

On the Use and Misuse of Child Height-for-Age Z-score in the Demographic and Health Surveys

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Abstract

This paper addresses the problem of model misspecification bias when estimating cohort-level determinants of child height-for-age z-score (HAZ) using data from the Demographic and Health Surveys (DHS). I show that the combination of DHS survey design and the biological realities of child health in developing countries create an artifact that can strongly bias regression estimates when identification relies on seasonal, annual or spatio-temporal variation associated with a subject's birth cohort. I formalize the econometric problem and show that flexible specifications of the HAZ-age profile can greatly mitigate the bias. When regression models can exploit within-cohort variation in the covariate of interest, appropriate fixed-effects models can effectively purge the bias. I also provide Monte Carlo evidence that DHS recommended inference strategies produce standard errors that are too small when estimating birth cohort determinants of HAZ.

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1 Introduction

I demonstrate how a statistical artifact in Demographic and Health Survey (DHS) data can bias regression estimates of cohort determinants of child height-for-age Z-score (HAZ). The short survey window of the DHS and the relatively long periods between survey rounds induce a mechanical, cyclical relationship in the data between birth cohort and age-at-measurement. Separately, the biological effects on children of chronic undernutrition and poor health cause a rapid loss of HAZ over the first few years of life, generating a strong, non-linear relationship between HAZ and age-at-measurement (the HAZ-age profile). Together, survey design and biology conspire to induce a spurious correlation in the data between birth cohort and HAZ that is driven by age-at-measurement and specifically determined by the particular dates on which the various survey rounds were conducted. The artifact is likely to bias coefficient estimates when the HAZ-age profile is misspecified in specific but common ways.

I refer to this relationship between birth cohort and HAZ as the *timing artifact*. As a general condition, the artifact arises when data is collected infrequently over a short period of time on an outcome variable that is itself a function of age. After providing the necessary background (Section 2), I formalize the econometric problem in the context of estimating cohort determinants of HAZ in the DHS, and interpret the bias in an omitted variables framework (Section 3). I consider two means of eliminating the bias: flexible specification of the HAZ-age profile, and identification based on within-cohort variation. Choice of strategy by the practitioner will depend primarily on the nature of the data the researcher is able to bring to bear on the question, but in general within-cohort estimates are preferable.

I then demonstrate the artifact and bias empirically in two environments (Section 4): estimating the effects of month of birth in Bangladesh and GDP per capita in year of birth in Zimbabwe. In the month of birth example, a regression with no control for age that ignores the relationship between month of birth and age-at-measurement estimates the peak-to-trough difference in seasonal effects at a statistically significant .245 standard deviations ($p = .017$). A more flexible model that specifies a spline control across age estimates a statistically insignificant peak-to-trough difference of .131 standard deviations ($p = .414$)¹, a decrease in magnitude of about 45%. In the GDP regressions, moving from no control for age to a spline specification reduces the implied effect of a 10% increase in GDP from .11 standard deviations to .03.

Following the empirical examples, I present results from a placebo-type Monte Carlo exercise simulating a “random weather shock” on data from Peru (Section 5). I find that cohort and region fixed-effects estimators that simultaneously harness geographic and temporal variation can fully purge the bias, but that estimators relying purely on temporal variation and flexible parametric specifications of the HAZ-age profile remain mildly biased.

The empirical and simulated exercises also provide evidence that DHS recommended standard error calculations, which account only for survey sampling and weighting (Madise et al., 2003), produce standard errors that tend to over-reject a true null hypothesis when estimating cohort effects². Suggestively, between 40% and 80% of other countries’ GDP time-series appear significantly correlated with the height of Zimbabwean children at the 95%

¹P-values are for F-stats testing Feb-Dec point estimates jointly 0 using DHS recommended inference accounting for weighting, stratification and cluster sampling.

²The simulation design and statistical theory closely follow Bertrand et al. (2004), which investigates standard error properties using the Current Population Survey in the United States.

level when DHS recommended standard errors are employed. These “rejection rates” decrease to between .03 and .09 when more conservative wild cluster-t bootstrap procedures are used to determine statistical significance. In the Monte Carlo environment, where rejection rates of .05 can be known to be expected, DHS standard error estimates lead to rejection rates of around .25 at 95% confidence. Alternative strategies that rely on cluster-robust analytic methods yield more appropriate rejection rates of around .08.

2 Background

Early life and in utero exposure to environmental and economic stresses has lasting effects on human capital development (Almond and Currie (2011), Uauy et al. (2011)). However, the mechanisms by which these early life conditions leave their mark are not well understood (Rasmussen, 2001). By tracking persistent birth cohort effects through all stages of life, researchers hope to flesh out the biological and socioeconomic mechanics behind the correlations between early life environment and later life economic success (Glewwe and Miguel, 2008). Methodologically, this literature has focused on accounting for important behavioral considerations like endogenous fertility, unobserved household characteristics, and/or non-random mortality (Strauss and Thomas (1995) ; Pitt (1997)). Previous research using the DHS has also mentioned the difficulty in effectively controlling for the HAZ-age profile (e.g. Paxson and Schady (2005), Agüero and Valdivia (2010)), though no prior work has provided a formal analysis of the problem.

This paper focuses on HAZ, an age- and gender-normalized measure of child height given in units of standard deviations and relative to the median age- and gender-conditional height distribution of a well-nourished, healthy

population of children. HAZ is of particular interest because it both captures the long-term cumulative effects of health throughout childhood and is known to be correlated with later life outcomes (Dewey and Begum (2011), Hoddinott et al. (2008)). If in utero shocks lead to permanent health effects, they should be measurable later in life via HAZ. In addition to the examples I investigate³, it has been used to estimate the effects of early life exposure to crop failure (Akresh et al., 2011), violence (Akresh et al., 2012), access to health facilities (Valdivia (2002)), agricultural price shocks (Cogneau and Jedwab (2012)), and a number of other birth cohort HAZ determinants.

HAZ is often described as “age-adjusted height”⁴. However, age may be the single best predictor of HAZ available for children in less developed countries. Figure 1 shows the HAZ-age profile for the three countries I analyze here - Bangladesh, Zimbabwe and Peru - with mean HAZ plotted across child age in months. In the first two years of life children in Bangladesh lose about 1.5 standard deviations of HAZ. By way of comparison, Bangladeshi children born to households in the highest asset index quintile are about 1 standard deviation taller than those born to parents in the lowest quintile. Bangladeshi children with mothers in the tallest height quintile are on average 1.1 standard deviations taller than those with mothers in the lowest quintile. HAZ as a measure of child health in developing countries may be “adjusted for age” in some sense, but it is also very strongly determined by age.

The DHS are natural datasets for investigating cohort determinants of HAZ. As of 2010, standard DHS surveys had been conducted 236 times in 84

³See Lokshin and Radyakin (2012), Agüero and Valdivia (2010) and Pörtner (2010) for papers examining month of birth, GDP per capita and weather shocks respectively.

⁴This analysis uses the 2006 WHO standards for HAZ, computed using the Stata package “haz06” and variables for age, height, and gender. I have also run most analyses using the previously favored NCHS/FELS/CDC standards. Choice of standardization does not qualitatively affect results.

countries, with 196 of the surveys similar to those analyzed here (Fabic et al., 2012). The costs of continual demographic surveillance are high, and so for most of the past two decades surveys have been implemented in participating countries only every four to five years, conducted over a period of just a few months. Continuity in cohorts between surveys is achieved by recording data on all children under the age of 5 found in sample households.

Use of the DHS has increased steadily. Fabic et al. (2012) finds 1177 journal articles published using DHS data between 1985 and 2010, and this count relies only on articles found on the measureDHS website and on PubMed. The rate at which articles using DHS data are published is also increasing, from less than 50 per year in 2000 to almost 150 per year in 2010. A little over half of these publications dealt with maternal and child health in general, and by 2010 there were around 25 papers per year being published on nutritional deficiencies, where the primary outcome is likely to be HAZ or some other anthropometric outcome.

3 Econometric Demonstration

This paper is focused on estimating the effects of birth cohort regressors on HAZ, but the econometric problem is not restricted to the particular outcome or dataset. Three ingredients are required for the timing artifact to bias regression estimates: 1) the data being analyzed comes from measurements of subjects taken over a short period of time at infrequent intervals; 2) the outcome variable of interest must be in part determined by the subjects' ages; and 3) the covariate of interest must be measured in relation to the subjects' birth cohorts, for example a birth year or birth quarter aggregate variable. The data collection method (1) ensures that cohort and age-at-measurement

are mechanically related in a manner determined by the dates on which the measurements were taken and the ages of the population of interest. The outcome-age profile (2) translates this age-cohort relationship into a spurious outcome-cohort relationship - the timing artifact. When estimating the effects of some cohort level variable (3), failure to adequately control for the outcome-age profile can allow residual, spurious variation in cohort HAZ (induced by the timing artifact) to induce an omitted variables bias on regression estimates. Intuitively, the least squares machinery associates the spurious timing artifact variation in HAZ with the time series variation in the birth cohort variable of interest. This problem is particularly worrisome when there are only a small number of cohorts (and thus no law of large numbers for “randomization” across cohorts) and/or there is temporal serial-correlation in the time series of the covariate of interest (since the spurious time series component of cohort HAZ induced by the timing artifact is also serially-correlated across time).

3.1 Intuition Using Month of Birth

The most basic form of the econometric problem is to estimate the effect of a birth cohort variable of interest β_T on some partially age-determined outcome Y , when data for estimating β_T comes from a random cross-section of the population of interest, and measurements are taken simultaneously and instantaneously on all sample subjects. Suppose real world data were generated in the following manner:

$$Y_{ia} = X'_{ia}\beta_x + \beta_T * T_a + f(age) + \epsilon_{ia} \quad (1)$$

Y , for individual i of age a , is determined by some set of covariates X , a covariate of interest (or treatment) T , the subject’s age at measurement, and

an iid error term ϵ . T is subscripted by ‘a’ because it is an aggregate measure common to some aspect of birth cohort, and in this framework cohort and age are perfectly collinear. I choose to subscript using a instead of c to emphasize that Y is partially a function of age, via $f()$, which is not restricted to be well behaved in any way. This stylized model is analogous to an empirical environment where one round of the DHS is employed, and the fact that the survey is conducted over several months is abstracted away.

3.1.1 Timing Artifact and Month of Birth

The first task in linking the timing artifact to estimates of β_T is to understand the relationship between T and age-at-measurement in the data. To make this relationship clear with an example, suppose T is a vector of dummy variables for month of birth. The mechanical relationship between age-at-measurement and month of birth makes the problem somewhat unique, but it also makes the timing artifact visible in a way that can be obscured once an additional layer of variation is introduced.

Assume for the moment that the survey in question was taken on January 1, 2000, at 12:01am, and that, like the DHS, it takes as a population of interest children under the age of 5 (all children born on or after January 1, 1995). In this case, all sample children who were born in December were measured at ages 1 month (born December 1999), 1 year 1 month (born December 1998), 2 years 1 month (born December 1997), etc. Similarly, all children born in June were measured at the ages of 6 months (born June 1999), 1 year and 6 months (born June 1998), etc. Because younger children have higher HAZ than older children (due to the HAZ-age profile), December-born children are not just younger, but also have higher HAZ than June-born children. To see

this graphically, consider Figure 1. In this setting, all children at the furthest left end of the graph would be December-born children. Six months to the right would be June-born children, who were measured at 6 months old and thus have lower mean HAZ than the December children who were measured at 1 month old. The same pattern repeats for children aged 1 year and 1 month (December-born) compared to children aged 1 year and 6 months (June-born), and again for ages of 2, 3 and 4 years, though for the older ages the exact trend is less visually apparent as the HAZ-age profile flattens⁵. Had the survey been conducted in summer, the reverse would be true.

Although this stylized environment represents only a single survey taken at a single point in time, the problem is easily generalized to include multiple survey rounds taken over several months each and several years apart. The left panel of Figure 2 shows survey timing in Bangladesh for four contiguous survey rounds, with the the year of the survey on the Y-axis and the months of the survey window (the months in which measurements were taken) plotted on the X-axis. Early rounds (1996 & 2000) were conducted mostly in winter, while the later rounds (2004 & 2007) were conducted in spring and summer. On aggregate, then, children born in late fall and winter were measured at younger ages than children born in spring. This is shown in the right panel of the figure, which is a bar graph depicting mean age-at-measurement by month of birth.

The effect of this imbalance of age across birth months on HAZ is represented in the top panel of Figure 3. This figure overlays a line graph of mean HAZ by birth month on the age-at-measurement bar graph shown in Figure 2.

⁵Even for 3 and 4 year olds, children are losing HAZ within any age-in-years bin. This is likely a result of both the standardizing process (de Onis et al., 2004) and age misreporting (Schoumaker, 2009), and it can be seen in Figure 6 which plots the estimated HAZ-age profile from a spline function with separate intercepts at round age in years.

The sinusoidal shape of the line graph strongly resembles the canonical season of birth graph. But the true relationship is clear from the bar graph: higher HAZ is apparent in months where children born in that month were measured at younger ages, and these months are the months immediately preceding the survey timing. An even more compelling example is shown in the bottom two panels. They show the same graph but with the sample restricted to children *measured* in either February (left panel) or November (right panel), and limited to the 1996 and 2000 survey rounds which both ran roughly November ('94/'99) to February ('96/'00). The February graph looks the same as the November graph shifted four months forward. The entire magnitude of the apparent birth month seasonality in child HAZ outcomes can be explained by the survey timing induced differential in age-at-measurement across birth months.

3.1.2 Bias

The correlation between HAZ and month of birth described above is an unconditional correlation. If we can fully control for the “natural” loss of HAZ over age, estimates of β_T should be unbiased. Common strategies to account for the HAZ-age profile include linear or polynomial specifications in age or dummy variables for age in months. However, misspecification of the HAZ-age profile can induce bias in the same manner as non-specification can. To formalize the problem econometrically, suppose that the researcher specifies a model of the following form using data generated by Equation 1, where $g(\text{age}_a; \gamma)$ is some parametric specification of $f(\text{age})$:

$$Y_{ia} = X'_{ia}\beta_x + \beta_T * T_a + g(\text{age}_a; \gamma) + \eta_{ia} \quad (2)$$

Unbiased estimation of Equation 2 requires that the error term η is uncorrelated with T . However, in the presence of misspecification of $g()$ such that $g()$ does not fully control for $f()$, analysis of the underlying data generating process shows this to be an unreasonable assumption. Returning to Equation 1, I add and subtract $g()$, producing:

$$Y_{ia} = X'_{ia}\beta_x + \beta_T * T_a + g(\text{age}_a; \gamma) + [f(\text{age}) - g(\text{age}_a; \gamma)] + \epsilon_{ia} \quad (3)$$

Define $\mu_a = f(\text{age}) - g(\text{age}_a; \gamma)$, the age-invariant systematic error in the estimate of $f()$ across age. Since both $f()$ and $g()$ are functions only of age, μ_a is then also solely a function of age and all subjects with the same age have the same value of μ . While η in the estimating equation was posited as being iid (as being ϵ), this exercise allows us to decompose η into two components: ϵ , the real-world iid error, and μ , the model misspecification error component that is the age-conditional difference between $f()$ and $g()$. Plugging this definition of η back into Equation 2, we see that the equation the researcher ought to be estimating is:

$$Y_{ia} = X'_{ia}\beta_x + \beta_T * T_a + g(\text{age}_a; \gamma) + \mu_a + \epsilon_{ia} \quad (4)$$

To simplify the analysis, assume that $f()$ is a linear combination of some set of age-transformations (age, its square, discontinuities, etc.), and that $g()$ does not contain one or more of these transformations. μ can now be understood as an omitted variable in the original specification of Equation 2. As the classical demonstration of omitted variables bias shows⁶, coefficient esti-

⁶see Angrist and Pischke (2008) pages 59-61 for proof and discussion.

mates on any variables included in the regression that are correlated with μ will be biased. Such a correlation does not require a particularly anomalous situation. In cases where $f()$ is smooth and $g()$ is specified to be smooth, μ will be smooth, and thus serially-correlated across age, and thus T and μ will likely move together (or against each other) in one fashion or another over at least some periods of time.

3.2 Two Strategies for Unbiased Estimation

In this subsection I discuss two approaches to purging regression estimates of timing artifact-induced bias. The choice of strategy by the researcher will depend largely on the types of identifying variation in the birth cohort variable the researcher can bring to bear on the problem. The first strategy is directly implied by the discussion above and the framing of the omitted variables bias interpretation: minimize the potential impact of μ through flexible specification of $g()$. This turns out, for reasons discussed in the following section, to be a more problematic approach than it first appears.

The second strategy hinges on the realization that model misspecification itself is not always problematic, even model misspecification that is invariant across age. The problem arises not only because μ is age-invariant, but also because age is correlated with T . In the presence of within-cohort variation in T , such as regional or subject-level quasi-experimental variation, it may be possible to separate out the cohort/age effect from the effect of T within each cohort, and thus achieve identification based on assumptions about the distribution of age (and hence the distribution of μ), *within cohort*, across T .

3.2.1 Flexible Specification of $g()$

One way to visualize the timing artifact is to plot out the HAZ “time-series” produced using DHS data by collapsing mean HAZ by date of birth. I show this in Figure 4. In the left panel I graph mean HAZ by birth date (aggregated to birth month), and in the right panel I provide the same graph disaggregated by survey round. While the left panel produces a graph that looks like a plausible, even if oddly sinusoidal, time-series, the right panel reveals the true dynamic. Most of the large variation seen in the aggregate time series is actually caused by overlapping survey windows. When the data is disaggregated, the graph appears in a more interpretable form - as the HAZ-age profile repeating (backwards) over time.

Figure 4 also raises a new modeling concern, the changing shape of this profile over time. To show this more clearly, Figure 5 overlays the sample mean HAZ-age profile from the first (1996) and last (2007) Bangladesh survey rounds. Over time, children are losing HAZ less rapidly during the first few years of life. This secular improvement in child-health affects both the *shape* and the *level* of the HAZ-age profile across survey rounds. Any attempt to absorb $f()$ via $g()$ must contend not only with the shape of the HAZ-age profile over age, but with the secular shifting of the HAZ-age profile over time.

To formalize these considerations, suppose data in the real world were generated in the following form, and gathered from a repeated cross section of S survey rounds, which are now each taken over several months, but taken several years apart:

$$Y_{iacs} = X'_{iacs} \beta_x + \beta_T * T_c + f(\text{age}_a, \text{birthdate}_c) + \epsilon_{iacs} \quad (5)$$

Y is determined for individual i of age a born into cohort c and surveyed

(measured) in round s . The incorporation of multiple survey rounds through the inclusion of a more refined timing-index is the first important difference between this model and the stylized model discussed above. The second difference is that $f()$ is no longer conceived as simply a function of age but also as a function of birthdate, to capture the changes in the HAZ-age profile over time.

Ignoring the shifts in the HAZ-age profile over time will generate a second type of model misspecification that may also induce a correlation between T and μ . Suppose that secular improvements in child health are raising the HAZ-age profile over time and that $g()$ is specified as static. μ now becomes a function of both age and time, the model misspecification associated with $g()$ improperly fitting the *shape* of the HAZ-age profile (μ in the stylized model), as well as the systematic error that is generated across survey rounds caused by the failure to account for the *level* changes over time. If not just the level but the shape of the profile is changing over time proper specification of $g()$ becomes even more difficult.

An obvious assumption that would allow the researcher to identify β_T is that $f()$ is separable into $f_1(\text{age})$ and $f_2(\text{birthdate})$, the assumption underlying the inclusion of both flexible controls for age and a set of secular calendar-time controls. These two options underlie the bulk of the previous empirical work referenced in this paper.

A second option is to assume that one can directly model the changes in $f()$ over time. For example, one could specify a spline function that shifts parametrically over time with different changes in the slopes of the spline segments for different age groups. Since very young children have HAZ much closer to the reference population median, there is little reason to believe that the mean HAZ of 1 month olds is likely to increase rapidly due to some secular

force. However, secular health improvements are quite likely to *slow the loss* of HAZ over the first few years of life.

3.2.2 Within-Cohort Variation in T

The second strategy for purging estimates of timing-artifact induced bias is to accept the problem of model misspecification but to specify the model in such a way that μ becomes orthogonal to T. In the environments described above, all children born at the same time had the same value of T. This forced identification of β_T to rely on assumptions about the shape of $f()$ and/or its movement over time. However, it is often the case that the researcher faces a problem where within-cohort variation in T is available or such variation can be constructed in some manner by bringing more refined data to the problem.

Suppose data were generated in the following form:

$$Y_{iacrs} = X'_{iacrs}\beta_x + \beta_T * T_{cr} + f(\text{age}_a, \text{birthdate}_c) + \epsilon_{iacrs} \quad (6)$$

This process is exactly the same as that described by Equation 5 except for the inclusion of subscript r denoting region of birth, and the fact that T now varies within birth cohort (across regions). In this situation a cohort and region fixed effects model can be specified. The cohort fixed effects are used to assure the similarity of the distribution of child ages between treatment and control. They are not functioning *as* a specification of $g()$, since the magnitude of μ is not a concern and the fit of the model to any individual child is not important. They are functioning *instead of* a specification of $g()$, rendering it unnecessary by exploiting variation such that μ becomes uncorrelated with T.

3.3 Inference

The bulk of this paper is concerned with the question of model-misspecification bias in estimates of $\hat{\beta}_T$, but the question of statistical inference is also important. Most work on standard error calculations and inference properties using the DHS has focused on accounting for survey design: sample selection (stratification and clustering) and population-probability weighting. Madise et al. (2003) demonstrates that failure to account for survey design can lead to greatly underestimated standard error estimates, giving researchers false confidence in the statistical significance of their results.

Madise et al. (2003) focuses on estimating standard errors for the effects of child, maternal and socioeconomic variables on child nutritional status. That context is different in theoretically important ways from the context of estimating standard errors for the effects of birth cohort aggregate variables. In this environment there is likely to be strong serial correlation in the regressor of interest within birth cohort or region. Bertrand et al. (2004) demonstrates that ignoring such serial correlation can lead to standard error estimates that are much too small.

Serial correlation is usually addressed spatially by clustering on the “cross-sectional” variable, in this case region. This is the environment in which Bertrand et al. (2004) and Cameron et al. (2008) operate. But in the context of birth cohort aggregate variables, the obvious serial correlation in the regressor of interest is temporal. I thus investigate both temporal and spatial correlation, “clustering” on either cohort or region groupings. Since the number of cohorts is generally small (less than 40), and since cluster-robust analytic estimators only converge in probability to the true standard error as the number of clusters goes to infinity, I follow Cameron et al. (2008) in

conducting a wild cluster-t bootstrap procedure to compute p-values when clustering on cohort or region is proper.

4 Empirical Examples

In this section I discuss two empirical contexts in which the timing artifact can induce bias on regression estimates of cohort variables. The first example, partially discussed in Section 3, is the effect of month of birth on HAZ. The second is the effect of GDP per capita in year of birth. Both examples demonstrate the fragility of point estimates to specification of $g(\text{age}, \text{birthdate})$, and in both cases the point estimates attenuate towards zero as the specification of $g(\cdot)$ becomes more flexible. The GDP example also lends itself nicely to investigation of the inference properties of various statistical significance estimators.

4.1 Data

In each empirical exercise, and for the data on which the placebo-test Monte Carlo simulations are run, I use four waves of DHS data from three countries, one country for each exercise - Bangladesh (month of birth) and Zimbabwe (GDP per capita in birth year) for the empirical examples, and Peru for the simulation exercise (weather shocks). All surveys were conducted between 1992 and 2011 (International, 2011). I aggregate all birthdates to month and year. Sample weights are re-normalized so that each survey is weighted equally, with relative sampling probability preserved within survey round. Results are robust to use of original DHS weights or no weights at all. Specifics of the sample selection and survey design are provided in Appendix A.

Table 1 displays unweighted summary statistics for the three example countries. Child HAZ is lowest in Bangladesh, where the mean HAZ hovers around the -2 standard deviations “stunting” level used internationally as a mark of moderate, chronic malnutrition. Children in Peru and Zimbabwe have slightly better HAZ measures, but remain more than a full standard deviation below the reference population of well nourished children. These numbers are consistent with many countries that face endemic disease and chronic nutritional shortfalls. Maternal age, education and height, and urban/rural composition are also consistent with much of the developing world (Ayad et al., 1997).

The covariates shown here, along with region dummies, constitute the set of X covariates used in the regressions that follow. Maternal education is entered in regressions as a set of five dummy variables⁷. Continuous variables other than child age and date of birth are entered into regressions linearly. Many papers that analyze child HAZ outcomes employ a much larger vector of covariates. I employ a sparse set to maintain sample size and increase transparency. Inclusion of larger covariate sets makes no qualitative difference to the results presented.

4.2 Month of Birth in Bangladesh

The first empirical question is whether or not the theoretical bias discussed in Section 3 actually affects point estimates to a meaningful degree after controlling for age. To determine the potential magnitude of the bias in an empirical setting, and to test the ability of increasingly flexible specifications of $g()$ to

⁷These are no education, primary school, secondary school, higher education, and a dummy for unknown educational status. Data definitions are provided by International (2012)

purge the bias, I estimate month of birth effects using data from Bangladesh. In this environment the timing artifact is likely to be quite strong since there is no mitigating factor between cohort and the regressor of interest, in the sense that we are not estimating something that varies by month of birth, but the actual effect of the month of birth itself. Beyond providing an intuitive environment for examining timing artifact-induced bias, month of birth has been shown to be a contributing determinant to all sorts of later-life outcomes, and various socio-biological mechanisms have been invoked to explain the correlation⁸. Other data artifacts can also induce the appearance of some month of birth effects⁹.

4.2.1 Month of Birth: Estimation

The estimating equation for month of birth, adapted from Equation 5 with month of birth replacing T and cohort (the interaction of month and year of birth) now omitted and subsumed by the birth month dummies, survey round dummies and age at measurement controls, is of the form:

$$HAZ_{iams} = X_{iams}\beta_x + MOB_m + g(age_a; \gamma) + \lambda_s + \eta_{iams} \quad (7)$$

The model conceives $f(age, birthdate)$ as separable: the shape of $f_1(age)$ is assumed to be constant across time, varying only in level shifts from survey to survey as captured by λ (the specification of $f_2(birthdate)$). I specify several

⁸Angrist and Krueger (1991) and Musch and Grondin (2001) use *institutional* explanations involving age-at-entry, while Buckles and Hungerman (2008) emphasize a *behavioral* mechanism of selection into birth timing. The development economics literature tends to focus on *biological* explanations, e.g. Lokshin and Radyakin (2012), which explains seasonal birth effects in India by in utero exposure to the monsoon rains

⁹Linn et al. (2011) show that estimating seasonal effects from data collected on births occurring over a fixed period of time can induce the appearance of seasonal differences in birthweight. Lewis (1989) shows that age-at-diagnosis can explain some seasonal birth patterns in schizophrenia.

forms of $g_1(\text{age}; \gamma)$: linear, quadratic, cubic and spline.

My base spline in this context is linear between knots, with the knots placed at round age-in-years (12 months, 24 months, 36 months and 48 months of age). Each knot is also allowed its own intercept, which is common across survey rounds, to account for standardization and age-reporting issues at round ages in years¹⁰. The fit of the spline is shown in Figure 6, which graphs out the estimated mean HAZ-age profile for Bangladesh from a regression of HAZ on the spline described with only the survey dummies also included in the regression. The left panel shows the fit of the spline (the line graph) on a scatter plot of mean HAZ across age, and the right panel shows the fit across birthdate. It is also possible to specify more flexible spline functions that directly estimate $f(\text{age}, \text{birthdate})$. The approach I take is to allow the slopes of the spline to shift parametrically or non-parametrically over time.

4.2.2 Month of Birth: Results

Figure 7 plots out the point estimates from four base specifications of $g(\text{age}; \gamma)$ - no age control, linear, quadratic, and the base spline. As the flexibility of the specification increases, the point estimates tilt towards 0 from both directions. The peak-to-trough difference within specification reduces from .245 without age controls, to .188 and .163 for the linear and quadratic specifications, to .131 with the base spline controls. F-stats and P-values for Feb-Dec jointly 0 and accounting for survey design are reported in the legend. All continuous specifications return statistically significant results ($p < .1$). The spline specification, however, returns statistically insignificant results ($p =$

¹⁰Since this specification is not continuous across age (due to the separate intercepts for each age-in-years), it may be more appropriately called a “piecewise linear” specification, with cubic splines called “piecewise polynomial”. I use the term “spline” for brevity.

.414).

Figure 7 shows that flexible specification of $g(\text{age}; \gamma)$ can mitigate timing artifact induced bias. Whether it eliminates the bias cannot be determined from this exercise. The separable model may be incapable of fully absorbing $f()$, or it may be that the secular trend is improperly modeled as survey round dummies, or it may be that the remaining seasonal pattern is real. A priori it is difficult to argue for one particular specification of the secular time trend or movement of the spline over another. It is possible, though, to test whether or not in this particular case some non-separable specification of $g(\text{age}, \text{birthdate}; \gamma)$ can make the effect disappear completely.

To test models that directly specify $g(\text{age}, \text{birthdate}; \gamma)$ I run a further series of regressions, collapsing month of birth into quarter of birth. The coarser definition of seasonality allows me to worry less about model flexibility compromising identification, and it simultaneously smooths out the timing artifact effect, giving the model the best chance of finding no effect. In Table 2, I show the results of regressions of HAZ on quarter of birth, the vector X of covariates, and several specifications of $g()$ that more explicitly model its movement over time.

Column 1 shows results using a simple linear in age specification including survey round fixed effects, while Column 2 adds in quadratic and cubic age adjustments. Column 3 shows the cubic spline results with a linear trend in birthdate, and Column 4 uses the same spline but adds a quadratic time trend in birthdate. Column 5 explicitly models $g(\text{age}, \text{birthdate}; \gamma)$, allowing each spline segment slope to follow a linear trend over time, while using survey round dummies to adjust for level changes in the HAZ-age profile. Column 6 allows the spline to have independently estimated slopes for each survey round and includes survey dummies. The bottom row of the table displays F-stats

and p-values testing the null hypothesis that quarters 2-4 are jointly 0 (with quarter 1 being omitted and set to 0), using standard errors accounting for survey design.

The primary row of interest is the row for the effect of Quarter 4 birth, as the point estimates increase in each quarter of birth and the highest quarter shows the effect of specification most clearly. Moving across Table 2 from left to right, and thus increasing the flexibility of the specification of $g()$, the coefficient on the Quarter 4 dummy drops by about 40%, from .1 standard deviations to .06. The point estimate on Quarter 4 is statistically significant across all specifications, but the F-stat testing Quarters 2-4 jointly 0 is only rejected at the 10% level for the first four columns. When the secular trend is allowed to be quadratic, or when the spline is allowed to parametrically bend over time, the F-stat indicates a failure to reject the null hypothesis of no effect. No specification makes the effect completely disappear, but increasing the flexibility of $g()$ consistently dampens the magnitude.

The relationship between month of birth and age creates a relationship between the covariate of interest and HAZ that is mechanically guaranteed. In the next section I present an example that investigates a more conventional time-series/cohort model.

4.3 GDP per Capita in Birth Year in Zimbabwe

The DHS is also a natural dataset to investigate the in utero and very early life effects of the business cycle on child health, e.g. Paxson and Schady (2005) and Agüero and Valdivia (2010) on Peru, Subramanyam et al. (2011) on India and Pongou et al. (2006) on Cameroon. The economic growth and HAZ literature is also the first that this author knows of to wrestle with the

econometric problem of separating out the effects of the HAZ-age profile from the time-series trends in the covariate of interest.¹¹

Figure 8 demonstrates the timing artifact graphically in the context of a GDP/HAZ time-series analysis. Using data from Zimbabwe, I overlay the GDP per capita time series on a disaggregated HAZ time-series similar to Figure 4 (which uses Bangladeshi data). The obvious problem is that any spurious correlation between the changes in GDP per capita and where the peaks/troughs of the HAZ-age profile hit in calendar time will cause bias in regression estimates when $g()$ is not properly specified.

4.3.1 GDP: Estimation

The model I specify to estimate the effects of (log) GDP per capita in year of birth, again derived from Equation 5, is:

$$HAZ_{iac} = X_{iac}\beta_x + \beta_{gdp} * \text{LogGDP}_c + g_1(\text{age}_a; \gamma_a) + g_2(\text{birthdate}_c; \gamma_c) + \eta_{iac} \quad (8)$$

This specification excludes survey-round effects, instead parametrically modeling movements in the HAZ-age profile over time. Since identifying variation necessarily comes from temporal changes in GDP, the specification of both $g_1(\text{age}; \gamma_a)$ and $g_2(\text{birthdate}; \gamma_c)$ are important.

To test the sensitivity of this model to specification of $g_1(\text{age}; \gamma_a)$ I test five specifications: unconditional on age, linear, quadratic, spline, and dummy variables for age-in-months. $g_2()$ is specified as linear or quadratic. Identifying

¹¹Paxson and Schady (2005) is the first to discuss the problem as one in which the object of interest is the entire HAZ-age profile. It is informative that, instead of regressing HAZ on GDP and estimating an elasticity, it simply compares HAZ-age profiles across survey rounds, and conceives of the problem as identifying changes in the shape of the HAZ-age profile over time

variation comes across time, and in the continuous specifications of $g()$ that includes both within-survey time variation (e.g. comparing children surveyed in 1996 and born in 1993 and 1994) along with variation in time across surveys (e.g. comparing the 1993 cohort with the 2005 cohort surveyed later). As the specification of $g_1()$ becomes more flexible, though, the models rely increasingly on between-survey variation, up to the point of the dummy variables for age in months which identify almost exclusively off between-survey variation. There are two concerns, though, with the dummy variable specification: first, the problem of implicitly comparing December-born and January-born children of the same age from the same survey who faced essentially the same economic conditions in utero but would have different values of GDP per capita in their birth year; and second the fact that, for any given age, there are only (S =) 4 values of GDP per capita identifying the effect. The increasingly flexible estimates utilize less and less of the available variation in GDP per capita.

An obvious compromise between a polynomial specification and the non-parametric specification is again a spline in age. Similar to the previous section, I set the spline to have internal knots at round ages in years with separate intercepts at each knot. Unlike in the month of birth example, here I can safely specify a cubic spline to allow the model to fit the non-linearity of the HAZ-age profile as well as possible.

4.3.2 GDP: Results

In Table 3 I present the results from a regression of HAZ on the same covariates used previously, a linear or quadratic trend in birthdate, and the specifications of $g_1(\text{age}; \gamma)$ discussed above. The effect of the age controls is clear, the point estimates dropping towards 0 with each increasingly flexible specification. The

unconditional regression estimates that a 10% increase in GDP per capita in birth year is associated with a .1 standard deviation increase in child height. Linear specifications only reduce the implied effect to about .085 standard deviations. The addition of age squared reduces the point estimate by almost another half, to .048 standard deviations. The spline and non-parametric specifications, my most flexible models, return estimated effects of around .03 standard deviations, 70% smaller than the estimate from the unconditional specification, and just over 35% less than the estimate from the quadratic specification.

The inclusion of quadratic controls for the secular trend leads to larger point estimates under the spline and non-parametric specifications. Point estimates in columns 6 and 7 are about 35% larger than the point estimates for the same specifications of $g_1()$ with linearly specified trends. One concern with the validity of these estimates is that, given the underlying data, there will always be a large increase in HAZ toward the very end of the time-series, since these cohorts are always composed of very young children. Thus, it may be that using a quadratic specification is unwise as it will become to some degree collinear with the HAZ-age profile at the far right end of the graph. This concern may or may not explain the increase in point estimates, and I cannot rule out the hypothesis that the quadratic specifications are improvements over the linear specifications.

All estimates of the effects of GDP in birth year on HAZ are statistically significant at the 95% level or lower when standard errors only accounting for survey design are computed. As discussed in Section 3.3, these standard errors do not account for serial-correlation in errors within cohort or region but outside of sampling cluster.

To provide initial evidence as to whether or not choice of inference strat-

egy can greatly affect our confidence level, I compute several alternative p-values and display them in the bottom rows of Table 3. The first row, labeled “DHS”, shows the p-value associated with the standard error estimates shown beneath the point estimates. This can be considered a baseline p-value against which to judge the alternatives. The second and third rows show p-values from standard errors that employ cluster-robust analytic specifications that cluster the data at the birth year-by-region level (210 clusters) and the survey-by-region level (40 clusters) respectively. P-values are based on t-distributions with degrees of freedom equal to the number of clusters minus 2. The bottom two rows provide p-values from the wild cluster-t bootstraps that are clustered on region and cohort separately¹².

There is a great deal of dissonance between the estimated confidence levels across inference strategies, particularly between the analytic and bootstrap estimators. All analytic clustering strategies, save for the region-by-survey clustering on the first dummy variable specification, return results that are statistically significant at the 10% level. However, both bootstraps fail to reject the null hypothesis at any standard confidence level for any of the specifications save the non-parametric specification with a quadratic trend in birthdate.

4.3.3 GDP: Informal Inference Test

The massive increase in the confidence interval (or, for the bootstraps, implied confidence interval) as clusters are defined at higher levels raises the possibility that the DHS recommended standard errors are inappropriately small. While I more formally test these inference strategies in the next section, I first pose

¹²For these bootstraps I use 500 repetitions, but in the alternate-country GDP regressions below I use only 300 bootstrap repetitions to save computational requirements.

two questions to the data. First, how often would a random GDP per capita time-series overlaid on the Zimbabwe DHS data generate results as extreme as those found in the real regression? Second, how often would we reject the null hypothesis of no effect when these placebo-GDPs are used? Neither of these can be answered perfectly, but I can provide some evidence in the empirical context.

Determining what, exactly, constitutes a placebo GDP time-series is difficult. As opposed to generating random GDP time-series, I take a less precise but more intuitive approach. I address both questions by estimating the implied effects of other country's GDP per capita time series on the Zimbabwean HAZ data. First, I take GDP per capita time-series data from the World Bank online reference for 183 countries with data on GDP from 1990 through 2011¹³. I then regress the Zimbabwe child height data on the GDPs of these other countries, using the base spline specification with a linear time trend and recording the point estimates and p-values.

Figure 9 graphs a kernel density plot of the GDP point estimates from other country's GDPs, and a dot for the estimated effect of Zimbabwe's own GDP. The vertical line shows the 95-th percentile of the distribution, and the Zimbabwe point estimate is smaller than this level. Overall, the "real" point estimate is less extreme (in terms of absolute value) than the point estimates obtained using 42 of 183 alternate GDPs (about 23%). This number is similar to the p-value obtained from the preferred birth year clustered bootstrap p-value (.188).

I then tabulate the rejection rates (5% level) and present them in Table 4. Using DHS inference methods, I find rejection rates between .39 and .84 depending on specification, with the more flexible specifications producing the

¹³Data was collected from data.worldbank.org

lower, but still too large, rejection rates. The cluster analytic standard errors reject the null at rates between .2 and .6, again with the more flexible specifications of $g()$ leading to lower rejection rates through the reduction of bias on the coefficient estimate. Of particular interest, though, are the rejection rates given by the wild cluster-t bootstraps that cluster on birth year, which reject the null hypothesis at a rate of around .03 to .09.

The GDP per capita exercise demonstrates two things. First, the timing artifact can bias results even when the right hand side variable of interest is not mechanically related to age, only measured in relation to it - the *value* of GDP per capita in birth year is associated with age-at-measurement, but it need not be. Second, standard errors calculated only to account for survey design may be too small. What is not clear from this exercise, or from the preceding exercise using month of birth, is whether the flexible specification of $g()$ is fully purging the bias and whether or not any of the strategies for estimating statistical precision produce reliable p-values.

5 Monte Carlo: “Weather Shocks” in Peru

This section presents a formal test of both proposed strategies for eliminating timing artifact-induced bias, as well a test of the quality of various precision estimators. To do this, I perform two Monte Carlo exercises based on data from the Peruvian DHS, where a placebo shock with known effect of 0 is randomly applied to children in some regions in each cohort but not to others. The use of a “shock” indicator for a particularly severe exposure of some type or another is common, and weather shocks exemplify the strand of the literature (del Ninno and Lundberg (2005)). The Peruvian data is used because of the relatively

large number of geographic regions provided in the data (25)¹⁴, sparing the need to define my own geographic regions and allowing for easier replication.

5.1 Simulation Design and Estimation

The spatio-temporal fixed-effects model I consider in this context is derived from Equation 6 but drops the age subscript since the model does not directly specify $g()$. It takes the following form:

$$HAZ_{ircs} = X_{ircs}\beta_x + \beta_{shock} * Shock_{rc} + \lambda_c + \gamma_r + \eta_{ircs} \quad (9)$$

If regional differences in child HAZ are essentially level differences in the HAZ-age profile, then, conditional on a dummy for region of birth, “control” (shock = 0) children should generate a reasonable counterfactual mean HAZ for children in their cohort in the “treated” (shock = 1) group. The additional identifying assumption of importance for this model is the equivalence of age distributions between treatment and control within cohort¹⁵.

This model is compared to models that directly specify $g()$ and follow Equation 8, but generalized to multiple regions. Identifying variation for these models can be conceived in exactly the same way as in the GDP example, since temporal, “within” variation is used to identify the coefficients. The variation is just repeated $R = \text{number of regions}$ times for each survey.

¹⁴There are only 13 regions in the 1992 data and 24 in 2000, but these same regions appear in other survey rounds, so I include them all. Since all regions are very likely to experience both “treatment” statuses in any given Monte Carlo repetition, this should not pose a problem.

¹⁵In the context of the DHS, this is a more stringent assumption than it may seem. Some cohorts are measured in two survey rounds, and thus a cohort of children could be measured at either a very young age or a relatively old one. Thus, the assumption requires not only within-survey-by-cohort equivalence in the distribution of child ages across treatment, but also similar proportions of observations in the cohort taken from each survey round over T .

The baseline simulation works in the following manner. First, each region-by-cohort cell is randomly assigned a shock value of 0 (control) or 1 (treatment) with equal probability. I then run three regressions on the placebo treatment data. The first regression includes a linear specification of the HAZ-age profile while the second includes the base cubic spline with internal knots and intercepts at round age in years. Both regressions include a linear trend in birthdate for $g_2(\text{birthdate}; \gamma)$. The third regression employs birth-year dummy variables (as represented in Equation 9). I then repeat this procedure 500 times, reassigning the placebo treatment variable through each repetition and saving the point estimates and p-values of $\hat{\beta}_{shock}$ for each regression.

In the case of this fully random shock, none of these specifications should return biased estimates in the mean of the repeated simulations. That is because, by construction, there is no correlation between age-at-measurement and the covariate of interest. A real test of the specifications must induce a correlation between cohort and T, preferably in a manner that is likely to occur in real world empirical investigations.

I induce the necessary correlation by increasing the probability of treatment in a single cohort. Call this year a “drought year”, it is a year when the shock hits more regions than in other years. The timing artifact will pressure the point estimates on T higher if the drought year corresponds to cohorts with high μ , and vice versa if μ is low for that cohort. I run the simulation 500 times treating the first cohort year as the drought year, and then repeat the procedure for all other cohorts.

5.2 Results: Independent Shocks

Figure 10 shows a kernel density estimation of the distribution of $\hat{\beta}_{shock}$ for the random (50-50) assignment of treatment status by region-cohort. As predicted, all three estimators produce unbiased estimates of $\hat{\beta}_{shock}$. The spline and fixed-effect models generate more precise estimates, likely driven by better performance when unlikely draws induce some timing artifact bias in some simulation repetitions.

The purely random placebo test demonstrates two things. First, it confirms the theoretical prediction that, when T is uncorrelated with age, the timing artifact does not infect the data. Second, it provides an excellent environment for testing the inference properties of the DHS recommended standard errors that account for survey design, and an analytic clustering strategy that accounts for serial correlation and heteroskedasticity. The particular form of the placebo-test leads to a natural clustering level of region-by-cohort, the level at which the shock is determined.

Through each replication of the placebo-test, I record whether or not the model rejects the null hypothesis of no effect at the 95% level. Table 5 displays the empirical rejection rates from the exercise. The linear specification using DHS recommended inference rejects a true null hypothesis of zero effect over 40% of the time. That is, if a researcher studying the effects of a one year drought used these models, they would find a statistically significant estimate 2 out of 5 times even if there was no real effect of drought, though only about half of those estimates would be both significant and of the sign anticipated. The spline and non-parametric specifications reject the true null about 30% of the time, still a rate about 6 times too often. Clustering at the region-by-birth year cell level greatly improves inference. All three specifications now reject

at a rate around .08.

5.3 Results: “Drought Years”

Now I induce a correlation between the shock and age-at-measurement by including in each simulation a “drought” year. If models misspecify $g()$ in the manner described in Section 3, the simulation should now induce a bias in point estimates of $\hat{\beta}_{shock}$.

The results are represented in Figure 11. The X-axis shows year, which refers to either year of drought (for the line graphs) or year of birth (for the bar graph). The bar graphs display mean age-at-measurement for children born in that year, with red bars indicating that measurements were taken in that year. The short-dash red line shows the mean estimate of $\hat{\beta}_{shock}$ when the HAZ-age profile is specified as linear. As predicted, the correlation between shock and cohort age induces bias in the improperly specified model. In the worst case, this bias is about .03 standard deviations, a small but economically significant amount, and that effect is generated from the addition of just a few extra “treated” regions in the birth year.

The other two lines show mean estimates of $\hat{\beta}_{shock}$ under the spline and fixed-effect specifications. The red long-dashed line representing the spline specification reveals the extent to which flexible specification can mitigate the bias induced by the timing artifact, but the specification does not completely purge the bias. The fixed-effect model, represented by the solid red line, stays firmly centered on zero, indicating that the model produces unbiased estimates regardless of survey- and shock-timing.

The relationship between the spline estimates and the linear estimates demonstrates three further points: First, the spline estimates are fairly close

to zero when the drought years hit cohorts that are not on the ends of the sample birthdates or just before or after a survey has taken place. Second, misspecification of $g()$ is most problematic when the drought hit the youngest cohort in a survey, showing the the discontinuity in the time-series at survey-round breaks can be particularly problematic. Finally, increased flexibility of $g()$ need not always simply reduce bias, but it can actually reverse the sign of the bias. The linear and spline estimates for the drought years near either end of the time period show coefficients that are getting more negatively biased as children grow younger (and thus are likely to have higher HAZ), opposite of the prediction for specifications unconditional on age.

Table 6 provides a summary of the regression results for several years in which the faux-drought induces large and small biases on the misspecified models. For each model and drought year I display the mean point estimate of $\hat{\beta}_{shock}$ and the rejection rates from the DHS and region-by-cohort clustered standard error calculations. 1996 is the drought year that induces the largest bias. A linear specification yields a mean point estimate bias of around .03 standard deviations. The linear model rejects the true null hypothesis of no effect at a rate of over .55 when using DHS recommended inference, and still around .15 when clustering at the cohort-by-region level. The mildly biased spline estimate and the unbiased fixed-effects estimate yield a rejection rate of over .25 using the DHS inference, but the rejection rate when clustering by cohort-region is down to between .06 and .09.

In general, the specifications that induce more bias tend to over-reject more than the unbiased estimators, simply because the t-statistics increase as the bias increases when the bias moves point estimates away from 0. Models that cluster at region-by-cohort levels, though, fair better under all specifications than models using DHS standard inference. However, cohort-region

clustering may still be too restrictive, since it fails to deal with the temporal serial-correlation in T that was problematic in the GDP context. I do not estimate the wild cluster- t bootstraps in this context due to the large computing requirements, but they may be appropriate for researchers operating in the spatio-temporal econometric environment when there is temporal serial-correlation in T .

6 Conclusion

Estimates of cohort determinants of child health can be greatly biased when the HAZ-age profile is not properly specified. The lesson to practitioners is clear: if possible and relevant, bring within-cohort variation in the covariate of interest to bear on the problem. If not, specify the HAZ-age profile as flexibly as possible. Researchers using other anthropometric or child health outcomes should examine the outcome-age profile before blindly specifying a functional form for their regression equations, and they should examine the movement of that profile over time when they are appending multiple rounds of the DHS. Standard error estimates should account for serial correlation and heteroskedasticity via analytic or bootstrap methods that cluster at higher levels than those accounting only for survey design. The extent to which these problems have led to the publishing of spurious results in the scientific literature requires further investigation.

I conclude with a note about the interpretation of coefficient estimates. In an OLS setting, the coefficient on the treatment variable is by definition modeled as a level shift in the HAZ-age profile. This is likely not the way that many inputs affect HAZ. Often times, it may be better to conceive of these inputs as bending the HAZ-age profile. While I hope this paper will convince

researchers to more carefully model the HAZ-age profile (and the outcome-age profile in general when dealing with child health outcomes), I also hope that it will spur new discussion about what exactly it is that we are trying to measure when we estimate treatment effects on HAZ.

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7 Figures and Tables

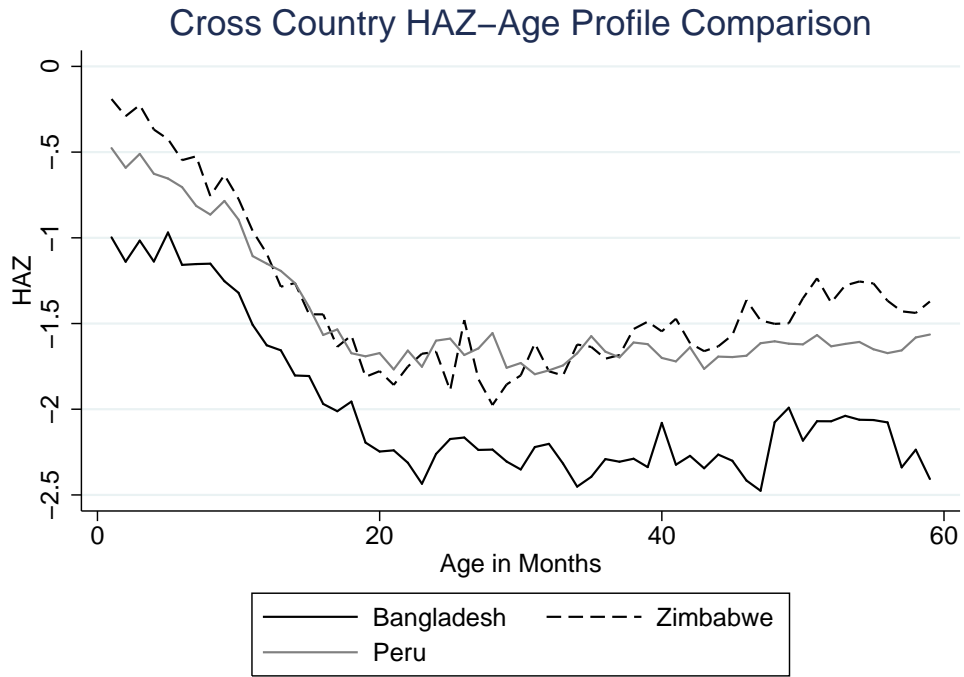


Figure 1: HAZ-age Profiles

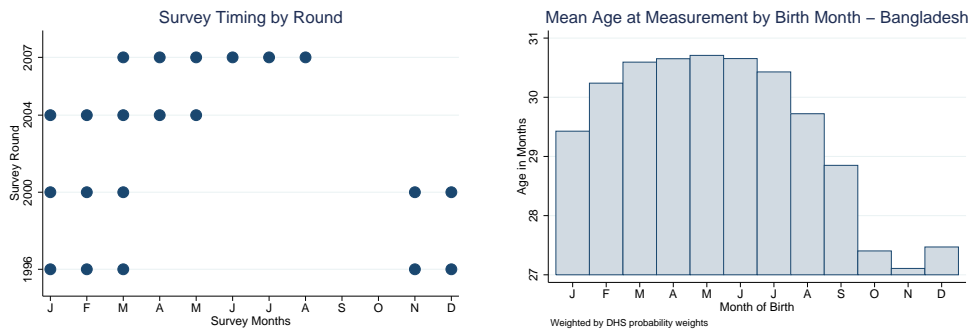


Figure 2: Survey Timing and Age at Measurement - Bangladesh

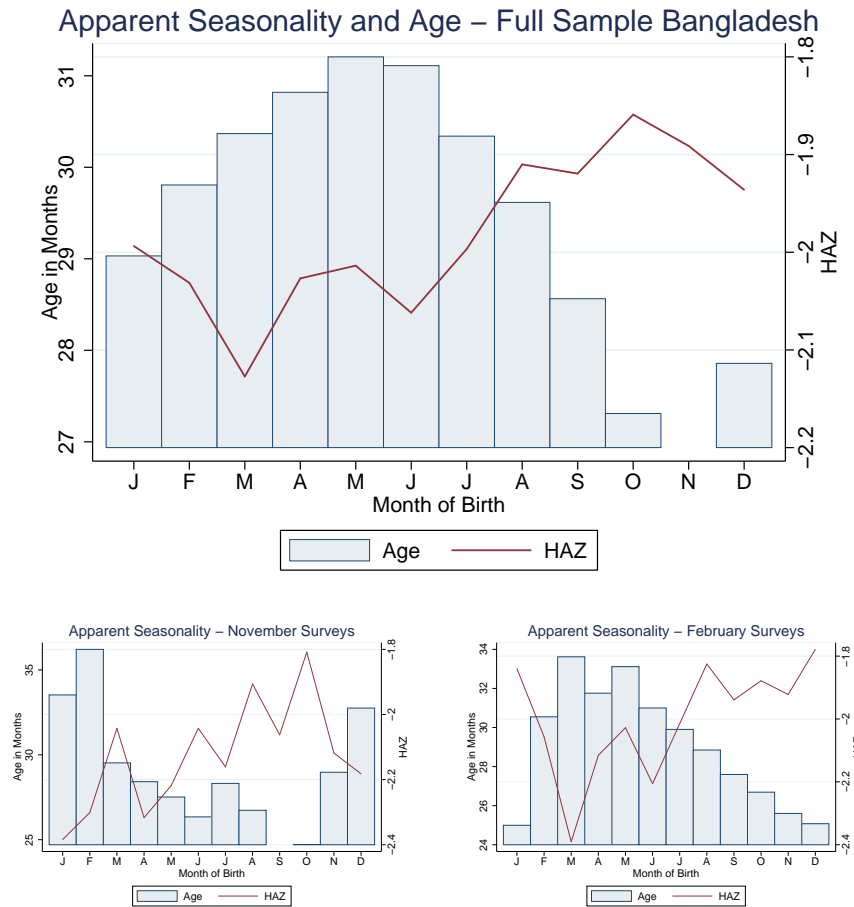


Figure 3: Survey Timing and Apparent Seasonality - Bangladesh

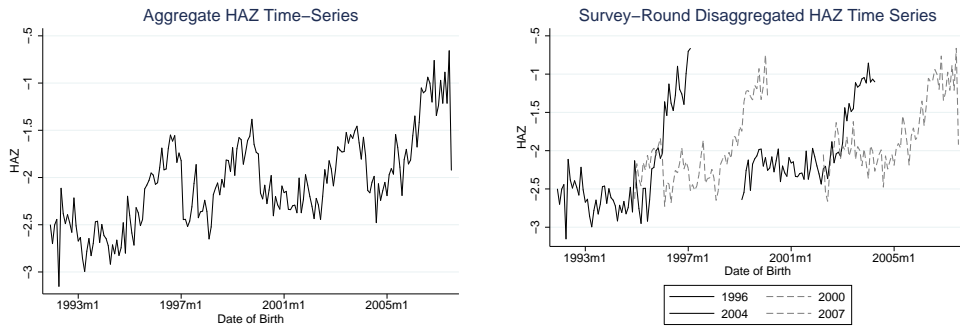


Figure 4: HAZ Time Series - Bangladesh

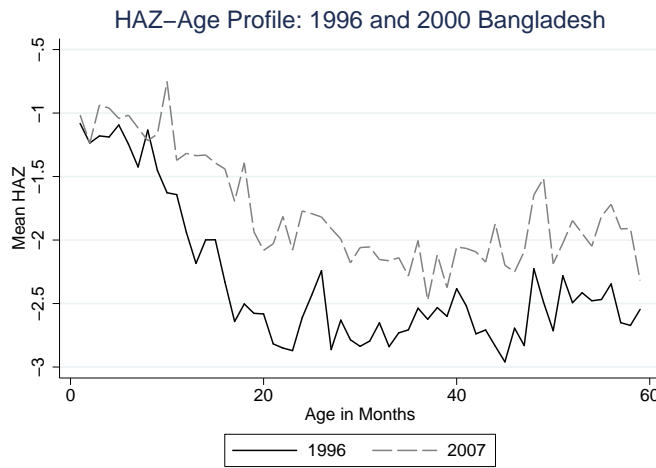


Figure 5: HAZ-age Profiles Over Time - Bangladesh

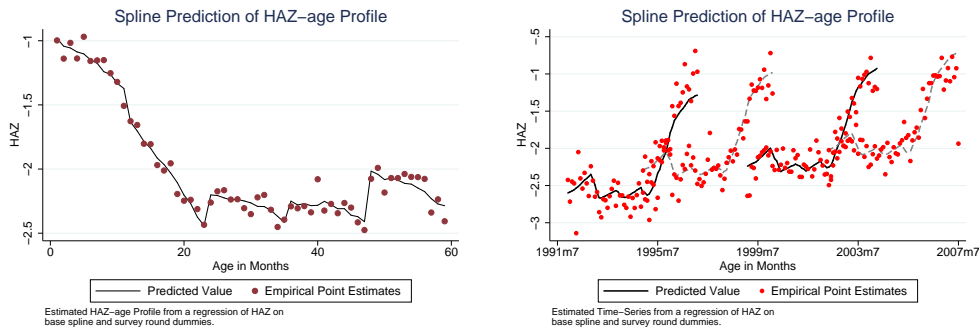


Figure 6: Spline Fit - Bangladesh

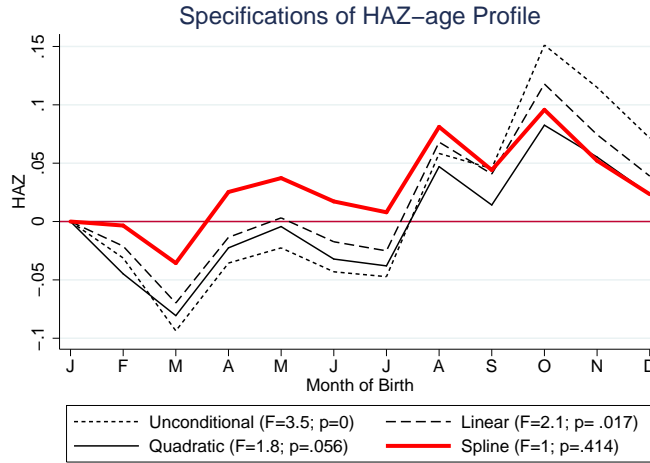


Figure 7: Flexible Specifications of $g()$ - Bangladesh

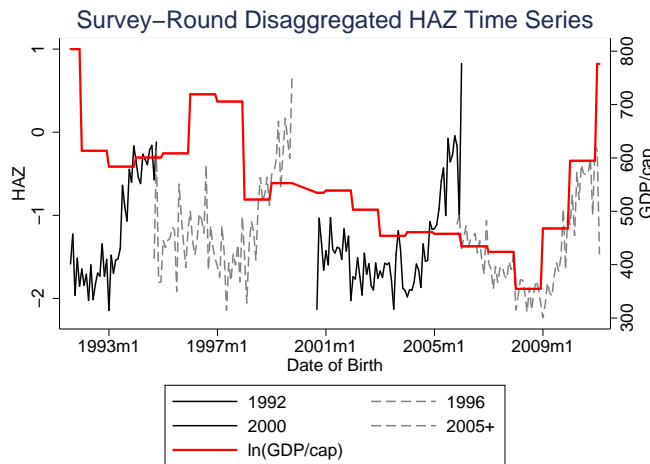


Figure 8: GDP and HAZ - Zimbabwe

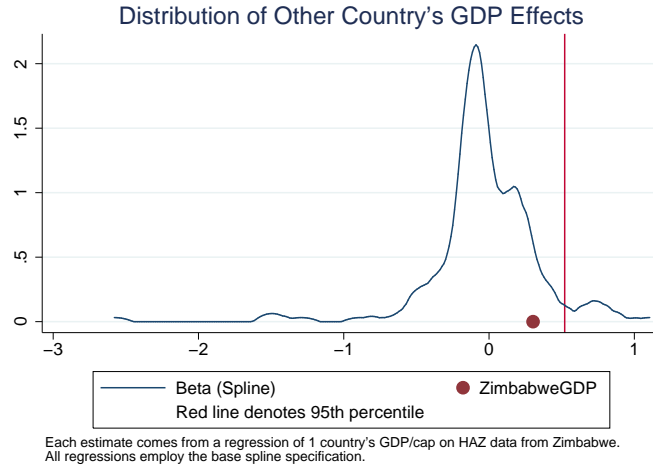


Figure 9: “Placebo” Estimates of $\hat{\beta}_{GDP}$ - Zimbabwe

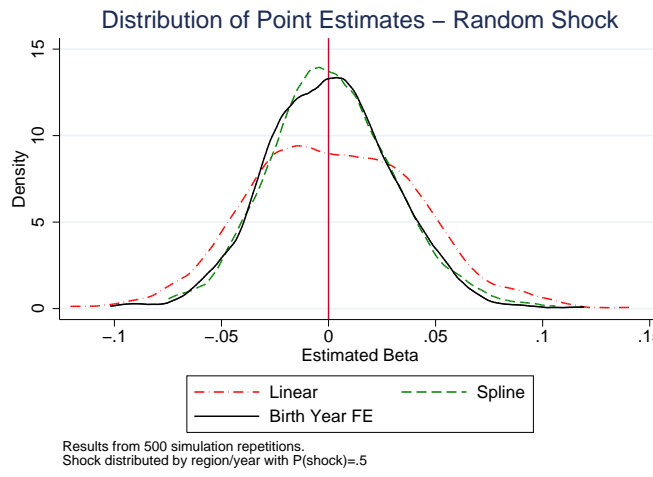


Figure 10: Distribution of $\hat{\beta}$ under Random Shock - Peru

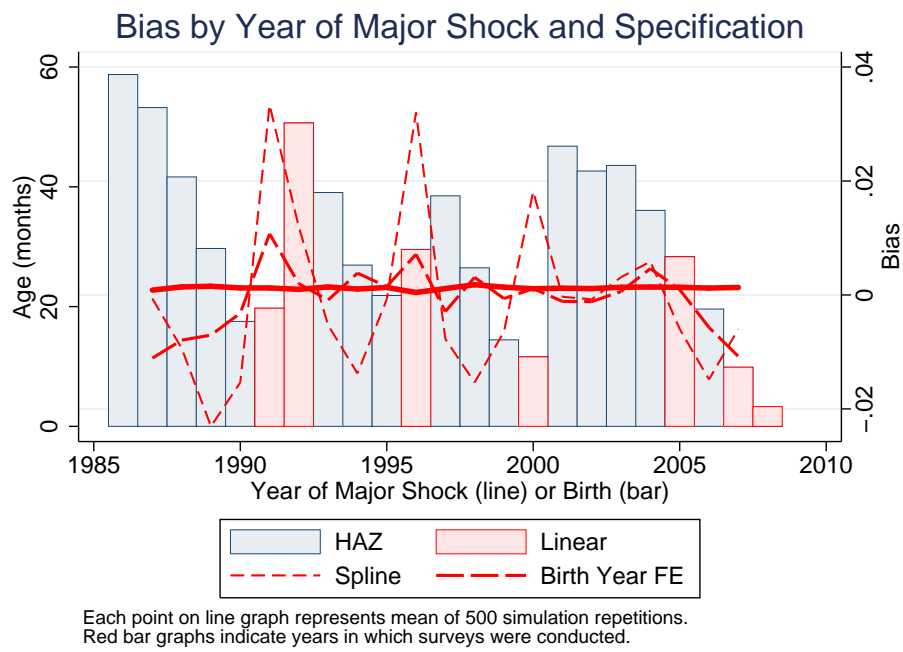


Figure 11: Drought and Bias - Peru

Table 1: Sample Summary Statistics

	Bangladesh	Peru	Zimbabwe
HAZ	-1.970	-1.454	-1.298
Child Age	29.14	30.00	25.98
Child Female	0.495	0.496	0.501
Birth Year	1999.4	1996.5	2001.8
Maternal HAZ	-220.5	-224.0	-64.41
Maternal Age	25.84	29.24	27.85
Secondary Ed.	0.298	0.484	0.533
Urban	0.263	0.517	0.239
N Surveys	4	4	4
Obs	21351	44205	13012

Table 2: Quarter of Birth and HAZ - Bangladesh

	(1)	(2)	(3)	(4)	(5)	(6)
	Linear	Cubic	Spline1	Spline2	Spline3	Spline4
	b/se	b/se	b/se	b/se	b/se	b/se
2nd Quarter	0.018 (0.03)	0.014 (0.03)	0.040 (0.03)	0.036 (0.03)	0.030 (0.03)	0.032 (0.03)
3rd Quarter	0.059* (0.03)	0.050* (0.03)	0.061** (0.03)	0.055* (0.03)	0.046 (0.03)	0.046 (0.03)
4th Quarter	0.104*** (0.03)	0.100*** (0.03)	0.070** (0.03)	0.065** (0.03)	0.060** (0.03)	0.060** (0.03)
Female	0.018 (0.02)	0.022 (0.02)	0.021 (0.02)	0.022 (0.02)	0.021 (0.02)	0.021 (0.02)
Urban	0.200*** (0.03)	0.205*** (0.03)	0.214*** (0.03)	0.211*** (0.03)	0.209*** (0.03)	0.209*** (0.03)
Maternal Age	0.012*** (0.00)	0.012*** (0.00)	0.012*** (0.00)	0.012*** (0.00)	0.012*** (0.00)	0.012*** (0.00)
Maternal HAZ	0.003*** (0.00)	0.003*** (0.00)	0.003*** (0.00)	0.003*** (0.00)	0.003*** (0.00)	0.003*** (0.00)
r2	0.158	0.188	0.199	0.200	0.202	0.203
Fstat	5.206	4.938	2.366	2.020	1.652	1.648
Pval	0.001	0.002	0.069	0.109	0.176	0.176

Point estimates and (standard errors) account for sample clustering and weights.

All regressions also include region dummies and five dummies for maternal education.

Table 3: Log GDP per cap and HAZ - Zimbabwe

	(1)	(2)	(3)	(4)	(5)	(6)	(7)
UC	Linear	Quad	Spline	NP	Spline-Q	NP-Q	
b/se	b/se	b/se	b/se	b/se	b/se	b/se	
loggdp	1.122*** (0.136)	0.853*** (0.127)	0.476*** (0.125)	0.301* (0.125)	0.277* (0.127)	0.411** (0.134)	0.387** (0.135)
bdate	0.001** (0.000)	0.000 (0.000)	-0.001* (0.000)	-0.001*** (0.000)	-0.001*** (0.000)	0.012* (0.005)	0.012* (0.005)
r2	0.041	0.078	0.126	0.134	0.139	0.135	0.139
P_values							
DHS	0.000	0.000	0.000	0.017	0.029	0.002	0.004
Cohort_Region	0.000	0.000	0.005	0.049	0.071	0.012	0.019
Survey_Region	0.002	0.000	0.010	0.077	0.115	0.047	0.069
Region_Wild	0.556	0.360	0.156	0.140	0.164	0.140	0.104
Cohort_Wild	0.388	0.244	0.112	0.280	0.248	0.140	0.048

UC is unconditional on age. Linear, quadratic (Quad) non-parametric (NP) refer to age controls. Spline refers to cubic spline with internal knots and intercepts at 12, 24, 36, and 48 months of age. All regressions also include region dummies, rural/urban and gender dummies, five dummies for maternal education and linear adjustments for child birthdate and maternal age and HAZ. Weighted using re-normalized sample probability weights. -Q indicates quadratic in birthdate included. Standard Errors in table account for weighting and sampling design (DHS) P-values below compare two clustering levels (Cohort- and Survey-by-Region) and two wild cluster-t bootstraps (Region and Cohort).

Table 4: Rejection Rates of Placebo GDP Effects

	DHS	Cluster	Bootstrap
Unconditional	0.841	0.590	0.0387
Linear	0.780	0.656	0.0884
Quadratic	0.473	0.311	0.0663
Spline	0.341	0.213	0.0387
NP	0.390	0.273	0.0552

Statistical significance determined at 95 percent level.

DHS refers to standard errors accounting for survey design.

Cluster uses SEs clustered at region by cohort level.

Bootstrap from wild cluster-t bootstraps clustered at cohort level.

Table 5: Random Shocks Rejection Rates (.05 level)

Linear:

DHS 0.416

Cluster 0.0720

Spline:

DHS 0.292

Cluster 0.0780

FE

DHS 0.250

Cluster 0.0820

Clustered at region-by-cohort level

Table 6: Drought Shocks Rejection Rates by Age Control (.05 level)

	1988	1992	1996	2000	2004
Linear:					
Beta (Linear)	-0.00935 (0.00165)	0.0119 (0.00168)	0.0321 (0.00162)	0.0181 (0.00167)	0.00587 (0.00169)
Survey Design	0.406 (0.0220)	0.412 (0.0220)	0.548 (0.0223)	0.430 (0.0222)	0.390 (0.0218)
Region-Cohort	0.0700 (0.0114)	0.0880 (0.0127)	0.150 (0.0160)	0.0980 (0.0133)	0.0740 (0.0117)
Spline:					
Beta (Spline)	-0.00793 (0.00122)	0.00198 (0.00123)	0.00715 (0.00119)	0.00111 (0.00122)	0.00464 (0.00123)
Survey Design	0.282 (0.0201)	0.262 (0.0197)	0.260 (0.0196)	0.268 (0.0198)	0.274 (0.0200)
Region-Cohort	0.0680 (0.0113)	0.0620 (0.0108)	0.0620 (0.0108)	0.0560 (0.0103)	0.0540 (0.0101)
FE:					
Beta (FE)	0.00142 (0.00128)	0.000987 (0.00129)	0.000435 (0.00130)	0.00110 (0.00127)	0.00140 (0.00129)
Survey Design	0.276 (0.0200)	0.272 (0.0199)	0.254 (0.0195)	0.268 (0.0198)	0.280 (0.0201)
Region-Cohort	0.0720 (0.0116)	0.0800 (0.0121)	0.0940 (0.0131)	0.0700 (0.0114)	0.0760 (0.0119)

Region-Cohort denotes clustering level

Standard Errors of Mean in parentheses

Columns refer to year with additional shocks

A Sample Construction

All regression samples are constructed using the same selection procedures. First, I append all four survey rounds of each country and drop all observations without height measurements, all children under the age of 1 month and all observations where child birthdate is imputed (5 in Bangladesh, 21 in Zimbabwe, and none in Peru). This leaves me with measurements on 22471 children in Bangladesh, 14524 in Zimbabwe and 46357 in Peru. Of these, I drop 785 observations from Bangladesh, 1265 from Zimbabwe and 1243 from Peru for having invalid measurements or measurements resulting in an HAZ score above 6 or below -6 standard deviations, which is the WHO definition of the range of valid HAZ scores and is common practice in the literature (Borghi et al., 2006). Excluding children with missing covariate measurements drops an additional 335 observations in Bangladesh, 247 in Zimbabwe and 909 in Peru. This leaves me with final sample sizes of: Bangladesh - 21351 ; Zimbabwe - 13012 ; and Peru - 44205 .

The DHS sampling method is partly unique to each survey but usually follows a similar procedure in most years and countries. Geographic regions are defined, and the country is stratified by region and urban/rural status. Clusters are chosen randomly from each strata, and households are sampled randomly (though with oversampling of some households) within cluster, though sampling is not uniform across time within strata¹⁶. Households are then assigned weights in order to make the weighted sample nationally representative, and these weights are normalized to sum to the sample size. All regressions in this work use re-normalized weights that sum to 1 within each survey round, implicitly weighting each round equally. Using the original DHS weights (which implicitly weights surveys by sample size), or using no weights at all, has no qualitative impact on the analyses presented.

¹⁶The fact that regions are surveyed at different times does lead to questions about the consistency of point estimates from region fixed effect models. The issue is not addressed in this work. The Monte Carlo simulations in Section 5 give some evidence that any bias induced by non-random sample timing across regions is likely to be of secondary concern in cases of region by birth year treatment assignment, but the effect may be problematic in other contexts