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Title: Derivation and validation of a total fruit and vegetable intake prediction model to identify targets for biomarker discovery using the UK National Diet and Nutrition Survey.

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List of abbreviations: AIC, Akaike information criterion; BIC, Bayesian information criterion; DINO, Diet In Nutrients Out; FV, fruit and vegetable; LR, likelihood ratio; MBPs, multi-metabolite biomarker panels; MG, modelling group; TFVpred, Total fruit and vegetable prediction; VG, validation group; VIF, variance inflation factor.

Abstract (297 words)

Background:

Dietary assessment in research and clinical settings is largely reliant on self-reported questionnaires. It is acknowledged that these are subject to measurement error and biases and that objective approaches would be beneficial. Dietary biomarkers have been purported as a complimentary approach to improve accuracy of dietary assessment. Tentative biomarkers have been identified for many individual fruit and vegetables (FV) but an objective total FV intake assessment tool has not been established.

Objective:

We aimed to derive and validate a prediction model of total FV intake (TFVpred) to inform future biomarker studies.

Methods:

Data from the National Diet and Nutrition Survey (NDNS) were used for this analysis. A modelling group (MG) consisting of participants aged >11 years from the NDNS years 5-6 was created (n=1746). Intake data for 96 FVs were analysed by stepwise regression to derive a model that satisfied three selection criteria: standard error of the estimate (SEE) ≤80, R^2>0.7, and ≤10 predictors. The TFVpred model was validated using comparative data from a validation group (VG) created from the NDNS years 7-8 (n=1865). Pearson's correlation coefficients were assessed between observed and predicted values in the MG and VG. Bland-Altman plots were used to assess agreement between TFVpred estimates and total FV intake.

Results:

A TFVpred model, comprised of tomatoes, apples, carrots, bananas, pears, strawberries and onions, satisfied selection criteria (R^2=0.761, SEE=78.81). Observed and predicted total FV

intake values were positively correlated in the MG (r=0.872, *P*<0.001, R^2=0.761) and the VG (r=0.838, *P*<0.001, R^2=0.702). In the MG and VG, 95.0% and 94.9% of TFVpred model residuals were within the limits of agreement, respectively.

Conclusions:

Intakes of a concise FV list can be used to predict total FV intakes in a UK population. The individual FVs included in the TFVpred model present targets for biomarker discovery aimed at objectively assessing total FV intake.

Keywords: fruit and vegetables, prediction model, dietary assessment, biomarkers, dietary questionnaires.

Background

Non-communicable diseases (NCDs) accounted for 71.3% of worldwide mortality in 2016 (1). The objective measurement of modifiable risk factors is vital in informing strategies to reduce the public health burden incurred by NCDs. Fruit and vegetable (FV) intake has been associated with a lower risk of cardiovascular disease (2–5), type 2 diabetes (6,7), and some forms of cancer (2,8). These NCDs accounted for approximately 28.6 million deaths in 2016, equating to half of global mortality (1), thus increasing FV consumption presents a potential opportunity to reduce the burden of disease.

Recent meta-analyses assessing the relationship between the quantity of FV intake and relative risk of all-cause mortality have produced equivocal results. Findings consistently indicate that relative risk of all-cause mortality is proportionately lower with increased consumption of FVs, yet the reported plateau in risk reduction ranges from 5 servings (5), to 10 servings of FV per day (2). This two-fold variation in the threshold of daily FV consumption at which there is the lowest relative risk of all-cause mortality is congruent with disparities in public health recommendations. The World Health Organization and Public Health England currently recommend the consumption of at least five servings (400 g) of FV per day (9,10), whereas the Danish Ministry of Food recommend the equivalent of 7.5 servings (600 g) per day (11). Findings from Aune *et al.* (2) infer that current recommendations, such as the UKs presented in the Eatwell Guide (9), may not sufficiently encourage higher levels of FV consumption that pertain to a lower risk of all-cause mortality. The evidence regarding optimal daily intake of FVs remains inconclusive, thus presenting a barrier toward informing public health recommendations, emphasising the necessity for further elucidation of the relationship between FV intake and NCDs.

Epidemiological studies aiming to determine diet-disease relationships assess dietary intake using self-report methods such as food diaries, 24-hour recalls and food frequency questionnaires (12–14). While necessary for obtaining data representative of habitual dietary intake, such methods are inherently subject to measurement error and biases and can be burdensome on participants (12,15–17). A more succinct method of intake data collection, i.e. reporting a single food group of interest could alleviate the burden on participants, while conversely reducing the utility of the data when the exploration of whole diet-disease associations is required. Appropriate study designs and methodologies can mitigate the measurement error and biases inherent to self-report methods (18). A combined approach, comprised of the simultaneous measurement of dietary biomarkers and self-report methods has been purported to improve the accuracy of dietary exposure measurements, thus facilitating the elucidation of diet-disease relations (18,19).

Candidate dietary exposure biomarkers for the objective measurement of total FV intake, including carotenoids and polyphenols (20,21), have been explored and shown to have limited utility. The establishment of an objective tool to assess total FV intake, rather than individual FV intake, has not yet proved efficacious or been validated (22). Untargeted metabolomic techniques are increasingly prevalent within the literature, making significant progress in the identification and quantification of specific dietary exposure biomarkers (23,24). The predominant focus of this research has been identifying single biomarkers for specific foods/food groups. Further to the identification of novel biomarkers, the use of a panel of biomarkers, by measuring a number of metabolites pertaining to a food/food group for a more accurate representation of dietary exposure, has been proposed (25). Multi-metabolite biomarker panels (MBPs) have been identified for the quantification of walnuts (26), bread

(27), cocoa (28), orange juice (29), wine (30) and whole dietary patterns (31,32), however a panel for total FV intake is yet to be established.

The National Diet and Nutrition Survey (NDNS) is a continuous, cross-sectional survey, designed to collect detailed quantitative information on the food consumption, nutrient intake and nutritional status of the UKs general population (33). Analysis of these data can provide novel insight into total FV eating habits. The aim of this research was to identify a concise number of FVs that are predictive of total FV intake. Identifying such FVs stands to direct future metabolomic biomarker studies that pursue the objective measurement of FV intake.

Methods

Study Design

This study analysed cross-sectional intake data of individuals from years 5-6 (2012/13 – 2013/14) and years 7-8 (2014-15 – 2015/16) of the NDNS rolling programme (33,34). The modelling dataset (years 5-6) and validation dataset (years 7-8) were retrieved from the UK data archive in September 2017 and January 2019, respectively.

Data Source

Full methodological details of the NDNS have been described elsewhere (35). In short, the full NDNS years 5 - 6 dataset was comprised of 2,546 participants (age 30 ± 24 years, mean ± SD) recruited from 323 postal sector random sampling units across the UK. Data were collected over 12 months to account for seasonal variation. Samples were stratified by country, ensuring proportional representation from England, Scotland, Wales and Northern Ireland. Following initial interviews to obtain background information and familiarise participants with the intake data collection method, 4-day food diaries were completed and participants over

the age of 4 years who consented to a nurse visit had anthropometric measurements (height, weight, waist and hip circumference, demi-span, blood pressure), and blood and urine samples taken. The modelling group (MG) dataset was obtained from this sample and included all participants > 11 years old (n = 1746).

Data Processing

The faction of NDNS data used in the current analysis consisted of food and drink consumption data collected using 4-day un-weighed food diaries (portions were quantified by household measures). Participants recorded the contents of all eating and drinking occasions over four consecutive days, including one weekend day. Food diaries were processed and coded using an adapted version of Health Nutrition Research's dietary assessment system DINO (Diet In Nutrients Out) (36). DINO disaggregates composite items and items that differ by preparation into individual foods with a unique code. The current analysis aggregated data of the same fruit/vegetable with differing codes, to form a daily intake value for individual FVs (g/day). Fruit juices, potatoes, and pulses (except for green beans, runner beans, and broad beans) were excluded from the analysis due to differences in nutrient composition from FV as included in the UK Eatwell Guide (9). We multiplied dried fruit intake by three, based on the respective water and micronutrient content, to standardize dried and non-dried FV intake (34). **Supplementary Table 1** outlines the details of individual FV intake data aggregation, FV consumption prevalence and mean daily intake in consumers only. Daily intake of 96 FVs were calculated and used as potential predictor variables. Individual FV intakes were summed to calculate total FV intake (g/day).

Statistical Analysis

All data were obtained and processed using IBM SPSS Statistics 24 (SPSS, Inc., Chicago, IL, USA) and analysed using Stata version 15 (College Station, TX: StataCorp LLC). The assumptions of multiple linear regression analysis were satisfied prior to analysis. Normality of residuals and homoscedasticity of the data were confirmed, and no transformations were applied to any variables. All potential predictors had a linear relationship with total FV intake.

We conducted automated forward stepwise regression analyses. Models began with an intercept and were iteratively constructed by selecting the predictor variable (individual FV intake) that accounts for the most unique variance in total FV intake. Subsequent models incorporated the individual fruit or vegetable that accounted for the most unique variance in total FV intake among the remaining predictor variables. Predictor variables were added with each model iteration until there was no longer an improvement in total FV intake variance accounted for by the model. Regression significance ($P < 0.05$) was taken to indicate that the independent variable predicts total FV intake. The variance inflation factor (VIF) was used to quantify correlation of predictors in a model, to detect any collinearity. Regression coefficients represent the mean change in outcome for one unit of change in the predictor variable and were used to compile regression the equation. The standard error of the estimates (SEE) was calculated and R^2 used to denote the proportion of variance in total FV intake explained by each model.

Model Selection Criteria

The rationale underpinning model selection criteria was to produce a regression equation that could be used to facilitate the discovery of FV biomarkers. The future utility of the model is dependent upon having few predictors to moderate the extent of biomarker measurement

required, while explaining a large proportion of the variance in predicted total FV intake. We established iterative models that satisfied three pragmatically determined selection criteria; a SEE ≤ an 80 g FV serving, variance in total FV intake (R^2) > 0.7, and the number of predictors in the model was capped at 10 to produce a concise assessment tool. Comparative assessment of regression models was facilitated by calculating adjusted R^2, Akaike information criterion (AIC), Bayesian information criterion (BIC) and penalised likelihood ratio (LR) testing. The aim of all comparative assessments was to ensure that all subsequent models were an improvement on the previous model.

Model Validation

Validation of the final total FV prediction model iteration (TFVpred) was conducted using a novel dataset from the NDNS years 7-8, with participants aged > 11 years. NDNS data collection methodologies were consistent with the years 5-6 used as the MG. The current analysis applied the same data processing procedure described above to the validation group (VG) dataset to obtain comparable FV intake data. The TFVpred equation was applied to the VG dataset to predict total FV intake (g/day). Pearson's r correlation coefficient was measured to determine linearity between observed and predicted total FV values. Correlational coefficient of determination (R^2) was calculated to measure the amount of variance in TFVpred estimated total FV intake explained by the observed total FV intake. Correlational analysis was conducted with observed and predicted FV intake in vegetarian and vegan subsets of the MG and VG to assess the validity of the prediction model in a subset of the population with known differences in FV consumption patterns. Bland-Altman plots were generated to assess the agreement between TFVpred estimates and observed total FV intake in modelling and validation groups. Limits of agreement were plotted at ± 1.96 SDs of the mean difference between the observed and predicted values of total FV intake.

Results

Multiple Linear Regression Models for prediction of total FV intake

In total, 4-day food diaries were analysed from 1746 participants in the MG, and 1865 participants in the VG. Forward stepwise regression model summaries are displayed in **Table 1**. Total FV prediction model 7 (TFVpred) was the first model iterated that met all model selection criteria, with an $R^2 > 0.7$, a SEE < 80 and contained ≤ 10 predictor variables. All seven models predicted total FV intake ($P < 0.05$). The proportion of variance explained by regression models (R^2) increased from 0.277 to 0.761 between models 1 and 7. Incremental reductions in SEE were observed with each regression model including a novel predictor. TFVpred, comprised of seven predictor FV coefficients and constant, is displayed in **Eq. 1**:

$$\mathbf{TFVpred} = \mathbf{1.773}(\mathbf{tomatoes}) + \mathbf{1.428}(\mathbf{apples}) + \mathbf{2.439}(\mathbf{carrots}) + \mathbf{1.211}(\mathbf{bananas}) + \mathbf{1.422}(\mathbf{pears}) + \mathbf{1.714}(\mathbf{strawberries}) + \mathbf{1.519}(\mathbf{onions}) + \mathbf{29.88}(\mathbf{constant}).$$

The TFVpred equation highlights the seven predictor FVs accounting for the most variance in total FV intake, namely tomatoes, apples, carrots, bananas, pears, strawberries and onions, thus presenting targets for intake biomarker discovery. Five FVs included in the TFVpred model (tomatoes, onions, carrots, bananas and apples) were within the top six most commonly consumed FVs (as per number of consumers), while strawberries and pears were within the top 15 and 24, respectively (**Supplementary Table 1**). All predictor variable FVs were within the top 40 FVs for mean daily intakes in consumers only.

Model Comparison

Comparison of regression models is shown in **Table 2**. The variance in total FV intake explained by models, when corrected for the number of predictors, incrementally increased with additional model iteration. The size of incremental augmentation in adjusted R^2

193 diminished as regression models progressed, with the maximum change being an increase of
194 0.174 from model 1 to model 2, and the smallest change was 0.028, observed between
195 models 6 and 7. Penalised-LR criteria, AIC and BIC, are presented for each model in Table 2.
196 AIC and BIC values were incrementally smaller as more predictors were added to the
197 regression models. LR tests for nested models were significant with all subsequent iterations,
198 indicating successive improvements in goodness of fit.

199 ***Model Validation***

200 In the MG, observed and predicted values of total FV intake were positively correlated (r =
201 0.872, $P < 0.001$) with an $R^2 = 0.761$ (**Figure 1A**). Observed and predicted total FV intake values
202 in the VG were also positively correlated (r = 0.838, $P < 0.001$) with an $R^2 = 0.702$ (**Figure 1B**).
203 Bland-Altman plots determined there was good agreement between observed and predicted
204 total FV intake values, with the MG (**Figure 2A**) and VG (**Figure 2B**) demonstrating 95.0% and
205 94.9% of residuals were within the limits of agreement, respectively. Observed and predicted
206 total FV intake values within vegetarian and vegan subsets were positively correlated in the
207 MG (r = 0.882, $P < 0.001$, $R^2 = 0.777$, **Supplementary Figure 1A**) and VG (r = 0.839, $P < 0.001$,
208 $R^2 = 0.704$, **Supplementary Figure 1B**).

209 ***Discussion***

210 To our knowledge, this is the first study to elucidate a concise group of individual FVs that are
211 predictive of total FV intake, accounting for 76.1% of total variance. The 7th model iteration,
212 TFVpred, was the first to satisfy predetermined selection criteria and was subsequently used
213 to predict total FV intake in the VG, using individual intake values of tomatoes, apples, carrots,
214 bananas, pears, strawberries and onions. Correlational analysis and Bland-Altman plots were
215 used to assess the efficacy of the TFVpred model when applied to the VG and demonstrated

strong agreement between observed and predicted values. TFVpred thus provides a potential assessment tool in estimating total FV intake, where valid measurements of seven individual FV intakes (tomatoes, apples, carrots, bananas, pears, strawberries and onions) are available. A multitude of comparisons between models were conducted to determine that TFVpred outperforms other models by AIC, BIC and LR test statistics, thereby the most appropriate model for estimating total FV intake (37). This research has the potential to consolidate the applicability of existing individual FV measurements obtained using dietary questionnaires. Furthermore, the identified FVs signify clear targets for novel biomarker discovery. Subsequent integration of validated biomarkers within the TFVpred equation provide additional utility as a potential tool for total FV intake estimation.

Dietary Questionnaires

Self-report methods of dietary intake assessment, such as food diaries, 24-hour recalls and food frequency questionnaires, have been a longstanding topic of debate in nutritional research (17,38), while remaining the most prevalent techniques to assess diet-disease relationships (4,39). Critics state that the reliance on memory and the influence of researcher/social-approval biases can incur random and systematic measurement errors, such as the over-reporting of FV intake (12–14,17). Furthermore, the accuracy of self-reported data may be influenced by the ability of individuals, or the sensitivity of the assessment method, to quantify the size and contents of a FV serving (40,41). Proponents of self-report methods acknowledge that while limitations exist, study design considerations and corrections for measurement error can be applied to gather insightful intake data, currently unobtainable using other means (42,43). The NDNS dataset used in the current study aimed to collect data accurately pertaining to the UK population by mitigating the effect of some of these limitations through appropriate study design. Daily food diaries were

completed over four consecutive days to minimise reliance on memory (42). Upon completion of food diaries, trained interviewers met with participants to aid the quantification of the food diary constituents, where original visual aids were insufficient (35). The NDNS dataset presents a useful source when compiling inferential statistical models, as in the present analysis. Given the robustness of the NDNS methodology, validation with an updated NDNS dataset was necessary and demonstrated the efficacy of the TFVpred model as a practical tool for total FV intake estimation.

Novel assessment of total FV intake using the TFVpred model could utilise existing methods of individual FV intake from dietary questionnaires. Measurements could be obtained via amended food frequency questionnaires, i.e. condensed to include only FV assessment, providing sufficient validation is conducted (39,44,45). Kristjansdottir *et al.* (44) reported that FV intake estimated using a combined 24-hour recall and food frequency questionnaire was associated with 7-day food diary reported intake, with a spearman's coefficient of 0.73 ($P < 0.001$). Furthermore, Block *et al.*(46) correlated FV intake obtained using 100-item food frequency questionnaires (47), and a single page screener questionnaire, reporting a spearman's coefficient of 0.71 ($P < 0.001$). Using a screener to assess FV intake could provide a time-effective alternative to a lengthy questionnaire and provide specific FV intake data. A practical application of the predictive FVs identified in the present analysis would be to incorporate these FVs in screener questionnaires or as prompts in multiple pass dietary assessment methods. Adopting such changes may increase the accuracy of dietary intake data, though amendments to validated dietary assessment tools would require subsequent validation. Incorporating measurements of the FVs identified in the TFVpred model within

existing dietary questionnaires presents an inexpensive tool for internal validation to improve the precision of dietary intake assessment.

Combining Dietary Questionnaires and Biomarkers

The prevailing recommendations from prominent research groups within the field of nutrition and dietary assessment include the combined assessment of diet using dietary questionnaires and biomarker quantification (18,19,25). A prospective application of the TFVpred model validated in the present analysis would be to integrate biomarker assessments for the seven FVs, providing an objective assessment tool that can be obtained from biological samples and be used to assess FV exposure alongside appropriately conducted questionnaires. The NDNS represents an example of how this may be achieved, due to the concurrent collection of self-report data and urine samples, however the assessment of a validated FV biomarker assessment panel is yet to be established (35). Systematic reviews exploring the efficacy of objective assessments of FV intake by dose-dependent concentration biomarkers have ascertained that no single candidate biomarker can accurately measure total FV intake (20,48). However, putative dose-dependent urinary biomarkers have been identified for some FVs including grapes (49), peas, apples, onions (50), red cabbage, strawberries and beetroot (31). Prevalent techniques aiming to identify a panel of biomarkers pertaining to individual foods/food groups include targeted and untargeted tandem high-performance liquid-chromatography mass-spectrometry, as well as proton nuclear magnetic resonance spectroscopy, with subsequent multivariate modelling (Principal Component-Discriminant Analysis, Partial Least Squares, and Random Forest Classification) (27,32,51). This has led to the identification of numerous metabolites purported as biomarkers of dietary exposure, although validation as dose-dependent biomarkers of intake, necessary prior to TFVpred

model integration, is less pervasive (49,52,53). The specificity of putative biomarkers ranges from individual foods (including FVs) to broad dietary pattern identification (32,54,55).

Potential confounding factors for biomarker identification include inherent genetic variance between individuals, physiological and lifestyle factors that may influence metabolism, biological sample handling and the analytical methodology (22). Future research should aim to negate some of these factors. For example, Garcia-Aloy *et al.* (25) propose the use of MBPs to provide an insight into dietary exposure. MBPs enable the simultaneous measurement of numerous metabolites that pertain to a specific food/food group, capturing a broader faction of dietary exposure. Once validated, prospective MBPs of individual FV intake could be integrated with the regression equation modelled in the present study as a method of estimating total FV intake. Dragsted *et al.* (56) identified a stringent set of post-discovery validity criteria for biomarkers, including assessments of: 1) biochemical plausibility and stability, 2) dose-dependency with low abundancy when intake is zero and saturation kinetics, 3) time-responsiveness to inform when biological samples can be collected, 4) robustness after co-ingestion with other foods, 5) reliability to ensure biomarkers are comparable to assessments from other questionnaire or biomarker measurements, 6) a reproducible analytical methodology. Meeting these standards is imperative if biomarkers are to improve the precision and accuracy of dietary assessment. Considerable work is necessary to elucidate in particular time-responsiveness and dose-dependency of putative FV biomarkers (25). At present, the limitations associated with both facets of dietary assessment cannot be fully alleviated by adopting sole usage of the alternate technique, thus combinations of dietary questionnaires and biomarker assessments should be explored (16,25).

Strengths & Limitations

FV servings of 80 g were used in the present analysis to compute regression models, thus FVs that deviated from the standard 80 g serving sizes, such as dried fruits, required numerical transformation prior to be considered a FV portion. This was conducted to prevent the potential exclusion of a subset of FVs that contribute to total FV intake, but do not constitute a regular FV serving. Some semi-dried fruits were not included in the current analysis due to the unknown composition of portion sizes. Consistent with other nutritional epidemiology research (57,58), children aged < 12 years (MG, n = 763; VG, n = 822) were excluded from the current analysis to mitigate the systematic error incurred by having dissimilar eating trends and serving sizes to adolescents and adults. As the current analysis was conducted using intake data from UK based participants ≥ 12 years, prospectively the TFVpred model should not be used to estimate total FV intake in children < 12 years. Deriving the TFVpred model using stepwise linear regression modelling and pragmatic predetermined selection criteria facilitated the formation of a model that included a combination of influential FVs that were predictive of total FV intake and frequently consumed in the population. TFVpred predictor FVs were among the most pervasively consumed in the MG and VG, indicating good suitability within a UK population. Future research should investigate the efficacy of the TFVpred model in other developed countries and further validation is required prior to use in non-UK based populations, as FV intake is variable between countries (59,60). A prominent challenge within the present study was producing a model with a small number of predictors that captured a substantial proportion of the variance in total FV intake, without including relevant cofactors such as socioeconomic status(61,62), food availability(63) and vegetarianism(64). The TFVpred model predictions were accurate for subsets of the population known to have different FV consumption patterns, as demonstrated by the correlation between observed

and predicted total FV intake in vegetarians and vegans. The TFVpred model also performed well across a broad variety of FV intakes, the small proportion of individuals that fall outside the upper LOA. Bland-Altman plots (Fig. 2) indicate that 4.70 % and 4.86 % of individuals in the MG and VG, respectively, fall outside the upper LOA, thus consuming a variety of FVs that are not accounted for by the model. The simultaneous assessment of cofactors of total FV intake and additional FVs would increase the accuracy of prediction models, however the aim of the present study was to identify a concise number of predictor FVs that can be integrated into dietary questionnaires to reliably estimate total FV intake in a UK population and identify targets for biomarker discovery, rather than establish a multifaceted prediction model of total FV intake.

Conclusions

The TFVpred model (Eq. 1) established in the current study provides a valuable tool for estimating total FV intake. Future utility of the TFVpred model would be improved with the integration of dose-dependent biomarkers/MBPs for the FVs that predict total FV intake (tomatoes, apples, carrots, bananas, pears, strawberries and onions). The identification of these FVs, through the establishment and validation of the TFVpred model provides a clear pathway for future research by identifying dose-dependent biomarker targets. Advances in biomarker identification and validation provide a valuable opportunity to obtain objective assessments of total FV intake that, in parallel with appropriate self-report techniques, could denote notable improvements in the accuracy of dietary assessment.

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Statement of authors' contributions to manuscript

EJO, TPD and LMO'C developed the research question and planned the analysis; EJO performed the analysis with support from SP and LMO'C; EJO drafted the manuscript with editorial support from OF, TPD, and LMO'C; and all authors read and approved the final manuscript.

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Tables

Table 1 – Multiple linear regression models using individual fruit and vegetable (FV) intake data from the National Diet and Nutrition Survey Rolling Programme years 5-6 to predict total FV intake (n = 1746).

Model	Predictor Variables	Regression *P value*	Constant	Regression Coefficient (β)	R^2	Standard Error of the Estimate	Variance Inflation Factor
1	Tomatoes	< 0.001	134.089	2.672	0.277	136.81	1.00
2	Tomatoes	< 0.001	104.069	2.352	0.451	119.24	1.02
	Apples	< 0.001		2.030			1.02
3	Tomatoes	< 0.001	69.595	2.277	0.567	105.92	1.02
	Apples	< 0.001		1.823			1.04
	Carrots	< 0.001		2.982			1.02
4	Tomatoes	< 0.001	46.973	2.091	0.664	93.26	1.04
	Apples	< 0.001		1.546			1.07
	Carrots	< 0.001		2.849			1.02
	Bananas	< 0.001		1.406			1.06
5	Tomatoes	< 0.001	45.125	2.060	0.702	87.91	1.04
	Apples	< 0.001		1.452			1.08
	Carrots	< 0.001		2.720			1.03
	Bananas	< 0.001		1.292			1.08
	Pears	< 0.001		1.362			1.05
6	Tomatoes	< 0.001	39.892	1.995	0.732	83.33	1.04
	Apples	< 0.001		1.453			1.08
	Carrots	< 0.001		2.673			1.03
	Bananas	< 0.001		1.250			1.08
	Pears	< 0.001		1.391			1.05
	Strawberries	< 0.001		1.762			1.01
7	Tomatoes	< 0.001	29.877	1.773	0.761	78.81	1.11
	Apples	< 0.001		1.428			1.08
	Carrots	< 0.001		2.439			1.05
	Bananas	< 0.001		1.211			1.08
	Pears	< 0.001		1.422			1.05
	Strawberries	< 0.001		1.714			1.01
	Onions	< 0.001		1.519			1.11

Table 2 – Comparison of multiple linear regression models using individual fruit and vegetable (FV) intake data from the National Diet and Nutrition Survey Rolling Programme years 5-6 to predict total FV intake (n = 1746).

Model	Cumulative Predictor Variables	Adjusted R^2	Change in adjusted R^2	Akaike information criterion	Bayesian information criterion	Likelihood Ratio Models Tested	Likelihood Ratio Test statistic	Likelihood Ratio Test *P*
1	Tomatoes	0.276	-	22133	22144	-	-	
2	Apples	0.450	0.174	21654	21670	1 and 2	481.13	< 0.001
3	Carrots	0.566	0.116	21241	21263	2 and 3	414.65	< 0.001
4	Bananas	0.664	0.098	20798	20825	3 and 4	445.38	< 0.001
5	Pears	0.701	0.037	20592	20625	4 and 5	207.35	< 0.001
6	Strawberries	0.732	0.031	20406	20445	5 and 6	187.86	< 0.001
7	Onions	0.760	0.028	20213	20256	6 and 7	195.84	< 0.001

Figures

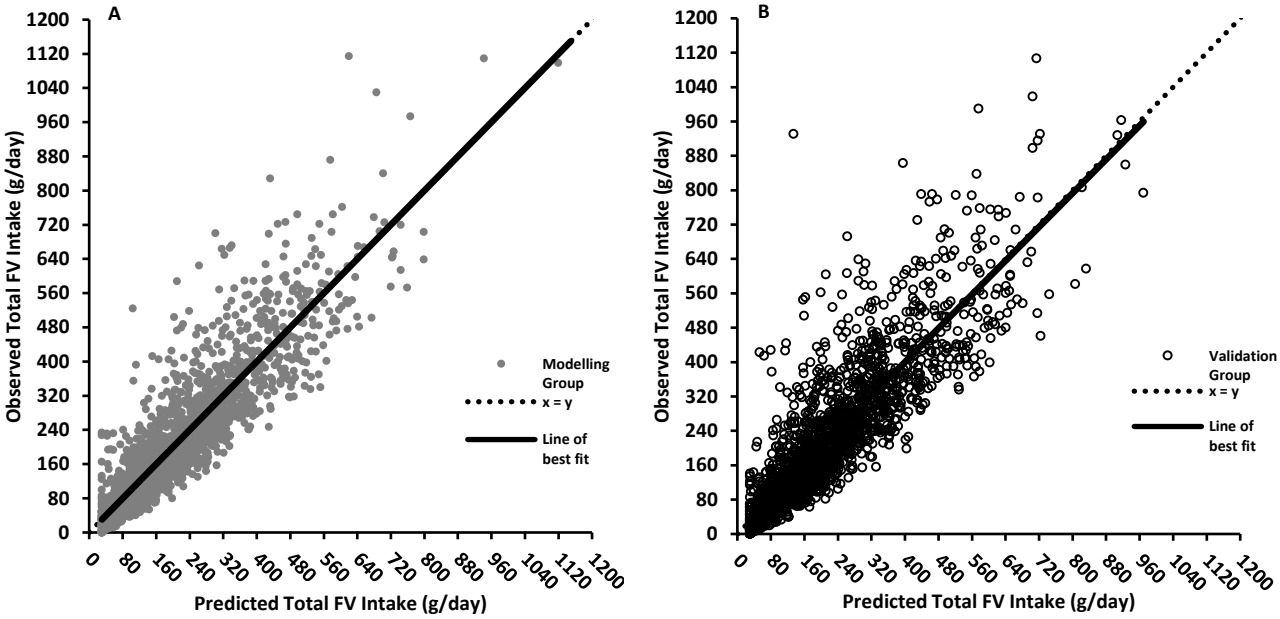

Figure 1 - Correlation between observed and predicted total FV intake using the TFVpred equation for the (A) modelling group (NDNS years 5-6, n = 1746) and (B) validation group (NDNS years 7-8, n = 1865). FV, fruit and vegetable.

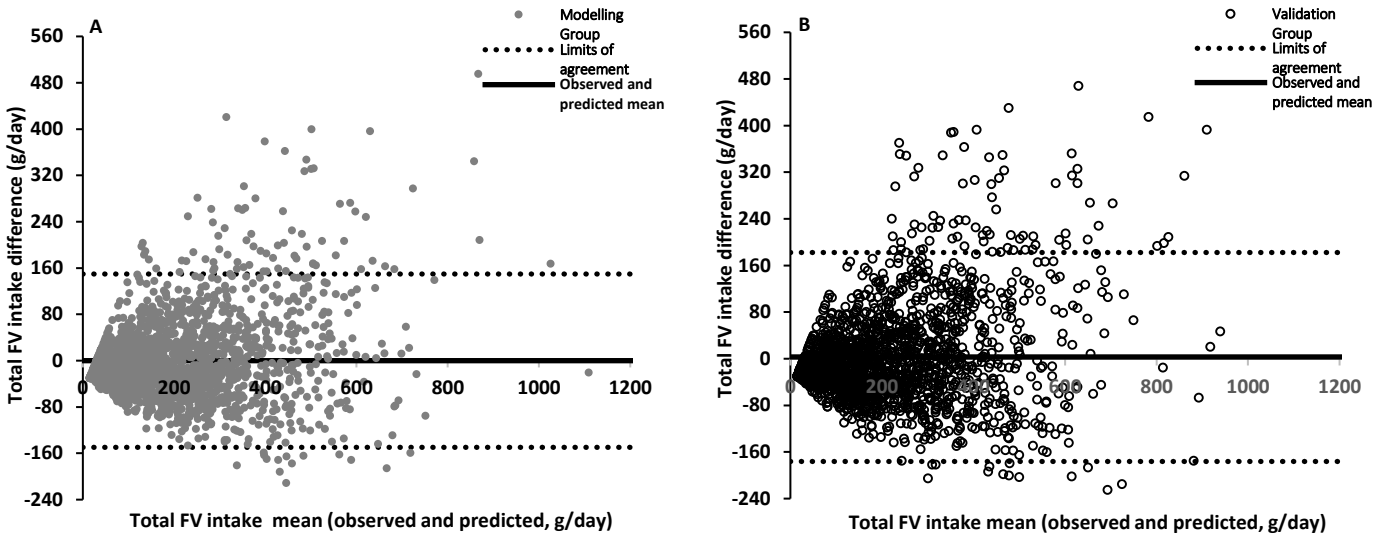

Figure 2 - Bland-Altman plots of total FV intake predictions in the modelling group (A, n = 1746) and validation group (B, n = 1865). Plots display the difference between total FV intake measured by the NDNS and total FV intake predicted by TFVpred model vs. the observed and predicted mean. Limits of agreement (dotted lines) are displayed at ± 1.96 SDs of the mean difference between the observed and predicted values of total FV intake. FV, fruit and vegetable.