**Predicting childhood overweight and obesity using maternal and early life risk factors: a systematic review**

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**Abbreviations**

WHO - World Health Organisation

NCMP - National Child Measurement Programme

DOHaD - Developmental origins of health and disease

MRH - Maternal resources hypothesis

IOTF - International Obesity Task Force

CDC - Centers for Disease Control

IQR – interquartile range

AUROC – Area under receiver operating curve

ALSPAC – Avon Longitudinal Study of Parents and Children

**Abstract**

**Background:** Childhood obesity is a serious public health challenge and identification of high-risk populations with early intervention to prevent its development is a priority. We aimed to systematically review prediction models for childhood overweight/obesity and critically assess the methodology of their development, validation and reporting.

**Methods:** Medline and Embase were searched systematically for studies describing the development and/or validation of a prediction model/score for overweight and obesity between 1 to 13 years of age. Data were extracted using the Cochrane CHARMS checklist for Prognosis Methods.

**Results:** Ten studies were identified that developed (one), developed and validated (seven) or externally validated an existing (two) prediction model. Six out of eight models were developed using automated variable selection methods. Two studies used multiple imputation to handle missing data.

From all studies, 30475 participants were included. Of 25 predictors, only seven were included in more than one model with maternal body mass index, birthweight and gender the most common.

**Conclusion**: Several prediction models exist but most have not been externally validated or compared to existing models to improve predictive performance. Methodological limitations in model development and validation combined with non-standard reporting restrict the implementation of existing models for the prevention of childhood obesity.

**Introduction**

The World Health Organisation (WHO) has identified childhood overweight and obesity as one of the most serious public health challenges of the 21st century with 42 million children aged under 5 years estimated as overweight globally in 20141. Data from the National Child Measurement Programme (NCMP) in England showed that in 2014/15, 22% of children in Reception (aged 4 to 5 years) and 33% in Year 6 (aged 10 to 11 years) were classified as overweight or obese with children in most deprived areas twice as likely than children in least deprived areas to be obese2. In 2012, the WHO published a report on population-based approaches to childhood obesity prevention, which identified improved government structures to support policy and intervention as well as population-based and community based interventions as actions to prevent childhood obesity3. In 2014, the European Union published a six-year action plan on childhood obesity with the goal of contributing to halting the rise in childhood overweight and obesity by 2020. In 2016, the UK Government published a plan for action for tackling childhood obesity with the aim to significantly reduce rates of childhood obesity within the next ten years by supporting healthier choices in children and engaging communities, schools and industry to make food and drink healthier4.

There is evidence that the in-utero environment induces a response in the foetus which can lead to enhanced susceptibility for diseases in later life5. This concept is described as the “developmental origins of health and disease (DOHaD)”. Developing foetuses adapt to an adverse in utero environment by undergoing structural, physiological and hormonal changes which are beneficial for short term survival, but at a cost for future health6 which could be transmitted through generations7. The “maternal resources hypothesis (MRH)” suggests that non-genetic evolution has led to a competitive dominance of adipocytes over other cell types in the acquisition and sequestering of energy in the body, which is maintained by the co-existence of excess maternal resources and sedentary behaviour during pregnancy leading to continued dysfunction in foetal metabolism8. Behavioural patterns are transmitted between generations through socially mediated learning9 and the postnatal environment could affect the behaviour of infants and young children based on that of the primary caregiver8. Thus, it has been suggested that DOHaD should include all aspects of environment and all sensitive windows (preconception, pregnancy, early childhood and any others yet to be identified)7.

Hence, the WHO Commission on ending childhood obesity considered it essential to address critical time periods in development including pre-conception and pregnancy as well as treating children identified as obese10. The increasing prevalence of obesity in women of reproductive age affects the health of the mother and puts the offspring at risk of developing childhood obesity and its consequences11. Given the lack of evidence on effective long term treatments, the focus of reducing childhood obesity rates should be on prevention12. Key to an effective prevention strategy is the ability to identify individuals at particular risk. There is increased risk of persistence of childhood weight status into adulthood13-16 particularly in children with two obese parents17-19 with a meta-analysis concluding a low probability of weight change without weight loss treatment20. Although this tracking of childhood BMI to adulthood was weaker in late adulthood21, the identification of high-risk populations and intervening as early as possible to prevent the development of overweight and obesity should be a priority22 due to the increased risk of adult morbidity and mortality associated with overweight and obesity in childhood and adolescence23. Once high risk populations are identified, mathematical models on childhood obesity trajectories that predict energy imbalance including excess energy intake underlying obesity24,25 and calculates the magnitude of intervention necessary to achieve change in weight25 can be used to guide the intervention.

The aim of this study was to systematically review studies of prediction models for childhood overweight and obesity using maternal and/or early life risk factors and critically assess the development and reporting of the methodology used to develop these models.

**Methods**

Medline and Embase were searched from their start dates to December 2016 using recommended filters and the bibliographies and citations of all included studies were hand searched (using Web of Science Core Collection). The outcome considered was overweight and obesity between 1 to 13 years of age. No criteria was defined for overweight and obesity as different criteria can be considered given the age under consideration. The following search strategy was used:

{Pediatric Obesity/ OR Fetal Macrosomia/ OR

[(child or childhood or children or p#ediatric\* or infant\* or toddler or embry\* or prenatal\* or neonat\*).mp. AND (obes\*.mp. OR overnutrition/ or obesity/ or overweight/ OR overweight.mp. OR over weight.mp.)]} AND

[exp causality/ OR ((Reinforc\* or Enabl\* or predispos\*) and factor\*).mp. OR (risk\* or predict\* or causal\* or prognos\* or causation).mp.] AND

[exp Maternal Behavior/ OR maternal.mp. OR mother\*.mp. OR early life.mp.]

*Eligibility criteria*

All studies that reported on one or more multivariable prediction models or scores that have been developed for individual risk estimation of future risk of childhood overweight and obesity were included. Studies that developed, developed and validated or just validated a risk score were not differentiated. The review was limited to studies conducted in humans and published in English. No limits were imposed on study timing or setting.

*Data extraction and critical appraisal*

The list of data extraction was based on the CHARMS checklist published by the Cochrane Prognosis Methods Group26. The TRIPOD statement was used to assess transparency in reporting27. NZ assessed all articles and extracted the data. Items extracted from studies describing model development included study design, study population and location, number of study participants, outcome and age of outcome if available, method of modelling, method of internal validation (random split of data, bootstrapping or cross-validation), number of predictors considered and included in the final model, model presentation and predictive performance including measures of discrimination and calibration where available.

For studies describing external model validation alone, items extracted included study design, study population and location, number of study participants and model performance. Predictors were checked to confirm that these were the same as the original model.

We have critically assessed the conduct and reporting of the methods used to develop these risk prediction models. However, a quantitative synthesis of the prediction models’ results was not performed as formal methods for meta-analysis of models are not yet fully developed and was beyond the scope of this review.

**Results**

From the 11867 articles identified by the search strategy, 143 full articles were reviewed of which nine articles were identified for inclusion in this review (Figure 1). An additional study was identified through hand searching the citations of the included studies. Eight of the studies developed a risk score, seven of which were internally (six) and/or externally (two) validated in the same publication, and two were external validation studies of two of the eight existing prediction models (Table 1).

*Study reporting*

Using the TRIPOD27 reporting recommendation, a median of 23 (interquartile range (IQR), 22 to 24) items out of 37 (31 for derivation or validation alone) were reported suggesting some shortcomings (Table 3). As this review assessed the extent of reporting, authors were not contacted to seek further information.

*Study designs, population and sample size*

Most of the studies used data from prospective birth cohorts and two studies used cross-sectional studies in childhood with retrospective data collection of maternal and early life factors. All the studies were in high income countries with the exception of data from Seychelles in the study that pooled cohort data from three studies.

*Outcomes, number of patients and events*

The outcome was overweight (three)28-30, obesity (three)31-33 or both (two)34,35 in the eight included studies that developed a score and the age at which this was predicted varied from one to 13 years of age in children. Sex- and age- specific BMI was calculated using the International Obesity Task Force (IOTF)29-31,34,35, Centers for Disease Control (CDC)32, WHO28 and UK90 growth chart33 criteria and appropriate thresholds for overweight or obesity applied.

The number of participants used to develop the prediction models was clearly reported in all studies. The number of participants was 30475 from all studies and the median number was 2015 (IQR 1644 to 5083) across the studies. Six29,30,32-35 out of eight studies reported the prevalence of the outcome in the study population of which two reported the prevalence of both overweight and obesity (12-23% overweight and 3-32% obesity). Where recorded, the median number of events that was used in model development was 821 (IQR 549 to 1374) for overweight and 133 (IQR 104 to 170) for obesity.

*Risk predictors*

Across the studies analysed, 57 putative predictors (Table 2) with a median of 11 risk predictors (IQR 8 to 19) were considered in the development models. These were defined a priori in six studies29,30,32-35, identified through previous multivariable regression31 or defined a priori for maternal predictors and through univariable regression for child predictors28. Only four of the six studies that defined predictors a priori provided the rationale or references for including these predictors.

Twenty-five predictors were included in the final risk prediction models. However, eighteen of these predictors were only included in one risk score model. The final reported prediction models included a median of six (IQR 5 to 6) predictors with maternal pre-pregnancy BMI, birthweight and infant gender included in seven out of eight scores (Table 2). Two studies assessed risk at birth (using preconception, antenatal and birth factors)29,35 whereas other scores incorporated weight gain in the first year of life30-34 predicting risk from the age of 12 months and over or childhood age- and sex- adjusted BMI at 5 years of age28 to predict risk at 10 years of age.

*Treatment of continuous risk predictors*

Four (50%) risk prediction models retained continuous predictors as continuous28,29,32,35, two (25%) categorised or dichotomised all continuous predictors and one (12.5%) retained some continuous predictors as continuous and categorised some predictors33. It was unclear how continuous risk predictors were treated in one study but a categorical score chart developed so it is likely that all continuous variables were categorised or dichotomised30.

*Missing data*

Four studies only included cases with complete data in model development28,29,33,34, two studies carried out multiple imputation32,35 and one study did not report the presence or handling of missing data31. The remaining study included participants with full anthropometric data at follow-up when outcome was assessed but it is unclear if there were missing data at previous data collection points and how this was handled30.

One of the studies that carried out multiple imputation had on average 1.7% (range 0 to 11.4%)35 missing data for each predictor whereas 17% of the other study32 participants had missing data for at least one predictor. Two of the studies that carried out complete case analysis, 23.8%29 and 27.2%28 of the sample were excluded due to the missing data but it is unclear what percentage of sample was excluded for missing data alone in the other studies33,34.

*Model building*

Six (75%) studies used automated variable selection (stepwise, backward deletion) to derive the final predictive model29,30,32-35.

All studies were clear on the method used to develop the prediction model - logistic regression was used in seven studies29-35 whereas linear regression was used in one study28. One study had selected predictor variables based on previous multivariable logistic regression analysis and only carried out univariable logistic regression to assign integer values to the categories of risk predictor variables without any further modelling31. Two models29,33 included interaction terms while modelling whereas there was no mention of interaction terms while modelling in the other studies.

*Predictive performance*

Model performance was assessed in all studies, seven of which used area under the receiving operator curve (AUROC) in either the derivation, validation or both cohorts. The other study tested for specificity and predictive value alone28. Although model performance was assessed and validated in all studies, only one study reported change in regression co-efficient post validation and updating the model29. Two studies from the UK used data from the same birth cohort (Avon Longitudinal Study of Parents and Children ALSPAC) for validation of the same outcome but at different ages (two33 and five36 years). Model development AUROC ranged from 0.64 to 0.91 (median 0.78, IQR 0.70 to 0.81). The AUROC of 0.91 was replicated in internal validation using bootstrapping and only decreased to 0.89 on external validation33.

Three studies29,32,35 carried out Hosmer-Lemeshow tests to test calibration, two of which did so during model development both achieving p>0.5. All studies assessed model classification (sensitivity and specificity) though one study31 did not present positive and negative predictive values.

*Internal Validation*

With the exception of two, all studies internally validated the models by random split of data30,34, random split followed by cross-validation28 or bootstrapping29,32,33. Of the studies that did not internally validate the model, one validated the model externally in two separate cohorts35 whereas the other was externally validated in a subsequent publication with overlapping authors in the development and validation papers31,37. Additionally one of the studies that internally validated the model using random split was also externally validated in a subsequent publication by the same authors30,36. Model validation AUROC ranging from 0.75 to 0.91 (median 0.78, IQR 0.77 to 0.81) was achieved and the original model was updated in one study only29. Of the studies that carried out Hosmer-Lemeshow test for calibration, one did not report the exact p-value but that p>0.5 was achieved32 whereas the other achieved p=0.30 on recalibration post validation29.

*External validation*

Only four of eight models have been externally validated - once for three models all of which used data from the same country for validation33,36,37 and twice for one model that was developed in Finland and validated in Italy and US35. Of the models validated using data from the same country, two studies calculated AUROC, which were 0.8936 and 0.6736. The only study that externally validated the model in two countries other than that in which it was developed35 found that AUROC (0.70, confidence intervals 0.63 to 0.77) and calibration (Hosmer-Lemeshow p=0.12) was satisfactory in one population but although AUROC (0.73, confidence intervals 0.67 to 0.80) was satisfactory in the other, calibration (Hosmer-Lemeshow p=0.02) was not. The predictors and model were then tailored to these populations by carrying out a replication analysis using stepwise logistic regression such that calibration achieved satisfactory levels. The initial model developed in Finland included six risk factors and reduced to three and five for the Italian and US cohort respectively with only two factors remaining consistent across all three models (maternal and paternal BMI). Ethnicity was introduced in the risk prediction score for the US and this was primarily because the birth cohort in Finland had high ethnic homogeneity. One of the external validation studies36 also developed a recalibrated model using multivariable logistic regression to apply a recalibrated algorithm reflecting the characteristics of the validation cohort, imputed model for missing risk factor prediction and a recalibrated imputed model which incorporated the two. This led to an increase in discrimination compared to the original model from 2% in the recalibrated to 25% in the recalibrated imputed model.

*Model presentation*

The complete regression formula (including all regression coefficients) was presented in six studies29,30,32-35 and two of these studies provided a decision rule/score chart or risk score algorithm29,30. Of the remaining two studies, one provided the regression coefficients28 whereas the other only provided a score chart31.

**Discussion**

To our knowledge, this is the first systematic review to examine prediction models for childhood overweight and obesity. Eight studies that developed prediction models were identified, however four of these prediction scores have been externally validated once or twice and there is no evidence of further validation or validation in populations outside of those in which this was developed. Additionally new models have been developed with no evidence of comparison with already existing models and none of the models have been compared with each other to assess predictive performance. There were inadequacies identified in reporting of the methodology of development of risk prediction models and there is no evidence of implementation of the risk scores. Whilst there is clear overlap between risk factors included in the prediction models, no single risk factor has been included in all prediction models with maternal pre-pregnancy BMI, infant gender and birthweight being the most commonly included. Thus, it is difficult to recommend the use of any one score, as there are no consistent predictors, no comparison between models and the outcome has been variable and predicted at different ages through childhood up to 13 years of age.

The question of predictors considered for inclusion in the model also needs to be considered. Although not included in the final prediction model, several predictors around infant temperament was considered. There are self-reported by parents and highly likely to be subjective. Additionally, these factors were identified a priori based on a previous systematic review but the conclusion of the review was that the evidence was inconclusive due to limited number of studies38.

Thirteen of the 25 risk factors identified were preconception and thus some of these could prove impactful in planned pregnancies such as maternal and paternal BMI whereas others are non-modifiable such as ethnicity. Although factors such as maternal education, occupation and income are modifiable, it is difficult to do so. Maternal smoking during pregnancy and hospital delivery were the only two antenatal risk factors identified and included in risk prediction. Eight of the ten early life risk factors identified can be broadly classified into weight gain particularly in first year of life and breastfeeding including weaning both of which are modifiable. The other two risk factors were gender and birthweight, of which gender is non-modifiable but birthweight can be monitored and is considered modifiable by factors known to affect foetal growth39.

Some key aspects of multivariable model development and validation need to be considered. These include handling missing data, method of treatment of continuous variables, selecting variables for inclusion in the model and methods of validation including assessing discrimination and calibration40. Missing data was identified in most studies which can introduce bias if inappropriately handled thus impeding the construction of a valid prediction model41. Multiple imputation minimises the effect of missing data provided data is missing at random42 and enables the use of all available data but was only done in 25% of studies included in this review. All other studies excluded participants with missing data which is an acceptable approach only if the amount of missing data is small43 however, these studies did not provide any indication of how much data was missing per individual and per variable to enable readers to reach their own judgement of the validity of the prediction.

At least three prediction models categorised some or all continuous variables for inclusion in the model. However, discarding information through categorisation of continuous variables to estimate a continuous relationship between a predictor variable and risk has been shown to lead to a substantial loss of power and precision44, thus reducing the efficiency of the analysis with increased probability of biased estimates45 and Type 146. In addition, a model that categorises continuous variables is unrealistic as individuals close to but on opposite side of the category cut-point will be characterised as having very different outcome when a very similar outcome is more likely47. It is recommended that continuous predictors are retained as continuous and suitable functions such as fractional polynomial are used47,48. Although this is true from a methodological point of view, the clinical practice in terms of implementation of any score needs to be considered. For example, the National Institute for Health and Care Excellence in the UK recommends action before, during and after pregnancy in women with BMI greater than 3049. Thus, including this categorisation could make the prediction rule easier to incorporate into clinical practice.

Although predictors shown to have little effect on the outcome should not be included in the prediction, the method of selection of predictor variables for inclusion is crucial. The majority of studies (75%) used an automated variable selection method which increases the likelihood that variables that do not truly predict the outcome will be identified as a predictor50. This is because it is a data-driven approach that cannot account for clinical relevance leading to biased regression estimates and poor predictions as true predictors could be excluded due to lack of power51,52. It also leads to loss of information due to inclusion of variables based on a binary decision. It has been suggested that a more reasonable reduction of variables using automated selection procedures could be achieved by using a liberal selection criteria such as p = 0.5052 instead of 0.05 which is more commonly used and has been used in all the prediction models included in this review that used this procedure. It could also be important to retain predictors known to be important from literature but does not achieve statistical significance in the model development dataset51.

Once developed, the performance of a model needs to be evaluated to demonstrate usability. Although a biased model could provide useful clinical separation into groups if the predictor information entered into the model is strong53, evidence is needed that the model performs well in populations other than that in which it was developed54. Validation can be internal or external using a completely different sample thus also examining the generalisability of the model54. Six studies (75%) internally validated the model through random split of the dataset (two), random split and cross-validation (one) or bootstrapping (three). Four studies (50%) externally validated the model, only one of which externally validated the model in cohorts from different countries. This was followed by replication analysis to rebuild the model in these two cohorts resulting in only two predictors being retained across all three models in this study (maternal and paternal BMI). As the use of random split sample decreases the precision of estimates and increases the frequency of missing important independent variable55, there is limited value in doing so unless the sample size is particularly large51. A non-random or chronological split has been suggested as a more precise approach but internal methods such as bootstrapping and cross-validation remain more informative53.

This review has been carried out with a systematic approach thus identifying all studies that have developed and/or validated a risk prediction model for childhood overweight and obesity. However, heterogeneity exists at many levels particularly the outcome (overweight, obesity or both) under consideration and age at which outcome is predicted. This heterogeneity combined with the deficiency of external validation limits the applicability of these scores. Additionally, poor reporting in aspects of development of the prediction models was observed with insufficient detail on steps involved in model building. Risk prediction models have nearly all been developed or validated in developed countries but almost half and one quarter of the estimated 42 million overweight children under the age of five years live in Asia and Africa respectively1. Models tailored to these countries are important, as associations are known to vary between ethnic groups.

**Conclusion**

Despite the existence of several models for the prediction of childhood overweight and obesity, most have not been externally validated or compared to existing models to assess predictive performance. Moreover as the outcome has been predicted at different ages, it may not be possible to combine or compare all models against each other. This review also highlights methodological limitations in model development and validation combined with non-standard reporting thus limiting the usability of these prediction models.

There remains a need to develop new methods for combining findings from existing prediction models and develop prediction models using robust methods of development followed by external validation and recalibrating to populations, which would then enable assessment of impact of the implementation of the score.

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Table 1: Summary of prediction models in the included studies

|  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Author, year | Druet2012 | Manios2013 | Manios2016 | Morandi2012 | Pei2013 | Redsell2016 | Robson 2016 | Santorelli2013 | Steur2011 | Weng2013 |
| N, derivation | 8236 | 2294 | - | 4032 | 1515 | - | 166 | 1868 | 1687 | 10810 |
| N, validation | 8236 | - | 5946 | 1503 | 757 | 980 |   | 867-880 | - | 2703 |
| Country | UK, Europe, America and Seychelles | Greece | Greece | Finland Validation – Italy, US | Germany | United Kingdom | USA (Latino cohort) | United Kingdom | The Netherlands | United Kingdom |
| Design | Meta-analysis of three birth cohorts  | Cross-sectional with retrospective data collection | Cross-sectional with retrospective data collection | Prospective birth cohort (Finland, US) Retrospective cohort (Italy) | Prospective birth cohorts | Prospective birth cohort | Birth cohort | Prospective birth cohorts | Prospective birth cohort | Prospective birth cohort |
| Outcome | Childhood obesity | Childhood obesity (9-13 years) | Childhood obesity (6-15 years) | Obesity and overweight at 7 years  | Overweight at age 10 years | Obesity at age 5 years | Obesity at 5 years | Obesity at age 1 year | Overweight at age 8 years | Overweight at age 3 years |
| Variables included | 4 | 5 | 5 | 6 | 5 | 7 | 10 (full model)5 (reduced model) | 4 | 6 | 7 |
| Derivation AUROC | - | 0.64 | - | 0.67 (overweight-obesity)0.78 (obesity) | - | - | 0.84 (full model)0.82 (reduced model) | 0.91 | - | 0.72 |
| Validation AUROC | 0.77 | - | 0.64 | 0.70, 0.73 | - | 0.67 (original model)0.93 (recalibrated) | - | 0.89 | - | 0.76 |
| TRIPOD | 21 | 19 | 20 | 28 | 23 | 23 | 24 | 29 | 24 | 23 |

Table 2: Predictor variables assessed (-) and included (+) in the models

|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Author, year | Druet2012 | Manios2013 | Morandi2012 | Pei2013 | Robson2016a | Santorelli2013 | Steur2011 | Weng2013 |
| Gender | + | + | - | + | + | + | + | + |
| Gestational age | - |  |  |  |  | - |  |  |
| Weight change 0-6 months |  |  |  |  | **+** |  |  |  |
| Weight gain 0-1 year (categorised) | - | + |  |  |  |  |  | + |
| Weight gain 0-1 year (continuous) | + |  |  |  |  | + |  |  |
| Weight gain 0-5 years (categorised)  |  |  |  | - |  |  |  |  |
| Standardised BMI at 60-64 months  |  |  |  | + |  |  |  |  |
| Birthweight | + |  | + | + | **+** | + | + | + |
| Maternal age |  |  |  |  | **+** |  |  | - |
| Maternal BMI | + | + | + |  | **+** | + | + | + |
| Maternal education |  | + |  |  |  |  | - | - |
| Pre-pregnancy maternal smoking |  |  | - |  |  |  |  |  |
| Maternal smoking during pregnancy |  | + | + | + |  | - | - | + |
| Maternal occupation |  |  | + |  |  |  |  |  |
| Maternal employment |  |  |  |  |  |  | - | - |
| Employment in pregnancy |  |  |  |  |  |  |  | - |
| Single parenthood/marital status |  |  | - |  |  |  |  | + |
| Gestational weight gain |  |  | - |  |  |  |  |  |
| Maternal alcohol consumption |  |  |  |  |  |  |  | - |
| Maternal feelings of depression |  |  |  |  |  |  |  | - |
| Maternal health |  |  |  |  |  |  |  | - |
| Maternal diabetes |  |  |  |  |  |  |  | - |
| Gestational diabetes |  |  |  |  |  | - |  |  |
| Hospital delivery |  |  |  |  |  |  | + |  |
| Delivery type |  |  |  |  |  |  | - | - |
| Number of household members |  |  | + |  |  |  |  |  |
| Obesity predisposing single-nucleotide polymorphisms  |  |  | - |  |  |  |  |  |
| Paternal BMI |  |  | + |  |  |  | + | + |
| Paternal education |  |  |  |  |  |  | - |  |
| Paternal employment |  |  |  |  |  |  | - |  |
| Family income (categorised) |  |  |  | + |  |  |  | - |
| Parental education (categorised) |  |  |  | + |  |  |  |  |
| Solids introduced at < or >6 months |  |  |  |  | + |  |  | - |
| Exclusive breastfeeding at 4-6 weeks |  |  |  |  | **+** |  |  |  |
| Any breastfeeding at 6 months |  |  |  |  | + |  | - |  |
| Ever breastfed in first year |  |  |  |  |  |  |  | + |
| Breastfeeding duration |  |  |  |  |  |  |  | - |
| Ever formula fed |  |  |  |  |  |  |  | - |
| First child/older siblings/number of own children |  |  |  |  | + |  | - | - |
| English language proficiency |  |  |  |  | + |  |  |  |
| Ethnicity |  |  |  |  |  | - | - | + |
| Smoking in the parental house |  |  |  |  |  |  | + |  |
| Living in a highly urbanized environment (≥2500 address/km2) |  |  |  |  |  |  | - |  |
| Maternal vegetable consumption during pregnancy |  |  |  |  |  |  | - |  |
| Premature birth of child |  |  |  |  |  |  | - |  |
| Region of birth |  |  |  |  |  |  | - |  |
| Financial status |  |  |  |  |  |  |  | - |
| Child care arrangements |  |  |  |  |  |  |  | - |
| Unhappy when feeding interrupted  |  |  |  |  |  |  |  | - |
| Makes a fuss going to sleep  |  |  |  |  |  |  |  | - |
| Makes a fuss after waking |  |  |  |  |  |  |  | - |
| Upset when not getting things |  |  |  |  |  |  |  | - |
| Does the infant sit up? |  |  |  |  |  |  |  | - |
| Does the infant stand? |  |  |  |  |  |  |  | - |
| Does the infant grab objects? |  |  |  |  |  |  |  | - |
| Does the infant hold objects? |  |  |  |  |  |  |  | - |
| Can the infant walk? |  |  |  |  |  |  |  | - |

**+** indicates predictors retained in the reduced model.

Table 3: TRIPOD items reported in the ten studies

|  |  |  |  |
| --- | --- | --- | --- |
| Title and abstract |  | TRIPOD item description | Reported |
| Title | 1 | Identify the study as developing and/or validating a multivariable prediction model, the target population, and the outcome to be predicted. | 8 |
| Abstract | 2 | Provide a summary of objectives, study design, setting, participants, sample size, predictors, outcome, statistical analysis, results, and conclusions. | 10 |
| Introduction |
| Background and objectives | 3a | Explain the medical context (including whether diagnostic or prognostic) and rationale for developing or validating the multivariable prediction model, including references to existing models. | 9 |
| 3b | Specify the objectives, including whether the study describes the development or validation of the model, or both. | 10 |
| Source of data | 4a | Describe the study design or source of data (e.g., randomized trial, cohort, or registry data), separately for the development and validation datasets, if applicable. | 10 |
| 4b | Specify the key study dates, including start of accrual; end of accrual; and, if applicable, end of follow-up. | 10 |
| Participants | 5a | Specify key elements of the study setting (e.g., primary care, secondary care, general population) including number and location of centres. | 10 |
| 5b | Describe eligibility criteria for participants. | 10 |
| 5c | Give details of treatments received, if relevant. | - |
| Outcome | 6a | Clearly define the outcome that is predicted by the prediction model, including how and when assessed. | 10 |
| 6b | Report any actions to blind assessment of the outcome to be predicted. | 0 |
| Predictors | 7a | Clearly define all predictors used in developing the multivariable prediction model; including how and when they were measured. | 8 |
| 7b | Report any actions to blind assessment of predictors for the outcome and other predictors. | 0 |
| Sample size | 8 | Explain how the study size was arrived at. | 10 |
| Missing data | 9 | Describe how missing data were handled (e.g., complete-case analysis, single imputation, multiple imputation) with details of any imputation method. | 4 |
| Statistical analysis methods | 10a | Describe how predictors were handled in the analyses. | 9 |
| 10b | Specify type of model, all model-building procedures (including any predictor selection), and method for internal validation. | 8 |
| 10c | For validation, describe how the predictions were calculated. | 8 |
| 10d | Specify all measures used to assess model performance and, if relevant, to compare multiple models. | 8 |
| 10e | Describe any model updating (e.g., recalibration) arising from the validation, if done. | 3 |
| Risk groups | 11 | Provide details on how risk groups were created, if done. | 0 |
| Development vs validation | 12 | For validation, identify any differences from the development data in setting, eligibility criteria, outcome, and predictors. | 2 |
| Results |
| Participants | 13a | Describe the flow of participants through the study, including the number of participants with and without the outcome and, if applicable, a summary of the follow-up time. A diagram may be helpful. | 6 |
| 13b | Describe the characteristics of the participants (basic demographics, clinical features, available predictors), including the number of participants with missing data for predictors and outcome. | 7 |
| 13c | For validation, show a comparison with the development data of the distribution of important variables (demographics, predictors, and outcome). | 1 |
| Model development | 14a | Specify the number of participants and outcome events in each analysis. | 4 |
| 14b | If done, report the unadjusted association between each candidate predictor and outcome. | 1 |
| Model specification | 15a | Present the full prediction model to allow predictions for individuals (i.e., all regression coefficients, and model intercept or baseline survival at a given time point). | 6 |
| 15b | Explain how to use the prediction model. | 6 |
| Model performance | 16 | Report performance measures (with CIs) for the prediction model. | 7 |
| Model updating | 17 | If done, report the results from any model updating (i.e., model specification, model performance). | 1 |
| Discussion |
| Limitations | 18 | Discuss any limitations of the study (such as non-representative sample, few events per predictor, missing data). | 10 |
| Interpretation | 19a | For validation, discuss the results with reference to performance in the development data, and any other validation data. | 3 |
| 19b | Give an overall interpretation of the results, considering objectives, limitations, results from similar studies, and other relevant evidence. | 10 |
| Implications | 20 | Discuss the potential clinical use of the model and implications for future research. | 10 |
| Other information |
| Supplementary information | 21 | Provide information about the availability of supplementary resources, such as study protocol, Web calculator, and datasets. | 6 |
| Funding | 22 | Give the source of funding and the role of the funders for the present study. | 9 |