

Estimating response propensity and nonresponse

bias for the 2022 Agricultural Production Census

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The Agricultural Production Census



- Undertaken every 5 years in partnership with the Ministry of Primary Industries
- Aims to provide a range of summary statistics on the agricultural industry in New Zealand (e.g., total number of cows).
- Target population is all businesses engaged in agricultural production activity during the year ended
- Despite being a survey, we have information on both responding an non responding farms.
- High-level strata consist of a combination of region and farm types
- The final stratification variable is the total land area of each farm measured in hectares.





The Agricultural Production Census



- In 2022 the APC had an atypical low response rate of 69%
- In comparison, in the 2017 census the response rate was 84%
- In addition to the general tendency of low response rates, Groundswell NZ called for all farmers and growers to boycott the APC
- Historically, nonresponse has been handled by donor imputation
- Low response rates can potentially introduce nonresponse bias.





Nonresponse bias



- Decreasing response rates may not always lead to nonresponse bias. Low response rates are not necessarily "bad" per se.
- Nonresponse bias occurs as a function of how correlated response propensity is to the attributes measured.
- Within the same survey, nonresponse bias can vary across different variables.
- To discern when nonresponse rates lead to nonresponse bias, we must understand how the

influences for and against participation are related to the survey measures.

Can nonresponse bias actually be quantified?



- Nonresponse bias is notoriously difficult to estimate because we do not know the nonrespondent's values.
- The bias estimation based on the sample respondents will not equal the population bias (i.e., you need the whole population)
- Bias approximations also need *Y* values for the nonrespondents (which we do not know), or some approximation of them (based on variables that correlate with them).
- Imputation assigns *Y* values to nonrespondents.
- All the expressions that relate response propensities to nonresponse bias are based on approximations because the estimators are nonlinear.
- While approximations are quite good in many cases, they may be less precise in some situations.

Response propensity



$$\phi_i = \phi(x_i) = \Pr(R_i = 1 | X = x_i)$$

- ϕ_i The response propensity for unit i
- x_i Auxilary data for unit i
- $R_i = 1$ If unit i responds

- Response propensities are unknown, we observe only the binary outcome of response or nonresponse
- We often have auxiliary data available for all sampled units that can be used to understand/adjust for non response
- We assume that $\phi_i > 0$ for all *i*
- Response propensities are often estimated by logistic regression, but probit and non parametric methods can also be used.
- Response propensities are dynamic and likely to vary with the recruitment protocol

Response propensity



• What causes a survey variable to be correlated to the likelihood to respond?



Figure 1. Three Relevant Causal Models Linking Response Propensity with Nonresponse Bias.

Response propensity weight adjustment



 $\hat{y} = \sum_{i} d_i \phi_i^{-1} y_i$

- An method to adjust for nonresponse
- The adjustment factor is the inverse of the estimated propensities of the respondents.
- The idea is to replace the unknown probability of response by an estimate

Research questions



- Which auxiliary variables best correlate with response propensities for the APC?
- Did response propensities change between 2017 to 2022?
- Is there evidence of nonresponse bias in key variables in the Agricultural Production Census?
- Does response propensity weight adjustment give different results to donor imputation?
- Can nonresponse bias decrease with response propensity weight adjustment?







 $\operatorname{response}_i \sim \operatorname{Binomial}(1, p_{i,j})$

$$p_{i,j} = \alpha_{\text{cell}[j]} + \beta_{\text{s,cell}[j]} \text{size}_i + \beta_{\text{r,cell}[j]} \text{past responses}_i$$
$$\begin{bmatrix} \alpha_{\text{cell}} \\ \beta_{\text{s,cell}} \\ \beta_{\text{r,cell}} \end{bmatrix} \sim \text{MVNormal} \left(\begin{bmatrix} \alpha \\ \beta_s \\ \beta_r \end{bmatrix}, \boldsymbol{\Sigma} \right)$$

- $\mathbf{\Sigma} = \mathbf{SRS}$
- $\alpha \sim \text{Normal}(0, 1.5)$
- $\beta_s \sim \text{Normal}(0, 0.5)$
- $\beta_r \sim \text{Normal}(0, 0.5)$
- $\sigma_{\alpha} \sim \text{Exponential}(1)$
- $\sigma_{\beta_s} \sim \text{Exponential}(1)$
- $\sigma_{\beta_r} \sim \text{Exponential}(1)$
- $\mathbf{R} \sim \text{LKJcorr}(2),$

- Multilevel Bayesian logistic model
- Imputation cell as a random effect (i.e., each cell had its own intercept and slope)
- Imputation cells are a combination of farm type, region and a range of sizes.
- Predictor variables : size (log transformed) and number of past responses.



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- Adjusted this model to 2017 and 2022 census data.
- Model comparison with LOOIC showed this is the best fit model for both sets of data compared to less complex models (e.g., only intercept)





- Posterior predictive checks showed this model is a relatively good fit to the data (figure shows 2022).
- Blue points are observed response rates while grey points and lines are median and 90% of the highest posterior density





Posterior draws of the expected value of the posterior predictive distribution for every observation in 2017 and in 2022









Bias within imputation cells





- Poststratification and imputation will reduce nonresponse bias if response propensities are homogeneous within strata
- And if there is little correlation between response propensities and the response variable

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Response propensity weight adjustment





Year

2017

2022

- Comparison between donor imputation and response propensity weight adjustment
- Dairy cows

Response propensity weight adjustment



• Beef cows



Main findings



- Main predictors of response propensity were size of the farm and the number of past responses.
- Response propensities decreased from 2017 to 2022 and were more widespread in 2022
- However, the patterns of response propensity remained similar to those observed in 2017
- For some imputation cells in 2022 response propensities were not homogeneous per strata and covaried with the number of animals, which likely introduced nonresponse bias at this level.
- At a regional level the results of the donor imputation and response propensity weighting methods were consistent





Moving forward



- Predicting response propensities is increasingly important to understand nonresponse bias.
 New covariates and predictive models can and should be explored if we want to accurately quantify nonresponse bias.
- As for the current release:
 - Some cells were divided to make them more homogeneous.
 - Suppression of outputs that derive from a high percentage of biased cells
- For future estimates:
 - New imputation methods can and should be explored under the assumption that response rates might not improve.
 - Multiple imputation
 - Weight calibration methods with response propensity
 - Model assisted estimates.







Thank you

Nonresponse bias



 $bias(y^{\overline{pst}}) \approx N^{-1} \sum \bar{\phi_h}^{-1} \sigma_{\phi_h} \sigma_{Y_h} \rho_{\phi_h,Y_h}$

- $\hat{y^0}$ estimated mean of the postratified estimator
- *h* denotes stratification classes

- Poststratification will reduce non response bias if the distributions of ϕ or *Y* are less variable within post strata than across post strata
- Or if their covariance is attenuated within post strata
- A good choice for a post stratification variable would be a variable highly correlated with the response propensities such that response propensities were constant with each level
- The total nonresponse bias is the sum of bias across all strata

Nonresponse bias



 $bias(\bar{y^0}) = \bar{\phi}^{-1}\sigma_{\phi}\sigma_{Y}\rho_{\phi,Y}$

- $\overline{y^0}$ The unadjusted estimator of the respondents mean
- $\bar{\phi}^{-1}$ The population mean of response propensities
- σ_{ϕ} The standard deviation of response propensities
- + σ_Y The standard deviation of the response variable
- $\rho_{\phi,Y}$ The correlation between response propensity and the response variable

- Stochastic representation of bias.
- It assumes that response is a random variable and the probability of response is like the probability in an additional phase of sampling
- However the probability for every unit in this phase is unknown, thus has to be *estimated*.
- The estimated respondent mean is unbiased if $\rho = 0$.

Nonresponse bias of the total



 $bias(\hat{y}^{pst}) \approx \sum_{h} \bar{\phi}_{h}^{-1} \sum_{i} Y_{hi}(\phi_{hi} - \bar{\phi}_{h})$

- Different estimators have different expressions of bias.
- The total bias of the estimate is the sum across all strata
- Imputation allows us to have Y values for respondents and nonrespondents

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- The bias estimation based on the sample respondents will not equal the population bias (i.e., you need the whole population)
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- Types of data that can produce estimates of nonresponse bias
 - Sample frame data (i.e., where records were available both on respondents and nonrespondents)
 - Supplemental data for both respondents and nonrespondents, linked to the sample data.
 - Follow up studies of nonrespondents, comparing the earlier respondent group to those former respondents
 - Reports of intentions to respond to a later survey, comparing those who report agreeing to respond with those who decline to respond
 - Screener interview data

Imputation cells



- The same variables used for forming selection cells are used to form imputation cells (region, farm type and farm size)
- With some minor adjustments for merging small cells
- Farm size is an imputation variable which strongly correlates with key response variables.
- For each nonrespondent the values for all variables to be imputed are copied from the next available donor in the cell
- Each unit can only be used as a donor up to 6 times
- Unlinking may occur
- Key farms are not imputed via donor imputation but by using past information
- If groups are homogeneous imputation will work.



Bias across imputation cells – Dairy cows





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Bias across imputation cells – Beef cows







Thank you