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Juggling complexity: Undertaking the first national linked data research on perinatal and maternal outcomes in Australia

ABSTRACT

Background

Data linkage offers a powerful mechanism for examining healthcare outcomes across populations. It has the potential to generate substantial robust datasets using routinely collected electronic data. However, it also presents methodological challenges, especially in Australia where many health datasets are maintained by eight separate states and territories. In a national study, we used linked data to investigate perinatal and maternal outcomes in relation to different places of birth. This study examined data from all eight jurisdictions regarding the outcomes from births planned in hospitals, birth centres and at home. Data linkage enabled the first compilation of an Australia-wide dataset on birth outcomes. However, jurisdictional differences in data collection created challenges in obtaining comparable cohorts for each birth setting comprised of women with similar low-risk pregnancies. The objective of this paper is to describe the techniques for managing linked data, and specifically for ensuring the resulting dataset contained only low-risk pregnancies.

Methods

This paper indicates the procedures for preparing and merging perinatal, inpatient and mortality data from different sources. It provides technical guidance to help address challenges arising in linked data study designs.
Results

We combined data from four collections of administrative healthcare and civil registration data from eight jurisdictions (states and territories). The merging process ensured that variables were consistent, compatible and relevant to study aims. To generate comparable cohorts across birth settings, we developed increasingly complex strategies to ensure that the dataset eliminated women with pregnancies at risk of complications during labour and birth. It was then possible to compare birth outcomes for comparable samples, enabling a specific examination of the impact of birth setting on maternal and infant safety across Australia.

Conclusions

Data linkage is a valuable resource to enhance knowledge about birth outcomes from different settings, notwithstanding methodological challenges. It is possible for researchers to develop and share practical techniques to address these challenges. This study has clear implications for jurisdictions to develop more consistent data collections to facilitate future data linkage.

KEYWORDS

Medical record linkage, pregnancy outcome, retrospective studies, pregnancy complications
ABBREVIATIONS

ABS       Australian Bureau of Statistics
ACT       Australian Capital Territory
APDC      Admitted Patient Data Collection
BIA       Birthplace in Australia
CHeReL    Centre for Health Record Linkage
CS        Caesarean section
DLU       Data linkage unit
HREC      Human Research Ethics Committee
ICD-AM    International Classification of Diseases – Australian modification
NICU      Neonatal intensive care unit
NSW       New South Wales
NT        Northern Territory
PDC       Perinatal Data Collection
PPN       Person project number
RBDM      Registry of Births, Deaths and Marriages
SA        South Australia
SCN       Special care nursery
WA        Western Australia
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BACKGROUND

Different countries provide maternity services in a variety of ways. In Australia, most births (97.5%) take place in public or private hospitals. Other settings include birth centres (either attached to a hospital or stand-alone) (1.8% of births) or at home (0.2%) [1]. In high-resource countries like Australia, healthy women giving birth generally have very good outcomes. Thus, given the small number of adverse events, determining which place of birth is safest is challenging. Combining data from multiple state and territories or across several years is more likely to provide evidence about safety. Generating such combined data, however, involves complex methodological and technical challenges to optimise the quality of the evidence from which to guide policy decisions.

This paper presents methodological experiences in the Birthplace in Australia: a population-based cohort study (BIA), examining the perinatal and maternal outcomes from births planned in hospital obstetric units, birth centres and at home (ref – authors). This research was a nation-wide retrospective cohort study, combining linked data from Australia’s eight jurisdictions (six states and two territories) for the period 2000-2012.

Similar large-scale studies examining outcomes by place of birth have been conducted in other high-income countries, including England [2], the Netherlands [3-6], Nordic countries [7, 8], Canada [9-12], the United States [13-15] and New Zealand [16, 17]. Although previous Australian research has investigated outcomes related to place of birth in single states [18-20], none has attempted to examine outcomes for women nation-wide. In Australia, the
data sets and variables are not uniform across the country, highlighting a challenge in creating a standard national data set.

Linking administrative data gathered over several years can generate the statistical power necessary to detect and compare rare outcomes such as perinatal mortality [21, 22] or examine health amongst vulnerable populations [23]. Data linkage combines electronic data from separate collections to amalgamate information about the same individual, facilitating research while maintaining privacy. However, its limitations include the time, technical intricacy and clerical burden involved, as well as concerns over the accuracy, consistency and comparability of data collected primarily for administrative purposes [24-31]. Increasing numbers of studies of maternal or perinatal health and wellbeing have utilised data linkage, in Australia [32-35] and elsewhere, particularly in Britain, the United States, and the Nordic countries [36].

Linking health-related data requires care, patience and expertise [23], especially linking maternal or neonatal datasets [22, 37, 38], and identifying and adjusting for errors and disparities [31, 32, 39-44]. Some researchers have described data linkage techniques for perinatal health research within one Australian state [23, 45, 46], although compiling and merging maternity data across jurisdictions is less common [40, 47].

In Australia, state and territory governments are responsible for much healthcare delivery (via public hospitals and community health), monitoring and administration. The BIA study used data from up to four comprehensive data collections in each of Australia’s eight jurisdictions, merging them into a robust dataset, containing information on multiple maternal and perinatal outcomes. However, the data screening and merging process spawned many methodological issues, related to the diversity of underlying data collections.
The BIA study required not only linking perinatal datasets, but also ensuring that the identified cohorts that were as similar as possible. This was imperative to eliminate potential variation from confounding factors other than the birth setting, especially differences in underlying risk factors that women may have which will influence their birth outcomes. It was important to exclude women with complex pregnancies, as they are likely to have higher rates of intervention and poorer outcomes than healthy women without any known risk factors. This stipulation added further complexity to the already intricate process of linking multiple datasets.

The objective of this paper is to describe data preparation techniques used to produce a national linked dataset on Australian women with low-risk pregnancies. It presents more detail on procedures than was feasible in the original BIA study report [authors]. Further, this paper aims to assist other researchers by highlighting and addressing problems encountered in managing and merging data from multiple sources in a complex healthcare system.

METHODS

The study retrospectively analysed routinely collected health data on women with low-risk pregnancies who gave birth between 2000 and 2012 in Australia. The BIA study was approved by the [University] Human Research Ethics Committee (HREC) (reference 2012000167).

Research questions

This paper addresses three research questions:

- How to merge already linked administrative data from eight states and territories?
• How to deal with inconsistencies in data collections?

• How to ensure a dataset that best represents women with low risk of obstetric complications, given the constraints of the data collections?

Data sources

In Australia, data on healthcare service delivery and civil registrations are gathered at state/territory level. To examine maternal and perinatal outcomes comprehensively, we requested linked data from four sources in each jurisdiction. These sources were:

• Perinatal Data Collections (PDC) collected by maternity providers

• Admitted Patient Data Collections (APDC) compiled by hospitals

• Registries of Births, Deaths and Marriages (RBDM) death registrations and

• Australian Bureau of Statistics (ABS) mortality data.

In Australia, data linkage uses probabilistic algorithms to match identifying data such as name, sex, date of birth and address across several databases. It allocates a match weight to similar pairs, representing the likelihood of a match [22, 31]. These identifiers are separated from the health record to preserve privacy, and the health records are ultimately coupled with a randomly-assigned anonymous person project number (PPN) [22, 48]. This method is necessary because Australians do not possess a unique identifier that can be matched across datasets, which would facilitate a deterministic data linkage process [46].

While there is some consistency in data collection and linkage across jurisdictions, limited coordination in data management creates complications for handling and merging the state-based databases into a readily accessible and efficient national database. Even though each jurisdiction recorded data on similar outcomes, they are stored in discrete systems, use
filenames of different formats, and comprise diverse variables with inconsistent value labels and contents. States also vary in quality assurance and linkage key protocols [40].

**Access to linked datasets**

Using a two-step process, we first applied to the Data Linkage Units (DLUs) in eight jurisdictions for linked data on specified variables from the four data collections. The Research Protocol assisted in completing disparate forms and ensuring that the data specifications remained consistent across the many data requests and ethics applications. We then applied for ethical approval from eight state-based HRECs (except in the Australian Capital Territory [ACT], where ethical approval needed to precede application to the data custodian). We used a national ethics application process while complying with state-specific requirements such as the Confidentiality Agreement (in Western Australia - WA) and the Victorian Specific Module.

The aim was to create a standardised national dataset, with consistent and comparable variables. Using the NSW dataset as a template for the master data collection, data were sequentially merged as they arrived from each state or territory, aligning them to the NSW variables. The filtering and screening processes further ensured that sample contained only women with low-risk pregnancies. This process took over two years and is described below.

**Process undertaken**

The following sections outline the approaches adopted to address the three research questions. Although presented here as three distinct and linear stages, in reality the process was more complicated and iterative. Data arrived from DLUs at various times and in widely
differing formats. We cleaned, screened and merged state-based data as they arrived and while we were awaiting datasets from other jurisdictions.

**Stage 1: Cleaning and validating linked data**

In order to develop an independent master file for each state, we extracted and merged data from the PDC, APDC, ABS and RBDM collections (or state-based equivalents). PDC data for 2000-2012 (or available years) were linked with APDC data on hospital admissions for the nine months prior to the birth (mother) and twelve months after the birth (mother and baby), and with ABS and RBDM data on deaths up to twelve months after the birth (mother and baby). One state provided mortality data for the first six months only.

This process generated two master files for each jurisdiction covering mothers and babies, although DLUs generally assigned PPNs for mothers and babies to enable matching. Except for two jurisdictions, the Mothers’ and Babies’ master files could be directly merged to form a state master file, using the process illustrated in Figure 1. For situations where a mother’s PPN and her baby’s PPN were unmatched, a unique phantom ID was assigned to each of the records in the Mothers Master file and Babies Master file to match them within the state master file.

<Insert Figure 1 here: The process of merging PDC, APDC, ABS and RBDM datasets into a state master file>

Basic cleaning during this stage involved validating and verifying demographic details across datasets, including age, gender, date of birth and date of death (some datasets only provided year and month). We compared consistency across all possible sources to minimise errors. For example, a particular PPN might contain gender or date of birth data
which were the same in all datasets except one; in this case, we decided on the data with most occurrences. We applied these data cleaning and validation processes throughout the given datasets for all variables of interest.

Missing data, such as age, arising from changed versions in a particular data collection system were reported accordingly. Data received from all sources were cleaned by eliminating duplicated (e.g. similar records within one unique personal number) and inaccurate or extreme cases (e.g. a child with several mothers). We validated data entries for discrepancies between sources (e.g. stillbirth with date of death), and determined value labels upon consensus among team members (e.g. regrouping of third degree and fourth degree perineal tears into a single variable) for subsequent merging into the master file. Owing to the large file size (often over 2GB), we generally commenced data screening after applying some selection criteria (e.g. spontaneous onset of labour, gestation weeks between 37 and 41) beforehand, to facilitate and streamline cleaning and checking processes.

Having collated the state master file, we performed more robust data validation and verification to minimise discrepancies across contents with similar variable types. The PPNs for mothers and babies were assigned in the DLUs to facilitate linkage. However, sometimes information on mothers and babies was independently stored in addition to the data with the PPN; this made merging the contents between mothers and babies more involved. Box 1 illustrates various possibilities, indicating reasons for excluding some records. It indicates that when data are linked and merged from several sources, some records may be repeated containing slightly different data.
Box 1: Potential scenarios in merging mother/baby data during validation

<table>
<thead>
<tr>
<th>PPN mother*</th>
<th>Birth event</th>
<th>PPN baby*</th>
<th>DOB baby MMM/YYYY</th>
<th>Included in study?</th>
<th>Reason for exclusion</th>
</tr>
</thead>
<tbody>
<tr>
<td>7654321</td>
<td>1</td>
<td>3456789</td>
<td>FEB/1999</td>
<td>No</td>
<td>Before 2000</td>
</tr>
<tr>
<td>7654321</td>
<td>2</td>
<td>5678934</td>
<td>JAN/2000</td>
<td>No</td>
<td>Multiple birth</td>
</tr>
<tr>
<td>7654321</td>
<td>2</td>
<td>6789543</td>
<td>JAN/2000</td>
<td>No</td>
<td>Multiple birth</td>
</tr>
<tr>
<td>7654321</td>
<td>3</td>
<td>4567893</td>
<td>DEC/2012</td>
<td>Yes</td>
<td>-</td>
</tr>
<tr>
<td>7654321</td>
<td>4</td>
<td>2345678</td>
<td>JAN/2013</td>
<td>No</td>
<td>After 2012</td>
</tr>
<tr>
<td>2345678</td>
<td>1</td>
<td>1234567</td>
<td>MAR/2004</td>
<td>No</td>
<td>Multiple mothers</td>
</tr>
<tr>
<td>3456712</td>
<td>1</td>
<td>1234567</td>
<td>MAR/2004</td>
<td>No</td>
<td>Multiple mothers</td>
</tr>
</tbody>
</table>

* Dummy PPNs used here as examples

We developed an approach for verifying records where one baby appeared to have several birth mothers using Algorithm 1. Initially, both variables PPN_Mum and PPN_Baby were sorted consecutively in ascending order, followed by an iterative one-step increment count for Mum_Num when the condition of PPN_Baby equal to previous PPN_Baby and PPN_Mum not equal to previous PPN_Mum was met.

Algorithm 1: Identifying the number of mothers per baby

*/Both PPN_Baby and PPN_Mum were consecutively sorted in ascending order.

SORT CASES BY PPN_Baby(A)PPN_mum(A).
COMPUTE MumNum=1.
IF (PPN_Baby = LAG(PPN_Baby)and PPN_mum<>LAG(PPN_mum))
Mum_Num=LAG(Mum_Num)+1.
EXECUTE.
FREQUENCIES VARIABLES=Mum_Num
/ORDER=ANALYSIS.
Then Algorithm 2 provided a more comprehensive approach to detecting records with the same birth mothers for the same babies, same birth mothers for different babies, different birth mothers for the same baby and different birth mothers for different babies. When a baby appeared to have more than one birth mother, DiffMum>1, then that record was omitted.

**Algorithm 2: Comprehensive combination between number of mothers and number of babies**

*/Both PPN_Mum and PPN_Baby were consecutively sorted in ascending order. Baby1 was assigned to count the number of similar PPN_Baby.
SORT CASES BY PPN_baby(UP)PPN_mum(UP).
COMPUTE baby1=1.
IF (PPN_Baby = LAG(PPN_Baby))baby1=LAG(baby1)+1.
VARIABLE LABELS baby1 'baby1 order'.
EXECUTE.

*/Both PPN_Baby and PPN_Mum were consecutively sorted in ascending order and baby2 was assigned to count the number of similar PPN_Baby.
SORT CASES BY PPN_mum(UP)PPN_Baby(UP).
COMPUTE baby2=1.
IF (PPN_Baby = LAG(PPN_Baby))baby2=LAG(baby2)+1.
VARIABLE LABELS baby2 'baby2 order'.
EXECUTE.

*/Crosstabulating baby1 and baby2 to view the uniqueness in PPNs between baby and mother.
CROSSTABS 
/TABLES=baby1 BY baby2 
/FORMAT=AVALUE TABLES 
/CELLS=COUNT 
/COUNT ROUND CELL.

*/If DiffMum>1, a baby has more than one mother, and the associated record has to be excluded.
IF (baby1 = 2 & baby2 = 1)DiffMum=2.
EXECUTE.
FREQUENCIES VARIABLES=DiffMum
The study methodology was initially pilot-tested within one jurisdiction – NSW, the most populous state which accounts for approximately one-third (31.9%) of Australian births [1]. This test used de-identified data from the NSW DLU, covering the four data collections (above) and the Register of Congenital Conditions. The original NSW master file sample comprised 258,161 women at low risk of complications who gave birth over an eight-year period (mid-2000 to mid-2008). Analysis of the linked data identified significant differences by planned place of birth in type of birth and interventions. The NSW dataset formed the master file for the national study, which adopted the NSW data variables and value labels.

**Stage 2: Merging data from different states and territories**

Using combined NSW data as the base for the national master file, data were sequentially merged from other state master files as they were received and verified (Figure 2).

*Fig 2: Development of national master file, indicating data sources and proportion of sample>*

Data collections varied between states and territories, often involving different variables or similar variables in different formats, requiring adjustment of the matching syntax to ensure values within data from every other state were compatible. For example, what is called ‘mode of birth’ in one state, is termed ‘birth method’ or ‘delivery’ in others, with varying constituent values (see Box 2 for examples of the syntax required). Even within NSW, the components of the dataset changed during the study period. For example, the variable ‘model of care’ was introduced to the PDC in 2006.
Box 2: Syntax required for merging data on mode of birth

In NSW, there were two versions of the mode of birth (‘delivery’) variable (deliv98 and deliv2011) with differing value labels. This called for a new set of value labels that would accommodate all options. We therefore developed the new variable deliv and proceeded to match similar values in data from other states or territories. For example, matching data to the value “6” for “Caesarean Section” required the following syntax to capture the data from other jurisdictions.

Northern Territory: ‘IF CHAR.INDEX(delivery_BB,"CS")>0 Y_deliv2015new=6.’

Queensland: ‘IF CHAR.INDEX(delivery_BB,"CS")>0 deliv=6.’

Tasmania: ‘IF (INDEX(BIRTH_MODE_P,"Caes")>0 or INDEX(ModeBirth_O,"Caes")>0) deliv=6.’

Victoria: ‘RECODE BirthMethod_Ds (‘6’=6) INTO deliv.’ and ‘RECODE BirthMethod_old (‘3’=6) (‘4’=6) INTO deliv.’

WA: ‘IF (INDEX(mid_method,"03")=1) deliv=6.’

For effective data management, we established various common strategies to avoid unnecessary reshuffling of data or other cumbersome processes. In practice, analysts devise their own syntax strategy. Box 3 outlines some examples of the many techniques that we developed to simplify data and procedures.
Box 3: SPSS syntax for simplifying data handling

<table>
<thead>
<tr>
<th>Tip</th>
<th>Syntax</th>
<th>Function</th>
</tr>
</thead>
</table>
| A   | COMPUTE id=$CASENUM.  
     | FORMAT id (F8.0).  
     | EXECUTE. | Keeping all records intact for quick reference if necessary. Assigned before any work was done on the original dataset. |
| B   | COMPUTE DiffY=(date_dth - dob) / (365.25 * time.days(1)).  
     | VARIABLE LABELS DiffY.  
     | VARIABLE LEVEL DiffY (SCALE).  
     | FORMATS DiffY (F8.2).  
     | VARIABLE WIDTH DiffY(8).  
     | EXECUTE. | Computing and validating the age provided in original dataset. |
| C   | STRING DiagP (A3).  
     | COMPUTE DiagP=CHAR.SUBSTR(Diag_codeP,1,3). | Assigning substring of a principal diagnosis code up to 3 characters. |
| D   | COMPUTE pdc_mumAgeOK1=TRUNC(pdc_mumAgeOK).  
     | VARIABLE LABELS pdc_mumAgeOK1 'Truncated to Year'.  
     | EXECUTE. | Truncating mother’s age into year to be consistent across all states. |
| E   | RECODE VAR1 (1=1) (2 thru 4=2) (5 thru 7=3) (8 thru Highest=4) INTO VAR2.  
     | EXECUTE. | Recoding a variable into another variable by regrouping the value contents. |
| F   | IF CHAR.INDEX(delivery_BB,"Forc")>0 deliv=2.  
     | EXECUTE. | Recoding values within a string variable into numeric values of a new variable. |

Tip A illustrates a technique for backtracking the original order of the dataset for quick reference on data entries; Tip B computes the mother’s age if age is not provided or validates the age if given age is in doubt; Tip C extracts the first three characters of a principal diagnosis code to help categorise risk status; Tip D truncates mother’s age into...
year of birth for consistency across all datasets (as maternal age data was presented inconsistently); Tip E recodes a series of value labels in one variable into another series of values in a new variable; Tip F captures contents that contain a certain string and assigns it into a new variable.

**Stage 3: Identifying low-risk cohorts**

The BIA study needed to minimise confounding variables in order to examine the specific impact of birth setting on perinatal safety and well-being. To ensure that the women in different birth place cohorts (hospital, birth centre, home birth) shared similar risk status, a three-phase process filtered out most pregnancies likely to have moderate or high risk of complication for labour and birth.

**Phase 1: Applying common inclusion criteria**

In this phase, the initial inclusion criteria specified:

- Birth between 2000 and 2012
- Singleton pregnancy
- Gestation between 37 weeks and 41 weeks + 6 days
- Spontaneous labour onset

The data received from DLUs varied considerably in the extent to which they addressed the risk factors requested. In one state 33.3% of data supplied met the study criteria for low-risk pregnancy, whereas in another 88.3% of records met them. This variation does not indicate that states and territories varied markedly in the proportion of pregnancies which are relatively low-risk; rather it demonstrates the extent to which DLUs could filter and extract relevant data.
As noted, we commenced the first phase of filtering data before merging the state master files into the national master file, not least to reduce the size of the unwieldy data sets and to facilitate computer execution speeds. This initial screening, either by data custodians or by the research team, resulted in eight individual data sets of women who met inclusion criteria which were cumulatively combined.

**Phase 2: Clarifying indicators of pregnancy risk**

The next stage eliminated cases with other factors that further contribute to the complexity of pregnancies, by closely examining variables provided and the related value labels. This phase excluded women who:

- Received no antenatal care
- Attempted a vaginal birth after one or more previous caesarean sections
- Had a breech presentation

**Phase 3: Excluding complicating medical conditions**

The third phase of screening for risk factors involved scrutinising APDC and PDC data (or equivalents) for International Classification of Diseases (ICD-10-AM) codes, to identify diagnoses and procedures related to pregnancy complications. We excluded women with high-risk O codes (Box 4) and/or infants with Q codes indicating congenital abnormalities in the current pregnancy.
**Box 4: ICD10\(^2\) codes indicating pregnancy complications**

<table>
<thead>
<tr>
<th>ICD-10-AM</th>
<th>Diagnosis</th>
</tr>
</thead>
<tbody>
<tr>
<td>O10</td>
<td>Pre-existing hypertension complicating pregnancy, childbirth and the puerperium</td>
</tr>
<tr>
<td>O11</td>
<td>Pre-eclampsia superimposed on chronic hypertension</td>
</tr>
<tr>
<td>O13</td>
<td>Gestational [pregnancy-induced] hypertension</td>
</tr>
<tr>
<td>O14</td>
<td>Pre-eclampsia</td>
</tr>
<tr>
<td>O15</td>
<td>Eclampsia</td>
</tr>
<tr>
<td>O24</td>
<td>Diabetes mellitus in pregnancy</td>
</tr>
<tr>
<td>O30</td>
<td>Multiple gestation</td>
</tr>
<tr>
<td>O31.2</td>
<td>Continuing pregnancy after intrauterine death of one fetus or more</td>
</tr>
<tr>
<td>O36.4</td>
<td>Maternal care for intrauterine death</td>
</tr>
<tr>
<td>O42</td>
<td>Premature rupture of membranes</td>
</tr>
<tr>
<td>O46</td>
<td>Antepartum haemorrhage</td>
</tr>
<tr>
<td>O75.5</td>
<td>Delayed delivery after artificial rupture of membranes</td>
</tr>
<tr>
<td>O75.7</td>
<td>Vaginal delivery following previous caesarean section</td>
</tr>
<tr>
<td>P95</td>
<td>Fetal death of unspecified cause</td>
</tr>
<tr>
<td>Q codes</td>
<td>Reportable congenital abnormalities</td>
</tr>
</tbody>
</table>


The iterative three-phase screening process is illustrated in Figure 3.
Box 5 indicates the SPSS syntax used in the three phases of filtering, identifying different indicators of potential obstetric risk, in order to generate a sample of women whose pregnancies were low-risk regardless of their planned place of birth.

### Box 5: SPSS syntax for identifying low-risk pregnancies

<table>
<thead>
<tr>
<th>Filtering</th>
<th>Syntax</th>
<th>Explanation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Phase 1</td>
<td>COMPUTE filter_$=((pdc_gestage &gt;= 37 &amp; pdc_gestage&lt;42) &amp; (Bdob_Yr &gt;= 2000 &amp; Bdob_Yr&lt;=2012) &amp; pdc_labons = 1 &amp; Non_Singleton_OK=0). VARIABLE LABELS filter_$ '(pdc_gestage &gt;= 37 &amp; pdc_gestage&lt;42) &amp; (Bdob_Yr &gt;= 2000 &amp; Bdob_Yr &lt;= 2012) &amp; pdc_labons = 1 &amp; Non_Singleton_OK=0 (FILTER)'. VALUE LABELS filter_$ 0 'Not Selected' 1 'Selected'. FORMATS filter_$ (f1.0). FILTER BY filter_$. EXECUTE.</td>
<td>Selected singleton born between 2000 and 2012, spontaneous onset in labour, inclusive 37 to 41 gestation weeks</td>
</tr>
<tr>
<td>Phase 2</td>
<td>IF Presentation = “Breech” or Delivery = “Breech” or prev_csbirth=”Yes” Exclude1=1.</td>
<td>Excluded vaginal breech and previous caesarean section</td>
</tr>
<tr>
<td>Phase 3</td>
<td>IF INDEX(Diag_code,&quot;O10&quot;)&gt;0 or INDEX(Diag_code,&quot;O11&quot;)&gt;0 or INDEX(Diag_code,&quot;O13&quot;)&gt;0 or INDEX(Diag_code,&quot;O14&quot;)&gt;0 or INDEX(Diag_code,&quot;O15&quot;)&gt;0 or INDEX(Diag_code,&quot;O24&quot;)&gt;0 or INDEX(Diag_code,&quot;O30&quot;)&gt;0 or INDEX(Diag_code,&quot;O31.2&quot;)&gt;0 or INDEX(Diag_code,&quot;O36.4&quot;)&gt;0 or INDEX(Diag_code,&quot;O42&quot;)&gt;0 or INDEX(Diag_code,&quot;O46&quot;)&gt;0 or INDEX(Diag_code,&quot;O75.7&quot;)&gt;0 or INDEX(Diag_code,&quot;P95&quot;)&gt;0 or INDEX(Diag_code,&quot;Q&quot;)&gt;0</td>
<td>Elimination of high-risk ICD codes: O10, O11, O13, O14, O15, O24, O30, O31.2, O36.4, O42, O46, O75.7, P95, all Q codes</td>
</tr>
</tbody>
</table>
In order to scrutinise outcomes on perinatal mortality, we conducted a line-by-line investigation of individual de-identified records of all deaths from births planned in birth centres and at home, and a random one-in-ten sample of planned hospital births that resulted in mortality. This highlighted a number of factors that indicated additional risk; some of these had not been detected by previous screening processes and were then added to the codes applied to the whole dataset. Given the complexity of the data and the recognised gaps in administrative data for the purposes of health research (especially for home births), it was essential to carry out manual validation to complement aggregate data preparation.

**RESULTS**

Over the period investigated by the BIA study, there were 3 171 800 births across Australia [49]. This figure represents all births in Australia 2007-2012, plus births 2000-2006 in NSW, Victoria, WA, SA, ACT and NT, and from 2005-2006 in Tasmania (Queensland and Tasmania provided data for limited time periods).

Overall, the BIA study received linked data about 2 524 329 births from the DLUs, which met the initial selection criteria to varying degrees. Figure 4 illustrates the effect of the various
phases of screening and filtering, demonstrating how the sample size decreased with the increasing specificity of requirements for a low-risk cohort.

*Figure 4 here: Refinement of sample size following three phases of data screening by state and territory>*

The various stages of cleaning, validating and screening records left 1,039,478 women remaining whose pregnancies were identified as low-risk by eliminating complicating conditions. The sample included data on outcomes from 1,251,420 births to women with full-term, singleton pregnancies without complications between 2000 and 2012 (or 2005-2012 in Tasmania and 2007-2012 in Queensland). The final low-risk cohort contained half the number of records originally received, following the multiple screening processes.

Table 1 presents the sources of data received from each jurisdiction and its contribution to the final master dataset.

**Table 1: Data sources, timeframe and contribution to national dataset, by state and territory**

<table>
<thead>
<tr>
<th>State or Territory</th>
<th>Data source</th>
<th>Years data available</th>
<th>Proportion of Australian births¹</th>
<th>Proportion of final dataset</th>
</tr>
</thead>
<tbody>
<tr>
<td>Australia Capital Territory (ACT)</td>
<td>ACT Death and Cause of Death Unit Record File</td>
<td>13 years (2000-2012)</td>
<td>1.9%</td>
<td>1.9%</td>
</tr>
<tr>
<td></td>
<td>ACT Perinatal Data Collection</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>ACT Admitted Patients Care</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>ACT Registrar of Births, Deaths and Marriages</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>New South Wales (NSW)</td>
<td>NSW Perinatal Data Collection</td>
<td>13 years</td>
<td>31.9%</td>
<td>40.5%</td>
</tr>
<tr>
<td></td>
<td>NSW Admitted Patient Data Collection</td>
<td></td>
<td></td>
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</tr>
<tr>
<td></td>
<td>NSW Register of Births, Deaths and Marriages</td>
<td></td>
<td></td>
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</tr>
<tr>
<td>State</td>
<td>Data Collection</td>
<td>Timeframe</td>
<td>Percentages</td>
<td></td>
</tr>
<tr>
<td>-----------------------------</td>
<td>---------------------------------------------------------------------------------</td>
<td>-----------------</td>
<td>-------------</td>
<td></td>
</tr>
<tr>
<td>Northern Territory (NT)</td>
<td>SA NT DataLink Perinatal Trends Inpatient Client Master Index Death Registry</td>
<td>13 years</td>
<td>1.3%</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>1.3%</td>
<td></td>
</tr>
<tr>
<td>Queensland</td>
<td>Queensland Perinatal Data Collection Queensland Health Admitted Patient Data Collection Queensland Registrar General Australian Bureau of Statistics - Deaths</td>
<td>6 years (2007-2012)</td>
<td>20.4%</td>
<td>9.1%</td>
</tr>
<tr>
<td>South Australia (SA)</td>
<td>SA NT DataLink Perinatal Maternal Family Link Birth, Death and Marriage, SA</td>
<td>13 years</td>
<td>6.6%</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>5.5%</td>
<td></td>
</tr>
<tr>
<td>Tasmania</td>
<td>Tasmanian Data Linkage Unit Cause of Death Unit Record File</td>
<td>8 years (2005-2012)</td>
<td>1.9%</td>
<td>1.6%</td>
</tr>
<tr>
<td>Victoria</td>
<td>Victorian Perinatal Data Collection</td>
<td>13 years</td>
<td>25.1%</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>29.6%</td>
<td></td>
</tr>
<tr>
<td>Western Australia (WA)</td>
<td>Hospital Morbidity Data System Midwives Notification System - Mortality</td>
<td>13 years</td>
<td>10.9%</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>10.5%</td>
<td></td>
</tr>
</tbody>
</table>

1 Hilder et al 2014. *Australia’s Mothers and Babies 2012*, AIHW

Table 1 indicates that the two states which provided data covering a shorter timeframe were under-represented in the final master file. Two other states were accordingly over-represented.

Figure 2 illustrates the complex process of developing the national master file from data sources from all states or territories. It indicates the variables for which data were available nationally.

**DISCUSSION**
The BIA study uniquely and ambitiously aimed to merge linked data from six states and two territories that used diverse methods for recording, storing and naming variables. The study entailed a complex and time-consuming process to verify, match, screen and clean data, and to ensure comparable cohorts. This paper has described several intricate procedures used to enhance the compatibility and integrity of data used to compare the outcomes for women who planned to give birth in three birth settings.

Data linkage is valuable for increasing the size and utility of datasets, especially to examine uncommon outcomes such as mortality [22] which are critical in evaluating the evidence on birthplace safety. However, the process is beset with challenges arising from datasets that are not always complete or compatible, and whose original purpose is not research [30].

Linked data offer enormous potential for exploring birth outcomes, albeit with an added degree of difficulty in a federation such as Australia where states and territories are responsible for relevant datasets [50]. To compare outcomes from different birth settings meaningfully, the BIA study endeavoured to distinguish compatible cohorts of women with similar low-risk pregnancies. Inconsistency in variables and data quality hampered this task. Eliminating obstetric complexity from the sample necessitated increasing the technical and analytical complexity to generate the most appropriate dataset.

Refining the Australia-wide master file over the stages of data preparation resulted in a final dataset of more than 1 million women with low-risk pregnancies, approximately two-fifths (39.5%) of all births during the study period [49]. This proportion highlights how the various phases of data screening greatly reduced the study denominator by limiting the sample to low-risk pregnancies as described.
As well as reducing the number of women in the sample, data linkage between eight jurisdictions also reduced the number of variables available for analysis. Differences between variables necessitated the most basic approach for the combined data. For example, not all jurisdictions distinguished between neonatal intensive care units and special care nurseries for unwell newborns, so we combined admissions to both facilities to ensure data consistency. Some states did not record data on specific items, meaning that we either did not report them or analysed a smaller sample for those variables; for example, maternal mortality outcomes were only available for six jurisdictions. At times, the process of adopting the lowest common denominator also potentially eroded the quality of data from some variables. Intended place of birth was a variable critical to our analysis as it is vital to exclude, for example, unintended home births without skilled birth attendants. However, in most states, women’s intended place of birth was recorded at an unspecified time in the pregnancy and not necessarily updated in administrative records. This is less ideal than documenting intention at onset of active labour given the potential for complicating conditions to occur throughout pregnancy. However, the other screening processes would have largely excluded those women who developed risk factors prior to labour onset and who required a hospital birth.

Limitations

This paper is limited in presenting only some of the techniques used to generate accurate and appropriate data for the BIA study. Other minor procedures were necessary to increase the integrity of the sample. However, this paper aimed to address several important considerations in using linked perinatal data in Australia and to provide guidance for future researchers attempting similar ventures.
The procedures to eliminate obstetric complexity in the dataset may have failed to detect all high-risk pregnancies as demonstrated by our final manual scrutiny of mortality data. A review of studies validating perinatal datasets suggested that diagnoses were less effectively recorded than procedures; it identified under-reporting of hypertension and diabetes amongst pregnant women [29]. Further, the selected ICD-10-AM codes for screening may have overlooked other complicating conditions.

Given these data are not collected primarily for research, they are dependent on the quality of data collection and entry by healthcare professionals and administrative staff.

**CONCLUSION**

This paper has illustrated the unique contribution that data linkage can make to understanding the impact of planned place of birth on maternal and perinatal outcomes, while expanding on the challenges involved. By offering practical guidance to help overcome common difficulties, this study aims to contribute to knowledge and research practice using this methodology.

The complexity and variability of data encountered in this study highlight the urgent need for more effective, transparent and uniform methods to collect and share healthcare data across Australian states and territories.
REFERENCES


