



# Go.Data Metadata Overview

## for Measles

### Goal of this document:

To describe the metadata elements within the core modules of Go.Data when using the default template outbreak **Measles**. This template outbreak is aligned with the most up-to-date guidance from [PAHO/WHO case investigation form](#) (2019).

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## Section 1. Background

Measles is a highly contagious, serious disease caused by a virus. Before the introduction of measles vaccine in 1963 and widespread vaccination, major epidemics occurred approximately every 2–3 years and measles caused an estimated 2.6 million deaths each year. More than 140 000 people died from measles in 2018 – mostly children under the age of 5 years, despite the availability of a safe and effective vaccine. Measles is caused by a virus in the paramyxovirus family and it is normally passed through direct contact and through the air. The virus infects the respiratory tract, then spreads throughout the body. Measles is a human disease and is not known to occur in animals.

This document presents with details the core components of Go.Data case creation including the case personal information, including location and address, epidemiology information, questionnaire and laboratory information. The last section is a demonstration of categorical data update in the Go.Data reference data section. Additionally, there is a section presenting the variables included in the creation of an event and a relationship between a case and exposures or contacts.

The personal information, epidemiology and laboratory sections are generic across all outbreaks for Go.Data. **The questionnaire section is strictly specific to Measles investigation and has been tailored to reflect relevant questions.**

Information not relevant to Measles should simply be ignored while collecting data.

**NB: Your country/area's location data (with geographic coordinates) needs to be configured after installing Go.Data, in addition to some specific categorical data categories.**

Additional documents and courses will support the adequate use of this document, including the [Go.Data open WHO course](#), [tutorial guide to update reference data](#), and the [Go.Data User Guide](#).

## Section 2. Personal

*NOTE: Cases, Contacts and Contacts of contacts modules have the same variables in the Personal tab; It can be added as many addresses as needed.*

### Data Elements – Personal (red indicates required field)

variable_code_godata	label	type	Notes, incl. relevance for EVD outbreak investigation
visualId	case or contact ID	freetext	Pattern for ID Masks set at the outbreak level.
firstName	First Name	freetext	
middleName	Middle Name	freetext	
lastName	Last Name	freetext	
age.years	Age - Months	numeric	Select Unit (Months or Years) from drop-down before entering number, then stored in database as age.years and age.months accordingly.
age.months	Age - Years	numeric	See above.
dob	Date of Birth	date	Option to enter dob or Age, but not both. If dob, calendar selector appears.
gender	Gender	Dropdown selector	SKIP LOGIC: If Female, Pregnancy variable appears.
pregnancyStatus	Pregnancy Status	Dropdown selector	Mapped to "Is the patient pregnant?" and "Weeks of pregnancy (01-42)"
occupation	Occupation	Dropdown selector	Option set configurable in reference data.
responsibleUserId	Assigned User	Dropdown selector	Select from drop-down list of existing users,
documents.type	Document type	Dropdown selector	Can use to capture other IDs like passport number, etc. Mapped to "Health Facility Case ID" for EVD investigation. Configure reference data to include this value (see Ref Data section).
documents.number	Document number	freetext	

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addresses.typeId	Address Type	Dropdown selector	Option set configurable in reference data. Default = Current Address. Must have at least one Current Address.
addresses.date	Address Date	date	
addresses.phoneNumber	Phone number	freetext	
addresses.emailAddress	Email address	freetext	
addresses.locationId	Location	Dropdown selector	Location drop-down, based on configured hierarchical reference data. Can select most granular level preferred.
addresses.city	City	freetext	
addresses.addressLine1	Address	freetext	Along with the address include the "Landmarks to locate the house"
addresses.postalCode	Postal code	freetext	
addresses.geoLocationAccurate	Coordinates are accurate	boolean	Toggle Switch to mark Yes or No (Default = NO) - in order to manually re-enter GPS vs. Using automatic GPS configured in Location Hierarchy.
geoLocationLat	Latitude	numeric	
geoLocation.Lng	Longitude	numeric	

## Section 3. Epidemiology

*NOTE: It can be added as many hospitalizations/isolations as needed.*

Search

Selected outbreak  
Measles

[Home](#) / [Cases](#) / [Create case](#)

Create case

Personal

Epidemiology

Questionnaire

Done

Details

Classification \*

Confirmed

Date of onset \*

2022-11-02

Date of onset approximate

Date of becoming case

2022-11-07

Date of infection

Case investigation status

Case investigation complete

Date of investigation completed

2022-11-11

Outcome

Alive

Date of outcome

2022-11-11

Refused transfer to treatment unit

Was burial performed safely?

Date of death

Location of death

Burial place name

Date of reporting \*

2022-11-07

Date of reporting approximate

Risk level

1 - Low

Risk reason

Vaccines received

Vaccine \*

Measles Rubella

Vaccine date

2017-10-11

Vaccine status \*

Fully Vaccinated

+ Add another vaccine

Hospitalization / Isolation details

Hospitalization / Isolation type \*

Hospitalization

From

2022-11-07

To

2022-11-10

Centre name

test1

Location

DEMO country

Comments

+ Add another hospitalisation

Data Elements – Epidemiology (red indicates required field)			
variable_code_godata	label	type	Notes, incl. relevance for EVD outbreak investigation
classification	Case classification	Dropdown selector	
dateOfOnset	Date of onset	date	Individual symptom date of onsets included in Questionnaire for fever and rash.
isDateOfOnsetApproximate	Date of onset approximate	boolean	Toggle Switch to mark Yes or No.
dateBecomeCase	Date of becoming a case	date	
dateOfInfection	Date of infection	date	
Caseinvestigationsatus	Case investigation status		
dateOfInvestigationcompleted	Date of investigation completed	date	
outcomeld	Outcome	Dropdown selector	Option set configurable in reference data (SKIP LOGIC: If Deceased, Burial variables appear). Please complete this value with the status of patient at time of this case report and update this information at the time of patient recovery and discharge from the hospital OR at the time of patient death.
dateOfOutcome	Date of outcome	date	
transferRefused	Refused transfer to treatment unit	boolean	Toggle Switch to mark Yes or No.
dateOfdeath	Date of Death	date	
locationOfdeath	Location of death	Dropdown selector	Location drop-down, based on configured hierarchical reference data. Can select most granular level preferred.

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burialPlaceName	Burial Place Name	freetext	
dateOfReporting	Date of reporting	date	Mapped to "date reported, local" and "date reported, national"
isDateOfReportingApproximate	Date of reporting approximate	boolean	Toggle Switch to mark Yes or No.
riskLevel	Risk Level	Dropdown selector	Can configure reference data to specify exposure risk categories, default = high/med/low.
riskReason	Risk Reason	freetext	
vaccinesReceived.vaccine	Vaccine	Dropdown selector	Configure reference data to include Measles/Rubella vaccines (see Ref Data section).
vaccinesReceived.date	Vaccine date	date	Mapped to "Date of last dose"
vaccinesReceived.status	Vaccine status	Dropdown selector	Configure reference data to include vaccines number of doses (see Ref Data section).
dateRanges.typeId	Hospitalization/isolation type	Dropdown selector	Option set configurable in reference data. Mapped to "Hospitalized?"
dateRanges.centerName	Centre Name	Dropdown selector	Option set configurable in reference data.
dateRanges.locationId	Location of hospital/facility	Dropdown selector	Location drop-down, based on configured hierarchical reference data. Can select most granular level preferred.
dateRanges.comments	Comments	freetext	Mapped to "Hospital record number"
dateRanges.startDate	From (Date)	date	
dateRanges.endDate	To (Date)	date	

## Section 4. Questionnaire (Case Investigation)

*NOTE: aligned with PAHO/WHO case investigation form (2019)*

Search

Selected outbreak  
Measles / Rubella

Home / Cases / Modify Name 1 Last name 1

Modify Name 1 Last name 1

CancelQuick actionsSave

PersonalEpidemiologyQuestionnaire

Health Care Interactions

1. Initial diagnosis

Select answer  
Measles

2. Reporting institution / Health service name

Write answer  
Institution 1

3. Reporting institution / Health service telephone

Write answer  
+99 999 9999

4. Type of provider reporting

Select answer  
Public

5. Date of consultation

Select date  
2022-11-02

6. Date of home visit

Select date  
2022-11-03

7. Detected by

Select answer  
Institutional search

Patient Information

8. Indicate household head or next of kin first and last name, if relevant

Write answer  
Name 2 Last name 2

9. Indicate the type of locality where the patient lives

Select answer  
Urban

10. Work or school address

Write answer  
School address 99



## Clinical information

### 11. Vaccinated against Measles/Rubella

Select answer  
Yes

**\*NOTE: After completing the questionnaire, please update case vaccination information in the "Epidemiology" tab within Case Module**

#### 11.1.1. Source of vaccination Information

Select answer  
Vaccination card

### 12. If the patient is pregnant, indicate the place where the birth will likely take place (After completing the questionnaire, please update case pregnancy information in the "Personal" tab within Case Module)

Write answer  
NA

### 13. If the patient is dead, indicate the primary cause of death (After completing the questionnaire, please update this information in the "Epidemiology" tab within the Case Module)

Write answer  
NA

### 14. Does the case present/has any signs and symptoms?

Select answer  
Yes

#### 14.1.1. Fever

Select answer  
Yes

##### 14.1.1.1. Date of fever onset

Select date  
2022-11-01

##### 14.1.1.2. Please indicate the temperature

Write numeric value  
38

#### 14.1.2. Rash

Select answer  
Yes

##### 14.1.2.1. Duration of the rash (in days)

Write numeric value  
5

##### 14.1.2.2. Date of rash onset

Select date  
2022-11-02

##### 14.1.2.3. Type of rash

Select answer  
Maculopapular

#### 14.1.3. Cough

Select answer  
Yes

#### 14.1.4. Conjunctivitis

Select answer  
Yes

#### 14.1.5. Coryza

Select answer  
No

14.1.6. Koplik Spots

Select answer  
No

14.1.7. Lymphadenopathy

Select answer  
No

14.1.8. Arthralgia

Select answer  
Yes

## Epidemiology and exposure

15. Were active case searches conducted?

Select answer  
Yes

**\*NOTE: After completing the questionnaire, please register identified cases in Go.Data**

15.1.1. Number of suspect cases detected during active case-search

Write numeric value  
4

16. Was the patient in contact with any pregnant woman?

Select answer  
Yes

**\*NOTE: After completing the questionnaire, please register the identified contact in Go.Data and complete pregnancy information in the 'Personal' tab within the Contact Module**

17. Are there other cases present in the case's municipality of residence?

Select answer  
Yes, with Rubella

**\*NOTE: After completing the questionnaire, please register identified cases in Go.Data**

18. Did the patient travel outside his/her province/state of residence 7-23 days before rash onset?

Select answer  
Yes

**Please add travel to each country below by first entering in "Select Date" the "Entry date", select "Add Travel History" and then fill subsequent fields (City/Country and Departure Date). You can add as many as needed**

19. If relevant, fill entry date, select "Add Travel History" and fill subsequent fields

19.1. Select date \*  
2022-10-26

Select answer  
Add travel history

19.1.1. City / Country

Write answer  
City 1, Country 1

19.1.2. Date of Departure

Select date  
2022-10-26

20. Setting where infected?

Select answer  
Community

^ 21. Ring vaccination?

Select answer

Yes

21.1.1. Date started

Select date

2022-11-03



21.1.2. Date ended

Select date

2022-11-04



21.1.3. Number of doses given during ring vaccination

Write numeric value

30

^ 22. Was rapid coverage monitoring done?

Select answer

Yes

22.1.1. What percentage of vaccinated persons was found?

Write numeric value

90

Outcome/Status

^ 23. FINAL CLASSIFICATION

Select answer

Measles

^ 24. Basis for confirmation

Select answer

Laboratory

^ 25. Basis for discarding

Select answer

^ 26. For confirmed cases, source of infection

Select answer

Unknown source

27. Classified by

Write answer

Technical officer 1

28. Date of final classification

Select date

2022-11-14



LEGEND Add answer Remove Download Browse

## Data Elements – Questionnaire (red indicates required field)

variable_code_godata	Label - EN	type	Notes
initial_diagnosis	Initial Diagnosis	single_answer	OPTIONS = Measles; Rubella; Dengue; Other rash illness; Other non-rash illness; Unknown
hcs_name	Reporting institution / Health service name	freetext	
hcs_telephone	Reporting institution / Health service telephone	freetext	
provider_type	Type of provider reporting	single_answer	OPTIONS = Public; Private; Other / Includes nested questions
consultation_date	Date of consultation	date/time	
homevisit_date	Date of home visit	date/time	
detected_by	Detected by	single_answer	OPTIONS = Spontaneous consultation; Laboratory; Institutional Search; Community Case Search; Contact investigation; Community Report; Other; Unknown
provider_specify	<i>IF OTHER: Specify</i>	freetext	
HH_Next_of_kin	Indicate household head or next of kin first and last name, if relevant	freetext	
locality_type	Indicate the type of locality where the patient lives	single_answer	OPTIONS = Urban; Periurban; Rural
work_school_address	Work or school address	freetext	
vaccination_status	Vaccinated against Measles/Rubella	single_answer	OPTIONS = yes; no; unk / Includes nested questions
	If yes, display markup: *NOTE: After completing the questionnaire, please update case vaccination information in the "Epidemiology" tab within Case Module		
vaccination_source	<i>IF YES: Source of vaccination Information</i>	single_answer	OPTIONS = Vaccination card; Health service record; Verbal
brith_place	If the patient is pregnant, indicate the place where the birth will likely take place	freetext	
cause_of_death	If the patient is dead, indicate the primary cause of death (After completing the questionnaire, please update this information in the "Epidemiology" tab within the Case Module)	freetext	
signs_and_symptoms	Does the case present/has any signs and symptoms?	single_answer	OPTIONS = yes; no; unk / Includes nested questions
symp_fever	<i>IF YES: Fever</i>	single_answer	OPTIONS = yes; no; unk / Includes nested questions
fever_onset	<i>IF YES: Date of fever onset</i>	date/time	
fever_temp	<i>IF YES: Please indicate the temperature</i>	numeric	
symp_rash	<i>IF YES: Rash</i>	single_answer	OPTIONS = yes; no; unk / Includes nested questions
rash_duration	<i>IF YES: Duration of the rash (in days)</i>	numeric	
rash_onset	<i>IF YES: Date of rash onset</i>	date/time	
rash_type	<i>IF YES: Type of rash</i>	single_answer	OPTIONS = Maculopapular; Vesicular; Other; Unk
symp_cough	<i>IF YES: Cough</i>	single_answer	OPTIONS = yes; no; unk
symp_conjunctivitis	<i>IF YES: Conjunctivitis</i>	single_answer	OPTIONS = yes; no; unk
symp_coryza	<i>IF YES: Coryza</i>	single_answer	OPTIONS = yes; no; unk
symp_koplik_spots	<i>IF YES: Koplik Spots</i>	single_answer	OPTIONS = yes; no; unk
symp_lymphadenopathy	<i>IF YES: Lymphadenopathy</i>	single_answer	OPTIONS = yes; no; unk
symp_arthralgia	<i>IF YES: Arthralgia</i>	single_answer	OPTIONS = yes; no; unk
active_searches	Were active case searches conducted?	single_answer	OPTIONS = yes; no; unk / Includes nested questions
	If yes, display markup: *NOTE: After completing the questionnaire, please register identified cases in Go.Data		
suspected_cases_detected	Number of suspect cases detected during active case-search	numeric	
contact_pregnant_woman	Was the patient in contact with any pregnant woman?	single_answer	OPTIONS = yes; no; unk / Includes nested questions
	If yes, display markup: *NOTE: After completing the questionnaire, please register the identified contact in Go.Data and complete pregnancy information in the "Personal" tab within the Contact Module		
additional_cases	Are there other cases present in the case's municipality of residence?	single_answer	OPTIONS = yes, with Measles; yes, Rubella; yes, with both; no; unk
	If yes, display markup: *NOTE: After completing the questionnaire, please register identified cases in Go.Data		
expo_travel	Did the patient travel outside his/her province/state of residence 7-23 days before rash onset?	single_answer	OPTIONS = yes; no; unk / Includes nested questions
	If yes, display markup: Please add travel to each country below by first entering in "Select Date" the "Entry date", select "Add Travel History" and then fill subsequent fields (City/Country and Departure Date). You can add as many as needed		
travel_history	If relevant, fill entry date, select "Add Travel History" and fill subsequent fields	single_answer	Add travel history

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city_country	<i>IF ADD TRAVEL HISTORY: City / Country</i>	single_answer	
departure_date	<i>IF ADD TRAVEL HISTORY: Date of Departure</i>	single_answer	
Setting_where_infected	Setting where infected?	single_answer	OPTIONS = Household contact; Community; Health Center; Unknown; Others
ring_vaccination	Ring vaccination?	single_answer	OPTIONS = yes; no; unk / Includes nested questions
date_started	<i>IF YES: Date started</i>	date/time	
date_ended	<i>IF YES: Date ended</i>	date/time	
ring_vacc_dosis	<i>IF YES: Number of doses given during ring vaccination</i>	Numeric	
rapid_coverage_monitoring	Was rapid coverage monitoring done?	single_answer	OPTIONS = yes; no; unk / Includes nested questions
rcm_vaccination_percentage	<i>IF YES: What percentage of vaccinated persons was found?</i>	Numeric	
final_classification	FINAL CLASSIFICATION	single_answer	OPTIONS = Measles; Rubella; Discarded
basis_for_confirmation	Basis for confirmation	single_answer	OPTIONS = Laboratory; Epidemiological Link; Clinical
basis_for_discarding	Basis for discarding	single_answer	OPTIONS = Measles/Rubella IgM-neg; Vaccine Reaction; Dengue; Parvovirus B19; Herpes 6; Allergic Reaction; Other Diagnosis / Includes nested questions
please_specify	IF Other Diagnosis: Please specify	Freetext	
source_of_infection	For confirmed cases, source of infection	single_answer	OPTIONS = Imported; Import-Related; Unknown source; Endemic / Includes nested questions
I_importing_country	<i>IF IMPORTED: Country of importation</i>	freetext	
IR_importation_country	<i>IFIMPORT-RELATED: Country of importation</i>	freetext	
classified_by	Classified by	freetext	
Final_classification_date	Date of final classification	date/time	

## Section 5. Laboratory

Add Lab details to a case/contact record (can also be bulk uploaded by lab personnel):

Search

Selected outbreak  
 Measles / Rubella

Home / Lab results / Cases / Name 1 Last name 1 / Name 1 Last name 1 Lab results / Modify lab result lab-ID-XX999

Cancel Quick actions Save

Details

### Details

Sample lab ID  
lab-ID-XX999

Sample taken \*  
2022-11-04

Sample delivered to lab  
2022-11-05

Sample tested  
2022-11-06

Date of the result  
2022-11-07

Lab name  
Health care center A

Sample type  
Serum

Test type  
IgM IF

Result  
Positive for targeted pathogen (if serology, e...)

Tested for  
Measles

Status  
In progress

Quantitative result

Notes

### Variant / Strain

Has variant / strain?

Date sample sent for determining variant / ...

Variant / Strain lab name

Variant / Strain result date

Variant / Strain result

Reason

### Data Elements – Laboratory (red indicates required field)

variable_code_godata	label	type	Notes
sampleIdentifier	Sample Lab ID	freetext	
dateSampleTaken	Date sample taken	date	
dateSampleDelivered	Date sample delivered to lab	date	
dateTesting	Date sample tested	date	
dateOfResult	Date of the result	date	
labName	Lab name	Dropdown selector	Option set configurable in reference data.
sampleType	Sample type	Dropdown selector	Option set configurable in reference data. Configure reference data to include the specific <i>sample type</i> collected for Measles/Rubella (see Ref Data section).
testType	Test type	Dropdown selector	Option set configurable in reference data. Option set configurable in reference data. Configure reference data to include the specific <i>test type</i> for Measles/Rubella (see Ref Data section).
result	Result	Dropdown selector	Option set configurable in reference data. Option set configurable in reference data. Configure reference data to include the specific <i>result type</i> for Measles/Rubella (see Ref Data section).
status	Status	Dropdown selector	Option set configurable in reference data.
testedFor	Tested for	freetext	Mapped to "Antigen", which includes: Measles, Rubella, Dengue, Parvovirus B19, Herpes 6, Enterovirus, Other.
quantitativeResult	Quantitative result	freetext	
notes	Notes	freetext	

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sequence.hasSequence	Has variant / strain?	boolean	Toggle Switch to mark Yes or No (skip logic: IF Yes, sequencing questions appear, IF No, only "Reason" appears).
sequence.dateSampleSent	Date sample sent for determining variant / strain	date	
sequence.labId	Variant / Strain lab name	Dropdown selector	
sequence.dateResult	Variant / Strain result date	date	
sequence.resultId	Variant / Strain result	Dropdown selector	
sequence.noSequenceReason	Reason	freetext	

## Section 6. Event Creation

*NOTE: can create exposure events and link cases to them. Address block is same to case creation.*

? Search

Selected outbreak  
Measles / Rubella

Home / Events / Create new event

### Create new event

1 Details
2 Done

#### Details

Name \*  
Birthday party

Date \*  
2022-10-26

Date of reporting \*  
2022-11-02

☐ Date of reporting approximate

Assigned user  
DEMO User

Category  
Party

End date  
2022-10-26

Description

#### Address

Type  
Current address

Date  
2022-11-14

Email address  
email@email.com

Phone number  
+99 999 9999

Location (Please specify the village or municipal area)  
DEMO country

City  
City 1

Postal code  
999

Address  
Address 1

Latitude \*  
1

Longitude \*  
1

☐ Coordinates are accurate

### Data Elements – Events (red indicates required field)

variable_code_godata	label	type	Notes
name	Name	freetext	Only name and date required, for each event, no Event ID
date	Date	date	Only name and date required, for each event, no Event ID
dateOfreporting	Date of reporting	date	
assignedUser	Assigned user	Dropdown selector	Option set configurable in reference data.
Category	Category	Dropdown selector	Option set configurable in reference data.
endDate	End date	date	
description	Description	freetext	Only name and date required, for each event, no Event ID
addresses.typeId	Address Type	Dropdown selector	Option set configurable in reference data. Default = Current Address. Must have at least one Current Address.
addresses.date	Address Date	date	
addresses.phoneNumber	Phone number	freetext	
addresses.emailAddress	Email address	freetext	
addresses.locationId	Location	Dropdown selector	Location drop-down, based on configured hierarchical reference data. Can select most granular level preferred.
addresses.city	City	freetext	
addresses.addressLine1	Address	freetext	
addresses.postalCode	Postal code	freetext	
addresses.geoLocationAccurate	Coordinates are accurate	boolean	Toggle Switch to mark Yes or No (Default = NO) - in order to manually re-enter GPS vs. Using automatic GPS configured in Location Hierarchy.
geoLocationlat	Latitude	numeric	
geoLocation.lng	Longitude	numeric	

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## Section 7. Relationship creation

*NOTE: can add relationships/exposures to source event or case or relationship to a contact where the case is the exposure*

?

Search

Q

Selected outbreak

✓

Measles / Rubella

▼

👤

Home / Cases / Name 1 Last name 1 / Exposures / Create relationship

Create relationship

1

Name 2 Last name 2

2 Done

Name 2 Last name 2

Date of first contact

2022-10-26

📅

Date of last contact

2022-10-26

📅

Is contact date estimated?

🔴

Certainty level

3 - High

▼

Exposure type

Direct physical contact

▼

Exposure frequency

1-5 Times

▼

Exposure duration

Medium (hours)

▼

Context of exposure

Home/household

▼

Cluster

▼

Relationship

cousins

Comment

📝

### Data Elements – Relationship creation (red indicates required field)

variable_code_godata	label	type	Notes
dateOfFirstContact	Date of first contact	date	
dateOfLastContact	Date of last contact	date	
contactDateEstimated	Is contact date estimated?	boolean	Toggle Switch to mark Yes or No
certaintyLevelId	Certainty level	Dropdown selector	Option set configurable in reference data.
exposureTypeId	Exposure type	Dropdown selector	Option set configurable in reference data.
exposureFrequencyId	Exposure frequency	Dropdown selector	Option set configurable in reference data.
exposureDurationId	Exposure duration	Dropdown selector	Option set configurable in reference data.
clusterId	Cluster	Dropdown selector	select from existing clusters within system
socialRelationshipTypeId	Context of exposure	Dropdown selector	Option set configurable in reference data.
socialRelationshipDetail	Relationship	freetext	
comment	Comments	freetext	

## Appendix: Reference data for categorical variables:

We have created some key reference data specifications for categorical variables in Go.Data that align with [Measles/Rubella \(PAHO/WHO – Case investigation form\)](#). These can be updated easily in the reference data section of Go.Data (see below).

*Default values not used in the context of Measles investigation can be deactivated. For more information see: [Tutorial for updating Go.Data Reference Data to align with a Disease-Specific Template](#)*

- Vaccine:

Configure reference data to include "Measles", "Rubella", "Measles Rubella", and "Measles Mumps Rubella" :

Search

Selected outbreak: Measles / Rubella

Home / Reference data / Vaccine

**Vaccine**

Quick actions + Add

14 results \*\*Filters are applied on this list (may include hidden fields)

Label	Code	Description	Icon	Colour	Order	Active	System Value
Measles	1		No icon	No colour	1	Yes	No
Rubella	2		No icon	No colour	2	Yes	No
Measles Rubella	3	MR	No icon	No colour	3	Yes	No
Measles Mumps Rubella	4	MMR	No icon	No colour	4	Yes	No

- Vaccine status:

Configure reference data to include "Zero dose", "One dose", "Two doses", "Three doses", etc.:

Search

Selected outbreak: Measles / Rubella

Home / Reference data / Vaccine Status

**Vaccine Status**

Quick actions + Add

10 results \*\*Filters are applied on this list (may include hidden fields)

Label	Code	Description	Icon	Colour	Order	Active	System Value
Zero dose			No icon	No colour	1	Yes	No
One dose			No icon	No colour	2	Yes	No
Two doses			No icon	No colour	3	Yes	No
Three doses			No icon	No colour	4	Yes	No
[...] doses			No icon	No colour	5	Yes	No
Unknown			No icon	No colour	6	Yes	No

- Type of sample:

Configure reference data to include "Nasopharyngeal aspirate/swab" and "Cerebrospinal Fluid" :

?

Search

Q

Selected outbreak

Measles / Rubella

▼

Home / Reference data / Type of sample

Type of sample

Quick actions

Add

14 results

\*\*Filters are applied on this list (may include hidden fields)

↺

☰

Label	Code	Description	Icon	Colour	Order	Active	System Value	
Serum			No icon	No colour	1	Yes	No	👁️ ✎️ ⋮
Nasopharyngeal aspirate/swab			No icon	No colour	2	Yes	No	👁️ ✎️ ⋮
Throat swab			No icon	No colour	3	Yes	No	👁️ ✎️ ⋮
Urine			No icon	No colour	4	Yes	No	👁️ ✎️ ⋮
Cerebrospinal Fluid			No icon	No colour	5	Yes	No	👁️ ✎️ ⋮
Other (specify in Notes field)			No icon	No colour	6	Yes	No	👁️ ✎️ ⋮

- Type of lab result:

Configure reference data to include "IgM EIA/Indirect" , " IgM EIA/Capture" , " Virus Isolation" , " IgM IF" , " IgG EIA/Capture" , " IgG IF" , and " HI" :

?

Search

Q

Selected outbreak

Measles / Rubella

▼

Home / Reference data / Type of lab result

Type of lab result

Quick actions

Add

15 results

\*\*Filters are applied on this list (may include hidden fields)

↺

☰

Label	Code	Description	Icon	Colour	Order	Active	System Value	
IgM EIA/Indirect			No icon	No colour	1	Yes	No	👁️ ✎️ ⋮
IgM EIA/Capture			No icon	No colour	2	Yes	No	👁️ ✎️ ⋮
Virus Isolation			No icon	No colour	3	Yes	No	👁️ ✎️ ⋮
PCR			No icon	No colour	4	Yes	No	👁️ ✎️ ⋮
IgM IF			No icon	No colour	5	Yes	No	👁️ ✎️ ⋮
IgG EIA/Capture			No icon	No colour	6	Yes	No	👁️ ✎️ ⋮
IgG IF			No icon	No colour	7	Yes	No	👁️ ✎️ ⋮
HI			No icon	No colour	8	Yes	No	👁️ ✎️ ⋮

- Lab test result:

Configure reference data to include "Inadequate specimen" , " Unknown (result not available)" :

Search

Selected outbreak  
Measles / Rubella

Home / Reference data / Lab test result

Lab test result

Quick actions ⌵ Add

6 results

\*\*Filters are applied on this list (may include hidden fields)

Label	Code	Description	Icon	Colour	Order	Active	System Value	
Negative for targeted pathogen			No icon	No colour	1	Yes	No	
Positive for targeted pathogen (if serology, enter titre in Qu...			No icon	No colour	2	Yes	No	
Inadequate specimen			No icon	No colour	3	Yes	No	
Inconclusive for targeted pathogen			No icon	No colour	4	Yes	No	
Unknown (result not available)			No icon	No colour	5	Yes	No	

Measles Reference Data Specifications		
variable_code	optionset_code	optionset_label
vaccinesReceived.vaccine	measles	Measles
vaccinesReceived.vaccine	rubella	Rubella
vaccinesReceived.vaccine	mr	Measles Rubella
vaccinesReceived.vaccine	mmr	Measles Mumps Rubella
vaccinesReceived.status	zero_dose	Zero dose
vaccinesReceived.status	one_dose	One dose
vaccinesReceived.status	two_doses	Two doses
vaccinesReceived.status	three_doses	Three doses
sampleType	nasopharyngeal_swab	Nasopharyngeal aspirate/swab
sampleType	cerebrospinal_fluid	Cerebrospinal Fluid
testType	lgm_eia_indirect	IgM EIA/Indirect
testType	lgm_eia_capture	IgM EIA/Capture
testType	Virus_isolation	Virus Isolation
testType	lgm_if	IgM IF
testType	lgg_eia_capture	IgG EIA/Capture
testType	lgg_if	IgG IF
testType	hi	HI
Result	inadequate_specimen	Inadequate specimen
result	unk	Unknown (result not available)