Association Between Preterm-Birth Phenotypes and Differential Morbidity, Growth, and Neurodevelopment at Age 2 Years
Results From the INTERBIO-21st Newborn Study

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IMPORTANCE The etiologic complexities of preterm birth remain inadequately understood, which may impede the development of better preventative and treatment measures.

OBJECTIVE To examine the association between specific preterm-birth phenotypes and clinical, growth, and neurodevelopmental differences among preterm newborns compared with term newborns up to age 2 years.

DESIGN, SETTING, AND PARTICIPANTS The INTERBIO-21st study included a cohort of preterm and term newborn singletons enrolled between March 2012 and June 2018 from maternity hospitals in 6 countries worldwide who were followed up from birth to age 2 years. All pregnancies were dated by ultrasonography. Data were analyzed from November 2019 to October 2020.

EXPOSURES/INTERVENTIONS Preterm-birth phenotypes.

MAIN OUTCOMES AND MEASURES Infant size, health, nutrition, and World Health Organization motor development milestones assessed at ages 1 and 2 years; neurodevelopment evaluated at age 2 years using the INTERGROWTH-21st Neurodevelopment Assessment (INTER-NDA) tool.

RESULTS A total of 6529 infants (3312 boys [50.7%]) were included in the analysis. Of those, 1381 were preterm births (mean [SD] gestational age at birth, 34.4 [0.1] weeks; 5148 were term births (mean [SD] gestational age at birth, 39.4 [0] weeks). Among 1381 preterm newborns, 8 phenotypes were identified: no main maternal, fetal, or placental condition detected (485 infants [35.1%]); infections (289 infants [20.9%]); preeclampsia (162 infants [11.7%]); fetal distress (131 infants [9.5%]); intrauterine growth restriction (110 infants [8.0%]); severe maternal disease (85 infants [6.2%]); bleeding (71 infants [5.1%]); and congenital anomaly (48 infants [3.5%]). For all phenotypes, a previous preterm birth was a risk factor for recurrence. Each phenotype displayed differences in neonatal morbidity and infant outcomes. For example, infants with the no main condition detected phenotype had low neonatal morbidity but increased morbidity and hospitalization incidence at age 1 year (odds ratio [OR], 2.2; 95% CI, 1.8-2.7). Compared with term newborns, the highest risk of scoring lower than the 10th centile of INTER-NDA normative values was observed in the fine motor development domain among newborns with the fetal distress (OR, 10.6; 95% CI, 5.1-22.2) phenotype.

CONCLUSIONS AND RELEVANCE Results of this study suggest that phenotypic classification may provide a better understanding of the etiologic factors and mechanisms associated with preterm birth than continuing to consider it an exclusively time-based entity.
Preterm birth is a heterogeneous syndrome, which has been previously described in terms of 3 factors: spontaneous or medically induced labor, presence or absence of preterm premature rupture of membranes, and mode of delivery. A phenotypic classification system, incorporating etiologically associated maternal, fetal, and placental characteristics as well as signs of parturition and pathway to delivery, has also been proposed. Twelve preterm birth phenotypes (ie, separate biological entities with specific risk factors, newborn anthropometry, and risk of neonatal morbidity and mortality) have previously been identified. The present INTERBIO-21st Newborn Study comprised phase 2 of the International Fetal and Newborn Growth Consortium for the 21st Century (INTERGROWTH-21st) Project, a population-based research initiative involving almost 70,000 mothers and infants worldwide that was conducted from 2008 to 2015. In this study, we investigated whether preterm birth phenotypes were associated with clinical, epidemiological, growth, and neurodevelopmental differences among preterm and term infants up to age 2 years.

Methods

The INTERBIO-21st Newborn Study included a cohort of preterm and term newborns who were enrolled from March 2012 to June 2018 and followed up from birth to age 2 years. The study was approved by the Oxfordshire Research Ethics Committee C, institutional research ethics committees at participating sites, and corresponding regional authorities. All mothers provided written informed consent. This study followed the Strengthening the Reporting of Observational Studies in Epidemiology (STROBE) reporting guideline for cohort studies.

Study Sites and Participants

The study was conducted in Pelotas, Brazil; Nairobi, Kenya; Kilifi, Kenya; Karachi, Pakistan; Soweto, South Africa; Mae Sot, Thailand; and Oxford, United Kingdom. The sites and study populations have been described elsewhere. In brief, the study enrolled preterm newborns who were live born at 23 weeks 0 days’ to 36 weeks 6 days’ gestation and term newborns who were live born at 37 weeks 0 days’ to 41 weeks 6 days’ gestation. All newborns were naturally conceived singleton with mothers who were age 18 years and older and residing in the catchment area of the participating hospital. Gestational age was estimated by ultrasonographic measurement of crown-rump length at less than 14 weeks 0 days’ gestation (75% of newborns) or head circumference at less than 24 weeks 0 days’ gestation (25% of newborns) using INTERGROWTH-21st standards.

Given the design of the INTERBIO-21st study, we selected a higher proportion of newborns with lower birth weight (ie, < third centile).

Deliveries were screened daily using a tablet computer (iPad; Apple), with software oversampling at lower gestational ages to increase statistical power for studying the highest-risk subgroups by producing a higher proportion of exposures and adverse neonatal outcomes. To correct for the oversampling of newborns who were small for gestational age (defined as birth weight <10th centile based on INTERGROWTH-21st standards) in the present analysis, we generated a term newborn group consisting of term infants who were appropriately grown for gestational age (defined as birth weight ≥10th centile based on INTERGROWTH-21st standards); approximately 10% of the term newborn group included randomly selected term infants who were small for gestational age to reflect the incidence of newborns who are small for gestational age in the general population.

Neonatal and Child Outcomes

The anthropometric measurement protocols, training materials, and quality control procedures were based on the World Health Organization (WHO) Multicentre Growth Reference Study, which produced the WHO Child Growth Standards. In brief, newborn measures were obtained within 12 hours of birth (and no later than 24 hours after birth) using identical equipment at all sites. An electronic scale (sensitivity, 10-20 g; Seca) was used to measure birth weight, and an infantometer (Harpenden; Chasmons) was used to measure recumbent length. Head circumference was measured using metallic nonextendable tape (Chasmons). The anthropometrists independently took all measures twice and compared values using maximum allowable differences of 50 g for weight, 7 mm for length, and 4 mm for head circumference. If any difference exceeded those values, both observers independently performed the relevant measurement a second time and, if necessary, a third time. The Oxford Anthropometric Standardization Unit regularly monitored staff performance and recommended retraining if measurements consistently exceeded the maximum allowable differences.

Standardized care and feeding practices were implemented using INTERGROWTH-21st protocols. Exclusive breastfeeding until age 6 months was encouraged, with standard supplementation for preterm newborns.

Each mother provided detailed information about the infant’s health and severe morbidities at birth, age 1 year, and age 2 years using project-specific forms. We used an unweighted composite neonatal outcome score comprising a severe neonatal morbidity index that included 1 or more of...
the following severe neonatal complications: bronchopulmonary dysplasia, hypoxic-ischemic encephalopathy, sepsis, anemia (requiring transfusion), periventricular hemorrhage or leukomalacia, retinopathy, necrotizing enterocolitis (Bell stage 2 or higher), and patent ductus arteriosus (requiring intervention).1,16-18 At ages 1 and 2 years, we assessed (1) hospital admission (≥1 admission); (2) diagnosis and treatment for infections requiring antibiotic medication (≥3 regimens), otitis media, pneumonia, bronchiolitis, parasitosis, diarrhea, vomiting, exanthema, skin disease, fever lasting 3 or more days (≥3 episodes), and meningitis; (3) neurologic disorders, including seizures and cerebral palsy; and (4) severe clinical conditions, including cardiovascular problems, gastroesophageal reflux, hemolytic conditions, trauma from injury, or any condition requiring surgery.

At birth, we measured weight, length, and head circumference following INTERGROWTH-21st protocols. At ages 1 and 2 years, we measured weight, length, and head circumference following WHO protocols.20 Motor development was assessed using the chronological age of the child by comparing parental information with 4 WHO developmental milestones: sitting without support, crawling on hands and knees, standing alone, and walking alone.19

We assessed neurodevelopment at age 2 years using the INTERGROWTH-21st Neurodevelopment Assessment (INTER-NDA), which measures multiple dimensions of early development among children aged 22 to 30 months.20 The INTER-NDA is implemented by nonspecialists21 and uses mixed methods to measure cognition, language development, fine and gross motor skills, and positive and negative mixed methods to measure cognition, language development, and separation. Clustering was considered satisfactory if the silhouette statistic was 0.6 or higher (range, −1.0 to 1.0).25 After the first analyses, we merged clusters associated with infections (extrauterine infection, chorioamnionitis, and perinatal sepsis) because each infection had low statistical power when analyzed separately (eTable 1 in the Supplement). Of 44 fetal anemia cases, cluster analysis categorized 28 cases (63.6%) as perinatal sepsis, 6 cases (13.6%) as mid to late pregnancy, 5 cases (11.4%) as fetal distress, 3 cases (6.8%) as preeclampsia, and 2 cases (4.5%) as congenital anomaly. Because most cases were associated with perinatal sepsis, we included those newborns in the infections group.

To assess associations between maternal risk factors and phenotypes, we performed multinomial logistic regression analyses to model nominal outcome variables. Odds ratios (ORs) for each phenotype (and for all preterm newborns) compared with the term newborn group were estimated based on multinomial modeling. Model 1 was adjusted for maternal age, height, first trimester and prepregnancy body mass index (calculated as weight in kilograms divided by height in meters squared), years of education, presence of prepregnancy mental illness, length of menstrual cycle, and smoking during pregnancy. Model 2 was adjusted for all variables included in model 1 plus the number of previous pregnancies, miscarriages, and terminations. Model 3 was adjusted for all variables included in model 2 plus the number of previous births, low birth weight (defined as <2500 g) and preterm infants, and neonatal deaths. We found no data that violated the independence of irrelevant alternatives assumption in our models.

For all newborns, we generated age- and sex-specific z scores for weight, length, head circumference, weight for length, and percentage of newborns lower than the 10th centile on the INTER-NDA and on visual tests using international standards.24

Statistical Analysis
To construct the preterm-birth phenotypes (eTable 1 in the Supplement), we were guided by the work of the INTERGROWTH-21st Consortium, which provided a conceptual frame and etiologic factors2-4 that were empirically tested.5 The same definitions of maternal, fetal, and placental conditions and the same analytical strategy using a 2-step cluster analysis (via the 2-step cluster algorithm in SPSS software, version 25; SPSS Statistics) were then applied to the data collected for the present study.5 This approach allowed us to compare the distribution of clusters and their proportional contribution across populations.

Quality was assessed based on silhouette measures of cohesion and separation. Clustering was considered satisfactory if the silhouette statistic was 0.6 or higher (range, −1.0 to 1.0).25 After the first analyses, we merged clusters associated with infections (extrauterine infection, chorioamnionitis, and perinatal sepsis) because each infection had low statistical power when analyzed separately (eTable 1 in the Supplement). Of 44 fetal anemia cases, cluster analysis categorized 28 cases (63.6%) as perinatal sepsis, 6 cases (13.6%) as mid to late pregnancy, 5 cases (11.4%) as fetal distress, 3 cases (6.8%) as preeclampsia, and 2 cases (4.5%) as congenital anomaly. Because most cases were associated with perinatal sepsis, we included those newborns in the infections group.

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For all newborns, we generated age- and sex-specific z scores for weight, length, head circumference, weight for length, and percentage of newborns lower than the 10th centile compared with INTERGROWTH-21st standards7 and preterm reference charts.26 At ages 1 and 2 years, similar z scores and centiles for weight, length, and head circumference were generated compared with the WHO Child Growth Standards.11 We used 1-way analysis of variance to test whether the means of each z score were different across the phenotypes and the term newborn group. Phenotypic analyses at age 2 years used chronological rather than corrected age.27

We performed logistic regression analysis to assess associations between phenotypes and neonatal, 1-year, and 2-year morbidity indices, presented as ORs with 95% CIs adjusted for study site. For the neonatal morbidity analyses, we compared individual preterm-birth phenotypes with the phenotype for no main maternal, fetal, or placental condition detected. Unadjusted and adjusted analyses were conducted for gestational age. Robust SEs were estimated in all logistic association models using the vce (cluster clustvar) package in Stata software, version 15 (StataCorp).

We used a Kruskal-Wallis nonparametric test to examine the equality of age distribution at the achievement of gross motor development milestones among phenotypes and the term newborn group. All analyses were performed using Stata software, version 15. Data were analyzed from November 2019 to October 2020.

Results
We prospectively enrolled 7540 women between 2012 and 2019; of those, 202 women were lost to follow-up or withdrew consent. After excluding 27 infants with missing birth weights, 7311 newborns (1381 preterm newborns and 5930 term newborns; 4633 newborns who were appropriately grown for
gestational age and 1297 newborns who were small for gestational age) remained (eFigure 1 in the Supplement). To restore the prevalence of infants who were small for gestational age to 10%, we randomly excluded 782 term newborns who were small for gestational age from the term newborn group. After all exclusions, a total of 6529 infants (3312 boys [50.7%]) were included in the final analysis. Of those, 1381 newborns were preterm births (mean [SD] gestational age at birth, 34.4 [0.1] weeks; mean [SD] maternal age, 29.3 [0.2] years) and 5148 newborns were term births (mean [SD] gestational age at birth, 39.4 [0.0] weeks; mean [SD] maternal age, 29.3 [0.1] years). The overall follow-up rate was 78% (71% for preterm newborns and 80% for term newborns) at 1 year and 67% (64% for preterm newborns and 68% for term newborns) at 2 years (eFigure 1 in the Supplement).

Each site’s contribution to the total study population ranged from 7.6% (Karachi, Pakistan) to 25.0% (Oxford, United Kingdom). For the preterm group, each site’s contribution ranged from 6.7% (Soweto, South Africa) to 27.0% (Oxford, United Kingdom). Sociodemographic characteristics were similar across groups from different sites.

Among 1381 preterm newborns, the largest phenotype was no main maternal, fetal, or placental condition detected (no main condition phenotype; 485 infants [35.1%]). Cluster selection also derived phenotypes with 1 main condition (with the exception of the severe maternal disease phenotype, comprising 85 infants [6.2%]), indicating that preterm birth was a heterogeneous syndrome. These phenotypes comprised infections (289 infants [20.5%]), preeclampsia (162 infants [11.7%]), fetal distress (131 infants [9.5%]), intrauterine growth restriction (110 infants [8.0%]), bleeding (71 infants [5.1%]), and congenital anomaly (48 infants [3.5%]) (eTable 2 in the Supplement). The main congenital anomalies diagnosed were in the face, heart, and limbs, with no observable common factor.

Cesarean deliveries were more common among preterm newborns compared with term newborns (696 of 1381 infants [50.4%] vs 1778 of 5148 infants [34.5%], respectively) and among preterm newborns with specific phenotypes, such as preeclampsia (124 of 162 infants [76.5%]), bleeding (61 of 71 infants [85.5%]), fetal distress (123 of 131 infants [93.9%]), and severe maternal disease (55 of 85 infants [64.7%]) (eTable 2 in the Supplement). Conditions such as preterm premature rupture of membranes were more common among those with preterm-birth phenotypes for infections (174 of 289 infants [60.2%]), severe maternal disease (32 of 85 infants [37.6%]), and congenital anomaly (17 of 48 infants [35.4%]). In addition, newborns with preterm phenotypes had lower birth weights (mean [SD], 2210 [15.9] g vs 3200 [6.2] g), shorter length at birth (mean [SD], 44.6 [0.1] cm vs 48.9 [0.0] cm), smaller head circumference at birth (mean [SD], 31.3 [0.1] cm vs 33.9 [0.1] cm), higher neonatal morbidity (eg, hospitalization in neonatal intensive care unit for >1 day, 755 of 1381 infants [54.7%] vs 380 of 5148 infants [7.4%]), and higher neonatal mortality (55 of 1381 infants [4.0%] vs 8 of 5148 infants [0.2%]) compared with the term newborn group.

For all preterm-birth phenotypes, a previous preterm birth was a risk factor for recurrence of preterm delivery (results for each risk factor assessed in each model are available in Figure 2 in the Supplement). Associations with preterm phenotypes were also identified, indicating a differential risk according to preterm phenotype. The no main condition phenotype was associated with young maternal age (OR, 1.6; 95% CI, 1.2-2.2), nulliparity (OR, 2.0; 95% CI, 1.3-3.1), short maternal stature (OR, 1.4; 95% CI, 1.1-1.8), fewer maternal years of education (OR, 1.5; 95% CI, 1.1-1.9), and 2 or more previous miscarriages (OR, 1.5; 95% CI, 1.1-2.2). The preeclampsia phenotype was associated with low maternal body mass index (OR, 2.3; 95% CI, 1.0-4.9) or overweight maternal status (OR, 3.0; 95% CI, 2.1-4.5), nulligravidity (OR, 1.9; 95% CI, 1.3-3.0), and delivery of a previous infant with low birth weight (OR, 2.8; 95% CI, 1.5-5.3). The infections phenotype was associated with both younger (OR, 1.9; 95% CI, 1.3-2.9) and older (OR, 1.4; 95% CI, 1.1-1.9) maternal age and previous termination of pregnancy (OR, 1.7; 95% CI, 1.1-2.8). The infections phenotype was associated with both younger (OR, 1.9; 95% CI, 1.3-2.9) and older (OR, 1.4; 95% CI, 1.1-1.9) maternal age and previous termination of pregnancy (OR, 1.7; 95% CI, 1.1-2.8). The intrauterine growth restriction phenotype was associated with short maternal stature (OR, 2.2; 95% CI, 1.3-3.8), smoking during pregnancy (OR, 5.0; 95% CI, 2.9-8.6), and previous miscarriages (OR, 2.2; 95% CI, 1.2-3.9).

Newborn Size and Postnatal Growth
Overall, 249 preterm newborns (18.0%) were small for gestational age and most were associated with the intrauterine growth restriction (73 infants [66.0%]), preeclampsia (42 infants [26.1%]), and fetal distress (33 infants [24.9%]) phenotypes (eFigure 3 in the Supplement).

Figure 1 presents anthropometric patterns from birth to age 2 years according to phenotype and for all preterm newborns and the term newborn group, expressed as mean z scores (actual z score values are presented in eTable 4 in the Supplement). Using z scores from birth to childhood was achievable because international standards, constructed using the prescriptive approach from the WHO,28 were available throughout the childhood growth period. Figure 1A shows birth weight distributed within 0 and −0.5 SD of the standards, with only the intrauterine growth restriction phenotype substantially lower than −1.0 SD. Beginning at age 1 year, 3 growth patterns emerged. The first pattern, which was close to the WHO Child Growth Standards 50th centile, consisted of the term newborn group and preterm newborns with the bleeding, preeclampsia, and severe maternal disease phenotypes. The second pattern consisted of all preterm newborns plus those with the no main condition, infections, and fetal distress phenotypes. The third pattern comprised newborns with the intrauterine growth restriction phenotype, which indicated a pattern that was increasing but remained lower than −1.0 SD of the WHO Child Growth Standards.11

Figure 1B presents the marked decrease in z scores for length and height at age 1 year, with a mixed pattern at age 2 years. Height at age 2 years appeared less affected in the term newborn group and in preterm newborns with some phenotypes. Newborns with the bleeding phenotype remained close to the 50th centile; those with the congenital anomaly and intrauterine growth restriction phenotypes were close to −1.0 SD.
of the WHO Child Growth Standards. Figure 1C shows a parallel pattern for postnatal head circumference growth among phenotypes. By age 2 years, different gradients were observed from newborns with the preeclampsia phenotype to the term newborn group and preterm newborns with other phenotypes, including the no main condition phenotype. Figure 1D shows patterns for birthweight, birthlength, and postnatal body mass index, expressed as $z$ scores. Unlike the 3 patterns in Figure 1A-C, similar trajectories were observed for all groups higher than the 50th centile (ie, children started being overweight for their height by age 1 year).

**Neonatal and Child Morbidity**

Figure 2 shows ORs for neonatal severe morbidity by preterm-birth phenotype compared with the no main condition phenotype. After adjusting for study site, all preterm-birth phenotypes had an increased risk of severe neonatal morbidity compared with the no main condition phenotype. The results remained similar after adjustment for gestational age.

An analysis of each phenotype separately revealed considerable heterogeneity. The highest risk of severe neonatal morbidity was found in newborns with the infections phenotype (OR, 9.7; 95% CI, 1.7-53.9); however, newborns with this phenotype had the highest risk of neonatal and postnatal growth restriction (Figure 1). There was minimal to no attenuation in ORs for preterm-birth phenotypes compared with the no main condition phenotype when adjusting for gestational age (eg, for the infections phenotype, OR, 37.5 [95% CI, 11.9-118.2]; for the bleeding phenotype, OR, 26.6 [95% CI, 7.8-90.7]; and for the congenital anomaly phenotype, OR, 19.9 [95% CI, 6.8-58.1]), indicating that underlying conditions were associated with increased morbidity independent of gestational age (Figure 2).

The phenotypes associated with different morbidities at age 1 year among preterm newborns compared with the term newborn group are shown in eTable 5 in the Supplement. Excluding the congenital anomaly phenotype, newborns with the no main condition (OR, 2.2; 95% CI, 1.8-2.7), infections (OR, 3.4; 95% CI, 2.3-5.2), and fetal distress (OR, 3.7; 95% CI, 2.4-5.9) phenotypes had a higher risk of hospitalization. Those with the preeclampsia (OR, 1.9; 95% CI, 1.5-2.6) and intrauterine growth restriction (OR, 2.0; 95% CI, 1.2-3.3) phenotypes had more infection-associated episodes. Newborns with the no main condition (OR, 4.6; 95% CI, 2.1-9.8), bleeding (OR, 3.5;
The Severe Neonatal Morbidity Index includes bronchopulmonary dysplasia, hypoxic-ischemic encephalopathy, sepsis, neonatal anemia (requiring transfusion), periventricular hemorrhage or leukomalacia, retinopathy of prematurity, necrotizing enterocolitis (Bell stage 2 or higher), and patent ductus arteriosus (requiring pharmacologic treatment or surgery). There were no cases of severe neonatal morbidity in newborns with the severe maternal disease phenotype.

The postnatal patterns varied across phenotype and age. For example, newborns with the intrauterine growth restriction phenotype had lower neonatal morbidity but more growth restriction and neurodevelopmental delay in childhood. Newborns with the no main condition phenotype who were not born group. Heterogeneity was observed, which would have been missed if all preterm newborns had been considered as a single entity. For example, excluding newborns with the congenital anomaly phenotype, those with the fetal distress phenotype had the highest risk of cognitive (OR, 5.1; 95% CI, 2.3-11.1), fine motor (OR, 10.6; 95% CI, 5.1-22.2), gross motor (OR, 3.9; 95% CI, 1.8-8.5), and language (OR, 6.3; 95% CI, 3.2-12.2) development problems. Newborns with the preeclampsia phenotype had a high risk of cognitive (OR, 2.7; 95% CI, 1.4-5.0), fine motor (OR, 4.7; 95% CI, 1.2-18.3), and gross motor (OR, 4.1; 95% CI, 2.4-7.0) development problems. Compared with term newborns, the highest risk of scoring lower than the 10th centile based on INTER-NDA normative values was observed in the fine motor development domain among newborns with the fetal distress (OR, 10.6; 95% CI, 5.1-22.2) and congenital anomaly (OR, 31.1; 95% CI, 3.2-300.3) phenotypes. The risk of fine motor delay was high among all preterm-birth phenotypes.

Discussion

The study's findings indicated that, among an international sample selected to maximize the inclusion of high-risk pregnancies, differential patterns existed with regard to maternal risk factors, the incidence of newborns who were small for gestational age, neonatal and childhood growth, neonatal and childhood severe morbidity, and neurodevelopment across preterm-birth phenotypes. These differential patterns are likely associated with specific pathologic factors that have implications for the growing fetus, as previously reported for early neonatal outcomes.5

The postnatal patterns varied across phenotype and age. For example, newborns with the intrauterine growth restriction phenotype had lower neonatal morbidity but more growth restriction and neurodevelopmental delay in childhood. Newborns with the no main condition phenotype who were not small for their gestational age had lower neonatal morbidity but increased morbidity at age 2 years. Those with the infections phenotype had lower gestational ages at birth and a low risk of being small for gestational age but a high risk of morbidity and postnatal growth restriction.

The association between preterm birth and the phenotype for intrauterine growth restriction, which was observed in 8.0% of all preterm infants, would not have emerged had the age at walking alone because this milestone is the least subject to maternal recall bias and because our data could be compared with the age ranges for achievement of WHO milestones.29 Newborns with the phenotypes for intrauterine growth restriction, bleeding, and congenital anomaly were considerably slower in reaching this milestone, with median ages at achievement of 15.0 months (25th-75th centile, 13.0-18.0 months), 15.0 months (25th-75th centile, 13.0-17.0 months), and 15.2 months (25th-75th centile, 13.5-18.0 months), respectively, representing a 3-month delay compared with the median age of 12 months reported by the WHO.29 Newborns with other phenotypes experienced a 1- to 2-month delay.

Table 2 presents the risk of scoring lower than the 10th centile of the normative population for the INTER-NDA domains24 according to phenotype and compared with the term newborn group. Heterogeneity was observed, which would have been missed if all preterm newborns had been considered as a single entity. For example, excluding newborns with the congenital anomaly phenotype, those with the fetal distress phenotype had the highest risk of cognitive (OR, 5.1; 95% CI, 2.3-11.1), fine motor (OR, 10.6; 95% CI, 5.1-22.2), gross motor (OR, 3.9; 95% CI, 1.8-8.5), and language (OR, 6.3; 95% CI, 3.2-12.2) development problems. Newborns with the preeclampsia phenotype had a high risk of cognitive (OR, 2.7; 95% CI, 1.4-5.0), fine motor (OR, 4.7; 95% CI, 1.2-18.3), and gross motor (OR, 4.1; 95% CI, 2.4-7.0) development problems. Compared with term newborns, the highest risk of scoring lower than the 10th centile based on INTER-NDA normative values was observed in the fine motor development domain among newborns with the fetal distress (OR, 10.6; 95% CI, 5.1-22.2) and congenital anomaly (OR, 31.1; 95% CI, 3.2-300.3) phenotypes. The risk of fine motor delay was high among all preterm-birth phenotypes.

Neurodevelopmental Outcomes

Figure 3 presents the median age (with 25th-75th centiles) of achievement for the WHO gross motor development milestone of age at walking alone because this milestone is the least subject to maternal recall bias and because our data could be compared with the age ranges for achievement of WHO milestones.29 Newborns with the phenotypes for intrauterine growth restriction, bleeding, and congenital anomaly were considerably slower in reaching this milestone, with median ages at achievement of 15.0 months (25th-75th centile, 13.0-18.0 months), 15.0 months (25th-75th centile, 13.0-17.0 months), and 15.2 months (25th-75th centile, 13.5-18.0 months), respectively, representing a 3-month delay compared with the median age of 12 months reported by the WHO.29 Newborns with other phenotypes experienced a 1- to 2-month delay.

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the syndrome's heterogeneity not been recognized. Notably, some preterm newborns with the intrauterine growth restriction phenotype were not included in that phenotype group for this study because they had other pathologic characteristics, such as preeclampsia or congenital anomalies. Other etiologic factors were likely associated with fetal growth restriction (eFigure 2 in the Supplement).

Our results support clustering 1 or more maternal, fetal, or placental conditions to construct phenotypes rather than relying on 1 associated factor. However, 35.1% of preterm births were clustered in the no main condition phenotype, which likely reflects reliance on the use of clinical parameters alone for classification or suggests more complex underlying associations between environmental and nutritional factors.

In this study, the dominant pathologic characteristics in the remaining phenotypes (3 maternal, 3 fetal, and 1 placental) commonly occurred with complications and comorbidities (particularly extraterine infection, chorioamnionitis, and perinatal sepsis) that were associated with the dominant condition and preterm birth (eTable 3 in the Supplement). Nevertheless, there are sufficient data to justify targeting the main putative conditions for the prevention and clinical management of these preterm-birth phenotypes.

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**Table 1. Morbidity Among Children at Age 2 Years According to Preterm-Birth Phenotype**

<table>
<thead>
<tr>
<th>Phenotype</th>
<th>Children with ≥1 hospitalization</th>
<th>Children diagnosed with or treated for ≥3 infections</th>
<th>Children diagnosed with or treated for neurologic disorders</th>
<th>Children diagnosed with or treated for severe clinical conditions</th>
</tr>
</thead>
<tbody>
<tr>
<td>No main condition detected</td>
<td>40/287 (13.9)</td>
<td>47/287 (16.4)</td>
<td>7/287 (2.4)</td>
<td>25/287 (8.7)</td>
</tr>
<tr>
<td>Preeclampsia</td>
<td>9/92 (9.8)</td>
<td>17/92 (18.5)</td>
<td>7/92 (7.6)</td>
<td>11/92 (12.0)</td>
</tr>
<tr>
<td>Infections</td>
<td>25/150 (16.7)</td>
<td>16/150 (10.7)</td>
<td>4/150 (2.7)</td>
<td>13/150 (8.7)</td>
</tr>
<tr>
<td>Bleeding</td>
<td>9/44 (20.5)</td>
<td>6/44 (13.6)</td>
<td>0/44 (0)</td>
<td>2/44 (4.5)</td>
</tr>
<tr>
<td>Intrauterine growth restriction</td>
<td>12/47 (25.5)</td>
<td>4/47 (8.5)</td>
<td>1/47 (2.1)</td>
<td>7/47 (14.9)</td>
</tr>
<tr>
<td>Fetal distress</td>
<td>17/76 (22.4)</td>
<td>12/76 (15.8)</td>
<td>3/76 (3.9)</td>
<td>10/76 (13.2)</td>
</tr>
<tr>
<td>Severe maternal disease</td>
<td>8/55 (14.5)</td>
<td>10/55 (18.2)</td>
<td>1/55 (1.8)</td>
<td>11/55 (20.0)</td>
</tr>
<tr>
<td>Congenital anomaly</td>
<td>7/29 (24.1)</td>
<td>3/29 (10.3)</td>
<td>1/29 (3.4)</td>
<td>12/29 (41.4)</td>
</tr>
<tr>
<td>All preterm newborns</td>
<td>127/780 (16.3)</td>
<td>115/780 (14.8)</td>
<td>23/780 (3.0)</td>
<td>91/780 (11.7)</td>
</tr>
<tr>
<td>Term newborn group</td>
<td>356/3224 (11.0)</td>
<td>326/3224 (10.1)</td>
<td>37/3234 (1.1)</td>
<td>315/3224 (9.8)</td>
</tr>
</tbody>
</table>

Abbreviations: NA, not applicable; OR, odds ratio.

a Adjusted for study site. 95% CIs are based on robust SEs.
b Infections include treatment with antibiotic medication (≥3 regimens), otitis media, pneumonia, bronchiolitis, parasitosis, diarrhea, vomiting, exanthema, skin disease, fever lasting 3 days or more (≥3 episodes), meningitis, and other infections that required antibiotic treatment.
c Neurologic disorders include seizures, cerebral palsy, and other neurologic disorders.
d Severe clinical condition includes cardiovascular problems, gastroesophageal reflux, any hemolytic condition, injury trauma, and any condition that required surgery.

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**Figure 3. Median Age for Developmental Achievement of Walking Alone According to Preterm-Birth Phenotype**

Horizontal lines represent interquartile ranges. For comparison, the 25th and 75th centiles of the WHO periods of achievement for the same developmental milestone are shown in the shaded area (with the median age represented by a broken vertical line). IUGR indicates intrauterine growth restriction; NMCD, no main condition detected; and WHO, World Health Organization.
The present study is distinct because (1) it has a firm conceptual basis; (2) it was designed to explore a priori hypotheses associated with the heterogeneity of preterm birth; (3) it accurately estimated gestational age in all newborns; (4) it oversampled newborns with low gestational ages to maximize the number of subgroups considered very high risk; (5) it achieved a 1:4 ratio for enrolled preterm and term newborns, substantially exceeding the 1:1 target; (6) it monitored growth, morbidity, and neurodevelopment from birth to age 2 years using standardized data collection systems; and (7) it assessed neurodevelopment compared with international normative values using a validated psychometric tool.

The study's results have implications for research and clinical practice. Preterm birth is a distinct syndrome in the medical field because it is defined by time (ie, gestational age at birth rather than etiologic, clinical, nutritional, or laboratory characteristics, as with other syndromes). Hence, it is important to estimate the gestational age as accurately as possible, consistent with the 2016 WHO guidelines, and to avoid the use of low birth weight (ie, <2500 g regardless of gestational age) as a factor. It has been asserted that low birth weight is a necessary factor to include because, in many regions of the world, gestational age cannot be reliably estimated. However, actions to minimize the incidence of preterm birth may nonetheless be taken and are warranted for a syndrome that occurs in at least 10% to 12% of all births worldwide and has serious implications for human developmental capacity. Governments, international organizations, and donors can work to improve antenatal care so that every woman receives adequate evaluation early in pregnancy.

### Limitations
This study has several limitations. The loss to follow-up at age 2 years was higher than that of previous studies, as some sites were located in rural or semiurban areas. This limitation was considered when the INTERBIO-21st study was designed, but we concluded that the loss of follow-up at such sites was offset by the enrollment of a high-risk population with accurate gestational ages and a higher number of more severe exposures (eg, infections and inadequate nutrition). Nevertheless, the risk of systematic loss to follow-up bias according to phenotype is unlikely because, with the exception of the intrauterine growth restriction phenotype, the follow-up rates across phenotypes were similar.

We could not include multiple births because we focused, for several logistical and sample size reasons, on singletons. In addition, we did not include stillbirths or late terminations because the childhood follow-up component was fundamental to the study. We also did not document the ways in which urinary tract infections were specifically diagnosed or record whether prophylactic antibiotic medications were prescribed for the treatment of group B streptococcus colonization. We categorized congenital anomaly as a separate phenotype to reinforce the integrity of preterm birth as a distinct syndrome. However, given the small sample of infants with the congenital anomaly phenotype (3.5% of all preterm newborns), we could not draw meaningful conclusions.

Although cluster methods are susceptible to nuances in the data, the fact that we obtained clusters that were similar to those acquired in a previous study, using identical methods but a data set with a different risk profile,
supports our hypothesis that these phenotypes occur systematically across populations. However, the phenotype prevalence changes depending on which factors predominate.

We recognize that there are competing etiologic characteristics and risk factors among the preterm birth, intrauterine growth restriction, and stillbirth syndromes,34 which were highlighted by Lee et al.35 Our study was designed to explore the preterm and intrauterine growth restriction syndromes in parallel. This overlapping approach was required because intrauterine growth restriction is an equally complex syndrome that has multiple etiologic characteristics and presentations. However, our focus on developmental outcomes precluded including stillbirths. Future research may explore these associations in detail.

Conclusions

In this study, the preterm birth syndrome was composed of well-defined phenotypes with differential neonatal morbidity, early childhood morbidity, growth, and neurodevelopment up to age 2 years. Therefore, the concept of preterm birth as an exclusively time-based entity may no longer be appropriate. Phenotypic classification of preterm newborns is likely to provide a better understanding of the etiologic factors and mechanisms associated with preterm birth. Although the study’s findings indicated that approximately 35% of phenotypes were not associated with distinct clinical conditions, newborns with preterm-birth phenotypes nevertheless remain at a higher risk of growth and developmental problems.

ARTICLE INFORMATION

Accepted for Publication: October 14, 2020.
Published Online: March 1, 2021.

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Obtained funding: Villar, Uauy, Kennedy.

Administrative, technical, or material support: Villar, Papageorghiou, Craik, Barsosio, Carvalho, Fernandes, Cheikh Ismail, Lambert, Norris, Stein, Tshivuila-Matala, Zondervan, Winsey, Nosten, Uauy, Bhutta.


Conflict of Interest Disclosures: Dr Villar reported receiving grants from the Bill and Melinda Gates Foundation during the conduct of the study and grants from the Bill and Melinda Gates Foundation, the European Research Council, and the National Institute of Health Research, personal fees from the British Journal of Obstetrics and Gynecology, Intelligent Ultrasound, the National Health Service of the United Kingdom, and Ultrasound Diagnostic Services; being a cofounder, shareholder, and senior advisor of Intelligent Ultrasound; being a paid scientific advisor for Oxford University Innovation; and being a deputy editor-in-chief of the British Journal of Obstetrics and Gynecology outside the submitted work. Dr Ochien reported receiving grants from the University of Oxford during the conduct of the study. Dr Lambert reported receiving grants from the Bill and Melinda Gates Foundation during the conduct of the study. Dr Tshivuila-Matala reported receiving a scholarship and stipend for living expenses from the Rhodes Trust during the conduct of the study. Dr Winsey reported receiving grants from the Nuffield Department of Women's and Reproductive Health during the conduct of the study. No other disclosures were reported.

Funding/Support: This study was supported by grant 49138 from the Bill and Melinda Gates Foundation to the University of Oxford.

Role of the Funder/Sponsor: The funding organization had no role in the design and conduct of the study; collection, management, analysis, and interpretation of the data; preparation, review, or approval of the manuscript; and decision to submit the manuscript for publication.


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REFERENCES


