Original article

Evidence of detrimental effects of prenatal alcohol exposure on offspring birthweight and neurodevelopment from a systematic review of quasi-experimental studies

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Abstract

Background: Systematic reviews of prenatal alcohol exposure effects generally only include conventional observational studies. However, estimates from such studies are prone to confounding and other biases.

Objectives: To systematically review the evidence on the effects of prenatal alcohol exposure from randomized controlled trials (RCTs) and observational designs using alternative analytical approaches to improve causal inference.


Selection criteria: RCTs of interventions to stop/reduce drinking in pregnancy and observational studies using alternative analytical methods (quasi-experimental studies e.g. Mendelian randomization and natural experiments, negative control comparisons) to determine the causal effects of prenatal alcohol exposure on pregnancy and longer-term offspring outcomes in human studies.

Data collection and analysis: One reviewer extracted data and another checked extracted data. Risk of bias was assessed using customized risk of bias tools. A narrative synthesis of findings was carried out and a meta-analysis for one outcome.
Main results: Twenty-three studies were included, representing five types of study design, including 1 RCT, 9 Mendelian randomization and 7 natural experiment studies, and reporting on over 30 outcomes. One study design–outcome combination included enough independent results to meta-analyse. Based on evidence from several studies, we found a likely causal detrimental role of prenatal alcohol exposure on cognitive outcomes, and weaker evidence for a role in low birthweight.

Conclusion: None of the included studies was judged to be at low risk of bias in all domains, results should therefore be interpreted with caution.

Systematic review registration: This study is registered with PROSPERO, registration number CRD42015015941

Key words: Alcohol, pregnancy, prenatal alcohol exposure, systematic review, quasi-experimental studies, negative control, Mendelian randomization, causal inference, neurodevelopment, FASD

Introduction
The effects of prenatal alcohol consumption have typically been studied using standard analytical approaches in observational studies.1 Systematic reviews have used these types of studies to determine the effects of prenatal alcohol exposure on several outcomes with a wide range and varying definition of alcohol intake including low-moderate to binge drinking. Outcomes such as central auditory disorders in children,2 orofacial clefts,3 speech and language4 and several birth outcomes including low birthweight, preterm birth and small for gestational age1,5,6 have been investigated. These have led to varying results from systematic reviews: an increased risk of detrimental outcomes at very heavy drinking levels,1,2 inconsistent evidence regarding effects of moderate, heavy, or binge drinking (5+ drinks on any occasion),3 inconsistent effects from low-moderate alcohol consumption (up to 83 g/week)5 and some evidence that even light prenatal alcohol consumption is associated with harmful birth outcomes (up to 32 g/week).6 However, estimates from such studies are prone to the effects of: (i) confounding by sociodemographic characteristics (age, ethnicity, education, socio-economic position) and behavioural factors (smoking and substance use) and (ii) measurement error, namely under-reporting of alcohol intake and/or recall bias. Therefore, the direction and size of any potential causal relationships cannot be determined without bias.

In recent decades, novel analytical approaches have been increasingly applied to data from observational studies in order to improve causal inference when assessing potential effects of prenatal alcohol exposure. These approaches include Mendelian randomization (MR),7 family-based designs such as paternal or sibling comparison studies8 and natural experiments.9 Their respective strengths and limitations are outlined in Box 1.

We conducted a systematic review of human studies that used experimental data (randomized controlled trials (RCTs)) or alternative analytical methods to improve causal inference applied to observational data, in order to determine the causal effects of maternal alcohol consumption in pregnancy on offspring outcomes at birth and later in life. Additionally, as is being recognised elsewhere,11–13 it is important in public health and in epidemiology to include work from other disciplines in order to avoid missing important contributions to the literature. We therefore present a co-citation analysis to evaluate whether studies of alcohol

Key Messages
• Systematic reviews of prenatal alcohol exposure effects generally only include conventional observational studies. However, estimates from such studies are prone to confounding and other biases.
• We conducted a comprehensive systematic review of experimental human data and alternative analytical approaches to improve causal inference based on observational data.
• We also developed customized risk of bias tools for Mendelian randomization, natural experiments and parental and sibling comparison, and applied them to studies with these designs.
• Our results showed a likely causal detrimental role of prenatal alcohol exposure on cognitive outcomes, and weaker evidence for a decrease in birthweight, confirming results from conventional observational studies.
• Guidance should continue to advise abstention from alcohol in pregnancy.
<table>
<thead>
<tr>
<th>Causal inference approach</th>
<th>Definition</th>
<th>Biases addressed</th>
<th>Strengths</th>
<th>Limitations</th>
</tr>
</thead>
<tbody>
<tr>
<td>Randomized controlled trial (RCT)</td>
<td>Subjects are randomly allocated to either exposure or control groups with assumption that there is no difference between the two groups except for the intervention they are receiving</td>
<td>Confounding, reverse causality, selection bias, loss-to-follow up bias (using intention-to-treat analysis), measurement error</td>
<td>Gold standard for estimating causal effects. Any effect is very likely to be causal if study has large number and trial is reliably performed</td>
<td>Generalizability may be questionable; impossible or unethical to randomize to certain exposures; can be expensive</td>
</tr>
<tr>
<td>Mendelian randomization (MR)</td>
<td>MR is the use of a genetic variant robustly associated with an exposure/risk factor of interest as an instrumental variable to test and estimate the causal effect of that exposure/risk factor with a disease or health related outcome</td>
<td>Confounding by shared genetics and environment, reverse causality, selection bias, measurement errors</td>
<td>Genetic instruments are not subject to confounding from environmental or lifestyle factors, are not influenced by the outcome, do not change over time and are measured with high accuracy</td>
<td>Low power, lack of instrumentation/weak instruments, horizontal pleiotropy, population stratification, inability to assess non-linear associations/dose–response estimation, assortative mating, not specific to the intrauterine period, developmental canalization, subject to dynastic effects bias</td>
</tr>
<tr>
<td>Sibling comparison</td>
<td>Compares outcomes when siblings are discordant for an exposure. If causal, then there will evidence of a difference in outcome in relation to discordant exposure levels within sibships</td>
<td>Confounding by genetics and environment, specificity of effect to intrauterine period</td>
<td>Improves causal inference of intrauterine exposures. Controls for familial background and related confounding factors</td>
<td>Assumes a stable family environment; potential for confounding by factors not perfectly shared by siblings; potential for measurement error of exposure; limited power</td>
</tr>
<tr>
<td>Natural experiment</td>
<td>Empirical study approach where a population is exposed to an external event or intervention at a specific time point. Associations are then compared with a similar cohort that was not exposed. The assumption is that exposure is caused by quasi-random assignment</td>
<td>Confounding by genetics and environment, reverse causality, specificity of effect to intrauterine period</td>
<td>Can include study settings which would be impractical or unethical to produce by researchers. Allows for long-term effects of the exposure of interest</td>
<td>Selection bias if treatment and control group are not sufficiently comparable, some unobserved confounding may remain, it may not be possible to study non-linear associations/dose–response estimation</td>
</tr>
<tr>
<td>Parental comparison</td>
<td>Maternal–child association is compared with paternal–child association for inferring causal effect of intrauterine exposure. If causal, maternal association is stronger than paternal association. Where associations are similar for both parents we assume that they are driven by genetic or postnatal environmental characteristics</td>
<td>Confounding by genetics and environment, specificity of effect to intrauterine period</td>
<td>Improves causal inference of intrauterine effect if exposures are measured in both parents at same time in pregnancy, and non-paternity is taken into account for phenotypic traits</td>
<td>Assumption that paternal exposures share same confounding structure as maternal exposures may not be correct; where parental associations are of similar magnitude this may be due to offsetting paternal pathways rather than shared confounding. Assumes no assortative mating</td>
</tr>
</tbody>
</table>
in pregnancy carried out in other disciplines, such as health economics, are currently being recognised in public health.

Methods

Selection criteria and search strategy

The protocol for this systematic review, carried out using PRISMA guidelines, is available from the PROSPERO systematic review register (registration number CRD42015015941; http://www.crd.york.ac.uk/PROSPERO/display_record.asp?ID=CRD42015015941.

We reported results from prospective observational studies on low-moderate consumption, adopting standard analytical approaches, in a separate manuscript. Here, we focus on RCTs and studies that used alternative analytical methods to improve causal inference (see Box 1). MR studies that only reported results of geneXenvironment analyses (i.e. stratified by levels of maternal alcohol consumption) were excluded, as these estimates may incur selection bias.

We adopted study specific definitions for all outcomes. Outcomes included the following. (i) Pregnancy outcomes: still birth [pregnancy loss after week 24, miscarriage, gestational length and preterm delivery (<37 weeks gestation)]; hypertensive disorders of pregnancy; gestational diabetes; small for gestational age (SGA, <10th percentile in weight or <−2 standard deviation scores) and birth size [weight (including low birth weight defined as <2500 g), length and head circumference]; low amniotic fluid (oligohydramnios); placenta previa; placental abruption; assisted delivery (including vacuum extraction, forceps delivery, Caesarean section); Apgar score at birth; admission to neonatal unit; congenital malformations. (ii) Features of fetal alcohol spectrum disorder (FASD): childhood growth restriction; cranial size and head circumference; developmental delays; behaviour problems; cognitive impairment and intelligent quotient (IQ); facial malformations.

The databases that were searched included: MEDLINE, PsychINFO, EMBASE on Ovid; the Cochrane Library including CENTRAL (the Cochrane Central Database of Controlled Trials) on Wiley Interscience; and Science Citation Index, Social Science Citation Index, on Web of Science from inception to 21 June 2018 (Supplementary Table 1, available as Supplementary data at IJE online). The search was limited to papers in English and excluded letters, animal studies, editorials and conference proceedings without corresponding full-text papers. Investigators tailored searches to each database. The search did not include grey literature and was focused on published medical literature. Additionally, we performed manual searches of the reference lists of: (i) papers included in recent systematic reviews of the effects of prenatal alcohol exposure on the outcomes of interest; and (ii) all recent papers citing those reviews.

Titles and abstracts, and full texts if necessary, were screened independently by two reviewers. Discrepancies were discussed between reviewers and resolved through consensus.

Data extraction

A custom-built Microsoft Access database was used to extract data. The following information from each study was extracted: title, authors, publication year, country/region, population characteristics (sample size, methods of sampling, age distribution, and ethnicity), study design, measures of exposure, assessment methods for outcomes (including whether this was derived from medical records, obtained via a research interview and the person reporting the outcome e.g. parent, teacher, health professional, researcher or child), model adjustments, and study results. If a study reported more than one result for each outcome, we extracted all of them (e.g. relative to different timing of exposure, model adjustments, etc.). Information from each included paper was extracted by the lead reviewer (L.M.) and subsequently checked for accuracy and completeness by another reviewer (H.B.E.). There were very few extraction errors and these were resolved through discussion between extractor and checker.

Data analysis

Odds ratios (OR) and 95% confidence intervals (CI) were derived from count data from individual studies, if they were not reported. Studies were meta-analysed if they used the same analytical approach and estimated the same outcome (e.g. MR analyses of the same genotype-outcome association, discordant siblings’ analyses looking at the same outcome, etc.). The I² statistic was used to determine percentage of variation due to heterogeneity. Where only two studies were available to meta-analyse, results were not pooled if they were very different from each other. Alternatively, a narrative summary of the results was given.

Risk of bias assessment

The Cochrane risk of bias tool was selected to explore risk of bias in eligible randomized control studies.

There are currently no widely accepted risk of bias assessment tools for the alternative observational study designs included in this systematic review (MR, sibling comparison, paternal comparison and natural experiments). We therefore considered the previous work in this
and pleiotropy. Because none of the studies are at low
genetic (two studies rated at high risk), genetic confound-

Having a weak instrument. Further concerns were non-
differential assessment of exposure was the main concern in
both studies. All MR studies were rated at moderate risk of
having a weak instrument. Further concerns were non-
genetic (two studies rated at high risk), genetic confound-

Results
A flowchart of the article review process is shown in Fig. 1.
A total of 5424 citation records were identified from
searching the four relevant databases. A manual search of
recent systematic reviews identified 34 additional articles.
After exclusions, 9 MR analyses, 6 negative control stud-
ies, 1 RCT and 7 papers based on natural experiments
were included, giving a total of 23 studies.

Risk of bias assessment
Table 1 shows the results of risk of bias assessments. No
study was rated low risk of bias in all domains. The RCT
was judged at low risk of bias in all except in the blinding
domain as participants were not blinded and self-reported
their alcohol use. For natural experiment studies the main
concerns with regard to validity were the differential trends
in outcome, instrument strength and selection bias. For pa-
ternal comparison studies potential for differential paternal
and maternal confounding and non-paternity were the key
threats to validity. In the 2 sibling-comparison studies dif-
ferential assessment of exposure was the main concern in
both studies. All MR studies were rated at moderate risk of
having a weak instrument. Further concerns were non-
genetic (two studies rated at high risk), genetic confound-

Mendelian randomization studies
We identified 9 MR studies examining the effects of prena-
tal alcohol exposure on pregnancy or offspring outcomes
(Table 2). All studies used known variants in alcohol
dehydrogenase (ADH) genes in mothers and/or offspring
as genetic proxies for the exposure: 5 employed a func-
tional variant in ADH1B, 21–23,27,29 2 a haplotype in
ADH1C, 24,25,28 and 2 a number of ADH variants com-
bined into an allele score 26,29 (Table 2). The ADH1B vari-

Figure 2 illustrates patterns of journal co-citations. It
shows four main journal clusters including (health) eco-

omics, clinical/alcohol research, genetics and epidemi-

ology. The journal with the highest citation is ‘Alcoholism:
Clinical and Experimental Research’. The two other jour-
nals with the highest tendency for co-citation are
genetics and epidemiology. The (health) economics cluster
has a weaker tendency for co-citation and is the most
isolated. The weak cross-disciplinary citation between health

economics and other public health/epidemiology/clinical
journals could be due to several reasons including differences
in the speed of publication as well as in the frequency of
citations.

Co-citation
data were collected from Web of Science. These data were analysed using VOSviewer version 1.6.5.
Weights/bubble size correspond to the strength of co-cita-

tion. The distant between bubbles corresponds to the num-
ber of times that journals are cited together in other
journals. The colours correspond to ‘communities’ (cluster-
ing) identified by the software, and not pre-specified sci-
entific disciplines.
gene alleles on oral cleft. Two case-control studies examined the risk of oral cleft, comparing faster with slower metabolizers according to ADH1C maternal and fetal genotype. A French study found evidence of lower risk of non-syndromic cleft for ADH1C*2*2 compared with 1*1 homozygotes, but did not report on whether genotype groups differed by alcohol consumption. The study from Norway found no evidence of association with either offspring cleft risk or maternal alcohol consumption (Table 2).

Pregnancy outcomes
A study of African American infants found no strong evidence of association between infant ADH1B genotype and measures of birth size and gestational age, but did not report levels of maternal alcohol use by genotype (Table 2).
Features of FASD

The US-based study by Stoler et al. found some evidence of higher odds of a FASD-like construct in offspring carrying the ADH1B*3 allele compared with *1*1 homozygotes (Table 2). The latter metabolize alcohol more slowly and were also reported to have been exposed to lower levels of alcohol in pregnancy. The same direction of effect was observed comparing offspring of mothers carrying ADH1B*3, and the evidence was stronger for those of black ethnicity. Another study on fetal alcohol syndrome (FAS), from South Africa, found evidence of lower risk comparing carriers of maternal (or fetal) (fast metabolizing and lower alcohol intake) ADH1B*2 with ADH1B*1*1 homozygotes (slower metabolizers and higher intake), and little evidence of an effect of ADH1B*3 on FAS, in a mixed-ancestry South African population (Table 2). This study did not report on genotype–alcohol use association.

Other outcomes

The four most recent (and by far the largest) MR studies reported on cognitive and behavioural childhood outcomes in the same UK-based cohort (Table 2). Two used multiple offspring ADH variants known to be expressed in fetal life. One of these found evidence of association with IQ at 8 years old, but not when using the maternal allele score; the effects were stronger for children of mothers reporting some alcohol consumption, but there was no evidence of association between the allele score and maternal alcohol use per se. The other study did not find an association between maternal genotype ADH1B*2 and an increased risk of children having early-onset-persistent...
<table>
<thead>
<tr>
<th>Study ID</th>
<th>Random sequence generation</th>
<th>Allocation concealment</th>
<th>Blinding of participants/personnel</th>
<th>Blinding of outcome assessment</th>
<th>Incomplete outcome data</th>
<th>Selective reporting</th>
<th>Other bias</th>
</tr>
</thead>
<tbody>
<tr>
<td>Tzilos 2011</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Some concerns (Moderate risk)</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Low risk</td>
</tr>
</tbody>
</table>

### Risk of bias assessment for Natural experiment studies

<table>
<thead>
<tr>
<th>Study ID</th>
<th>Confounding 1—similar populations</th>
<th>Confounding 2—different trends in outcome</th>
<th>Confounding 3—population-level cointervention</th>
<th>Instrument strength</th>
<th>Assessment bias—outcome</th>
<th>Selection bias</th>
</tr>
</thead>
<tbody>
<tr>
<td>Barreca 2015</td>
<td>Low risk</td>
<td>High risk</td>
<td>Low risk</td>
<td>Unclear risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
</tr>
<tr>
<td>Evans 2016</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Unclear risk</td>
<td>Low risk</td>
<td>High risk</td>
</tr>
<tr>
<td>Fertig 2009</td>
<td>Low risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
</tr>
<tr>
<td>Nilsson 2017</td>
<td>Low risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
</tr>
<tr>
<td>Zhang 2010</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
</tr>
<tr>
<td>Zhang 2011</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
</tr>
<tr>
<td>Cil 2017</td>
<td>Low risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Low risk</td>
</tr>
</tbody>
</table>

### Risk of bias assessment for Parental Comparison studies

<table>
<thead>
<tr>
<th>Study ID</th>
<th>Confounding 1—paternal and maternal confounding similar</th>
<th>Confounding 2—timing of both parents’ exposure</th>
<th>Confounding 3—nonpaternity</th>
<th>Confounding 4—paternal effect</th>
<th>Assessment bias—exposure</th>
</tr>
</thead>
<tbody>
<tr>
<td>McCormack 2018</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
</tr>
<tr>
<td>Alati 2008</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Low risk</td>
</tr>
<tr>
<td>Alati 2013</td>
<td>High risk</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Low risk</td>
</tr>
<tr>
<td>Zuccolo 2016</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Low risk</td>
</tr>
</tbody>
</table>

### Risk of bias assessment for Sibling Comparison Studies

<table>
<thead>
<tr>
<th>Study ID</th>
<th>Confounding 1—birth order</th>
<th>Confounding 2—individual level</th>
<th>Assessment bias—exposure</th>
<th>Assessment bias—outcome</th>
<th>Selection Bias</th>
</tr>
</thead>
<tbody>
<tr>
<td>D’Onofrio 2007</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>High risk</td>
<td>Moderate risk</td>
<td>Low</td>
</tr>
<tr>
<td>Eliertsen 2017</td>
<td>Low risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low</td>
</tr>
</tbody>
</table>

### Risk of bias assessment for Mendelian Randomization studies

<table>
<thead>
<tr>
<th>Study ID</th>
<th>Weak instrument bias</th>
<th>Genetic confounding</th>
<th>Non-genetic Confounding (e.g. lifestyle factors)</th>
<th>Pleiotropy—additional direct effects btw IV &amp; outcome</th>
<th>Selection Bias</th>
</tr>
</thead>
<tbody>
<tr>
<td>Arfsten 2004</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>High risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
</tr>
<tr>
<td>Lewis 2012</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
</tr>
<tr>
<td>Boyle 2010</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Low risk</td>
</tr>
<tr>
<td>Chevrier 2005</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Low risk</td>
</tr>
<tr>
<td>Stoker 2002</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>High risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
</tr>
<tr>
<td>Viljoen 2001</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
<td>Moderate risk</td>
</tr>
<tr>
<td>Zuccolo 2013</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Moderate risk</td>
<td>Low risk</td>
<td>Low risk</td>
</tr>
</tbody>
</table>

(Continued)
behavioural problems, however this may be due to lack of statistical power (Table 2).\textsuperscript{29} The other two studies both used the functional ADH1B variant, and found some evidence that the offspring of mothers genetically predisposed to consuming less alcohol had better academic performance at ages 7, 11, 14 and 16, but no association between offspring genotype and their educational outcomes,\textsuperscript{28} nor was there evidence for an effect of genotype on IQ.\textsuperscript{27} Both studies reported lower alcohol consumption in mothers carrying the rare ADH1B*2 allele compared with the ADH1B*1*1 homozygotes.

Sibling comparison studies
Two sibling-comparison studies compared behavioural outcomes in siblings differentially exposed to alcohol in utero (Table 2).

Features of FASD
The study from the USA examined externalizing problems (measured through the Behaviour Problem Index) at ages 4–11 and found evidence that siblings exposed to moderate levels of prenatal alcohol had higher rates of conduct problems compared with their unexposed siblings, however there was no evidence of differences in attention or impulsivity problems.\textsuperscript{30} The more recent study from Norway compared differentially exposed siblings in terms of their attention-deficit hyperactivity disorder (ADHD) at 5 years of age.\textsuperscript{31} Results differed slightly depending on the ADHD scale used, with evidence of increased prenatal alcohol exposure being associated with higher ADHD levels according to the revised Conner’s Parent Rating Scale, but less strong evidence for the Child Behaviour Rating Checklist.\textsuperscript{31}

Parental comparison studies
Four maternal–paternal comparison studies met our inclusion criteria. These investigated the effects of prenatal alcohol exposure on neurocognitive domains in offspring: childhood educational achievement,\textsuperscript{33} IQ,\textsuperscript{32} cognitive development\textsuperscript{35} and head circumference\textsuperscript{34} (Table 2).

Features of FASD
Two reports from the same UK-based study found no evidence of association between regular maternal alcohol use in pregnancy and either school results at 11\textsuperscript{33} or IQ at 8 years of age.\textsuperscript{32} One of the studies did find some evidence that increased levels of maternal binge drinking in pregnancy (consuming 32+g alcohol/occasion) were associated with decreased school results at age 11 years, whereas paternal exposure was associated with improved school results.\textsuperscript{33} The other report did not find the same level of
Table 2. Quasi-experimental studies examining the effects of prenatal alcohol exposure (PAE) on pregnancy and childhood outcomes. *3 Denotes rare allele found in sub-Saharan African populations; *2 denotes rare allele found in European ancestry populations; *1 denotes common allele found worldwide

<table>
<thead>
<tr>
<th>Study (year)</th>
<th>Country</th>
<th>Sample size</th>
<th>Exposure definition (proxy for PAE)</th>
<th>Exposure categories</th>
<th>Age at outcome assessment (child)</th>
<th>Outcomes</th>
<th>Summary of results &amp; conclusions as presented in the paper</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mendelian randomization studies</td>
<td>Viljoen et al. (2001)21</td>
<td>South Africa</td>
<td>56 Mother–child pairs</td>
<td>MR</td>
<td>5–9 years</td>
<td>Fetal alcohol syndrome (FAS)</td>
<td>Fetal genotype: <em>3</em> vs <em>1</em> OR 0.85 (95% CI: 0.26, 2.72)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>178 Controls</td>
<td>Khoisan–Caucasian mixed ancestry</td>
<td></td>
<td></td>
<td>Maternal genotype: <em>3</em> vs <em>1</em> OR 0.85 (95% CI: 0.26, 2.72)</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Stoler et al. (2002)22</td>
<td>USA</td>
<td>173 White women</td>
<td>MR</td>
<td>At birth</td>
<td>Blinded physician assessment of a composite trait including: growth restriction, microcephaly or 4 of 6 predefined facial features typical of FAS</td>
<td>Maternal genotype (White women): OR <em>3</em> vs <em>1</em>: 6.68 (95% CI: 0.89, 49.90)</td>
</tr>
<tr>
<td></td>
<td></td>
<td>85 Black women</td>
<td>MR</td>
<td></td>
<td></td>
<td>Maternal genotype (Black women): OR <em>3</em> vs <em>1</em>: 2.62 (95% CI: 0.97, 7.04)</td>
<td>Fetal genotype (White children): OR <em>3</em> vs <em>1</em>: 4.75 (95% CI: 0.35, 64.74)</td>
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<td></td>
<td>Arfsten et al. (2004)23</td>
<td>USA</td>
<td>305 African American infants</td>
<td>MR</td>
<td>Birth weight (g)</td>
<td>Birth weight (g)</td>
<td>Fetal genotype (Black children): OR <em>3</em> vs <em>1</em>: 3.75 (95% CI: 1.00, 14.02)</td>
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<td>GA (weeks)</td>
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<td>LBW (&lt;2500g)</td>
<td>Birth weight (g)</td>
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<td>SGA</td>
<td>Birth weight (g)</td>
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<td>Chevrier et al. (2005)24</td>
<td>France</td>
<td>205 Case children + parents (195 fathers and 211 mothers)</td>
<td>MR</td>
<td>Birth weight (g)</td>
<td>Birth weight (g)</td>
<td>Fetal genotype: <em>3</em> vs <em>1</em>: 3.211 vs 31.96, P=0.83</td>
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<td>115 Control children and 54 mothers</td>
<td>MR</td>
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<td>GA (weeks)</td>
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<td>SGA</td>
<td>Birth weight (g)</td>
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<td>Boyles et al. (2010)25</td>
<td>Norway</td>
<td>Mother–child pairs (483 cases 503 controls)</td>
<td>MR</td>
<td>Birth weight (g)</td>
<td>Birth weight (g)</td>
<td>Fetal genotype: <em>3</em> vs <em>1</em>: 3.96 vs 39.4, P=0.37</td>
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<td>SGA</td>
<td>Birth weight (g)</td>
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<td>Lewis et al. (2012)26</td>
<td>UK</td>
<td>4 167 Mother–child pairs</td>
<td>MR</td>
<td>Birth weight (g)</td>
<td>Birth weight (g)</td>
<td>Fetal genotype: <em>3</em> vs <em>1</em>: 0.93 (95% CI: 0.50, 1.90)</td>
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<td>SGA</td>
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<td>Zucolo et al. (2013)27</td>
<td>UK</td>
<td>6268 Mother–child pairs for KS2 analysis</td>
<td>MR</td>
<td>Birth weight (g)</td>
<td>Birth weight (g)</td>
<td>Fetal genotype: <em>3</em> vs <em>1</em>: 0.93 (95% CI: 0.50, 1.90)</td>
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<td>von Hinke Kessler Scholder et al. (2014)²⁸</td>
<td>UK</td>
<td>4061 pairs for WISC analysis</td>
<td>Mother-child pairs</td>
<td>MR</td>
<td>Gene: ADH1B SNP-rs number: rs1229984</td>
<td><em>1</em>1 Slow metabolizers (ref. category) <em>2</em> Fast metabolizers</td>
<td>7–16 years</td>
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<td>Murray et al. (2016)²⁹</td>
<td>UK</td>
<td>8621 Sibling comparison</td>
<td>Maternal genotype analysis: 3114 mother-child pairs</td>
<td>MR</td>
<td>Gene: ADH1B SNP-rs number: rs1229984</td>
<td><em>1</em>1 Slow metabolizers (ref. category) <em>2</em> Fast metabolizers</td>
<td>4–13 years</td>
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<tr>
<td>D’Onofrio et al. (2007)³⁰</td>
<td>USA</td>
<td>8621 Sibling comparison</td>
<td>Siblings discordant for PAE Moderate alcohol exposure vs none</td>
<td>Moderate alcohol exposure vs none</td>
<td>4–11 years</td>
<td>Conduct problems Attention/impulsivity problems: Moderate vs no alcohol intake: MD 0.05 SE 0.02 Moderate vs no alcohol intake: MD 0.03 SE 0.02 (β = 0.017, 95% CI: 0.005, 0.030) (β = 0.011, 95% CI: -0.002, 0.024) Attention-deficit hyperactivity disorder (ADHD) Scales: Moderate vs no alcohol intake: MD 0.05 SE 0.02 Moderate vs no alcohol intake: MD 0.03 SE 0.02</td>
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<tr>
<td>Eikelen et al. (2017)³¹</td>
<td>Norway</td>
<td>34 283 Sibling comparison</td>
<td>Siblings discordant for PAE Not available</td>
<td>Not available</td>
<td>5 years</td>
<td>Conduct problems Attention/impulsivity problems: Moderate vs no alcohol intake: MD 0.05 SE 0.02 Moderate vs no alcohol intake: MD 0.03 SE 0.02 (β = 0.017, 95% CI: 0.005, 0.030) (β = 0.011, 95% CI: -0.002, 0.024) Attention-deficit hyperactivity disorder (ADHD) Scales: Moderate vs no alcohol intake: MD 0.05 SE 0.02 Moderate vs no alcohol intake: MD 0.03 SE 0.02 (β = 0.017, 95% CI: 0.005, 0.030) (β = 0.011, 95% CI: -0.002, 0.024)</td>
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<tr>
<td>Alati et al. (2008)³²</td>
<td>UK</td>
<td>43 32 Maternal-paternal comparison</td>
<td>Mother and partner alcohol intake</td>
<td>Mother and partner alcohol intake</td>
<td>First trimester (regular use): never</td>
<td>IQ: Wechsler Intelligence Scale for Children (WISC) 1st trimester regular alcohol use (pdiff = 0.43) MD per increase in maternal category: 0.03 (−0.58, 0.65) MD per increase in paternal category: 0.40 (−0.01, 0.02)</td>
<td>Parental control studies</td>
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<td>Mother and partner alcohol intake</td>
<td>First trimester (regular use): never</td>
<td>11 years</td>
<td>Academic achievement: Key Stage 2 scores (standardized)</td>
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<td>1–4 drinks/occasion</td>
<td>5–10 drinks/occasion</td>
<td>10+ drinks/occasion</td>
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<td>MD per increase in maternal category: –0.45 (–1.32, 0.43)</td>
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<td>MD per increase in paternal category: 0.10 (–0.36, 0.56)</td>
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<tr>
<td>Zuccolo et al. (2016)34</td>
<td>Norway</td>
<td>46 178</td>
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<td>Mother and partner alcohol intake</td>
<td>Non-drinker (ref)</td>
<td>At birth and at 3 months post-partum</td>
<td>Head circumference</td>
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<td>&lt; 1 unit/week</td>
<td>1–2 units/week</td>
<td>3–4 units/week</td>
<td>5+ units/week</td>
<td>At birth: Fully and mutually adjusted model:&lt;br&gt; 1st trimester regular Alcohol use (PDiff = 0.406)</td>
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<td>Mother: mean difference (SD) 0.00 (95% CI: –0.02, 0.02)</td>
<td>Father: mean difference (SD) = 0.00 (95% CI: –0.05, 0.04)</td>
<td>1–2 units- Mother: mean difference (SD) = –0.02 (95% CI: –0.05, 0.01)</td>
<td>Father: mean difference (SD) = 0.01 (95% CI: –0.03, 0.05)</td>
<td>MD per increase in maternal category: –0.68 (–1.03, –0.33)</td>
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<td>3–4 units- Mother: mean difference (SD) = 0.06 (95% CI: 0.02, 0.11)</td>
<td>Father: mean difference (SD) = 0.01 (95% CI: –0.03, 0.05)</td>
<td>5+ units- Mother: mean difference (SD) = 0.01 (95% CI: –0.04, 0.06)</td>
<td>Father: mean difference (SD) = –0.01 (95% CI: –0.04, 0.03)</td>
<td>MD per increase in paternal category: 0.27 (0.07, 0.46)</td>
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<td>At 3 months post-partum: fully and mutually adjusted model:&lt;br&gt; &lt; 1 unit- Mother: mean difference (SD) = 0.02 (95% CI: –0.00, 0.05)</td>
<td>Father: mean difference (SD) = –0.02 (95% CI: –0.08, 0.04)</td>
<td>1–2 units- Mother: mean difference (SD) = –0.02 (95% CI: –0.05, 0.02)</td>
<td>Father: mean difference (SD) = –0.03 (95% CI: –0.08, 0.02)</td>
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<td>3–4 units- Mother: mean difference (SD) = 0.04 (95% CI: –0.02, 0.10)</td>
<td>Father: mean difference (SD) = –0.04 (95% CI: –0.09, 0.01)</td>
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<td>McCormack et al. (2018)</td>
<td>Australia</td>
<td>2030</td>
<td>Material–paternal comparison</td>
<td>Mother and partner alcohol intake</td>
<td>First 6 weeks: Abstinent, Low, Moderate, Binge, Heavy; Second 6 weeks: Abstinence</td>
<td>12 months</td>
<td>Infant cognitive development (Bayley Scales for Infant Development, third edition)</td>
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<td>Fertig and Watson (2009)</td>
<td>USA</td>
<td>All women: 16 165 747; Native White women: 11 426 203</td>
<td>Changes in minimum legal drinking age (MLDA)</td>
<td>Lower (18 years) vs higher (19–21 years) MLDA</td>
<td>At birth</td>
<td>Low birth weight (&lt;2500g)</td>
<td>MLDA of 18 main effect: -0.17% (SE 0.07)</td>
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<td>Preterm birth (&lt;37 weeks)</td>
<td>MLDA of 18 vs mother ≥7 years of age interaction: 0.30% (SE 0.18)</td>
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<td>Congenital anomalies</td>
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<td>Native Black women</td>
<td>USA</td>
<td>303 108</td>
<td>MLDA of 18 x mother 18–20 years of age interaction; 0.26% (SE 0.10)</td>
<td>MLDA of 18 main effect; -0.35% (SE 0.09)</td>
<td>Mean of outcome: 7.5</td>
<td>MLDA of 18 x mother 18–20 years of age interaction; 0.26% (SE 0.10)</td>
<td>MLDA of 18 main effect; -0.35% (SE 0.09)</td>
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<td>Martin et al. (2010)</td>
<td>USA</td>
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<td>MLDA of 18 x mother 18–20 years of age interaction; 0.26% (SE 0.10)</td>
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<td>USA</td>
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<td>Effects of different MLDA (18–21 years), when woman is 14 (e.g. proxying for different alcohol availability)</td>
<td>MLDA of 18 main effect; -0.19% (SE 0.08)</td>
<td>MLDA of 18 x mother 18–20 years of age interaction; 0.26% (SE 0.10)</td>
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<td>Effects of different MLDA (18–21 years), when woman is 14 (e.g. proxying for different alcohol availability)</td>
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<td>Barreca and Page (2015)</td>
<td>USA</td>
<td>14–17 years at conception 3 314 000</td>
<td>Differences in MLDA Lower (18 years) vs higher (19–21) MLDA</td>
<td>At birth Low birthweight (&lt;2 500 g)</td>
<td>Effects of different MLDA (18–21 years), when woman is 14 (e.g. proxying for different alcohol availability)</td>
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<td>21–24 years at conception 10 178 000</td>
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<td>Evans et al. (2016)</td>
<td>USA</td>
<td>1 704 191 (Education sample) 198 518 (Obesity/height sample)</td>
<td>Effect of state alcohol prohibitions</td>
<td>States with alcohol prohibitions vs states without prohibitions</td>
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<td>Exposed children had: ( \text{MLDA of 18 x mother 18–20 years of age interaction; 0.04% (SE 0.09)} ) Mean of outcome: 10.7 ( \text{MLDA of 18 main effect; 0.39% (SE 0.32)} ) ( \text{MLDA of 18 x mother 14–17 years of age interaction; 0.46% (SE 0.53)} ) ( \text{MLDA of 18 x mother 18–20 years of age interaction; 0.19% (SE 0.26)} ) Mean of outcome: 90.1 4</td>
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<td>Cal (2017)</td>
<td>USA</td>
<td>Birth weight: 60 914 264 Pre-term: 53 276 541 FAS: 28 371 025 APGAR: 48 291 613</td>
<td>Effect of point-of-sale warnings about risks of drinking during pregnancy</td>
<td>States with warnings vs states without warnings (including pre-post-intervention within states)</td>
<td>At birth</td>
<td>Low birth weight (&lt;2500 g) Very low birth weight (&lt;1500 g) Pre-term (&lt;37 weeks) Very pre-term (&lt;32 weeks) FAS Low APGAR (&lt;7)</td>
<td>( \text{0.115% (SE 0.082) decreased odds of low birth weight (P &gt; 0.1)} ) ( \text{0.047% (SE 0.023) decreased odds of very low birth weight (P &lt; 0.05)} ) ( \text{0.065% (SE 0.14) decreased odds of pre-term birth (P &gt; 0.1)} ) ( \text{0.052% (SE 0.029) decreased odds of very pre-term birth (P &gt; 0.1)} ) ( \text{0.003% (SE 0.002) decreased odds of FAS (P &gt; 0.1)} ) ( \text{0.014% (SE 0.047) decreased odds of low APGAR (P &gt; 0.1)} ) Exposed children had: ( \text{APGAR; 48 291 613} )</td>
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<td>Nilsson (2017)</td>
<td>Sweden</td>
<td>35 3742</td>
<td>Relancing the regulation of alcohol sales</td>
<td>Children born in counties with relaxed regulation</td>
<td>Mean age 32 years</td>
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<td>Randomised controlled trial</td>
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<td>Intervention group 27</td>
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<td>Children born to mothers receiving intervention on prenatal alcohol use vs to mothers receiving standard care</td>
<td>At birth</td>
<td>Birthweight</td>
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<td>Intervention group: mean = 3189.6, SD = 328.0</td>
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<td>Control group: mean = 2965.3, SD = 387.7</td>
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</table>

LBW, low birth weight; ELBW, extremely low birthweight; GA, gestational age; APGAR, appearance, pulse, grimace, activity, respiration; MLDA, minimum legal drinking age; NICU, neonatal intensive care unit; OR, odds ratio; CI, confidence interval; MD, mean difference; SD, standard deviation; SGA, small for gestational age; IQ: intelligence quotient; RR, relative risk.

a Unweighted allele score composed of the four fetal SNPs.

b Adjustments: mother’s ancestry principle components from Genome wide association studies (GWAS) analysis.

c Adjustments: sex, other parent’s alcohol consumption, maternal age, parity, socio-economic position, ethnicity, and, maternal and paternal education and smoking.

d Test for trend maternal and paternal alcohol intake in full model: $P = 0.267$ and $P = 0.201$.

e Test for trend maternal and paternal alcohol intake in full model: $P = 0.390$ and $P = 0.124$.

f Test for trend maternal and paternal alcohol intake in full model: $P = 0.545$ and $P = 0.056$.

g Test for trend maternal and paternal alcohol intake in full model: $P = 0.178$ and $P = 0.090$.

h Adjustments: state fixed effects, year-month fixed effects, maternal age fixed effects, state-specific time trends and birth characteristic controls.

i Adjustments: state fixed effects, year fixed effects, mother’s education, age, marital status, smoking during pregnancy, real income per capita and real beer taxes (federal plus state level).

j Adjustments: state fixed effects, year-by-month fixed effects, age fixed effects, state-specific trends, age-by-year fixed effects, state-by-age fixed effects and state-by-year fixed effects.
evidence to support an association of prenatal binge drinking with offspring IQ at age 8 years. In a large Norwegian cohort, there was no evidence of association between maternal or paternal alcohol use during or before pregnancy and head circumference at birth or 3 months. In the same study, odds of microcephaly increased with higher paternal but not maternal alcohol consumption prior to pregnancy and in the first trimester. A recent Australian study showed no consistent evidence of association between maternal alcohol use in different trimesters of gestation and cognitive function in children aged 1 year (Bayley Scales of Infant Development), and even scatter evidence for partner alcohol intake.

Natural experiments
Seven reports analysed data from natural experiments involving changes in government laws that effected the availability or affordability of alcohol, or required point-of-sale warnings about the risks of drinking alcohol during pregnancy (Table 2). The studies by Fertig and Watson and Barreca and Page were based on US-wide birth data and estimated the association between MLDA and low birthweight (<2500 g), preterm delivery (<37 weeks) and congenital anomalies, with the latter additionally examining Apgar scores. Both used a triple difference approach (Supplementary Material, available as Supplementary data at IJE online) and substantially the same data, although the latter study ran additional analyses with more covariates and interaction terms to check the robustness of the model to some of its assumptions. When running similar age-specific analyses, the second study replicated the first study’s results of an increase in both preterm deliveries and low birthweight corresponding to a lowering of MLDA, more marked for babies conceived to younger (<18 year old) compared with older (18–20 year old) women. In more fully adjusted analyses, the negative association with birthweight was still found to be robust for younger mothers (<18 years). However, no consistent evidence of association was found for other age groups in the main effects analyses, or for other adverse fetal outcomes including gestational age, congenital abnormalities and Apgar score. Neither study reported data on actual population-level alcohol use. The third study, by Zhang and Caine (2011), investigated the same outcomes (low birthweight, preterm delivery and Apgar scores) in relation to a State’s MLDA at the time a woman is 14 years old. The difference with

Pregnancy outcomes
Three US-based studies used reductions in the minimum legal drinking age (MLDA) to proxy for prenatal alcohol exposure, under the assumption that a lower MLDA would increase alcohol availability to young women (Table 2).
respect to the two previous studies was that the ‘exposed’
status is assigned based on MLDA at the time the women
are 14 years, regardless of what it is when she is older and
pregnant. The authors hypothesize that the drinking envi-
ronment at age 14 sets a woman’s future ‘drinking propen-
sity’ including binge drinking behaviour, but no data were
reported to confirm this. The estimates were derived from
difference-in-difference specifications, but with additional
controls for State-specific effects. The authors presented
evidence that women who lived in a State where the
MLDA was 18 years at the time they themselves were 14
years, compared with those in States with higher MLDA,
had higher chances of giving birth to low birthweight
babies with lower Apgar scores, but no association with
prematurity.38

A fourth paper examined the effect of within-State
changes in alcohol taxation in the US and within-State var-
iation in birthweight and Apgar scores37 (Table 2). The
authors found evidence that increases in alcohol taxes are
associated with increases in birthweight and Apgar scores.
The authors also tried to validate their assumptions that
changes in taxation are a valid proxy for alcohol consump-
tion and therefore prenatal alcohol exposure, by regressing
several alcohol drinking variables from a federal behaviou-
rnal survey on alcohol taxation. They found some evidence
of reduced binge drinking behaviour among pregnant
women, corresponding to increases in alcohol taxes,
however no evidence that the quantity consumed was sen-
titive to alcohol pricing.37

Another US-based study explored the impact of State
laws requiring point-of-sale warnings about the risks of
drinking alcohol during pregnancy on outcomes including
birthweight, pre-term birth, FAS and Apgar scores.41
There was evidence that the warnings reduced the chances
of very low birth weight babies (<1500 g), but no evidence
of association with the other outcomes. The authors vali-
dated their assumption that alcohol warning signs would
reduce prenatal alcohol exposure by regressing several al-
cohol drinking variables on whether the State prescribed
health warnings or not, using both individual birth and na-
tional survey data. They found that adoption of the law
was associated with a reduction in alcohol consumption
and binge drinking among pregnant women.

Features of FASD

Two studies looked at long-term offspring outcomes
(Table 2). Based on data from World War II US enlistees,
the first study used different timings of prohibition imple-
mentation in different States to proxy for reduced likeli-
hood of prenatal alcohol exposure as a result of reduced
availability to women, and examined attained education
and height in adult offspring.40 The authors report an in-
crease in years of education associated with the introduc-
tion of prohibition, but no evidence of an effect on height.

| Table 3. Summary of direction of association of prenatal alcohol exposure with selected outcomes (cognitive/brain development and birthweight), in the context of expected and observed differences in prenatal alcohol exposure in each study |

<table>
<thead>
<tr>
<th>Outcome</th>
<th>Study</th>
<th>Direction PAE—outcome</th>
<th>Direction Exposure proxy—PAE</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cognition, brain develop-</td>
<td>Lewis et al. (2012) 26</td>
<td>↓</td>
<td>NA (Not Applicable) NA</td>
</tr>
<tr>
<td>ment</td>
<td>Zuccolo et al. (2013) 27</td>
<td>↑</td>
<td>↓</td>
</tr>
<tr>
<td></td>
<td>von Hinke Kessler Scholder et al. (2014) 28</td>
<td>↑</td>
<td>↓</td>
</tr>
<tr>
<td></td>
<td>Zuccolo et al. (2016) 24</td>
<td>↑</td>
<td>↓</td>
</tr>
<tr>
<td></td>
<td>McCormack et al. (2018) 35</td>
<td>↑</td>
<td>↓</td>
</tr>
<tr>
<td></td>
<td>Alati et al. (2008) 32</td>
<td>↑</td>
<td>↑</td>
</tr>
<tr>
<td></td>
<td>Alati et al. (2013) 33</td>
<td>↑</td>
<td>↑</td>
</tr>
<tr>
<td></td>
<td>Nilsson (2017) 42</td>
<td>↑</td>
<td>↑</td>
</tr>
<tr>
<td></td>
<td>Evans et al. (2016) 40</td>
<td>↑</td>
<td>↑</td>
</tr>
<tr>
<td>Birthweight</td>
<td>Arfsten et al. (2004) 23</td>
<td>↔</td>
<td>NA NA</td>
</tr>
<tr>
<td></td>
<td>Fertig and Watson (2009) 36</td>
<td>↓</td>
<td>↑</td>
</tr>
<tr>
<td></td>
<td>Barreca and Page (2015) 39</td>
<td>↓</td>
<td>↑</td>
</tr>
<tr>
<td></td>
<td>Zhang and Caine (2011) 38</td>
<td>↓</td>
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</tr>
<tr>
<td></td>
<td>Zhang (2010) 37</td>
<td>↑</td>
<td>↓</td>
</tr>
<tr>
<td></td>
<td>Cil (2017) 41</td>
<td>↑</td>
<td>↑</td>
</tr>
<tr>
<td></td>
<td>Tzilos et al. (2011) 43</td>
<td>↑</td>
<td>↓</td>
</tr>
</tbody>
</table>

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However, there were no estimates of actual alcohol consumption in States introducing prohibition.  

A Swedish study compared earnings, education and welfare dependency rates in children born in counties that did and did not relax the regulation of alcohol sales in 1967. The relaxation of alcohol policy, used as a proxy for increased prenatal alcohol exposure, was shown to be related to reduced earnings, years of schooling and high school completion rates, as well as to a higher proportion of individuals on welfare. The author reported some evidence of increased consumption of alcohol for the counties during the period where the more liberal policy applied, but no results specifically for pregnant women.

Randomized controlled trial

We included one RCT feasibility study with a small sample size (control group 23 women, intervention group 27 women; Table 2).

Pregnancy outcomes

In the RCT feasibility study, 50 pregnant women who screened positive for risky drinking were randomized: 27 pregnant women in the intervention group received a 20-min computer-based, self-administered program intended to motivate them to reduce their drinking, whereas 23 pregnant women in the control group received a questionnaire about television preferences. Follow-up after 1 month (average 33 days) showed no difference in alcohol use between the intervention and control groups but some evidence of higher birthweight for infants born to women in the intervention group compared with the control group. As there was no strong evidence of a difference in alcohol consumption between the randomized groups this does not support any causal effect of alcohol on birthweight but may suggest bias in the RCT, some pathways (other than change in alcohol) from the intervention to birthweight that might counter any effect of alcohol and/or too little power to detect effects on alcohol robustly.

Discussion

Summary of the evidence

Our systematic review of the literature found a limited number of studies addressing the effects of prenatal alcohol exposure using experimental designs or alternative analytical strategies to improve causal inference in observational studies, which we described in narrative format. Twenty-three reports were included, representing five types of study design, with MR and natural experiments the most common designs (9 and 7 studies, respectively). Cognitive outcomes were the most commonly reported (by 9 studies), followed by birthweight (7 studies). The overall picture that emerges from this review is that moderately strong evidence exists for detrimental effects of prenatal alcohol exposure on cognitive outcomes (Table 3). For cognitive outcomes and birth weight outcomes, we found the highest degree of consistency across study types (MR,26–28 parental comparisons33 and natural experiments exploiting different policy changes40,41) as well as with the direction of association predominantly reported in conventional epidemiological studies. Based on natural experiments36–39 and one feasibility RCT,43 some evidence was also found for reduced birthweight following higher prenatal alcohol exposure (Table 3), in line with recent reviews6 and pooled analyses of observational studies.49

Only one outcome-study design combination had more than one result that could be combined into a meta-analysis. For the rest, we described results in narrative format. We also developed and deployed customized risk of bias (RoB) assessment tools for the different types of study design. None of the studies scored ‘low’ RoB in all domains, therefore we recommend caution in interpreting the results of any one study as ‘causal’, since it is impossible to predict the overall direction of bias affecting each result.

Results of our co-citation analysis showed that the field of (health) economics is relatively isolated compared with the other clusters. It also shows a limited number of studies in public health. This shows that the findings published in health economics journals are not well recognised in the fields of epidemiology and public health, although the evidence they contribute should be considered alongside that from more traditional epidemiological studies when updating public health guidance on alcohol use, as evidenced by our reviewing efforts.

Strengths and limitations of alternative study designs

An extensive literature exists exploring the strengths and limitations of the observational study designs and analytical strategies19 included in this review, especially when applied to the study of intergenerational effects such as here.50,51 In theory, all study types attempt to minimize confounding by shared genetic and environmental factors by design, all but MR and some of the natural experiments address the specificity of the effect to the intrauterine period (i.e. not confounded by postnatal alcohol use), and MR and natural experiments avoid reverse causality (Box 1). In practice, sources of bias varied both across and within each study-type category, as evidenced by our customized RoB tools showing some of the included studies
being at higher risk of bias than others. For example, data availability may restrict the extent to which one can test and/or account for potential differential trends in studies exploiting natural experiments such as MLDA. Similarly, data availability may restrict the extent to which one can explore whether (in particular historic) policies affected prenatal alcohol consumption. This is also true for many of the (particularly older) MR studies that did not report genotype associations with maternal alcohol use. Furthermore, ensuring that the analytical strategy identifies effects that are specific to the intrauterine period may be difficult. For example, a reduction in the MLDA in the year of birth is likely to be related to alcohol exposure in that year, but potentially also in the year after. This is less of an issue in studies that exploit temporary changes in alcohol exposure, such as Nilsson, as temporary policies are more likely to only affect alcohol exposure at that point in time only.42 In MR studies, one analytical strategy that improves specific attribution of effects to the intrauterine period is using alcohol metabolizing genotypes in the offspring (not just the mothers) as proxy for prenatal alcohol exposure. This is because maternal genotype in theory predisposes to lower or higher alcohol use in pregnancy as well as before and after (therefore it is not specific to the intrauterine period). Additionally, MR studies of intrauterine exposures that do not account for both offspring and maternal genotype can suffer from bias because of violation of the exclusion restriction assumption.52 On the other hand, offspring genotype (conditional on maternal genotype) is more specific, since children do not consume alcohol themselves and the only time in early life where they are exposed to alcohol is in utero. Therefore, different alcohol metabolizing genotypes in the offspring could modulate prenatal alcohol exposure, independently of maternal alcohol use. This strategy of presenting results for offspring genotype adjusted for maternal genotype was only adopted by a couple of the included MR studies and has the additional advantage of minimizing dynastic effects bias.

An additional strength of some of the natural experiments included here is that they investigated possible mechanisms for the observed effects of prenatal alcohol exposure, in particular through a postulated increase in unplanned pregnancies (also known as ‘compositional changes’). This was explored through, e.g. sensitivity analyses to test whether MLDA changes resulted in more unplanned pregnancies. The idea is that, if MLDA led to an increase in unplanned pregnancies, this may have particularly affected mothers with a systematically different e.g. socio-economic position, whose children also have systematically different outcomes. But these effects are then driven by socio-economic confounding, not (necessarily) only by intrauterine toxicity. This was done by Fertig and Watson36 examining the percentage of births recorded with missing paternal information, with the analysis confirming evidence of effect for this in black women, and stronger effects in younger girls (<18 years), thus providing a possible partial explanation for the birthweight effects in their study. Compositional changes or changes in the demographics of mothers giving birth, are also thought to play a role in explaining some of the effect on adverse pregnancy outcomes observed in the study by Zhang.37 Specifically, since an increase in alcohol taxes appeared to lead to a reduction in pregnancies amongst younger and less educated mothers, who are more likely to experience adverse pregnancy outcomes, maternal age and education (over and above alcohol consumption per se), may explain some of the apparent effect of alcohol. The study by Nilsson42 was able to avoid potential bias due to possible compositional changes by focusing on children who were conceived prior to the start of the relaxation of alcohol policy. Hence, his study did not include children who were conceived due to the change in alcohol policy.

Another study by Barreca and Page39 additionally investigated the presence of an early selection effect that intrauterine alcohol exposure could have on the least healthy foetuses, by examining gender ratio of live births as a marker of early fetal loss. The authors’ interpretation, although highly speculative, is that this selection indeed is present and could explain the unexpected direction of effect for the main effect analyses in their study.

Small sample sizes in many of the studies (especially for the earlier studies) means that estimates were often imprecise. This was particularly true for the MR studies, some of which were among the first ever to be conducted, and none of which adopted a multi-cohort approach to increase sample size, or multiple genetic variants to improve the variance explained in alcohol consumption, as is recommended and customary in recent times.33

Another limitation of the MR and natural experiment studies that were included in this review is the inability to provide dose–response estimates. Instead, they provide estimates of the effect of prenatal alcohol exposure around mean levels of consumption in the study sample. This falls short of the most interesting research question which is whether the effects are linear or whether there is a threshold at low levels of drinking under which alcohol is not harmful to the fetus.

Additionally, for MR studies, only ADH variants have been used and there is a possibility that acetaldehyde is both the deterrent to drinking and the cause of damage, which could lead to null results. Many more loci affecting alcohol intake are now available for future studies, although their effect on prenatal alcohol use will require validation in studies of pregnant women.
Strengths and limitations of this systematic review

This systematic review is the first of its kind to explicitly search for and integrate the evidence from different study designs and analytical approaches in a true triangulation framework.\(^5\) Efforts were made to include studies from different disciplines for the first time, as evidenced by the results of our co-citation analysis (Fig. 2). Alongside this triangulation approach, strengths of this review include the pre-registered protocol http://www.crd.york.ac.uk/PROSPERO/display_record.php?ID=CRD42015015941, and the thorough assessment of RoB through deployment of customized RoB tools.

This literature could be affected by publication bias. Given the lack of sufficient numbers of studies to meta-analyse, we could not investigate publication bias through funnel plots, so it remains speculative whether further unpublished (negative) studies could exist. On the other hand, we notice a trend for a number of studies to attempt replication of certain (positive) seminal papers (e.g. Eilertsen et al.\(^3\)) replicating D’Onofrio et al.\(^2\) Barreira and Page\(^3\) replicating Fertig and Watson,\(^3\) Boyles et al.\(^2\) replicating Chevrier et al.\(^3\) and occasionally failing to replicate the original results. Being able to capture these failures to replicate is a strength of the current review.

The main limitation of the review derives from the nature of the evidence we found, the paucity and heterogeneity of which prevented us from pooling effects through meta-analysis. Instead, we systematically grouped results by outcome and study type and examined them for consistency.

Another important limitation is that we cannot infer causal dose–response relationships based on this body of evidence. For example, the effect estimates from MR studies using maternal genotypes effectively refer to the average difference in alcohol to which the offspring are exposed. Therefore, what we can infer is that (often small) increases in prenatal alcohol exposure are associated with lower neurocognitive outcomes and to a lesser extent lower birth-weight, based on studies that minimize confounding. The most pressing question of relevance to public health remains whether the recommendation to abstain from alcohol in pregnancy is backed by solid evidence, as opposed to being purely precautionary. We have already explored this extensively with our previous review of observational studies on the effects of low levels of drinking in pregnancy, which concluded that the abstinence advice was mainly a precaution.\(^6\) The research studies brought together by the current review add to this in a significant way, tipping the balance towards a more solid evidence-base, in particular for neurocognitive and behavioural outcomes.

Public health/policy and research implications

This review seeks to address an area of great public health impact. Alcohol use in pregnancy is still widespread worldwide,\(^5\) despite claims that it causes the most common neurodevelopmental impairments, included under the umbrella diagnosis of FASD.\(^5\) The claims of causality implicit in the diagnostic definition of FASD, however, have occasionally been disputed (e.g. McLennan et al.\(^5\)) due to lack of robust evidence of specific alcohol effects on different domains in the child, and whether thresholds apply. This review highlights the need for more studies using a variety of analytical approaches to establish the extent to which prenatal alcohol exposure causes specific neurobehavioral outcomes in the offspring. Studies simultaneously addressing multiple sources of bias are particularly needed (e.g. MR exploiting trio data from fathers, mothers and offspring, to conduct both negative control MR with paternal effects, and analyses accounting for transmitted and un-transmitted alleles,\(^5\) and 2-samples MR using recently developed approaches to study intrauterine effects),\(^5\) as are studies allowing for more sophisticated dose–response estimation (e.g. Silverwood et al.\(^9\)).

This evidence will then feed into a revised and improved definition of FASD. Results from this review will also inform future reviews of guidelines on alcohol use in pregnancy. The current UK guidelines for example, revised down to abstinence in 2016,\(^6\) are heavily based on the precautionary principle. The present review of alternative study designs to improve causal inference will strengthen the evidence base for the abstinence recommendation, as well as highlighting the considerable gaps in evidence and quality of studies needed to move the field forward and draw firm conclusions.

Conclusion

Our understanding of the specific causal effects of alcohol in pregnancy, especially at low levels of exposure, is limited due to biases affecting traditional observational methods and the practical and ethical obstacles to conducting an RCT. Alternative study designs such as MR and natural experiments make an important contribution to our understanding of these effects, as they overcome some of the limitations of traditional methods. Currently, these comprise a modest body of evidence suggestive of a detrimental effect on cognitive outcomes and infant birthweight, which corroborate findings of conventional epidemiological studies. The studies included in this review do not provide evidence on whether the effect of alcohol exposure is linear, or whether there is a safe threshold for drinking in pregnancy, although many of them compare groups of
offspring with at most small differences in their prenatal alcohol exposure. Although it remains true that the only way to avoid alcohol-related risks to the fetus is to abstain from alcohol during pregnancy, it is also important to communicate both to mothers-to-be and healthcare professionals that there remains uncertainty in the evidence base for this recommendation, although we welcome the fact that more and more studies with complementary strengths and weaknesses are emerging in this field.

Supplementary data
Supplementary data are available at IJE online.

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Authors’ contributions
L.M.: drafting the protocol, drafting the risk of bias assessment tools, preparing the data extraction database, data screening and collection (lead reviewer), data analysis, figures, tables, data interpretation, and drafting the manuscript. T.J.: data extraction, data quality checking, and full text screening. S.I.: risk of bias reviewer, data quality checking, and provided methodological advice for the review and meta-analysis conduct. H.B.E.: screening, data collection, data quality checking. J.S.: protocol development/study design (including the development of searching, article screening and data collection strategies), oversaw and managed the review process and the review team, provided methodological advice for the review and meta-analysis conduct, and critically revised the manuscript. T.H.M.M.: advising on the protocol, preparing the protocol, screening title and abstract. G.D.S.: obtained funding, formulation of project plan, review of the manuscript. J.L.D.: obtained funding and contributed to drafting the manuscript and approved the final version. V.L.: data extraction. D.A.L.: risk of bias reviewer, contributed to data interpretation and review of earlier manuscript drafts, approved the final version that has been submitted. S.J.L.: contributed to data interpretation and review of earlier manuscript drafts, approved the final version that has been submitted. S.vH.: risk of bias reviewer, revised one of the risk of bias assessment tools, completed risk of bias reviews and review of manuscript draft. A.F.: risk of bias reviewer, contributed to the study design, reviewed manuscript draft. L.Z.: contributed to the study protocol, study design, data interpretation, drafted parts of the manuscript, designed and revised the risk of bias assessment tools, completed risk of bias reviews. L.M.has full access to all the data in the study and had final responsibility for the decision to submit for publication. The manuscript is an honest, accurate and transparent account of the study being reported, no important aspects of the study have been omitted and any discrepancies from the study as planned (and, if relevant, registered) have been explained.

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