

# Born in Bradford Data Dictionary

Eclipse neonatal data

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## Background

This document is a data dictionary for Eclipse neonatal data. It describes 27 variables from 1 source. This document was built from Born in Bradford database version BUILD-JAN2018.

## Born in Bradford

Born in Bradford is a longitudinal multi-ethnic birth cohort study aiming to examine the impact of environmental, psychological and genetic factors on maternal and child health and wellbeing. Bradford is a city in the North of England with high levels of socio-economic deprivation and ethnic diversity. Women were recruited at the Bradford Royal Infirmary at 26-28 weeks gestation. For those consenting, a baseline questionnaire was completed. The full BiB cohort recruited 12,453 women and 3353 of their partners across 13,776 pregnancies and 13,858 children between 2007 and 2010. The cohort is broadly characteristic of the city's maternal population. Mean age of the mothers at study recruitment was 27 years old. Researchers are looking at the links between the circumstances of a child's birth, the context in which they grow up, their health and well-being and their educational progress. Ethical approval for the data collection was granted by Bradford Research Ethics Committee (Ref 07/H1302/112).

## Study identifiers

Study identifiers are standardised across Born in Bradford data sources to enable linking of data from different sources.

Variable	Variable Label	Details
<b>ChildID</b>	BiB Child ID	Unique ID assigned to each child at birth. Where birth outcome is unknown for a given pregnancy, ChildID will be blank and there is no child recruited to the study from that pregnancy. Use MotherID with ChildID to link siblings together. Note that twins have separate ChildIDs but the same PregnancyID.
<b>FatherID</b>	BiB Father ID	Unique ID assigned to partners post-recruitment. Use FatherID with PregnancyID to link fathers across pregnancies. Where FatherID matches across two PregnancyIDs, but those PregnancyIDs are associated with different MotherIDs, this is a father with two separate pregnancies in the cohort with different mothers. Likewise, where MotherID matches across two PregnancyIDs, but those PregnancyIDs are associated with different FatherIDs, this is a mother with two separate pregnancies in the cohort with different fathers.
<b>MotherID</b>	BiB Mother ID	Unique ID assigned to each mother post-recruitment. MotherID should be used when looking for pregnancies or children associated with the same mother. Data collected at pregnancy level will duplicate for MotherIDs that are in the study for more than one pregnancy.
<b>PregnancyID</b>	BiB Pregnancy ID	Unique ID assigned to each mother at recruitment. It is named PregnancyID because a mother can enrol for more than one pregnancy. If a mother returns to enrol for a second or third pregnancy, she is assigned a new PregnancyID. Children and partners from that pregnancy can be linked to the mother by the PregnancyID

## eClipse Baby Measures

Database ID for source: ec1bby

This source is measured at the **child** level. It contains data from 13525 children with one observation per child. There are 27 variables with a total of 13525 observations.

### Description

eClipse maternity electronic record: baby measures

Variable	Variable Label	Details
<b>ec1bdcirc</b>	Abdominal circumference (cm)	Routine Healthcare: Continuous value <hr/> Range 3.1 to 53.5 Mean 31.19 11686 non-missing values
<b>ec1bapgar1m</b>	APGAR at 1 minute	Routine Healthcare: Integer value <hr/> Range 0 to 10 Mean 8.47 13285 non-missing values
<b>ec1bapgar5m</b>	APGAR at 5 minutes	Routine Healthcare: Integer value <hr/> Range 0 to 10 Mean 9.05 13272 non-missing values
<b>ec1bbabysex</b>	Sex of baby	Routine Healthcare: Categorical value <hr/> 13525 non-missing values <hr/> Coding [ec1bbyec1bbabysex]: 1 = Male 2 = Female
<b>ec1bbirthwt</b>	Birth weight (g)	Routine Healthcare: Integer value <hr/> Range 480 to 5800 Mean 3205.46 13524 non-missing values

Variable	Variable Label	Details
<b>eclbrtasst</b>	Assistance during birth	Routine Healthcare: Categorical value <hr/> 13515 non-missing values <hr/> Coding [eclbbyeclbrtasst]: 1 = None 2 = Forceps 3 = Ventouse 4 = Forceps and ventouse
<b>eclbrthocm</b>	Outcome of birth	Routine Healthcare: Categorical value <hr/> 13525 non-missing values <hr/> Coding [eclbbyeclbrthocm]: 1 = Livebirth 2 = Stillbirth
<b>eclbrthord</b>	Birth order	Routine Healthcare: Integer value <hr/> Range 1 to 3 Mean 1.01 13525 non-missing values
<b>eclbrtpres</b>	Presentation at birth	Routine Healthcare: Categorical value <hr/> 13525 non-missing values <hr/> Coding [eclbbyeclbrtpres]: 1 = Cephalic 2 = Cephalic brow 3 = Cephalic face 4 = Cephalic vertex 5 = Breech 6 = Breech extended 7 = Breech flexed 8 = Breech footling 9 = Breech frank 10 = Breech dorso-anterior 11 = Breech complete 12 = Compound 13 = Oblique 14 = Transverse 15 = Unknown
<b>eclheadcir</b>	Head circumference (cm)	Routine Healthcare: Continuous value <hr/> Range 20.9 to 43 Mean 34.20 12372 non-missing values

Variable	Variable Label	Details
<b>ecllgaukwho</b>	Large for gestational age (>90th centile)	<p>Derived: Categorical value</p> <hr/> <p>Large for gestational age is Yes if birthweight is above 90th percentile on UK WHO fetal growth charts for sex and gestational week at birth. Only calculated for singletons, missing for multiple births.</p> <hr/> <p>13194 non-missing values</p> <hr/> <p>Coding [ec1bbyyesno]: 0 = No 1 = Yes</p>
<b>eclmidarmc</b>	Mid-arm circumference (cm)	<p>Routine Healthcare: Continuous value</p> <hr/> <p>Range 2 to 32.5 Mean 10.65 11668 non-missing values</p>
<b>eclrtbirth</b>	Route of birth	<p>Routine Healthcare: Categorical value</p> <hr/> <p>13525 non-missing values</p> <hr/> <p>Coding [ec1bbyeclrtbirth]: 1 = Vaginal 2 = Caesarean</p>
<b>eclrupbrth</b>	Time from membrane rupture to birth	<p>Routine Healthcare: Text value</p> <hr/> <p>1947 unique values 13525 non-missing values</p>
<b>eclsdystoc</b>	Shoulder dystocia	<p>Routine Healthcare: Categorical value</p> <hr/> <p>12419 non-missing values</p> <hr/> <p>Coding [ec1bbyyesno]: 0 = No 1 = Yes</p>

Variable	Variable Label	Details
<b>eclsgaukwho</b>	Small for gestational age (<10th centile)	<p>Derived: Categorical value</p> <hr/> <p>Small for gestational age is Yes if birthweight is below 10th percentile on UK WHO fetal growth charts for sex and gestational week at birth. Only calculated for singletons, missing for multiple births.</p> <hr/> <p>13194 non-missing values</p> <hr/> <p>Coding [ec1bbyyesno]:  0 = No  1 = Yes</p>
<b>eclsubscap</b>	Subscapular skinfold thickness (cm)	<p>Routine Healthcare: Continuous value</p> <hr/> <p>Range 0.8 to 10.4  Mean 4.72  9230 non-missing values</p>
<b>ecltmpbrth</b>	Temperature at birth	<p>Routine Healthcare: Continuous value</p> <hr/> <p>Range 3.7 to 39.9  Mean 36.86  11632 non-missing values</p>
<b>ecltriceps</b>	Triceps skinfold thickness (cm)	<p>Routine Healthcare: Continuous value</p> <hr/> <p>Range 1.2 to 10.8  Mean 5.08  9257 non-missing values</p>
<b>eclvitk1st</b>	Vitamin K first dose	<p>Routine Healthcare: Text value</p> <hr/> <p>43 unique values  13525 non-missing values</p>
<b>eclvitkcon</b>	Vitamin K consent	<p>Routine Healthcare: Categorical value</p> <hr/> <p>13348 non-missing values</p> <hr/> <p>Coding [ec1bbyeclvitkcon]:  0 = Not indicated  1 = Consent declined  2 = Consent obtained  3 = Given with consent  4 = Given without consent in an emergency  5 = Offer being considered</p>

Variable	Variable Label	Details
<b>flgabdcirc</b>	Abdominal circ. flag	<p>Routine Healthcare: Categorical value</p> <hr/> <p>Flag variables indicate a value was changed during QC</p> <hr/> <p>11689 non-missing values</p> <hr/> <p>Coding [ec1bbyflag1b1]:            1 = Value checked and is correct            2 = Value was incorrect and has been changed            3 = Value cannot be verified            4 = Value not checked</p>
<b>flgbirthwt</b>	Birth weight flag	<p>Routine Healthcare: Categorical value</p> <hr/> <p>Flag variables indicate a value was changed during QC</p> <hr/> <p>50 non-missing values</p> <hr/> <p>Coding [ec1bbyflag1b1]:            1 = Value checked and is correct            2 = Value was incorrect and has been changed            3 = Value cannot be verified            4 = Value not checked</p>
<b>flgheadcir</b>	Head circ. flag	<p>Routine Healthcare: Categorical value</p> <hr/> <p>Flag variables indicate a value was changed during QC</p> <hr/> <p>129 non-missing values</p> <hr/> <p>Coding [ec1bbyflag1b1]:            1 = Value checked and is correct            2 = Value was incorrect and has been changed            3 = Value cannot be verified            4 = Value not checked</p>
<b>flgmidarmc</b>	Midarm circ. flag	<p>Routine Healthcare: Categorical value</p> <hr/> <p>Flag variables indicate a value was changed during QC</p> <hr/> <p>149 non-missing values</p> <hr/> <p>Coding [ec1bbyflag1b1]:            1 = Value checked and is correct            2 = Value was incorrect and has been changed            3 = Value cannot be verified            4 = Value not checked</p>

Variable	Variable Label	Details
<b>flgsubscap</b>	Subscap. skinfold thickness flag	<p>Routine Healthcare: Categorical value</p> <hr/> <p>Flag variables indicate a value was changed during QC</p> <hr/> <p>86 non-missing values</p> <hr/> <p>Coding [ec1bbyflag1b1]:</p> <p>1 = Value checked and is correct</p> <p>2 = Value was incorrect and has been changed</p> <p>3 = Value cannot be verified</p> <p>4 = Value not checked</p>
<b>flgtricep</b>	Tricep skinfold thickness flag	<p>Routine Healthcare: Categorical value</p> <hr/> <p>Flag variables indicate a value was changed during QC</p> <hr/> <p>67 non-missing values</p> <hr/> <p>Coding [ec1bbyflag1b1]:</p> <p>1 = Value checked and is correct</p> <p>2 = Value was incorrect and has been changed</p> <p>3 = Value cannot be verified</p> <p>4 = Value not checked</p>