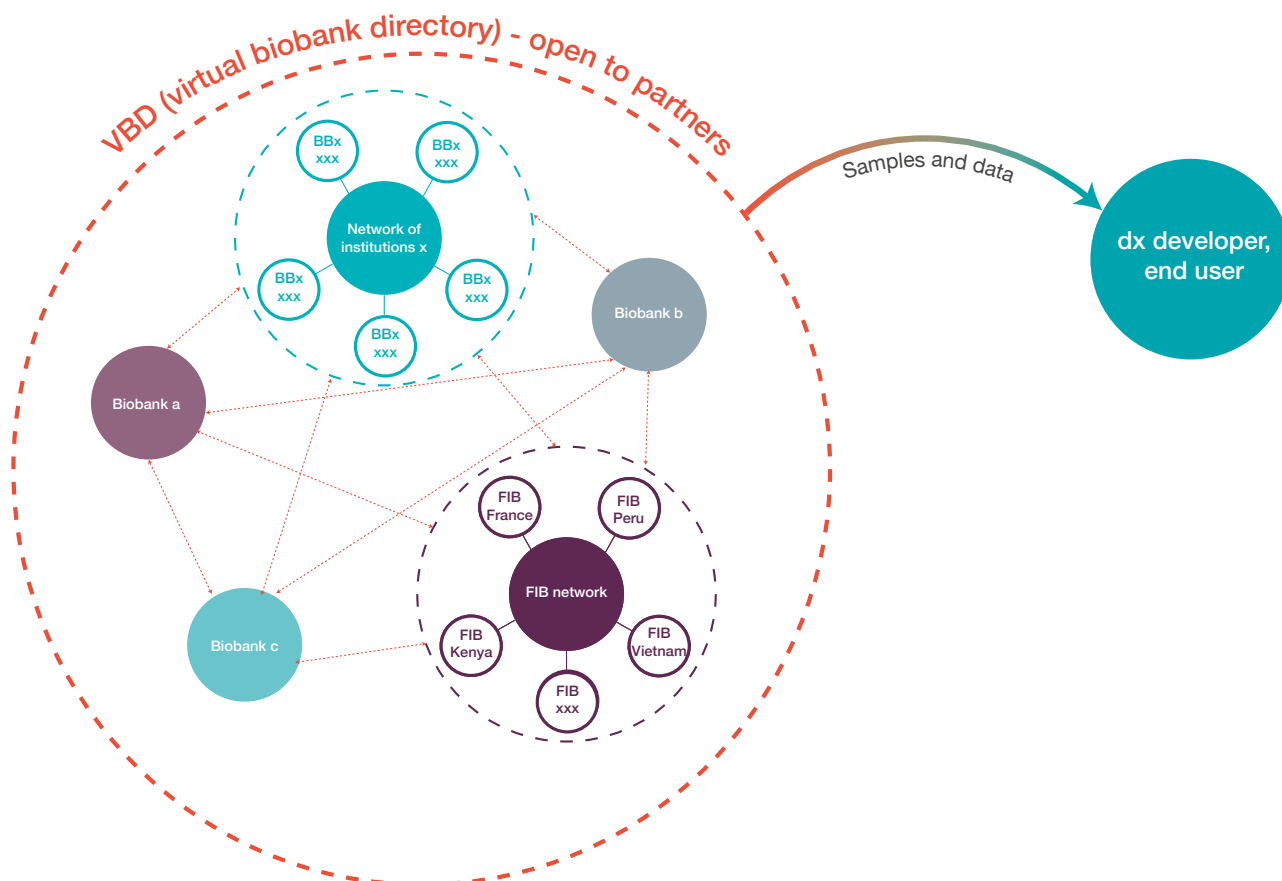
## SAMPLE SET

- Sex
- Age range
- Number of donors

## MATERIAL PRESERVATION DETAILS

- Material type
- Storage temperature
- Microscopic assessment

# VIRTUAL BIOBANK DIRECTORY INFRASTRUCTURE:

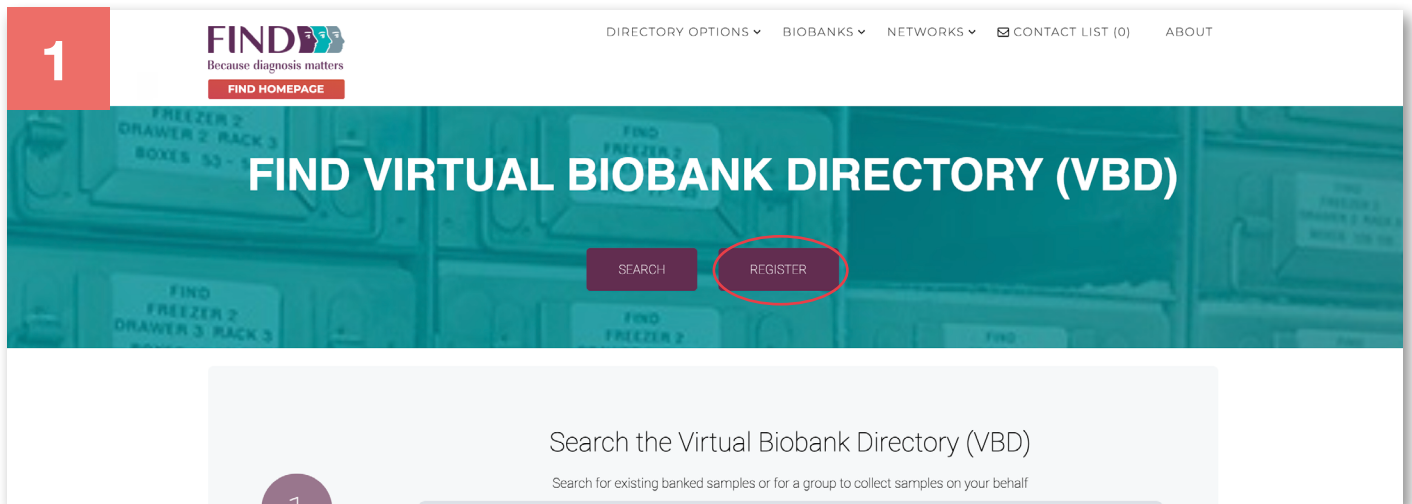


# STEP ONE:

## CREATE AN ACCOUNT

Visit: <https://vbd.finddx.org/>

Click: Register



➤ You can register as a biobank or a network by clicking on the corresponding item.

## 2

### REGISTER A BIOBANK

A biobank is any infrastructure that holds or can collect and distribute human samples and data (e.g. biobanks, bioresources, biorepositories, cohorts and clinical trials).

For more information on registering please click [here](#).

[REGISTER A NEW BIOBANK](#)

### REGISTER A NETWORK

A network is a group of biobanks that have come together with some common objective or agreed standard.

If you are unsure about whether to register a biobank or network then please [contact us](#).

[REGISTER A NEW NETWORK OF RESOURCES](#)

➤ You will be redirected to a new page to fill in preliminary information.

## Register a new biobank

If you have a collection of biological fluid samples or the ability to collect samples for diagnostic evaluations then you can add an overview to our database by registering below.

|   |   |
|---|---|
| Biobank name *                          | <input type="text" value="My Biobank"/>             |
| Admin name *                            | <input type="text" value="John Smith"/>             |
| Admin email *                           | <input type="text" value="john.smith@example.com"/> |
| <input type="button" value="Register"/> |   |

- Add the name of your biobank or network.
- Add the admin name. This is the person responsible for managing the biobank/network information.
- Add an email for the admin. It is essential that you have a direct access to this email account as it will be used for further registration steps and login to the FIND VBD.
- Later in the registration process you will have the possibility to add additional email addresses associated with your biobank/network.
- A validation link will be emailed to you.
- Click on the link to validate your account.

# STEP TWO:

## CREATE A PROFILE

Complete general information about your Biobank.  
Mandatory information is indicated by a red asterisk.

**1** [Home](#) / [FIB UCT](#) / [Sample resource details](#) / [Edit Biobank details](#)

### Edit Biobank details

Name \* ⓘ

Description \* ⓘ

URL ⓘ

Contact email \* ⓘ

Contact phone number ⓘ

Logo

## No Logo

Address Details ⓘ

Address \*

City \*

Country \*

Postcode \*

## Governance Compliance

Institution \* 

School / Department

Ethics Committee Approval 

## Services

- Infectious strain isolation
- Ability to recontact
- Access to the full pathology archive
- Cell culture
- Data analytics
- Digital imaging
- Molecular Extractions
- Immunohistochemistry - scoring
- Immunohistochemistry - staining
- Nucleic acid extraction

## Services

- Infectious strain isolation
- Ability to recontact
- Access to the full pathology archive
- Cell culture
- Data analytics
- Digital imaging
- Molecular Extractions
- Immunohistochemistry - scoring
- Immunohistochemistry - staining
- Nucleic acid extraction

## Data sharing settings

- Opt out of sharing data with other directories

## Reasons for Registering

- Condition for local ethics approval
- Condition for funding
- Requirement set by host institution
- Desire to make sample collections visible
- Desire to make collection service visible

Other

Cancel

Update details

# STEP THREE :

## ADD COLLECTIONS

You can add several collections to your biobank/network. For each collection added, complete the corresponding information as shown below.

**1**

Home / FIB UCT / Collections

Biobank Details Admins Funders Collections Capabilities Network Acceptance Annual Stats

**COLLECTIONS**

[Add collection](#)

*There are no collections here yet. A collection is the highest level of detail. [Add a collection now.](#)*

➤ Click on “add collection”

**2**

Home / FIB UCT / Collections / Add collection

### Add collection

**⚠** We expect you to ensure that this information can be shared in the public domain.

Disease status **?** \*

Title **?**

Description **?**

Year started **?** \*

Access condition **?** \*

|  |  |
|--|--|
| <input type="radio"/> Open to applicants                 | <input type="radio"/> Data access only             |
| <input type="radio"/> Open in response to specific calls | <input type="radio"/> Access restricted at present |
| <input type="radio"/> Open only through collaboration    | <input type="radio"/> Closed to access             |

Collection type **?**

|  |  |
|--|--|
| <input type="radio"/> Disease specific | <input type="radio"/> Longitudinal     |
| <input type="radio"/> Case-control     | <input type="radio"/> Population-based |
| <input type="radio"/> Cohort           | <input type="radio"/> Quality control  |
| <input type="radio"/> Cross-sectional  | <input type="radio"/> Clinical Trial   |

Collection status **?** \*

Not-started

In progress

Completed

Collection point **?** \*

|  |  |
|--|--|
| <input type="radio"/> Pre-diagnosis    | <input type="radio"/> Post-diagnosis                     |
| <input type="radio"/> During diagnosis | <input type="radio"/> Multiple points in patient pathway |



Consent restriction ⓘ \*

- Genetic analysis restriction
- Human genetic analysis restriction
- Scope of use restriction
- Export restriction
- Disease area restriction
- Project specific restriction
- Commercial restriction

Associated data ⓘ

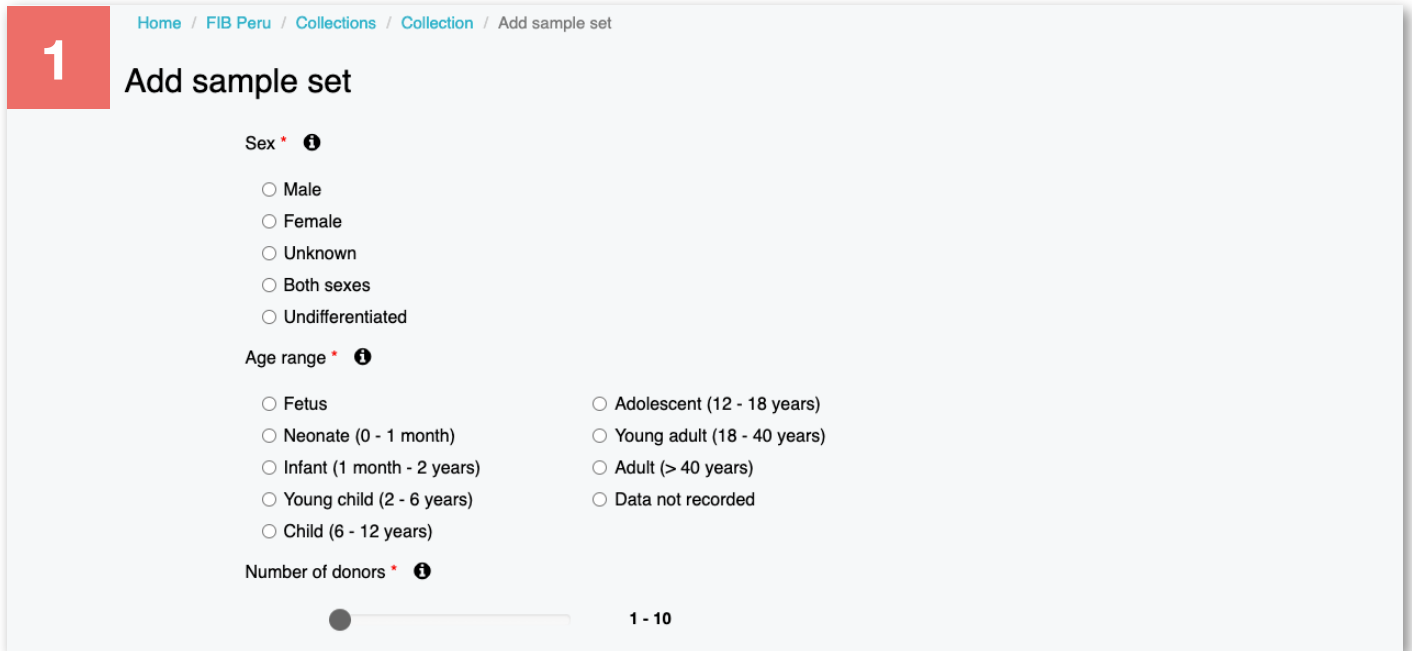
| Data type   | Provision time (months)  |
|---|--|
| Quality Data  |  |
| ⓘ <input type="checkbox"/> Freezer temperature logs               | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Quality indicators                     | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| Research Data   |  |
| ⓘ <input type="checkbox"/> Biomarker datasets                     | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Genomic datasets                       | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| Annotation Data   |  |
| ⓘ <input type="checkbox"/> Clinical records                       | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Followup records                       | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Genealogical records                   | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Imaging data                           | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> National registries                    | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Pre-analytical data                    | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Physiological/biochemical measurements | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Primary care records                   | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Survey data                            | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Treatment records                      | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Psychological data                     | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Participant Ethnicity                  | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| ⓘ <input type="checkbox"/> Microbiological results                | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |

[Cancel](#) [Create collection](#)

# STEP FOUR:

## ADD SAMPLE SET DETAILS

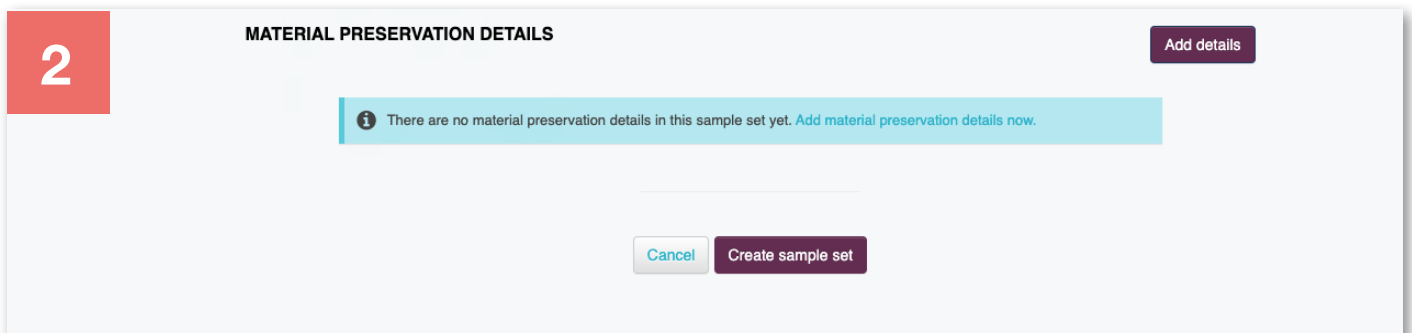
➤ For each collection, add and describe the sample sets.



The screenshot shows a web application interface for adding sample set details. At the top, there is a breadcrumb trail: Home / FIB Peru / Collections / Collection / Add sample set. A large red square with the number '1' is on the left. The main heading is 'Add sample set'. Below this, there are three sections of form fields:

- Sex \*** (with an information icon):
  - Male
  - Female
  - Unknown
  - Both sexes
  - Undifferentiated
- Age range \*** (with an information icon):
  - Fetus
  - Neonate (0 - 1 month)
  - Infant (1 month - 2 years)
  - Young child (2 - 6 years)
  - Child (6 - 12 years)
  - Adolescent (12 - 18 years)
  - Young adult (18 - 40 years)
  - Adult (> 40 years)
  - Data not recorded
- Number of donors \*** (with an information icon):
  - A horizontal slider bar with a dark grey circle on the left and a line extending to the right.
  - The value '1 - 10' is displayed to the right of the slider.

➤ You should also give details about the preservation of the samples, by clicking on the “Add details” button.



The screenshot shows a web application interface for material preservation details. At the top, there is a heading 'MATERIAL PRESERVATION DETAILS' and a dark purple button labeled 'Add details'. Below this, there is a light blue information box with an information icon and the text: 'There are no material preservation details in this sample set yet. Add material preservation details now.' At the bottom, there are two buttons: a light blue 'Cancel' button and a dark purple 'Create sample set' button.

➤ A new window will pop up, where you will be asked to fill the material preservation details (i.e. Material Type, Storage Temperature, Microscopic Assessment).

## Add sample set

### Sex \*

- Male
- Female
- Unknown
- Both sexes
- Undifferentiated

### Age range \*

- Fetus
- Neonate (0 - 1 mo)
- Infant (1 month - 2 years)
- Young child (2 - 6 years)
- Child (6 - 12 years)

### Number of donors \*



### MATERIAL PRESERVATION

There

### Add material preservation detail

Please answer the following questions for each material type available for this collection. To facilitate data entry, you can copy your current answers for re-use.

#### Material type \*

Please select the type of sample you have stored; if is not there please get in touch.

- |  |  |
|--|--|
| <input type="radio"/> Serum  | <input type="radio"/> Amniotic fluid   |
| <input type="radio"/> Bronchoalveolar lavage                               | <input type="radio"/> Blood (whole)  |
| <input type="radio"/> Breast milk  | <input type="radio"/> Buccal cells   |
| <input type="radio"/> Buffy coat   | <input type="radio"/> Cells from non-blood specimen type (non-viable)          |
| <input type="radio"/> Cells from non-blood specimen type (viable)          | <input type="radio"/> Cells from fine needle aspirate                          |
| <input type="radio"/> Cord blood   | <input type="radio"/> Cerebrospinal fluid                                      |
| <input type="radio"/> Disrupted tissue (non viable)                        | <input type="radio"/> Dried whole blood  |
| <input type="radio"/> Nasal washing  | <input type="radio"/> Isolated peripheral blood mononuclear cells (non viable) |
| <input type="radio"/> Isolated peripheral blood mononuclear cells (viable) | <input type="radio"/> Nails  |
| <input type="radio"/> Plasma (single spun)                                 | <input type="radio"/> Placenta   |
| <input type="radio"/> Red Blood Cells                                      | <input type="radio"/> Plasma (double spun)                                     |
| <input type="radio"/> Semen  | <input type="radio"/> Saliva   |
| <input type="radio"/> Sputum   | <input type="radio"/> Solid tissue (non viable)                                |
| <input type="radio"/> Swab medium  | <input type="radio"/> Stool  |
| <input type="radio"/> Urine (first morning)                                | <input type="radio"/> Urine (random)   |
| <input type="radio"/> Isolated pathogens (Derivative)                      | <input type="radio"/> Other  |
| <input type="radio"/> Host cDNA (Derivative)                               | <input type="radio"/> Host genomic DNA (Derivative)                            |
| <input type="radio"/> Pathogen DNA (Derivative)                            | <input type="radio"/> Host ccfRNA (Derivative)                                 |
| <input type="radio"/> Pathogen protein extracts (Derivative)               | <input type="radio"/> Pathogen RNA (Derivative)                                |
|  | <input type="radio"/> Cell lines (Derivative)                                  |

#### Storage Temperature \*

Select how the samples are currently stored. If there is something missing please contact us.

- |                                      |                                       |
|--------------------------------------|---------------------------------------|
| <input type="radio"/> RT             | <input type="radio"/> -85°C to -60°C  |
| <input type="radio"/> -35°C to -18°C | <input type="radio"/> <-135°C         |
|                                      | <input type="radio"/> Liquid nitrogen |

#### Microscopic Assessment \*

For samples examined by a pathologist. Please select if the sample type selected above contains (affected) or does not contain (non-affected) the disease referenced in this collection. If the sample is not examined by a pathologist, please mark N/A.

- Affected
- Non-affected
- Not applicable

# STEP FIVE:

## ADD CAPABILITIES

Complete information about your ability to prospectively collect samples, based on disease, consent, how many sample donors you expect approximately at a usual timeframe of a collection, other associated data and the amount of time necessary to provide that data.

Home / FIB Peru / Capabilities

Biobank Details Admins Funders Collections Capabilities Network Acceptance Annual Stats

### CAPABILITIES

Add capability

There are no capabilities here yet. Add a capability now

➤ Fill the requested information at the Add capability step.

Home / FIB Peru / Capabilities / Add capability

### Add capability

Disease status ⓘ \*

Protocols ⓘ  Bespoke consent form  
 Bespoke SOP

Annual donor expectation ⓘ \* e.g. 120

Associated data ⓘ

| Data type   | Provision time (months)  |
|---|--|
| Quality Data                                      |  |
| <input type="checkbox"/> Freezer temperature logs | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| <input type="checkbox"/> Quality indicators       | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| Research Data                                     |  |
| <input type="checkbox"/> Biomarker datasets       | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |
| <input type="checkbox"/> Genomic datasets         | <input type="radio"/> Immediate <input type="radio"/> 0-3 <input type="radio"/> 3-6 <input type="radio"/> >6 |

Annotation Data

- Clinical records  Immediate  0-3  3-6  >6
- Followup records  Immediate  0-3  3-6  >6
- Genealogical records  Immediate  0-3  3-6  >6
- Imaging data  Immediate  0-3  3-6  >6
- National registries  Immediate  0-3  3-6  >6
- Pre-analytical data  Immediate  0-3  3-6  >6
- Physiological/biochemical measurements  Immediate  0-3  3-6  >6
- Primary care records  Immediate  0-3  3-6  >6
- Survey data  Immediate  0-3  3-6  >6
- Treatment records  Immediate  0-3  3-6  >6
- Psychological data  Immediate  0-3  3-6  >6
- Participant Ethnicity  Immediate  0-3  3-6  >6
- Microbiological results  Immediate  0-3  3-6  >6

REFERENCE GUIDE TO:  
**FIELD DESCRIPTIONS**

| BIOBANK DETAIL   | ADDITIONAL INFORMATION  | EXAMPLE  |
|--|---|--|
| Name   | What is the name of the resource you are registering? It is up to you to present your biobank the way you want.                                       | Tuberculosis biobank   |
| Description  | Use this for general information that is not captured in other fields. You can use as much space as needed.   | A collection of human bodily fluids from Tuberculosis positive patients. |
| URL  | If your biobank has a website or page you can include it here.  | www.biobanktb.com  |
| Contact email  | The email address that people will use to contact you.  | contact@biobanktb.com  |
| Contact phone number   | The phone number that people will use to contact you.   | Tel: country code, city code, number                                     |
| Logo   | If your biobank has a logo you can put it here and it will be displayed on the FIND VBD website.  |  |
| Address  | Use the address where access to the samples is based rather than where they are stored.   | 9 Chemin des Mines, 1202, Geneva, Switzerland                            |
| Institution  | The organization where your governance is managed.  | Foundation of Innovative New Diagnostics (FIND)                          |
| Ethics Committee Approval  |   |  |
| Services <ul style="list-style-type: none"> <li>• Infectious strain isolation</li> <li>• Ability to recontact</li> <li>• Access to the full pathology archive</li> <li>• Cell culture</li> <li>• Data analytics</li> <li>• Digital imaging</li> <li>• Molecular extractions</li> <li>• Immunohistochemistry - scoring</li> <li>• Immunohistochemistry - staining</li> <li>• Nucleic acid extraction</li> </ul> | Check if your collections are approved by an Ethics Committee (i.e. IRB)<br><br>Does your biobank offer any extra services? If so indicate that here. |  |
| Reasons for registering <ul style="list-style-type: none"> <li>• Conditions for local ethics approval</li> <li>• Condition for funding</li> <li>• Requirement set by host institution</li> <li>• Desire to make sample collections visible</li> <li>• Desire to make collection service visible</li> </ul>   | Select the reasons for registering your sample resource with the directory. You may select multiple reasons.  |  |
| Other  | If there is any other useful information you would like to communicate you can put it here.   |  |

| ADD COLLECTION   | ADDITIONAL INFORMATION   | EXAMPLE   |
|--|--|---|
| Disease status   | We are seeking to harmonize the terms used to describe diseases and have adopted <a href="#">SNOMED CT</a> in this effort. We are seeking to discuss with the domain experts what terms are most relevant for certain diseases. Therefore, if you do not see a relevant term presented on the form, please contact us. If you are collecting from healthy volunteers, use the term 'Fit and Well'. | Tuberculosis  |
| Title  | This is an optional field in case your collection has a specific title.  | TB in country X   |
| Description  | Add a description of your collection.  | Sputum samples from <i>Mycobacterium tuberculosis</i> sputum positive patients from country X |
| Year started   | The year when the first samples were collected.  | 2012  |
| Access condition:<br><ul style="list-style-type: none"> <li>• Open to applicants</li> <li>• Open in response to specific calls</li> <li>• Open only through collaboration</li> <li>• Data access only</li> <li>• Access restricted at present</li> <li>• Closed to access</li> </ul>   | Please choose one of the access conditions applicable to your facility   |   |
| Collection type:<br><ul style="list-style-type: none"> <li>• Disease-specific</li> <li>• Case-control</li> <li>• Cohort</li> <li>• Cross-sectional</li> <li>• Longitudinal</li> <li>• Population-based</li> <li>• Quality control</li> <li>• Clinical trial</li> </ul>   | Select the category that best describes your collection. If you think a category is missing, please contact us.  |   |
| Collection status:<br><ul style="list-style-type: none"> <li>• Not started</li> <li>• In progress</li> <li>• Completed</li> </ul>  | Indicate your collection status.   |   |
| Collection point:<br><ul style="list-style-type: none"> <li>• Pre-diagnosis</li> <li>• Post-diagnosis</li> <li>• During diagnosis</li> <li>• Multiple points in patient pathway</li> </ul>   | Select the category that best describes the point in the diagnosis pathway at which your samples were collected.   |   |
| Consent restriction:<br><ul style="list-style-type: none"> <li>• Genetic analysis restriction</li> <li>• Human genetic analysis restriction</li> <li>• Scope of use restriction</li> <li>• Export restriction</li> <li>• Disease area restriction</li> <li>• Project specific restriction</li> <li>• Commercial restriction associated data</li> </ul> | Please indicate any areas of research for which you do not have consent.   |   |
| Associated data  | See table <a href="#">ADD ASSOCIATED DATA TO COLLECTION AND CAPABILITY on page 18</a>  |   |



| ADD SAMPLE SET   | ADDITIONAL INFORMATION   | EXAMPLE                   |
|--|--|---------------------------|
| Sex: <ul style="list-style-type: none"> <li>• Male</li> <li>• Female</li> <li>• Unknown</li> <li>• Both sexes</li> <li>• Undifferentiated</li> </ul>   | Choose gender category based on the options provided.  | Both sexes                |
| Age range: <ul style="list-style-type: none"> <li>• Fetus</li> <li>• Neonate</li> <li>• Infant</li> <li>• Young child</li> <li>• Child</li> <li>• Adolescent</li> <li>• Young adult</li> <li>• Adult</li> <li>• Data not recorded</li> </ul> | Select the age category that best reflects the sample set.                                     | Young adult (18-40 years) |
| Number of donors   | Use the slide bar to select the approximative range of the number of donors in the sample set. | 1001-3000                 |

| ADD MATERIAL PRESENTATION DETAILS   | ADDITIONAL INFORMATION   | EXAMPLE                 |
|---|--|-------------------------|
| Material type   | Divide your sample set into the different types of material you have sampled. If you think a material type, is missing <a href="#">please contact us</a> .   | Serum<br>-35°C to -18°C |
| Storage temperature   | Choose appropriate temperature category based on the options provided.   |                         |
| Microscopic assessment <ul style="list-style-type: none"> <li>• Affected</li> <li>• Non-affected</li> <li>• Not applicable</li> </ul> | If samples were examined by a pathologist, indicate if the sample type contains (affected) or does not contain (non-affected) the disease referenced in this collection. If the sample was not examined by a pathologist, select “Not applicable”. |                         |

| ADD CAPABILITY  | ADDITIONAL INFORMATION  | EXAMPLE      |
|---|---|--------------|
| Disease status  | We are seeking to harmonize the terms used to describe diseases and have adopted <a href="#">SNOMED CT</a> in this effort. We are seeking to discuss with the domain experts what terms are most relevant for certain diseases. Therefore, if you do not see a relevant term, <a href="#">please contact us</a> . If you are collecting from healthy volunteers, use the term ‘Fit and Well’. | Tuberculosis |
| Protocols <ul style="list-style-type: none"> <li>• Bespoke consent form</li> <li>• Bespoke SOP</li> </ul> | Select these if you can offer custom-made SOPs and consent procedures.  |              |
| Annual donor expectation  | Provide an estimate of the number of patients from whom you collected samples and data in a year.   | 120          |
| Associated data   | See table <a href="#">ADD ASSOCIATED DATA TO COLLECTION AND CAPABILITY on page 18</a>   |              |

## ADD ASSOCIATED DATA TO COLLECTION AND CAPABILITY

## DESCRIPTION

These fields will permit you to show what data you have and how long it would take you to provide it. Provision times are: immediate, 0 to 3 months, 3 to 6 months, or more than 6 months. If you are unable to provide any of the data listed here, do not tick anything.

### Quality data:

- Freezer temperature logs
  - Quality indicators
- If you keep records of freezer temperatures over time.
  - If you keep any records that would indicate the quality of the sample or if a QMS system is followed, tick this field. It could, for example, be following an ISO protocol or checking a random sample for yield.

### Research data:

- Biomarker datasets
  - Genomic datasets
- Results of an assay of anything that can be used as an indicator of a particular disease state or some other physiological state.
  - Results of any genome analysis.

### Annotation data:

- Clinical records
  - Follow up records
  - Genealogical records
  - Imaging data
  - National registries
  - Pre-analytical data
  - Physiological/biochemical measurements
  - Primary care records
  - Survey data
  - Psychological data
  - Participant ethnicity
  - Microbiological results
- Data obtained from accessing clinical records.
  - Data recorded about a patient after the sample was taken.
  - Data recorded about the lines of family descent.
  - Access to images or data recorded from images.
  - Data obtained from national registries such as NCRAS.
  - Data obtained from pathology records.
  - Assay results.
  - Data obtained from accessing primary care records.
  - Results from any surveys the participants were involved in
  - Data obtained from accessing treatment records.
  - Data obtained from accessing psychological records.
  - Ethnicity of the participant.
  - Microbiological indicators associated to the samples.