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## Article

# Effect of Probiotics on Pregnancy Outcomes in Gestational Diabetes: Systematic Review and Meta-Analysis

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**Abstract:** Background: Gestational diabetes is a common complication during pregnancy that can lead to numerous adverse outcomes. Some studies suggest that probiotics may be used to treat gestational diabetes, however, the results remain controversial. We conducted a systematic review and meta-analysis to evaluate the effect of probiotics on blood glucose and pregnancy outcomes in women with gestational diabetes. Methods: A systematic search of PubMed, Embase, and Cochrane databases was performed (start date to August 22, 2023). Primary outcomes included fasting blood sugar (FBS), fasting serum insulin (FSI), homeostasis model assessment of insulin resistance (HOMA-IR), and quantitative insulin sensitivity check index (QUICKI). Secondary outcomes included pregnancy and neonatal outcomes. Results: 15 articles (n = 1006 women) met the inclusion criteria and were included in the analysis. Compared to a placebo, probiotics can decrease FBS (MD -2.58, 95% CI -4.38 to -0.79,  $p < 0.01$ ), FSI (MD -2.29, 95% CI -3.40 to -1.18,  $p < 0.01$ ), HOMA-IR (MD -0.56, 95% CI -0.81 to -0.32,  $p < 0.01$ ), birthweight (MD -101.20, 95% CI -184.62 to -17.77,  $p = 0.02$ ), neonatal intensive care unit (NICU) (RR 0.60, 95% CI 0.40 to 0.89,  $p = 0.01$ ), and hyperbilirubinemia (RR 0.31, 95% CI 0.16 to 0.61,  $p < 0.01$ ), alongside higher QUICKI (MD 0.01, 95% CI 0.00 to 0.01,  $p < 0.01$ ). However, no other significant results were obtained. Conclusion: Probiotics may improve blood glucose indicators and reduce neonatal hyperbilirubinemia, NICU admissions, and birth weight in women with GDM.

**Keywords:** probiotics; pregnancy; GDM; gestational diabetes; meta analysis

## 1. Introduction

The most common metabolic complication of pregnancy is gestational diabetes mellitus (GDM), the incidence of which has been increasing significantly in the last decade, accounting for 12-18% of all pregnant patients [1]. Studies have shown that genetic, epigenetic and environmental factors can combine to cause GDM. Poor glycemic control during pregnancy may lead to adverse maternal and infant outcomes, such as maternal preeclampsia, hypertension, neonatal macrosomia, and hypoglycemia [2].

Therefore, the cornerstone of GDM management is glycemic control. Lifestyle interventions are used as the initial treatment for GDM and include medical nutrition therapy and daily exercise. Patients are asked to check their blood glucose levels frequently at home to ensure that blood glucose goals are being met. If these measurements do not meet glycemic goals, medication should be initiated [3].

Although women with gestational diabetes still make insulin, their bodies are not sensitive to it and do not produce enough insulin to maintain blood glucose control. Insulin is often the treatment of choice for women who are unable to maintain their blood glucose treatment goals through medical nutrition therapy or other pharmacological therapies. Insulin may be an alternative for women who cannot tolerate the side effects of oral antidiabetic medications such as metformin [4]. However, some serious side effects of insulin and oral hypoglycaemic drugs, such as hypoglycaemia and gastrointestinal symptoms, limit their use, and sometimes these treatments are more dangerous for mothers and babies.

The definition of probiotics is live microorganisms that provide health benefits when consumed in sufficient quantities. For decades, it has been discovered that probiotics can treat acute gastroenteritis, *Clostridium difficile*-associated diarrhoea, irritable bowel syndrome, and digestive issues, among others [5]. Some studies have shown that the pathogenesis of GDM may be related to abnormalities in the composition of the gut flora of pregnant women. Dietary supplements can play a role by regulating the intestinal microbiota. Therefore, the use of dietary supplements, such as probiotics, to target the intestinal flora and regulate the disordered intestinal flora would be a potential method for the prevention and treatment of GDM [6].

However, some relevant randomized controlled trials (RCTs) have shown that probiotics have no effect on fasting blood sugar (FBS), fasting serum insulin (FSI), and neonatal weight in patients with gestational diabetes [7,8]. Some meta-analyses of GDM include RCTs involving obese or overweight pregnant women, rather than those with GDM. Therefore, considering these inconsistent effects, we conducted a novel meta-analysis of RCTs to assess the utility of probiotics on glycaemic control, maternal and infant outcomes in patients with gestational diabetes.

## 2. Methods

This meta-analysis was conducted in accordance with the Preferred Reporting Items for Systematic Reviews and Meta-Analysis (PRISMA) [9]. This study was registered at PROSPERO (CRD42023456637).

### 2.1. Literature Search

A comprehensive literature search was conducted using PubMed, Embase and the Cochrane Library. The search was limited to the English language (up to 22 August 2023). Furthermore, we conducted a search on <http://www.clinicaltrials.gov/> to find information on registered RCTs in order to identify trials that were completed but had not yet published results. We primarily used the following search terms: "Gestational Diabetes Mellitus," "Probiotics," and "randomized controlled trials." We combined specific keywords and free text items with Boolean operators to create our search strategy.

### 2.2. Inclusion Criteria And Exclusion Criteria

The meta-analysis comprised studies that met the following criteria: fully open randomised controlled studies conducted according to English-language criteria; patients were women diagnosed with GDM in accordance with the American Diabetes Association, the World Health Organization, the Carpenter and Kushan Guidelines, and the Australian Pregnancy Association Consensus Guidelines; and interventions consisted of probiotics alone or therapies of probiotic-accessorised other substances; and the full text was available for review. The following studies were excluded: Ongoing trials whose results had not been presented or published; repetitive publications; and review articles, case reports, conference abstracts and guidelines.

### 2.3. Risk of Bias Assessment

Two independent investigators assessed the risk of bias in the literature. The recommended "risk of bias" tool from the Cochrane Collaboration was used.

### 2.4. Data Extraction

Each included study was extracted independently by two researchers. Any disagreements were discussed or negotiated with a third researcher. The main information we extracted included: first author, year of publication, country, sample size of each trial group, dose of probiotics and number of weeks of duration. We focused on three outcomes: glycaemic control, maternal outcomes and neonatal outcomes. Glycaemic outcomes included FBS, FSI, homeostasis model assessment of insulin resistance (HOMA-IR), and quantitative insulin sensitivity check index (QUICKI); maternal outcomes included pre-eclampsia, caesarean section, preterm labour, induction of labour and

polyhydramnios; neonatal outcomes included birth weight, neonatal length, neonatal head circumference, neonatal intensive care unit (NICU) admission, 1-min Apgar score, 5-min Apgar score, hyperbilirubinemia, macrosomia, large for gestational age (LGA), and neonatal hypoglycaemia.

### 2.5. Quality Assessment

Two researchers evaluated the risk of bias of the included clinical trials independently according to the the Cochrane Collaboration Tool [10]. We use Cochrane Handbook for Systematic Reviews of Interventions version 6.3 for quality evaluation. The following quality assessments were included: selection bias, performance bias, detection bias, attrition bias, reporting bias, and other bias. The results were classified as "low risk", "high risk" and "unclear risk". Disagreements at the time of quality assessment were resolved by discussion or referred to a third evaluator for adjudication.

### 2.6. Statistical Analysis

Probiotic efficacy was assessed by pooling prognostic indicators. Dichotomous data are expressed as relative risk ratios (RR) and 95% confidence intervals (CI); continuous data are expressed as mean difference (MD) and 95%CI, statistically significant differences were deemed as such when  $p < 0.05$ .

Statistical heterogeneity was assessed by chi-squared test and  $I^2$  statistic. When statistically significant heterogeneity was found ( $I^2 > 50\%$  or  $p < 0.1$ ), a random-effects model was used [11]. If not, fixed effects models are selected. All statistical calculations were conducted with Review Manager version 5.4 (Cochrane Collaboration, Software Update, Oxford, UK). The risk of bias was assessed with the Egger statistical method using Stata software 12.0 (StataCorp, based in College Station, Texas, USA). We assessed bias only for studies included in a sample of 10 or more. For outcomes with greater heterogeneity, we conducted subgroup analyses examining the effects of both country and duration.

## 3. Results

### 3.1. Study Selection and Characteristics

A total of 221 relevant articles were retrieved from various databases and 122 studies were obtained after removing duplicates. Subsequently, 104 irrelevant articles were removed because some of them were of the review type, non-RCTs, etc. The remaining 18 articles were reviewed in full text, of which 3 had insufficient data. Therefore, a total of 15 RCT articles[8,12–25] were finally selected for meta-analysis. **Error! Reference source not found.** depicted the search process, and



Table 1. Basic characteristics of eligible studies.

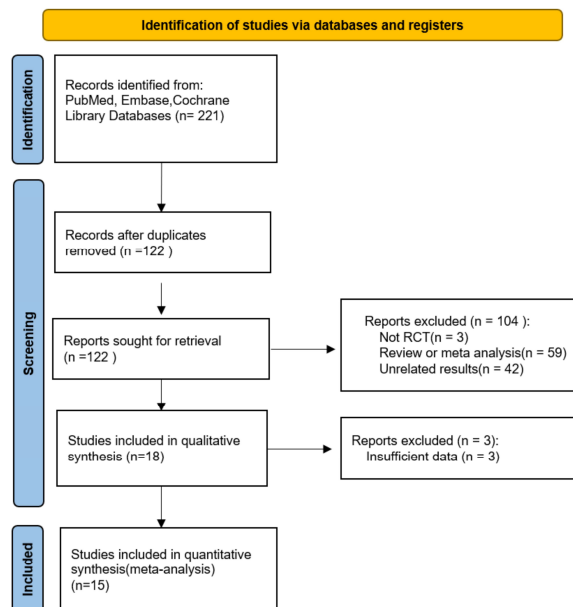
First authors	Year	Country	Probiotic Intervention	Control Intervention	Probiotic Dose	Probiotic group (N )	Control group ( N)	Duration (weeks)
Dolatkhah	2015	Turkey	<i>Lactobacillus acidophilus</i> LA-5, <i>Bifidobacterium</i> BB-12, <i>Streptococcus thermophilus</i> STY-31 and <i>Lactobacillus delbrueckii bulgaricus</i> LBY-27	placebo	4 biocap>4 × 109 CFU	29	27	8
Lindsay	2015	Ireland	<i>Lactobacillus salivarius</i> UCC118	placebo	1×109 CFU/g	48	52	8
Karamali	2016	Iran	<i>L. acidophilus</i> , <i>L. casei</i> and <i>B. bifidum</i> strains	placebo	2 × 109 CFU/g each	30	30	6
Jafarnejad	2016	Iran	VSL#3 ( <i>Streptococcus thermophilus</i> , <i>Bifidobacterium breve</i> , <i>Bifidobacterium longum</i> , <i>Bifidobacterium infantis</i> , <i>Lactobacillus acidophilus</i> , <i>Lactobacillus plantarum</i> , <i>Lactobacillus paracasei</i> , and <i>Lactobacillus delbrueckii</i> subsp. <i>bulgaricus</i> )	placebo	112.5 × 109 CFU	37	35	8
Ahmadi	2016	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus casei</i> , <i>Bifidobacterium bifidum</i> plus 0.8 g inulin	placebo	2 × 109 CFU/g each	35	35	6
Nabhani	2018	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus plantarum</i> , <i>Lactobacillus fermentum</i> , <i>Lactobacillus gasseri</i>	placebo	<i>L. acidophilus</i> (5 × 1010 CFU/g), <i>L. plantarum</i> (1.5 × 1010 CFU/g), <i>L. fermentum</i> (7 × 109 CFU/g), <i>L. gasseri</i> (2 × 1010 CFU/g)	45	45	6

Badehnoosh	2018	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus casei</i> and <i>Bifidobacterium bifidum</i>	placebo	2 × 10 <sup>9</sup> CFU/g each	30	30	6
Karamali	2018	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus casei</i> and <i>Bifidobacterium bifidum</i> strains plus 800 mg inulin	placebo	2 × 10 <sup>9</sup> CFU/g each	30	30	6
Hajifaraji	2018	Iran	<i>L.acidophilus</i> LA-5, <i>Bifidobacterium</i> BB-12, <i>Streptococcus thermophilus</i> STY-31 and <i>Lactobacillus delbrueckii bulgaricus</i> LBY-27 plus dextrose anhydrous filler and magnesium stearate lubricant	placebo	4 biocap >4×10 <sup>9</sup> CFU	29	27	8
Babadi	2019	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus casei</i> , <i>Bifidobacterium bifidum</i> , and <i>Lactobacillus fermentum</i>	placebo	2 × 10 <sup>9</sup> CFU/g each	24	24	6
Sahhaf Ebrahimi	2019	Iran	Probiotic yoghurt containing <i>Lactobacillus acidophilus</i> and <i>Bifidobacterium lactis</i>	placebo	300 g/day of probiotic yoghurt(contained 10 <sup>6</sup> CFU <i>Lactobacillus acidophilus</i> and <i>Bifidobacterium lactis</i>	42	42	8
Kijmanawat	2019	Thailand	<i>Lactobacillus acidophilus</i> and <i>Bifidobacterium bifidum</i>	placebo	1 × 10 <sup>9</sup> CFU/g each	28	29	4
Jamilian	2019	Iran	<i>Lactobacillus acidophilus</i> , <i>Bifidobacterium bifidum</i> , <i>Lactobacillus reuteri</i> , and <i>Lactobacillus fermentum</i>	placebo	8 × 10 <sup>9</sup> CFU/day	29	28	6

Amirani	2022	Iran	<i>Lactobacillus acidophilus</i> , <i>Bifidobacterium bifidum</i> , <i>Bifidobacterium lactis</i> <i>Bifidobacterium longum</i> Additionally , selenium	placebo	2 × 109 CFU/day each	26	25	6
Yefet	2022	Israel	<i>Bifidobacterium bifidum</i> , <i>Bifido-</i> <i>bacterium lactis</i> , <i>Lactobacillus</i> (L) <i>acidophilus</i> , <i>L. paracasei</i> , <i>L.rhamnosus</i> and <i>Streptococcus thermophilus</i>	placebo	2 capsules/day ( >6×109 CFU/capsule)	41	44	2

Abbreviations: CFU, colony-forming units.

listed the basic characteristics of eligible trials. At baseline, age, height and weight were not significantly different between the probiotic and control groups. A total of 1006 participants were included in our quantitative analyses, 503 in the probiotic group and 503 in the placebo group. All included trials were published between 2015 and 2022, and all 15 trials were conducted in 5 countries: 11 in Iran, 1 in Ireland, 1 in Israel, 1 in Thailand and 1 in Turkey. This shows that Iran was the main country studied. The findings of Egger's test for FBS, FSI, and HOMA-IR were 0.749, 0.022, and 0.020, respectively. The results suggest a possible publication bias in the probiotics' impact on the FSI and HOMA-IR variables.



**Figure 1.** Included in the flow chart.

### 3.2. Glycemic outcomes

#### 3.2.1. FBS

Results from 12 trials with a total of 805 GDM patients included FBS (**Error! Reference source not found.A**). In the meta-analysis, the mean difference in FBS levels (MD -2.58, 95% CI -4.38 to -0.79,  $p < 0.01$ ) implies that probiotics statistically significantly improved FBS levels in pregnant women with GDM.

#### 3.2.2. FSI

The findings from 10 studies with 661 GDM patients consistently incorporated FSI (**Error! Reference source not found.B**). The studies indicated that the probiotic group exhibited lowered FSI levels (MD -2.29, 95% CI -3.40 to -1.18,  $p < 0.01$ ).

#### 3.2.3. HOMA-IR

HOMA-IR measurements from 661 expectant mothers with GDM across ten studies were analysed (**Error! Reference source not found.C**). The findings indicate that the HOMA-IR levels in the intervention group were lower than those in the placebo group (MD -0.56, 95% CI -0.81 to -0.32,  $p < 0.01$ ).

#### 3.2.4. QUICKI

The QUICKI test results from seven studies including 432 pregnant women diagnosed with GDM are shown (**Error! Reference source not found.D**). The results of the studies propose that

QUICKI levels in the intervention group (MD 0.01, 95% CI 0.00 to 0.01,  $p < 0.01$ ) were higher than those in the placebo group.



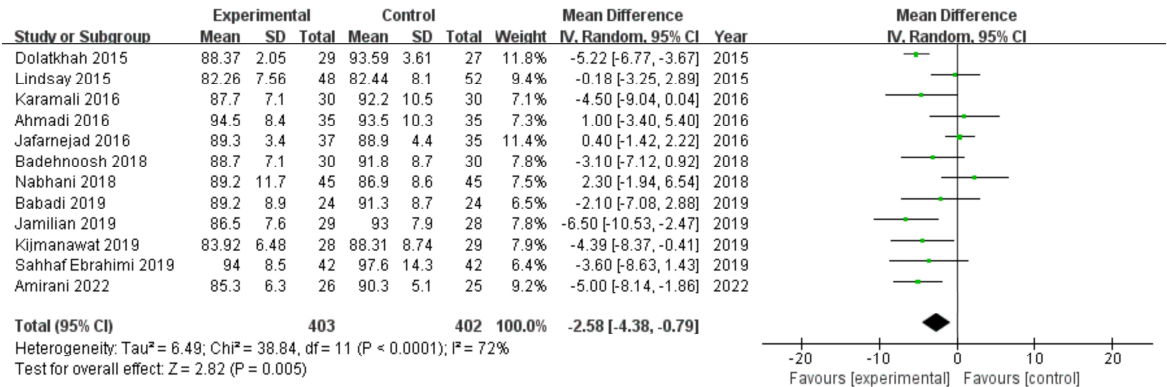
Table 1. Basic characteristics of eligible studies.

First authors	Year	Country	Probiotic Intervention	Control Intervention	Probiotic Dose	Probiotic group (N )	Control group ( N)	Duration (weeks)
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Lindsay	2015	Ireland	<i>Lactobacillus salivarius</i> UCC118	placebo	1×10 <sup>9</sup> CFU/g	48	52	8
Karamali	2016	Iran	<i>L. acidophilus</i> , <i>L. casei</i> and <i>B. bifidum</i> strains	placebo	2 × 10 <sup>9</sup> CFU/g each	30	30	6
Jafarnejad	2016	Iran	VSL#3 ( <i>Streptococcus thermophilus</i> , <i>Bifidobacterium breve</i> , <i>Bifidobacterium longum</i> , <i>Bifidobacterium infantis</i> , <i>Lactobacillus acidophilus</i> , <i>Lactobacillus plantarum</i> , <i>Lactobacillus paracasei</i> , and <i>Lactobacillus delbrueckii</i> subsp. <i>bulgaricus</i> )	placebo	112.5 × 10 <sup>9</sup> CFU	37	35	8
Ahmadi	2016	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus casei</i> , <i>Bifidobacterium bifidum</i> plus 0.8 g inulin	placebo	2 × 10 <sup>9</sup> CFU/g each	35	35	6
Nabhani	2018	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus plantarum</i> , <i>Lactobacillus fermentum</i> , <i>Lactobacillus gasseri</i>	placebo	<i>L. acidophilus</i> (5 × 10 <sup>10</sup> CFU/g), <i>L. plantarum</i> (1.5 × 10 <sup>10</sup> CFU/g), <i>L. fermentum</i> (7 × 10 <sup>9</sup> CFU/g), <i>L. gasseri</i> (2 × 10 <sup>10</sup> CFU/g)	45	45	6

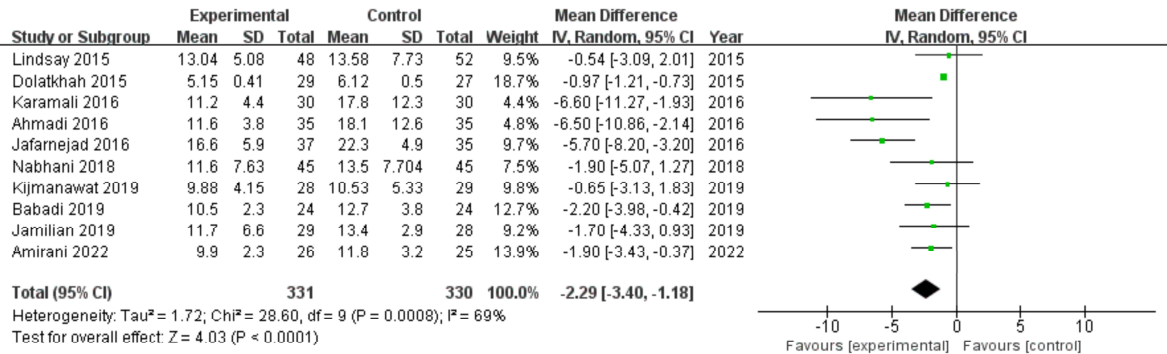
Badehnoosh	2018	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus casei</i> and <i>Bifidobacterium bifidum</i>	placebo	2 × 10 <sup>9</sup> CFU/g each	30	30	6
Karamali	2018	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus casei</i> and <i>Bifidobacterium bifidum</i> strains plus 800 mg inulin	placebo	2 × 10 <sup>9</sup> CFU/g each	30	30	6
Hajifaraji	2018	Iran	<i>L.acidophilus</i> LA-5, <i>Bifidobacterium</i> BB-12, <i>Streptococcus thermophilus</i> STY-31 and <i>Lactobacillus delbrueckii bulgaricus</i> LBY-27 plus dextrose anhydrous filler and magnesium stearate lubricant	placebo	4 biocap >4×10 <sup>9</sup> CFU	29	27	8
Babadi	2019	Iran	<i>Lactobacillus acidophilus</i> , <i>Lactobacillus casei</i> , <i>Bifidobacterium bifidum</i> , and <i>Lactobacillus fermentum</i>	placebo	2 × 10 <sup>9</sup> CFU/g each	24	24	6
Sahhaf Ebrahimi	2019	Iran	Probiotic yoghurt containing <i>Lactobacillus acidophilus</i> and <i>Bifidobacterium lactis</i>	placebo	300 g/day of probiotic yoghurt(contained 10 <sup>6</sup> CFU <i>Lactobacillus acidophilus</i> and <i>Bifidobacterium lactis</i>	42	42	8
Kijmanawat	2019	Thailand	<i>Lactobacillus acidophilus</i> and <i>Bifidobacterium bifidum</i>	placebo	1 × 10 <sup>9</sup> CFU/g each	28	29	4
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Amirani	2022	Iran	<i>Lactobacillus acidophilus</i> , <i>Bifidobacterium bifidum</i> , <i>Bifidobacterium lactis</i> <i>Bifidobacterium longum</i> Additionally , selenium	placebo	2 × 10 <sup>9</sup> CFU/day each	26	25	6
Yefet	2022	Israel	<i>Bifidobacterium bifidum</i> , <i>Bifido-</i> <i>bacterium lactis</i> , <i>Lactobacillus</i> (L) <i>acidophilus</i> , <i>L. paracasei</i> , <i>L.rhamnosus</i> and <i>Streptococcus thermophilus</i>	placebo	2 capsules/day ( >6×10 <sup>9</sup> CFU/capsule)	41	44	2

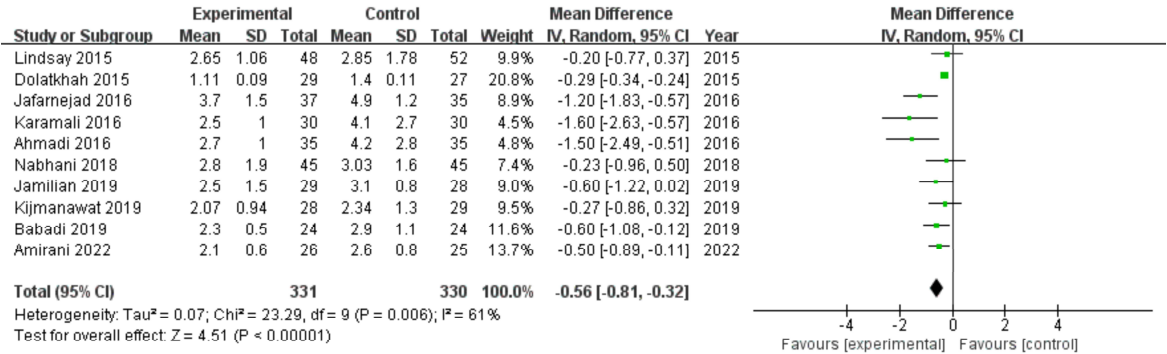
Abbreviations: CFU, colony-forming units.



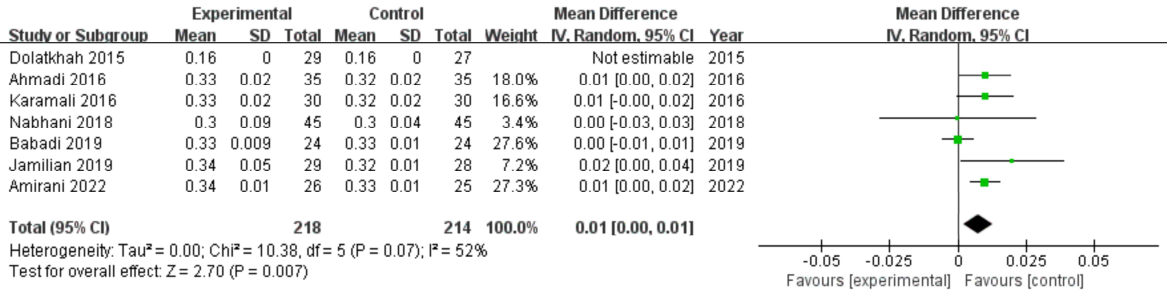
(A)



(B)



(C)



(D)

**Figure 2.** Effect of probiotic on glucose control in pregnant women with GDM: (A) FBS (mg/dL), (B) FSI (mU/L), (C) HOMA-IR, and (D) QUICKI. FBS: fasting blood sugar; FSI: fasting serum insulin; HOMA-IR: the homoeostatic model assessment for insulin resistance; QUICKI: quantitative insulin sensitivity check index.

3.3. Maternal outcomes

3.3.1. Preeclampsia

The preeclampsia incidence was included in all 4 studies of 313 pregnant women (**Error! Reference source not found.A**). The probiotic group did not display any statistically significant incidence of pre-eclampsia in comparison to the placebo group (RR 1.14, 95% CI 0.56 to 2.31,  $p = 0.71$ ).

3.3.2. Cesarean delivery

The forest plot (**Error! Reference source not found.B**) displays a merged analysis of the outcomes of caesarean sections in the experimental and control groups. The combined effect size estimations indicated that the differentiation between the two groups was not statistically noteworthy (RR 0.88, 95% CI 0.48 to 1.64,  $p = 0.70$ ).

3.3.3. Preterm delivery

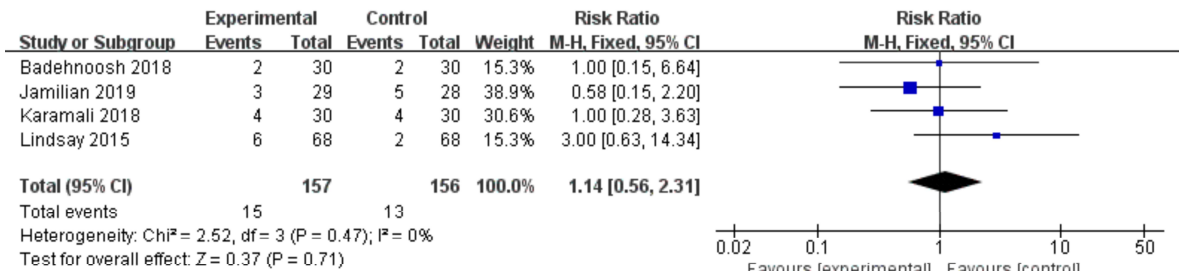
Three studies included information from 177 participants who had a caesarean section (**Error! Reference source not found.C**). There was no statistical difference in the risk of preterm birth between the probiotic and placebo groups (RR 0.99, 95% CI 0.23 to 4.24,  $p = 0.99$ ).

3.3.4. Induction of labor

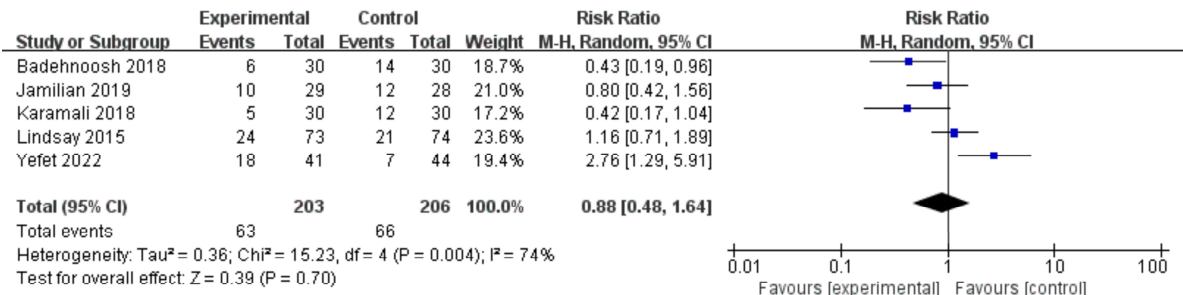
Information from 212 participants who underwent a caesarean section was analysed in the two included studies (**Error! Reference source not found.D**). The probiotic group did not show a statistically significant difference in the risk of caesarean section when compared to the placebo group (RR 0.83, 95% CI 0.34 to 2.03,  $p = 0.69$ ).

3.3.5. Polyhydramnios

In addition, we found 3 publications that included a combined sample of 177 subjects with polyhydramnios during pregnancy (**Error! Reference source not found.E**). Nevertheless, the difference between the two groups was not statistically significant (RR 0.46, 95% CI 0.13 to 1.56,  $p = 0.21$ ).

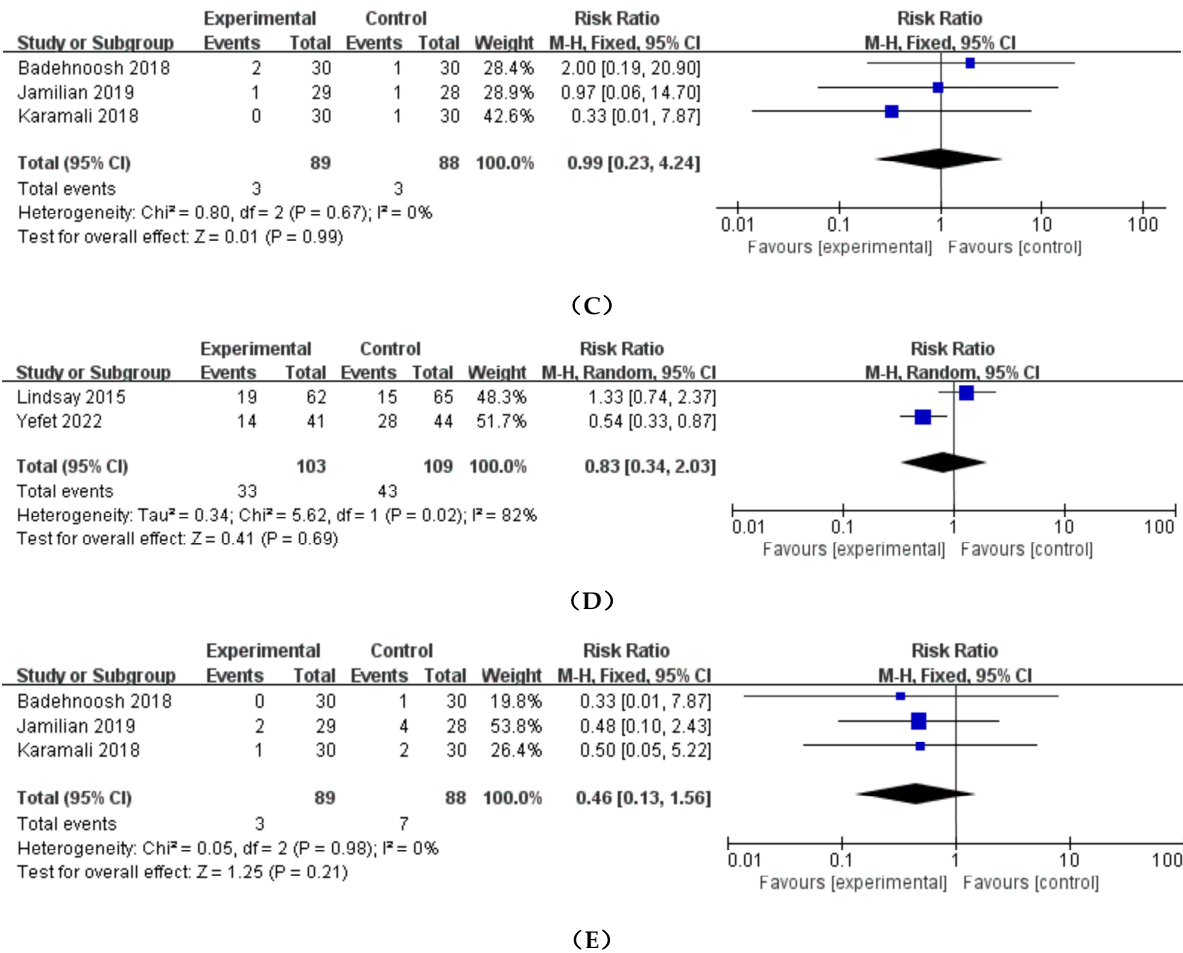


(A)



(B)





**Figure 3.** Effect of probiotic on pregnancy outcomes in pregnant women with GDM: (A) Preeclampsia, (B) Cesarean delivery, (C) Preterm delivery, (D) Induction of labor, and (E) Polyhydramnios.

3.4. Neonatal outcomes

3.4.1. Birthweight, Neonatal length and head circumference

Birth weight was compared between the two groups in seven studies. Meta-analysis (**Error! Reference source not found.A**) showed that among all included studies, probiotic group (MD -101.20, 95% CI -184.62 to -17.77, p = 0.02) had a significant reduction in newborn weight compared to placebo group.

For neonatal length in a total of 5 studies (**Error! Reference source not found.B**), there was no statistically significant difference between the probiotic intervention group and the placebo group for neonatal length (MD -0.44 , 95% CI -1.01 to 0.12, p = 0.12).

There were 6 studies on Neonatal head circumference (**Error! Reference source not found.C**). There was no statistically significant difference in the results of the comparison between the probiotic and placebo groups (MD -0.02 , 95% CI -0.36 to 0.31, p = 0.89).

3.4.2. NICU

A total of six studies were included (**Error! Reference source not found.D**), and a forest plot showed a reduced risk of NICUs compared to placebo (RR 0.60, 95% CI 0.40 to 0.89, p = 0.01).

3.4.3. Apgar score

We retrieved Apgar scores, including scores taken at one-minute and five-minute intervals (**Error! Reference source not found.E-F**). For both the one-minute Apgar score (MD 0.03, 95% CI -

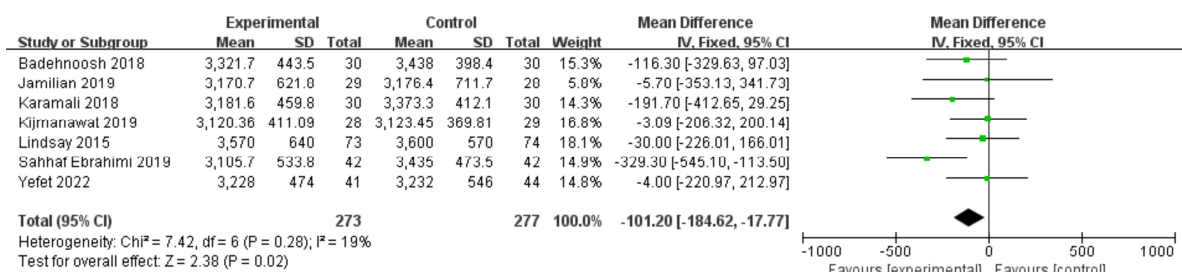
0.06 to 0.11,  $p = 0.51$ ) and the five-minute Apgar score (MD 0.04, 95% CI -0.03 to 0.12,  $p = 0.23$ ), there was no significant difference detected between the probiotic group and the placebo group.

### 3.4.4. Hyperbilirubinemia

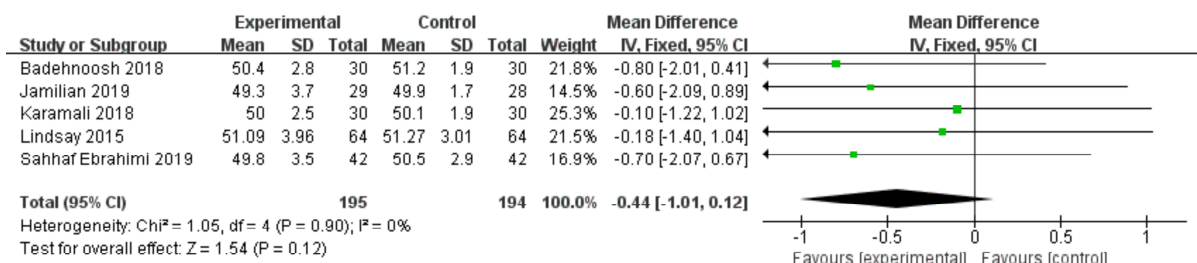
Four studies were included in the analysis (Error! Reference source not found.G), revealing a significant difference between the two groups. The probiotic group exhibited a lower risk of hyperbilirubinaemia in neonates compared to the placebo group (RR 0.31, 95% CI 0.16 to 0.61,  $p < 0.01$ ).

### 3.4.5. Other neonatal outcomes

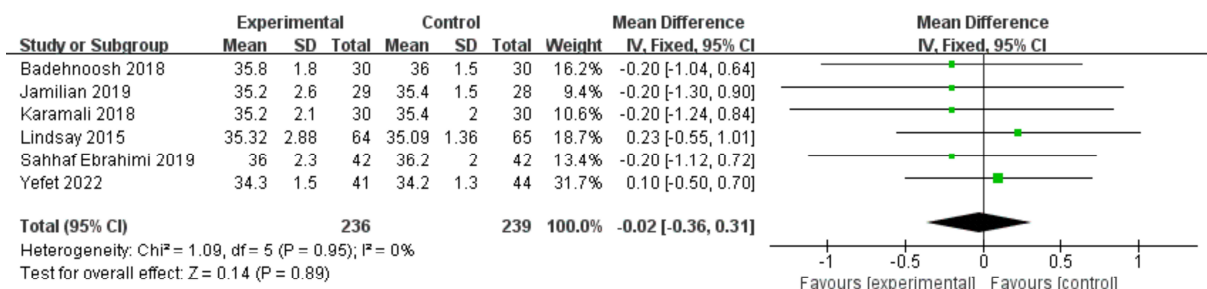
Furthermore, we summarised data on macrosomia, LGA and neonatal hypoglycaemia and found that none of the three outcomes were statistically significant (Error! Reference source not found.H-J). For Macrosomia, we collected five papers comprising 408 individuals. These studies showed that the probiotic group did not reduce the incidence of macrosomia (RR 0.37, 95% CI 0.12 to 1.15,  $p = 0.08$ ). Similarly, no reduction was observed in the incidence of LGA (RR 0.72, 95% CI 0.38 to 1.36,  $p = 0.31$ ) or neonatal hypoglycaemia (RR 0.80, 95% CI 0.43 to 1.51,  $p = 0.49$ ).



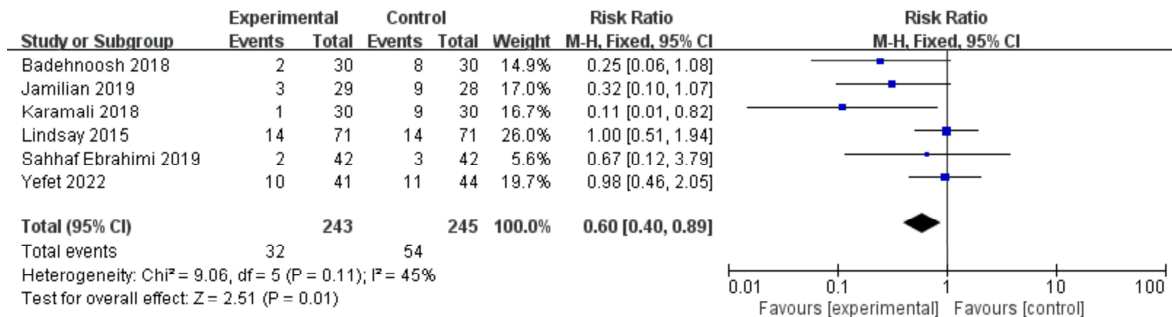
(A)



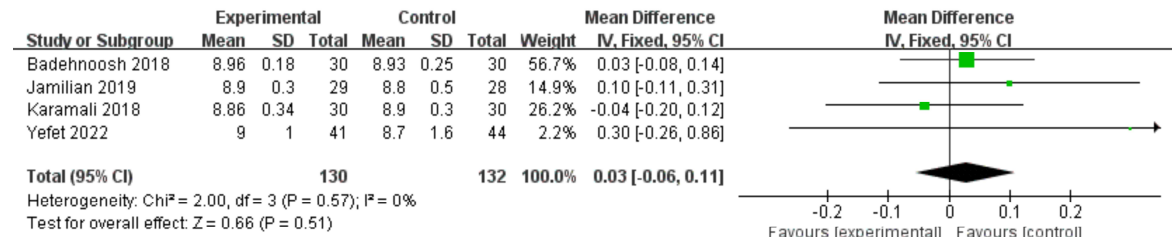
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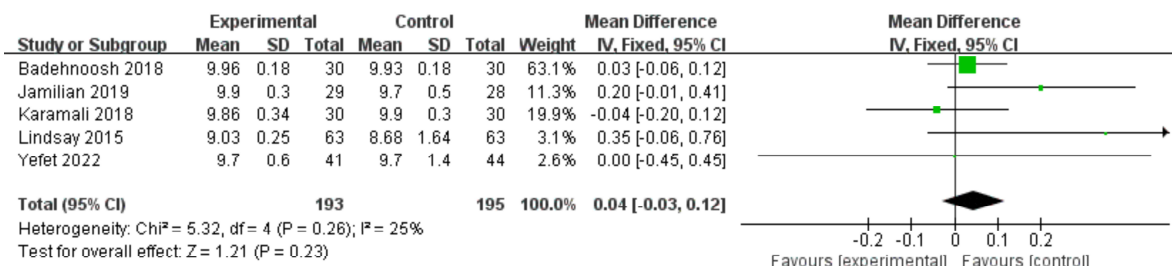
(C)



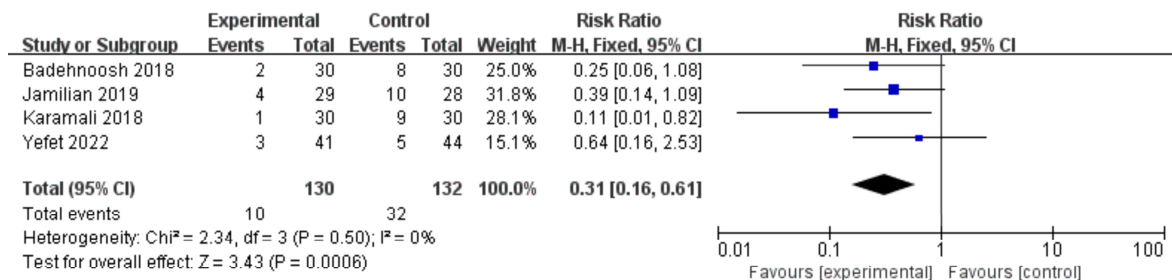
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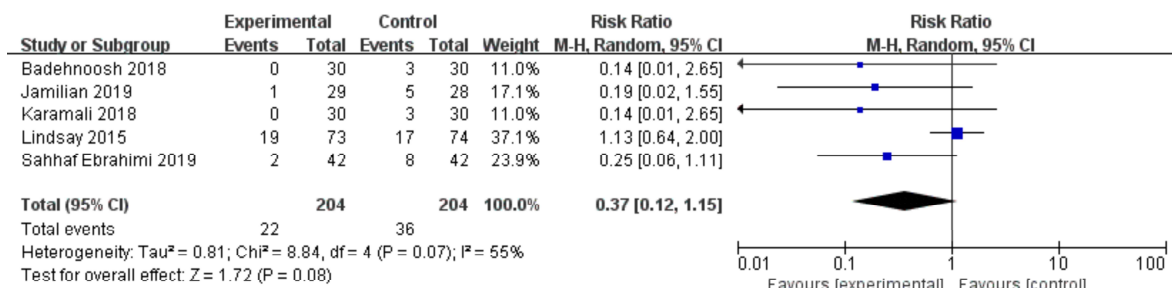
(E)



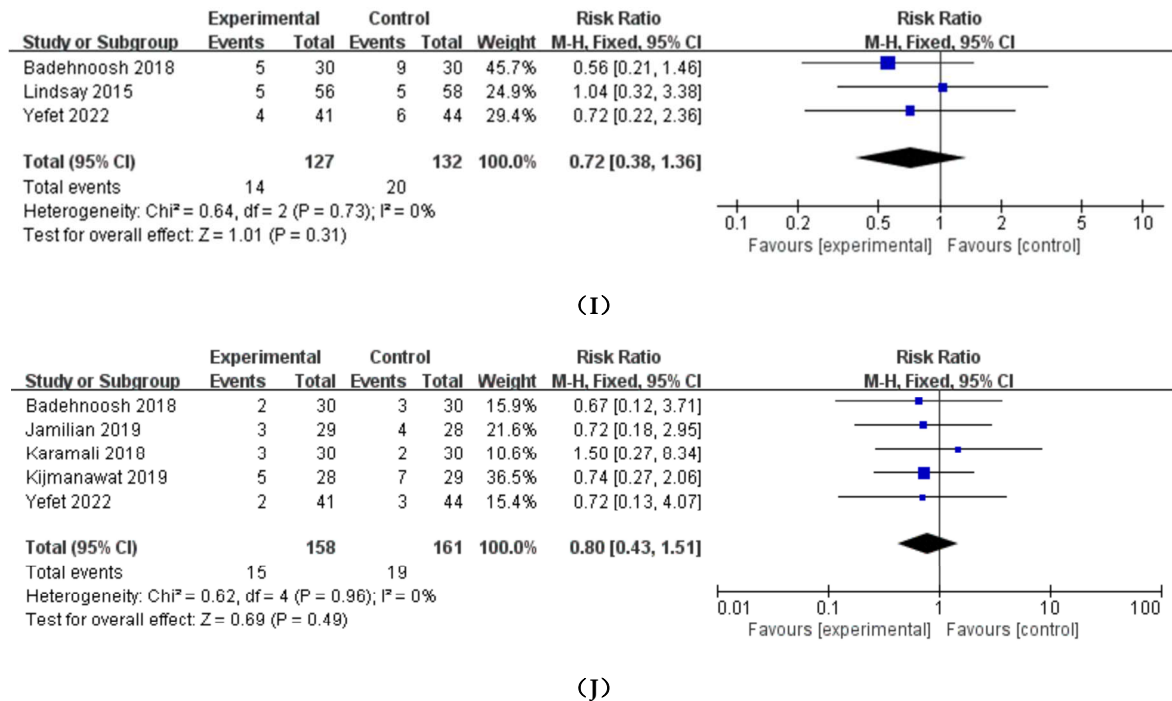
(F)



(G)



(H)



**Figure 4.** Effect of probiotic on neonatal outcomes in pregnant women with GDM: (A) Birthweight (g), (B) Neonatal length (cm), (C) Neonatal head circumference (cm), (D) NICU, (E) 1-min Apgar score, (F) 5-min Apgar score, (G) Hyperbilirubinemia, (H) Macrosomia, (I) LGA, and (J) Neonatal hypoglycaemia. NICU: neonatal intensive care unit; LGA: large for gestational age.

### 3.5. Subgroup Analysis

Due to FBS ( $I^2 = 72\%$ ), FSI ( $I^2 = 69\%$ ), HOMA-IR ( $I^2 = 61\%$ ), QUICKI ( $I^2 = 52\%$ ), cesarean delivery ( $I^2 = 74\%$ ) and macrosomia ( $I^2 = 55\%$ ), these outcomes existed with large heterogeneity and were therefore analyzed in subgroups, however, there were too few experiments on the outcome of induction of labour ( $I^2 = 82\%$ ), so subgroup analyses were not performed. The duration and country subgroups were grouped to examine potential reasons for heterogeneity. Different groups' p-values and heterogeneity are shown in **Error! Reference source not found.** Results suggest that country could have contributed to heterogeneity in FSI, HOMA-IR and macrosomia and did not explain heterogeneity in other outcomes; whereas duration did not explain heterogeneity in any outcomes.

**Table 2.** Results of subgroup analyses based on duration and country.

Subgroup	Studies	Participants	MD/RR (95% CI)	Heterogeneity ( $I^2\%$ )	P
<b>FBS</b>					
Iran	9	592	-2.26 (-4.35, -0.17)	64	0.03
Non Iran	3	213	-3.37 (-6.64, -0.10)	76	0.04
Duration $\geq$ 8weeks	4	312	-2.56 (-3.64, -1.48)	87	< 0.00001
Duration < 8weeks	8	493	-3.10 (-4.54, -1.66)	52	< 0.0001
<b>FSI</b>					
Iran	7	448	-2.81 (-3.71, -1.91)	55	< 0.00001
Non Iran	3	213	-0.96 (-1.20, -0.72)	0	< 0.00001
Duration $\geq$ 8weeks	3	228	-1.01 (-1.25, -0.77)	85	< 0.00001

Duration < 8weeks	7	433	-2.16 (-3.05, -1.26)	33	< 0.00001
<b>HOMA-IR</b>					
Iran	7	448	-0.70 (-0.93, -0.48)	44	< 0.00001
Non Iran	3	213	-0.29 (-0.34, -0.24)	0	< 0.00001
Duration ≥ 8weeks	3	228	-0.30 (-0.35, -0.24)	76	< 0.00001
Duration < 8weeks	7	433	-0.58 (-0.80, -0.36)	35	< 0.00001
<b>QUICKI</b>					
Iran	6	376	0.01 (0.00, 0.01)	52	0.0001
Non Iran	1	56	Not estimable	Not estimable	Not estimable
Duration ≥ 8weeks	1	56	Not estimable	Not estimable	Not estimable
Duration < 8weeks	6	376	0.01 (0.00, 0.01)	52	0.0001
<b>Cesarean delivery</b>					
Iran	3	177	0.57 (0.36, 0.89)	1	0.01
Non Iran	2	232	1.70 (0.73, 3.97)	72	0.22
Duration ≥ 8weeks	1	147	1.16 (0.71, 1.89)	Not estimable	0.55
Duration < 8weeks	4	262	1.16 (0.71, 1.89)	79	0.62
<b>Macrosomia</b>					
Iran	4	261	0.20 (0.07, 0.56)	0	0.002
Non Iran	1	147	1.13 (0.64, 2.00)	Not estimable	0.67
Duration ≥ 8weeks	2	231	0.85 (0.51, 1.42)	72	0.53
Duration < 8weeks	3	177	0.16 (0.04, 0.71)	0	0.02

Abbreviations: FBS, fasting blood sugar; FSI, fasting serum insulin; HOMA-IR, homeostatic model assessment for insulin resistance; QUICKI, the quantitative insulin sensitivity check index.

### 3.6. Assessment of study quality

All studies provided detailed explanations of the generation of randomised sequences and were assessed to be at low risk of bias. Ten studies clearly explained the allocation concealment process, and the remaining four studies which lacked this information were categorised as being at an unclear risk of bias. One study, which was an open-label trial, was deemed to have a high risk of bias, whereas 14 studies mentioned the blinding of assessors and were thus considered to be at low risk of bias. Based on a review of the study protocols, it was observed that all studies were at low risk of bias for follow-up and other biases. This is supported by the risk of bias graph (**Error! Reference source not found.**) and the risk of bias summary (**Error! Reference source not found.**). Consequently, the studies met the quality criteria.



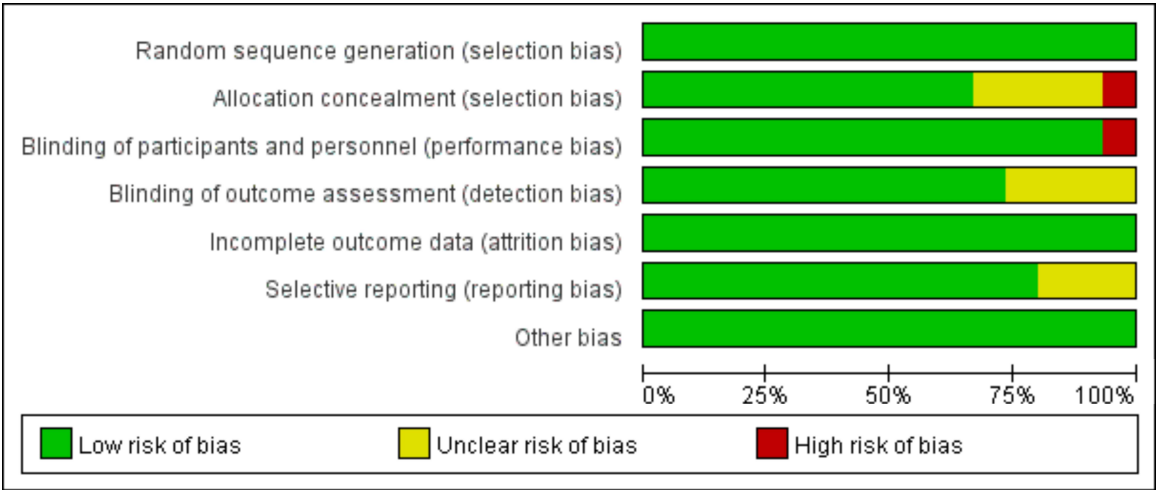


Figure 5. The risk of bias graph.

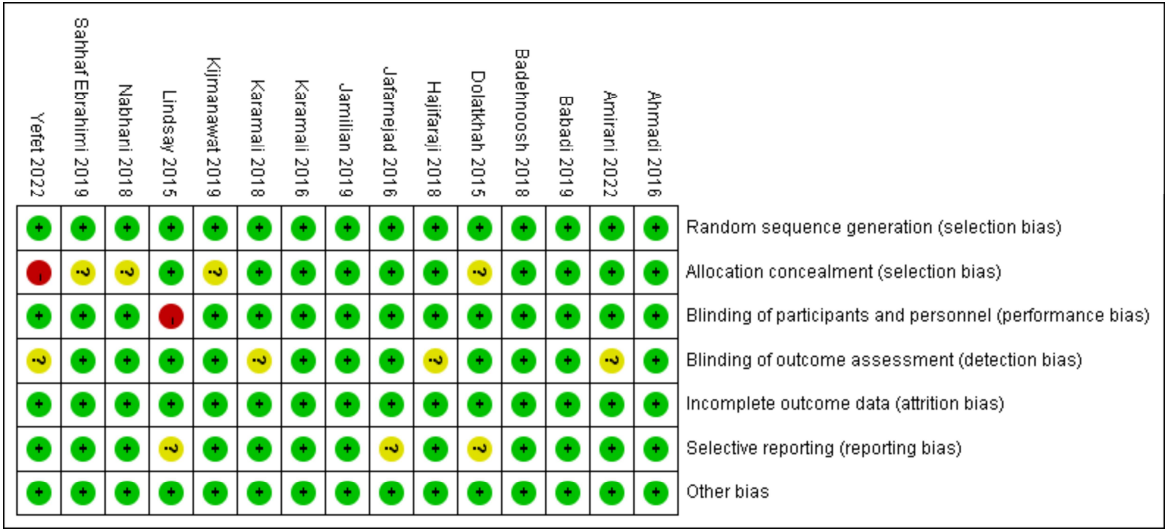


Figure 6. The risk of bias summary.

4. Discussion

The objective of this meta-analysis was to evaluate the glycaemic impact of probiotics compared to placebo among patients with gestational diabetes, along with presenting maternal and neonatal outcomes. Our meta-analysis demonstrates that regarding glycaemic outcomes, probiotics resulted in decreased levels of FBS, FSI, HOMA-IR, and increased levels of QUICKI. In relation to maternal outcomes, probiotics were found to have a significant impact on our set of preeclampsia, cesarean delivery, preterm delivery, induction of labour, and polyhydramnios indicators. However, they did not show any statistically significant results for other relevant outcomes. Concerning neonatal outcomes, probiotics were noted to result in reduced birthweight, NICU admissions and hyperbilirubinemia. Subgroup analyses indicated that the country was able to explain the heterogeneity of FSI, HOMA-IR, and macrosomia.

The most prevalent complication during pregnancy in present times is GDM. The incidence of diagnosed hyperglycaemic and even overtly diabetic young women is increasing. The primary risk factors for GDM are obesity or being overweight, a previous history of GDM, later age at childbearing, and a family history of type 2 diabetes. GDM works by altering glucose regulation during pregnancy to provide nutrients to the developing fetus[26]. Insulin sensitivity is reduced in late pregnancy compared to pre-pregnancy, while basal endogenous glucose production is increased

[27,28]. In pregnant women with normal blood glucose, pancreatic  $\beta$ -cells produce more insulin to maintain normal glucose levels. Diagnosis is usually made using an oral glucose tolerance test (OGTT), but a non-fasting glucose provocation test is also used in some places to assess the need for a woman to undergo a complete oral glucose tolerance test, and one of the commonly used criteria includes the International Diabetes Federation (IDF), the American Diabetes Association (ADA) or the World Health Organisation (WHO). Treatment for GDM involves a healthy diet, exercise regimen, and appropriate medication. The Academy of Nutrition recommends a balanced diet for GDM, providing adequate macronutrients and micronutrients to limit blood sugar fluctuations in pregnant women and support the nutritional requirements of the foetus[29,30]. Carbohydrate intake can have an impact on blood glucose levels in pregnant women, so diets should be comprehensive and diverse, with particular attention paid to the carbohydrate content. In addition, physical activity requires energy consumption, and this energy comes from the oxidative breakdown of sugar. Therefore, appropriate exercise can consume sugar, thus reducing blood glucose levels, and can assist expecting mothers in maintaining their blood glucose levels better. Common exercises include walking, yoga, and dancing[31,32]. However, when blood glucose levels remain abnormal, treatment with insulin medication is necessary[33]. In some countries, oral hypoglycaemic agents, primarily metformin and glibenclamide, are also employed. Nevertheless, medication comes with risks and potential adverse effects. For instance, oral hypoglycaemic drugs carry the risk of hypoglycaemia to the foetus, while insulin injections may cause discomfort and skin reactions at the injection site. The pharmacological treatment entails choosing the appropriate drug, adjusting the dosage, and so on, which complicates diabetes management. Moreover, it must be used in strict compliance with the physician's advice and monitoring guidance. Medication is typically administered to manage blood glucose levels, however, gestational diabetes often resolves spontaneously postpartum. Therefore, personalized treatment is required. Hence, the necessity of pharmacotherapy must be balanced against its use throughout the entirety of pregnancy. Each pregnant woman's physical state and pancreatic function is distinctive, and their response to various drugs can vary. Thus, there is a need to find more convenient and safer treatment options.

A mounting body of research indicates a crucial connection between glucose intolerance and dysbiosis of gut microflora (GM) during pregnancy[34]. As per the analyzed data, probiotics exhibit a beneficial impact on glycaemic control and might have a role in preventing and treating GDM, thus constituting a promising strategy to lessen GDM occurrence[35,36]. Evidence suggests that probiotics may reduce adverse pregnancy outcomes in GDM by modulating gut microbial composition and improving metabolism. The corresponding mechanisms are believed to involve the secretion of short-chain fatty acids, modulation of hormone secretion, and reduction of inflammatory responses[6].

One study discovered that the addition of probiotics during pregnancy did not seem to decrease the possibility of diabetes or enhance other neonatal and maternal outcomes[37]. The Shahriari et al. study aimed to investigate the impact of probiotic supplementation during pregnancy on the likelihood of GDM and other maternal and neonatal outcomes. Mothers underwent assessment for GDM presence via a 75 g OGTT between weeks 24-28 of pregnancy. The probiotic group, comprising of 507 pregnant women, were randomly assigned. Each 500 mg probiotic capsule contained a blend of *Lactobacillus acidophilus* LA1, *Bifidobacterium longum* sp54 cs, and *Bifidobacterium bifidum* sp9 cs, while the control group was given placebo. The study found that the incidence of GDM was similar in both the probiotic group (41.9%) and the control group (40.2%) ( $p = 0.780$ ). Additionally, there were no significant differences in FBS (88.68 vs. 89.61 mg/dL;  $p = 0.338$ ), OGTT-1h (163.86 vs. 166.88 mg/dL;  $p = 0.116$ ), and OGTT-2h (138.39 vs. 139.27 mg/dL;  $p = 0.599$ ) between the probiotic and control groups. Yet another study hints at a potential decrease in the occurrence of LGA[38]. The objective was to ascertain if probiotics (*Lactobacillus rhamnosus* and *Bifidobacterium animalis* *Lactis* subspecies) consumed by overweight and obese women from mid-pregnancy prevented GDM, as 411 women were randomly allocated. The frequency of GDM was 12.3% among the placebo group and 18.4% among the probiotic group ( $p = 0.10$ ). For the outcomes of preeclampsia, the probiotic group had a rate of 9.2%, while the placebo group had a rate of 4.9% ( $p = 0.09$ ). The incidence of LGA was 2.4% in the probiotic group and 6.5% in the placebo group ( $p = 0.042$ ). Furthermore, there were no significant

differences in other pregnancy and neonatal outcomes. Both studies have demonstrated that probiotics do not prevent the emergence of GDM. Nonetheless, there is a contentious debate over the results regarding maternal and infant outcomes.

In previous studies, fewer meta-analyses focused on pregnancy outcomes and neonatal outcomes, and instead mainly collected data on metrics of glycemic control, lipid metabolism and inflammation[39,40]. Unlike previous studies, we exclusively selected patients with gestational diabetes to evaluate the influence of probiotics on GDM patients. Additionally, the preceding research featured obese and overweight pregnant women, who were not diagnosed with GDM – this could have introduced biased results[7]. Due to the restricted number of outcomes, we gathered some typical glycaemic outcomes, as well as maternal and infant outcomes. Our findings revealed no significant statistical differences in maternal outcomes, however, neonatal outcomes such as birthweight, NICU, and hyperbilirubinemia were significant.

The outcome of macrosomic babies in our study showed no statistically significant difference between the probiotic group and the placebo group, contrary to a previous study[41]. Furthermore, a prior meta-analysis[42] suggested that probiotics could elevate the risk of pre-eclampsia, but this did not impact neonatal birthweight. However, our results contrasted those of our counterparts, likely due to the inclusion of additional RCTs in our study.

In our study, the subanalyses of various countries produced divergent endpoints, likely due to the majority of studies being conducted in Iran. These findings underline the possibility that women across different nations may have varying gut microbiota influenced by distinct dietary and lifestyle choices, resulting in dissimilar gut microbiota. Additionally, differences in genetic and environmental backgrounds may cause variations in the roles of relevant and metabolic pathways in vivo. In future studies, greater consideration should be given to country-specific effects when determining appropriate probiotic supplements for various nations. Furthermore, subgroup analyses were performed for duration, and none of the subgroups with an 8-week cut-off could clarify the heterogeneity of the outcomes of interest. As a result, a different duration may need to be selected for subgroup analyses in the future.

Meta-analysis has a number of strengths. Firstly, combining data from 15 articles enhances the sample size ( $N = 1006$ ) and boosts the dependability of the outcomes. Second, the majority of the integrated randomised controlled trials were high-quality studies that employed randomised sequence generation and double-blind methods, thus the resulting data had a high level of reliability. Thirdly, the study collected not only glycaemic indicators but also relevant maternal and infant outcomes in women with gestational diabetes, making the results more representative. These strengths render the findings of this study of high scientific worth and practical use. However, this study has several limitations. Firstly, the number of participants in the included randomised controlled trials was relatively small, ranging from 40 to 100. Secondly, because of restricted data, only one study has reported on postprandial glycaemia. This study evaluated solely fasting glycaemia and did not examine the impact of probiotics on postprandial glycaemia. Furthermore, only one study investigated small for gestational age (SGA). Thirdly, data on maternal and infant outcomes were insufficient, as they failed to present a comprehensive range of outcomes and consider the long-term effects on mothers and offspring. Furthermore, our subgroup analyses did not group the type and dose of probiotics together appropriately, thereby obscuring the extent of heterogeneity. Therefore, more randomised controlled trials are required to achieve greater precision in the results. Overall, the data indicates a positive effect of probiotics for patients with GDM, as demonstrated by improvements in glucose-related markers and certain neonatal markers.

## 5. Conclusions

In conclusion, this meta-analysis, based on short-term data, indicates that probiotics have the potential to improve glycaemic control and reduce the risk of some neonatal outcomes compared with placebo. Therefore, probiotics may be a safe and effective treatment for GDM. However, due to the limited number of RCTs, further research is required to evaluate the long-term effects of probiotics on pregnant women and neonates.

**Author Contributions:** R.W., J.J.H., H.L., and Z.J.L. all made significant contributions to the content and structure of the study. R.W. and J.J.H. reviewed each experiment for eligibility and extracted and tabulated the relevant data. R.W. analysed the data. R.W. drafted the article. H.L. checked the format of the article. R.W., J.J.H., H.L., and Z.J.L. commented on the content of the article. All authors have read and agreed to the published version of the manuscript.

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