Maternal genetic risk factors for spontaneous preterm birth: a systematic review and meta-analysis

Tea Mladenić¹, Anita Barišić², Nina Pereza¹, Saša Ostojić¹, Borut Peterlin³, and Sanja Dević Pavlić¹

¹Sveuciliste u Rijeci Medicinski fakultet ²Klinicki bolnicki centar Rijeka ³Univerzitetni klinicni center Ljubljana

April 19, 2024

Abstract

Background: Despite various genomic approaches used in prior studies investigating association of maternal genetic variability with spontaneous preterm birth (sPTB), results show inconsistency and contradictions. Objectives: To: conduct a systematic review of studies analysing the association between maternal genetic variants and sPTB; evaluate retrieved studies based on selection criteria; classify studies into hypothesis-based and hypothesis-free; perform a meta-analysis to identify the strongest associations. Search Strategy: PubMed, Scopus and reference lists were searched until October 2023. Selection Criteria: English-language case-control, cross-sectional and prospective cohort studies examining the association between maternal genetic variations and sPTB were included. Data collection and Analysis: Data on authors, publication year, ethnicity, genes/variants, P-values, study type, sample size, inclusion criteria and methods were collected. The association strength was estimated using odds ratios with 95% confidence intervals. Main Results: 81 studies met eligibility criteria: 72 utilized a hypothesis-based and 9 a hypothesis-free approach. 34 studies qualified for a meta-analysis revealing a significant association in $TNF-\alpha$ (rs1800629) gene for alleles, additive and recessive genetic models (P < 0.05). From the hypothesis-free approach, 7 variants in 5 genes (EBF1, EESEC, HSPA1L, ASTN1, MAST1) reached global significance ($P < 5 \times 10^{-8}$). Conclusions: No specific genes or variants were clearly associated with the risk of sPTB. Among hypothesis-based studies, limited gene overlap indicates inconsistent SNP associations. TNF- α (rs1800629) emerges as the only with a modest signal for future analyses. Additional 5 genes from the hypothesis-free approach showed a globally significant association. Funding: / Keywords: Preterm Birth, Genetic Association Study, Genome-Wide Association Study, Exome Sequencing

Hosted file

BJOG manuscript upload.docx available at https://authorea.com/users/771656/articles/856013maternal-genetic-risk-factors-for-spontaneous-preterm-birth-a-systematic-review-andmeta-analysis

Hosted file

Figure 1 BJOG.docx available at https://authorea.com/users/771656/articles/856013-maternal-genetic-risk-factors-for-spontaneous-preterm-birth-a-systematic-review-and-meta-analysis

Hosted file

Table 1 BJOG.docx available at https://authorea.com/users/771656/articles/856013-maternal-genetic-risk-factors-for-spontaneous-preterm-birth-a-systematic-review-and-meta-analysis

Hosted file

Table 2 BJOG.docx available at https://authorea.com/users/771656/articles/856013-maternal-genetic-risk-factors-for-spontaneous-preterm-birth-a-systematic-review-and-meta-analysis

Hosted file

Table 3 BJOG.docx available at https://authorea.com/users/771656/articles/856013-maternal-genetic-risk-factors-for-spontaneous-preterm-birth-a-systematic-review-and-meta-analysis

Hosted file

Supp. mat. BJOG.docx available at https://authorea.com/users/771656/articles/856013-maternal-genetic-risk-factors-for-spontaneous-preterm-birth-a-systematic-review-and-meta-analysis

\backslash	Describe any methods used to assess certainty (or confidence) in the body of evidence for an outcome.	15	Certainty assessment	
6-7	Describe any methods used to assess risk of bias due to missing results in a synthesis (arising from reporting biases).	14	Reporting bias assessment	
200	Describe any sensitivity analyses conducted to assess robustness of the synthesized results.	13f		
U 1 an	Describe any methods used to explore possible causes of heterogeneity among study results (e.g. subgroup analysis, meta-regression).	13e		
5 1 00	Describe any methods used to synthesize results and provide a rationale for the choice(s). If meta-analysis was performed, describe the model(s), method(s) to identify the presence and extent of statistical heterogeneity, and software package(s) used.	13d		
5-2	Describe any methods used to tabulate or visually display results of individual studies and syntheses.	13c		
5,2	Describe any methods required to prepare the data for presentation or synthesis, such as handling of missing summary statistics, or data conversions.	13b		
5-20	Describe the processes used to decide which studies were eligible for each synthesis (e.g. tabulating the study intervention characteristics and comparing against the planned groups for each synthesis (item #5)).	13a	Synthesis methods	
4-8	Specify for each outcome the effect measure(s) (e.g. risk ratio, mean difference) used in the synthesis or presentation of results.	12	Effect measures	
6-7	Specify the methods used to assess risk of bias in the included studies, including details of the tool(s) used, how many reviewers assessed each study and whether they worked independently, and if applicable, details of automation tools used in the process.	1	Study risk of bias assessment	
5-6	List and define all other variables for which data were sought (e.g. participant and intervention characteristics, funding sources). Describe any assumptions made about any missing or unclear information.	10b		
2-6	List and define all outcomes for which data were sought. Specify whether all results that were compatible with each outcome domain in each study were sought (e.g. for all measures, time points, analyses), and if not, the methods used to decide which results to collect.	10a	Data items	
0)	Specify the methods used to collect data from reports, including how many reviewers collected data from each report, whether they worked independently, any processes for obtaining or confirming data from study investigators, and if applicable, details of automation tools used in the process.	Q	Data collection process	
5	Specify the methods used to decide whether a study met the inclusion criteria of the review, including how many reviewers screened each record and each report retrieved, whether they worked independently, and if applicable, details of automation tools used in the process.	œ	Selection process	
Figure 1, Tada	Present the full search strategies for all databases, registers and websites, including any fitters and limits used.	7	Search strategy	
S	Specify all databases, registers, websites, organisations, reference lists and other sources searched or consulted to identify studies. Specify the date when each source was last searched or consulted.	o	Information sources	
5,6	Specify the inclusion and exclusion criteria for the review and how studies were grouped for the syntheses.	J	Eligibility criteria	
C	Provide an explicit statement of the objective(s) of question(s) the review addresses.	4	Objectives	
10		-	Rationale	
2			INTRODUCTION	
2,3	See the PRISMA 2020 for Abstracts checklist.	2	Abstract	
			ABSTRACT	
-	Identify the report as a systematic review.	-	Title	
			TITLE	
Location where item is reported	Checklist item	ltem	Section and Topic	
	PRISMA 2020 Checklist	MA 20	PRIS	

1,____

de 51,5-6

Section and Topic	ltem #	Checklist item	Location where item
RESULTS			
Study selection	16a	Describe the results of the search and selection process, from the number of records identified in the search to the number of studies included in the review, ideally using a flow diagram.	ncluded in
	16b	Cite studies that might appear to meet the inclusion criteria, but which were excluded, and explain why they were excluded.	& Figure 1
Study characteristics	17	Cite each included study and present its characteristics.	Table 1-3,51, 8-11
Risk of bias in studies	18	Present assessments of risk of bias for each included study.	9-10, ST
Results of individual studies	19	For all outcomes, present, for each study: (a) summary statistics for each group (where appropriate) and (b) an effect estimate and its precision (e.g. confidence/credible interval), ideally using structured tables or plots.	orecision Table 1-3
Results of	20a	For each synthesis, briefly summarise the characteristics and risk of bias among contributing studies.	8-11, ST Table 2
syntheses	20b	Present results of all statistical syntheses conducted. If meta-analysis was done, present for each the summary estimate and its precision (e.g confidence/credible interval) and measures of statistical heterogeneity. If comparing groups, describe the direction of the effect.	on (e.g.
	20c	Present results of all investigations of possible causes of heterogeneity among study results.	-4-
	20d	Present results of all sensitivity analyses conducted to assess the robustness of the synthesized results.	-14-
Reporting biases	21	Present assessments of risk of bias due to missing results (arising from reporting biases) for each synthesis assessed.	9-10,S
Certainty of evidence	22	Present assessments of certainty (or confidence) in the body of evidence for each outcome assessed:	
DISCUSSION			
Discussion	23a	Provide a general interpretation of the results in the context of other evidence.	47-14
	23b	Discuss any limitations of the evidence included in the review.	11-14
	23c	Discuss any limitations of the review processes used.	4
	23d	Discuss implications of the results for practice, policy, and future research.	14-15
OTHER INFORMATION	TION		
Registration and	24a	Provide registration information for the review, including register name and registration number, or state that the review was not registered.	red. PLOSPERC
protocol	24b	Indicate where the review protocol can be accessed, or state that a protocol was not prepared.	
	24c	Describe and explain any amendments to information provided at registration or in the protocol.	
Support	25	Describe sources of financial or non-financial support for the review, and the role of the funders or sponsors in the review.	15
Competing interests	26	Declare any competing interests of review authors.	15
Availability of data, code and other materials	27	Report which of the following are publicly available and where they can be found: template data collection forms; data extracted from included studies; data used for all analyses; analytic code; any other materials used in the review.	cluded

From: Page MJ, McKenzie JE, Bossuyt PM, Boutron I, Hoffmann TC, Mulrow CD, et al. The PRISMA 2020 statement: an updated guideline for reporting systematic reviews. BMJ 2021;372:n71. doi: 10.1136/bmj.n71

A STATE

PRISMA 2020 Checklist