

# Template Map for the Case Import File

Last Updated: April 22, 2025

# **Table of Contents**

INSTRUCTIONS	3
INFANT DEMOGRAPHIC INFORMATION	4
SCREENING INFORMATION	7
INITIAL SPECIMEN COLLECTION INFORMATION	8
SUBSEQUENT SPECIMEN COLLECTION INFORMATION	8
POINT-OF-CARE TEST INFORMATION	g
INTERVENTION, FOLLOW-UP, AND DIAGNOSIS	g
FINAL DIAGNOSIS	10
If condition is Presence of Other Hb Variant	13
If condition is Critical congenital heart disease – CCHD and finalDiagnosis is CCHD	13
If condition is "Spinal Muscular Atrophy – SMA	14



#### **INSTRUCTIONS**

This template map provides variable names and acceptable values for the <u>case import file</u>. This import file is one of the two options for newborn screening programs to enter individual cases into the NewSTEPs Repository. The other option is to use the <u>online webform</u>.

The <u>case import file</u> contains the common demographic and screening variables that are asked for all conditions. It also contains final diagnosis for certain conditions. General instructions to use the import file include:

- Required fields are indicated below; these variables must have an acceptable value entered in order for the import to work
- For fields that are not required, the variable or column is also not required
  - For non-required variables/columns included in the CSV file, enter an acceptable value or leave empty
- Variables/columns may be in any order
- Each row is unique to the case/baby; please be sure to select the correct condition, this
  includes secondary conditions

Download the <u>case import file</u>, enter the data that is being reported, and save the document as a CSV file to your desktop. To import the file into the repository, select **Choose File** on the right-hand side of the screen. The File Explorer for your desktop will appear and the desired file can be selected. Next, select **Submit CSV** to import the file. If data isn't formatted correctly, the import will not be accepted.

Common errors in import files include:

- Abbreviation of the state or territory name; please spell out
- Conditions not spelled correctly or use the correct format; it is suggested that you copy and paste directly from this template map and only abbreviate conditions found on page
- NULL versus true zero: only enter zero when the value is a true zero, otherwise leave the cell empty



#### INFANT DEMOGRAPHIC INFORMATION

**state** - name of the state/territorial newborn screening program, REQUIRED\* Acceptable values:

- Alabama
- Alaska
- American Samoa
- Arizona
- Arkansas
- California
- Colorado
- Connecticut
- Commonwealth of the Northern Mariana Islands
- Delaware
- District of Columbia
- Florida
- Georgia
- Guam
- Hawaii
- Idaho
- Illinois
- Indiana

- lowa
- Kansas
- Kentucky
- Louisiana
- Maine
- Maryland
- Massachusetts
- Michigan
- Minnesota
- Mississippi
- Missouri
- Montana
- Nebraska
- Nevada
- New Hampshire
- New Jersey
- New Mexico
- New York
- North Carolina
- North Dakota

- Ohio
- Oklahoma
- Oregon
- Pennsylvania
- Puerto Rico
- Rhode Island
- South Carolina
- South Dakota
- Tennessee
- Texas
- US Virgin Islands
- Utah
- Vermont
- Virginia
- Washington
- West Virginia
- Wisconsin
- Wyoming

birthYear - The year in which the birth occurred, REQUIRED\*

stateUniqueId - The unique identifier assigned to the case by the state, REQUIRED\*

**condition** - Name of condition, REQUIRED\* Acceptable values:

- 2,4 Dienoyl-CoA reductase deficiency DE RED
- 2-Methyl-3-hydroxybutyric aciduria 2M3HBA
- 2-Methylbutyrylglycinuria 2MBG
- 3-Hydroxy-3-methyglutaric aciduria HMG
- 3-Methylcrotonyl-CoA carboxylase deficiency 3-MCC
- 3-Methylglutaconic aciduria 3MGA
- Argininemia ARG
- Argininosuccinic aciduria ASA
- Beta-Ketothiolase deficiency BKT
- Biopterin defect in cofactor biosynthesis BIOPT (BS)
- Biopterin defect in cofactor regeneration BIOPT (RG)
- Biotinidase deficiency BIOT
- Carbamoyl phosphate synthetase I deficiency CPS
- Carnitine acylcarnitine translocase deficiency CACT
- Carnitine palmitoyltransferase type I deficiency CPT IA
- Carnitine palmitoyltransferase type II deficiency CPT II



- Carnitine uptake defect/carnitine transport defect CUD
- Citrullinemia, type I CIT
- Citrullinemia, type II CIT II
- Classic galactosemia GALT
- Classic PKU & Hyperphe
- Congenital Toxoplasmosis TOXO
- Congenital adrenal hyperplasia CAH
- Congenital hypothyroidism CH
- Critical congenital heart disease CCHD
- Cystic fibrosis CF
- Cytomegalovirus CMV
- Duchenne Muscular Dystrophy DMD
- Ethylmalonic encephalopathy EME
- Fabry
- Formiminoglutamic acidemia FIGLU
- Galactoepimerase deficiency GALE
- Galactokinase deficiency GALK
- Gaucher
- Glucose-6-phosphate dehydrogenase deficiency G6PDD/G6PD
- Glutaric acidemia type I GA1
- Glutaric acidemia type II GA2
- Guanidinoacetate Methyltransferase GAMT
- Hb No structural variant
- Hearing loss HEAR
- Holocarboxylase synthetase deficiency MCD
- Homocystinuria HCY
- Human Immunodeficiency Virus HIV Exposure
- Hypermethioninemia MET
- Hyperornithinemia with Gyrate Deficiency Hyper ORN
- Hyperornithinemia-hyperammonemia-homocitrullinemia syndrome HHH
- Isobutyrylglycinuria IBG
- Isovaleric acidemia IVA
- Krabbe Disease
- Long-chain L-3 hydroxyacyl-CoA dehydrogenase deficiency LCHAD
- Malonic acidemia MAL
- Maple syrup urine disease MSUD
- Medium-chain acyl-CoA dehydrogenase deficiency MCAD
- Medium-chain ketoacyl-CoA thiolase deficiency MCKAT
- Medium/short-chain L-3-hydroxyacl-CoA dehydrogenase deficiency M/SCHAD
- Methylmalonic acidemia (cobalamin disorders) Cbl A,B
- Methylmalonic acidemia (methylmalonyl-CoA mutase) MUT
- Methylmalonic acidemia with homocystinuria Cbl C,D
- Mucopolysaccharidosis I MPS I
- Mucopolysaccharidosis II MPS II
- Niemann Pick



- Nonketotic Hyperglycinemia NKH
- Ornithine transcarbamylase deficiency OTC
- Pompe
- Presence of Hb S
- Presence of Other Hb Variant
- Prolinemia Type I/ Type II PRO
- Propionic acidemia PROP
- Pyroglutamic acidemia 5-OXO
- Severe Combined Immunodeficiencies SCID
- Short-chain acyl-CoA dehydrogenase deficiency SCAD
- Spinal Muscular Atrophy SMA
- T-cell related lymphocyte deficiencies
- Trifunctional protein deficiency TFP
- Tyrosinemia, type I TYR I
- Tyrosinemia, type II TYR II
- Tyrosinemia, type III TYR III
- Very long-chain acyl-CoA dehydrogenase deficiency VLCAD
- X-linked Adrenoleukodystrophy
- Zellweger Syndrome

Note: The following condition abbreviations can be used instead of using the entire **condition** name:

_	2	N A	$\sim$	$\sim$
•	3-	IVI	יטו	U

ASA

BIOT

BKT

CAH

• Cbl A,B

CCHD

• CF

CH

CIT

CUD

GA1

GALT

GAMT

HCY

HEAR

HMG

• IVA

LCHAD

MCAD

• MCD

MPS I

MPS II

• MSUD

• MUT

Pompe

PROP

• TFP

TYR I

VLCAD

**gestationalAge** - the gestational age in weeks (please use whole numbers only)

birthWeight - the birth weight in grams

biologicalGender - the biological gender of the infant



Acceptable values: FEMALE, MALE, UNSPECIFIED, UNKNOWN

ethnicity - The ethnicity of the infant

Acceptable values:

- HISPANIC\_LATINO\_OR\_SPANISH
- NOT HISPANIC LATINO OR SPANISH
- NOT REPORTED
- UNKNOWN

Note: only one value should be specified

race - the race of the infant

Acceptable race values:

- ISLANDER
- ASIAN
- NATIVE AMERICAN
- BLACK OR\_AFRICAN\_AMERICAN
- WHITE
- UNKNOWN
- NOT REPORTED

Note: If more than one value applies, separate each value with a colon (e.g., ISALNDER:WHITE) Note: ISLANDER = Native Hawaiian or other Pacific Islander

#### **SCREENING INFORMATION**

**screeningIdentifyingRisk** - The screening result which indicated this infant was at risk for the disorder. Acceptable values:

- Initial Screen
- Second Required Screen
- Subsequent Screen

**prenatalTestForRisk** - Was prenatal testing done that indicated that this infant was at risk for this disorder? Acceptable values: TRUE, FALSE, UNKNOWN

**familyHistoryRisk** - Was there a family history that indicated that this infant was at risk for this disorder? Acceptable values: TRUE, FALSE, UNKNOWN

**diagnosedAfterNewbornScreening** - Was this individual identified outside of newborn screening? Acceptable values: TRUE, FALSE, UNKNOWN

**missedDiagnosisReason** - The reason this diagnosis was not identified by newborn screening. *Note:* should only be answered if diagnosedAfterNewbornScreening is TRUE Acceptable values:

- Parental Refusal
- Lost to follow-up after unsatisfactory specimen
- Biologic false negative / result within normal range
- Did not have a valid screen due to error



#### Other

**otherMissedDiagnosisReason** - Text description of the missed diagnosis reason up to 254 characters long. *Note: should only be answered if missedDiagnosisReason is OTHER* 

#### INITIAL SPECIMEN COLLECTION INFORMATION

**birthToInitialSpecimenCollection** - hours between birth and initial specimen collection. Integer value. *Not specified for conditions "Critical congenital heart disease - CCHD", "Hearing loss - HEAR"* 

**birthToInitialSpecimenCollectionIncludesTime** - Acceptable values: TRUE, FALSE *Note:* TRUE signifies that the data available for the calculation of elapsed time included time as well as date

**birthToInitialReceiptByLab** - Time elapsed from birth until the initial NBS specimen was received by the lab, in days (as measured by 24-hour periods since the birth). Integer value. Not specified for conditions "Critical congenital heart disease - CCHD", "Hearing loss - HEAR"

**birthToInitialReceiptByLabIncludesTime** - Acceptable value: TRUE, FALSE *Note: TRUE* signifies that the data available for the calculation of elapsed time included time as well as date

**birthToInitialResultRelease** - Time elapsed from birth until the release of out-of-range results as a result of the initial screen, in days (as measured by 24-hour periods since the birth). *Not specified for conditions "Critical congenital heart disease - CCHD", "Hearing loss - HEAR"* 

**birthTolnitialResultReleaseIncludesTime** - Acceptable value: TRUE, FALSE. *Note: TRUE* signifies that the data available for the calculation of elapsed time included time as well as date

#### SUBSEQUENT SPECIMEN COLLECTION INFORMATION

**birthToSubsequentSpecimenCollection** - Time elapsed from birth until the subsequent NBS specimen was collected, in days (as measured by 24-hour periods since the birth). *Not specified for conditions "Critical congenital heart disease - CCHD", "Hearing loss - HEAR"* 

**birthToSubsequentSpecimenCollectionIncludesTime** - Acceptable value: TRUE, FALSE *Note:* TRUE signifies that the data available for the calculation of elapsed time included time as well as date



**birthToSubsequentReceiptByLab** - Time elapsed from birth until the subsequent NBS specimen was received by the lab, in days (as measured by 24-hour periods since the birth). Not specified for conditions "Critical congenital heart disease - CCHD", "Hearing loss - HEAR"

**birthToSubsequentReceiptByLabIncludesTime** - Acceptable value: TRUE, FALSE *Note:* TRUE signifies that the data available for the calculation of elapsed time included time as well as date

**birthToSubsequentResultRelease** - Time elapsed from birth until the release of out-of-range results as a result of the subsequent screen, in days (as measured by 24 hour periods since the birth). *Not specified for conditions "Critical congenital heart disease - CCHD", "Hearing loss - HEAR"* 

**birthToSubsequentResultReleaseIncludesTime** - Acceptable value: TRUE, FALSE. *Note:* TRUE signifies that the data available for the calculation of elapsed time included time as well as date

# POINT-OF-CARE TEST INFORMATION

**birthToPointOfCareTestInterval** - Time elapsed from birth in hours until the point of care screening test was performed. *Only specified for conditions "Critical congenital heart disease - CCHD", "Hearing loss - HEAR"* 

**birthToPointOfCareTestIntervalIncludesTime** - Acceptable value: TRUE, FALSE *Note:* true signifies that the data available for the calculation of elapsed time included time as well as date. Only specified for conditions "Critical congenital heart disease - CCHD", "Hearing loss - HEAR"

# INTERVENTION, FOLLOW-UP, AND DIAGNOSIS

**birthToIntervention** - Time elapsed from birth until intervention by an appropriate medical provider occurred, in days (as measured by 24-hour periods since the birth)

**birthToDiagnosisConfirmation** - Time elapsed from birth until confirmation of the diagnosis occurred, in days (as measured by 24-hour periods since the birth)

**treatmentInOtherState** - Is infant receiving treatment/care out-of-state? Acceptable values: TRUE, FALSE, UNKNOWN

**treatmentState** - state where infant receives treatment/care? *Note: should only be answered if treatmentInOtherState is TRUE*Acceptable values: see list provided for **state** 

**diagnosisReversed** - Is this diagnosis reversed? Note: this does not refer to the therapeutic interventions to address a condition (i.e., surgery, treatment, therapy, etc)
Acceptable values: TRUE, FALSE, UNKNOWN



**diagnosisReversedYear** - year diagnosis reversed (*note: enter four-digit year*) Note: should only be answered if diagnosisReversed is TRUE

### **FINAL DIAGNOSIS**

**finalDiagnosis** - final diagnosis as determined by the medical provider performing the clinical diagnostic workup, REQUIRED\*

Note: not all conditions require a final diagnosis; please use the table to see what conditions need a final diagnosis and the associated acceptable values. The final diagnosis categories do NOT include any of the secondary or other conditions listed on the RUSP. These should be entered as a separate case (see **conditions**).

Condition	Acceptable Values
3-Methylcrotonyl-CoA carboxylase deficiency - 3-MCC	<ul> <li>3-Methylcrotonyl-CoA Carboxylase Deficiency</li> <li>3-MCC</li> <li>Maternal MCC deficiency</li> <li>MT-ATP6 related mitochondrial disorders</li> <li>Unknown</li> </ul>
Argininosuccinic aciduria - ASA	<ul><li>Argininosuccinic Acidemia/ Aciduria (ASA)</li><li>Pyruvate carboxylase deficiency</li><li>Unknown</li></ul>
Biotinidase deficiency - BIOT	<ul><li>Profound Biotinidase deficiency</li><li>Partial Biotinidase deficiency</li><li>Unknown</li></ul>
Citrullinemia, type I - CIT	<ul><li>Citrullinemia, Type I</li><li>Pyruvate carboxylase deficiency</li><li>Unknown</li></ul>
Carnitine uptake defect/carnitine transport defect - CUD	<ul> <li>Carnitine Uptake Deficiency (CUD)</li> <li>Maternal Carnitine Deficiency (primary and secondary)</li> <li>Unknown</li> </ul>
Classic PKU & Hyperphe	<ul> <li>Classic phenylketonuria - PKU</li> <li>Benign hyperphenylalaninemia - H-PHE</li> <li>HyperPhe diet controlled</li> <li>Dihydropterine reductase deficiency (DHPR)</li> <li>DNAJC12</li> <li>Parenteral nutrition</li> <li>Maternal PKU</li> <li>Unknown</li> </ul>
Classic galactosemia - GALT	<ul><li>Classic Galactosemia</li><li>Duarte variant galactosemia</li><li>Unknown</li></ul>
Congenital hypothyroidism - CH	<ul> <li>Primary Congenital Hypothyroidism</li> <li>Secondary Congenital Hypothyroidism</li> <li>TBG Deficiency (Thyroxine Binding Globulin) or other protein binding defect</li> <li>Transient Congenital Hypothyroidism</li> <li>Unknown</li> </ul>



Condition	A acceptable Malves
Condition	Acceptable Values
	Classic 21-Hydroxylase Deficiency- Salt     Wasting
	Classic 21-Hydroxylase Deficiency- Simple
Congenital adrenal hyperplasia - CAH	Virilizing
	Other Adrenal disorder
	Unknown
	CCHD
Critical congenital heart disease - CCHD	Non critical CCHD
Children congenital mount discuss Comb	Other
	• Unknown
	CFTR-Related Metabolic Syndrome (CRMS)
Cystic fibrosis - CF	CFTR-Related Disease
	Typical Cystic Fibrosis (CF)
	Unknown
	Holocarboxylase synthetase deficiency (MCD)
	Maternal 3-methylcrotonyl-CoA carboxylase
Holocarboxylase synthetase deficiency - MCD	deficiency
	MT-ATP6 related mitochondrial disorders
	Other biotin disorder
	Unknown
	Classic Homocystinuria
	Methionine Adenosyltransferase (MAT I/III      Deficiency)
Homocystinuria - HCY	Deficiency)
	<ul><li>Glycine n-methyltransferase (GNMT)</li><li>Adenosylhomocysteine Hydrolase Deficiency</li></ul>
	Macrosymomocysteme nydrolase Deliciency     Unknown
	Isovaleric Acidemia/ Aciduria (IVA)
	Short/branched chain acyl-CoA
Isovaleric acidemia - IVA	dehydrogenase Deficiency (SBCAD) or 2-
	methylbutyrl CoA dehydrogenase deficiency
	Unknown
	Infantile Onset Krabbe Disease
Krabbe Disease	<ul> <li>Later Onset Krabbe Disease</li> </ul>
Nabbe Disease	Uncertain Type/Onset
	Unknown
	Classic
	Intermediate
Maple syrup urine disease - MSUD	Thiamine-response
,y ansassss	Hydroxyprolinemia
	Unclassified
	• Unknown
Methylmalonic acidemia (methylmalonyl-CoA mutase) - MUT	• Mutase (-) (mut-)
	Mutase (0) (mut0)
	Maternal vitamin B12 deficiency
	Succinate-CoA ligase deficiency
	Unclassified
Mathedra I and a side and a few to the standard of the standar	Unknown     Oak alassis A staffician ass (Obl.A)
Methylmalonic acidemia (cobalamin disorders) -	Cobalamin A deficiency (Cbl A)     Cobalamin B deficiency (Cbl B)
Cbl A,B	Cobalamin B deficiency (Cbl B)



Condition	Acceptable Values
Condition	Cobalamin Dv2 deficiency (Cbl Dv2)
	Maternal vitamin B12 deficiency
	Succinate-CoA ligase deficiency
	Unclassified
	• Unknown
	Cobalamin C deficiency (Cbl C)
	Cobalamin D deficiency (Cbl D)     Cobalamin D deficiency (Cbl D)
	Cobalamin F deficiency (Cbl F)  Cobalamin Pud deficiency (Cbl Pud)
Methylmalonic acidemia with homocystinuria - Cbl	<ul><li>Cobalamin Dv1 deficiency (Cbl Dv1)</li><li>Cobalamin J deficiency (Cbl J)</li></ul>
C,D	<ul> <li>Cobalamin J deficiency (Cbl J)</li> <li>Maternal vitamin B12 deficiency</li> </ul>
	Succinate-CoA ligase deficiency
	Other cobalamin deficiency
	Unclassified
	Unknown
	MPS I - severe
	MPS I - severity not determined
Mucopolysaccharidosis I - MPS I	MPS I - attenuated
	Uncertain Type/Onset
	Unknown
Mucopolysaccharidosis II - MPS II	Severe
	Attenuated
	Uncertain Type/Onset
	Unknown
	Infantile Onset (IO) Pompe Disease
Pompe	Late Onset (LO) Pompe Disease
1 onipe	Uncertain Type/Onset
	Unknown
	S,S disease (Sickle cell anemia) - Hb SS
	S, Beta 0-thalassemia - Hb S/B0Th
Presence of Hb S	S, Beta + thalassemia - Hb S/B+ Th
	S,C disease - Hb S/C
	• S, Other
	Unknown
	Hemoglobin C disease
	<ul><li>Hemoglobin D disease</li><li>Hemoglobin E disease</li></ul>
Presence of Other Hb Variant	Library and all the O. Amerika discussion
	Hemoglobin O-Arab disease     Other hemoglobin disorder
	Unknown
	Alpha thalassemia major (Fetal Hydrops)
Hb - No structural variant	Beta thalassemia major (Cooley's anemia)
	Hgb H disease
	Unknown
	Propionic Acidemia (PROP)
	Maternal vitamin B12 deficiency
Propionic acidemia - PROP	Succinate-CoA ligase deficiency
	Unknown
Severe Combined Immunodeficiencies - SCID	Classic SCID
	Leaky SCID
	Omenn Syndrome
	Jillotti Oyttarottio



Condition	Acceptable Values
	Unknown
	Tyrosinemia, Type I (hepatorenal)
Tyrosinemia, type I - TYR I	Transient Tyrosinemia of the neonate (TTN)
	Unknown
	X-Linked Adrenoleukodystrophy (in Males)
	X-Linked Adrenoleukodystrophy (in Females)
	Contiguous ABCD1 DXS1357E deletion
	syndrome (CADDS)
	Peroxisomal Disorder
Y linked Adronaloukodystrophy	Acyl-CoA Oxidase Deficiency
X-linked Adrenoleukodystrophy	D-Bifunctional Protein Deficiency
	Dyamin-like protein 1 (DLP1)
	ABDC5
	Non-peroxisomal Disorder
	Uncertain Type/Onset
	Unknown

**otherFinalDiagnosisName -** Specify the name for the other final diagnosis when the value "OTHER" is entered for *finalDiagnosis* 

### If condition is Presence of Other Hb Variant

alphaThalassemiaPresent- Alpha thalassemia present?

Acceptable values: TRUE, FALSE, UNKNOWN

Note: must only be entered when condition is "Presence of Other Hb Variant"

# If condition is Critical congenital heart disease - CCHD and finalDiagnosis is CCHD

**cchdFinalDiagnosesDetails-** Specify type of CCHD diagnosed. Acceptable values:

- TRUNCUS ARTERIOSUS
- TOTAL ANOMALOUS PULMONARY VENOUS CONNECTION
- TETRALOGY OF FALLOT
- PULMONARY\_ATRESIA
- EBSTEIN\_ANOMALY
- HYPOPLASTIC LEFT HEART SYNDROME
- SINGLE\_VENTRICLE
- TRICUSPID\_ATRESIA
- TRANSPOSITION OF GREAT ARTERIES
- DOUBLE\_OUTLET\_RIGHT\_VENTRICLE
- COARCTATION OF AORTA
- INTERRUPTED AORTIC ARCH
- AORTIC VALVE DISEASE



Note: must only be entered when CCHD FinalDiagnosis is CCHD; can add multiple selections by using a colon to separate each acceptable value (e.g., TRUNCUS ARTERIOSUS:PULMONARY ATRESIA:SINGLE VENTRICLE)

# If condition is Spinal Muscular Atrophy - SMA

**newbornSMN2MolecularTest** - newborn screen molecular test for SMN2? Acceptable values: TRUE, FALSE, UNKNOWN Note: only enter if condition is "Spinal Muscular Atrophy - SMA"

**newbornSMN2MolecularTestValue** - SMN2 copy number? Acceptable values: ONE, TWO, TWO\_OR\_MORE, UNKNOWN Note: only enter if condition is Spinal Muscular Atrophy - SMA" and newbornSMN2MolecularTest is TRUE

**postNewbornSMN2MolecularTest** - post-newborn screen molecular test for SMN2? Acceptable values: TRUE, FALSE, UNKNOWN Note: only enter if condition is "Spinal Muscular Atrophy - SMA"

**postNewbornSMN2MolecularTestValue** - SMN2 copy number? Acceptable values: ONE, TWO, TWO\_OR\_MORE, UNKNOWN Note: only enter if condition is Spinal Muscular Atrophy - SMA" and newbornSMN2MolecularTest is TRUE

#### **CHANGE LOG**

# Modifications made from August 2023 version

- Removed all diagnostic variables
  - Note: Collecting diagnostic information was well intended to standardize the identification and classification of disorders, aligning with the <u>public health</u> <u>surveillance case definitions</u> for newborns in the United States. However, it was decided that the cons (i.e., staff time, poor feedback loop with clinicians, inability to obtain all diagnostic information to feed into the classification system) outweigh the benefits of collecting this information. NewSTEPs will continue to collect demographic and newborn screening information for individual cases, which will continue to help inform our birth prevalence, health equity, and quality improvement practices.

# Modifications made from January 2024 version

 Updated spelling from Holocarboxylase synthase deficiency - MCD to Holocarboxylase synthetase deficiency - MCD

#### Modifications made from March 2024 version



Updated language for diagnosedAfterNewbornScreening to clarify that the diagnosis
was made outside of newborn screening. Note: this was just a change to the variable
label and does not impact queries.

# **Modifications from May 2024 version**

- Final diagnosis category is now a required field
- Unknown was added to the final diagnosis for all conditions that have a final diagnosis option

### **Modifications from June 2024 version**

- In the case template map, separated NOT\_REPORTED and UNKNOWN for ethnicity. This was a typo only in the case template map as these two categories were accidentally on the same line.
- Updated spelling from Cobalamin Dv2 (CblDv2) to Cobalamin Dv2 deficiency (CblDv2).
- Updated spacing in the following final diagnosis options
  - Classic 21-Hydroxylase Deficiency- Salt Wasting
  - Classic 21-Hydroxylase Deficiency- Simple Virilizing
  - Cobalamin A deficiency (Cbl A)
  - Cobalamin B deficiency (Cbl B)
  - Cobalamin C deficiency (Cbl C)
  - Cobalamin Dv2 deficiency (Cbl Dv2)
  - Cobalamin D deficiency (Cbl D)
  - Cobalamin F deficiency (Cbl F)
  - Cobalamin Dv1 deficiency (Cbl Dv1)
  - Cobalamin J deficiency (Cbl J)
  - o S,C disease- Hb S/C
  - S,S disease (Sickle cell anemia) Hb SS
- Updated formatting
  - o Argininosuccinic aciduria ASA
  - Beta-Ketothiolase deficiency BKT
  - Biotinidase deficiency BIOT
  - Carbamovl phosphate synthetase I deficiency CPS
  - Carnitine acylcarnitine translocase deficiency CACT
  - Carnitine uptake defect/carnitine transport defect CUD
  - Citrullinemia, type I CIT
  - Citrullinemia, type II CITII
  - Classic galactosemia GALT
  - Congenital Toxoplasmosis TOXO
  - Congenital adrenal hyperplasia CAH
  - Congenital hypothyroidism CH
  - o Critical congenital heart disease CCHD
  - Cystic fibrosis CF
  - Cytomegalovirus CMV
  - Ethylmalonic encephalopathy EME
  - Formiminoglutamic acidemia FIGLU
  - o Galactoepimerase deficiency GALE
  - Galactokinase deficiency GALK
  - Guanidinoacetate Methyltransferase GAMT
  - Hb No structural variant
  - Hearing loss HEAR



- Holocarboxylase synthetase deficiency MCD
- Homocystinuria HCY
- Hypermethioninemia MET
- Spinal Muscular Atrophy SMA

# **Modifications from September 2024 version**

- Added clarification that gestational age should be in whole numbers.
- Updated spelling for SMA molecular variables from UKNOWN to UNKNOWN.

#### **Modification from December 2024**

- Updated language from Was this individual not identified by newborn screening to Was this individual identified outside of newborn screen?
- Add Duchenne Muscular Dystrophy DMD as an acceptable condition.
- Updated final diagnosis for Maple syrup urine disease by removing Type IA III, and adding classic, intermediate, thiamine-response.
- Added final diagnosis of unclassified to Maple syrup urine disease, Methylmalonic acidemia (methylmalonyl-CoA mutase) - MUT, Methylmalonic acidemia (cobalamin disorders) - Cbl A, B, and Methylmalonic acidemia with homocystinuria – Cbl C,D.
- Added final diagnosis of uncertain type/onset to Mucopolysaccharidosis I MPS I, Pompe, and X-linked Adrenoleukodystrophy.
- Added final diagnoses to Krabbe Disease of infantile onset Krabbe disease, later onset Krabbe disease, uncertain type/onset, and unknown.
- Added final diagnoses to Mucopolysaccharidosis II MPS II of severe, attenuated, uncertain type/onset, and unknown.

