

# **Testing for Heterogeneous Treatment Effects in Experimental Data: False Discovery Risks and Correction Procedures**

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# **Testing for Heterogeneous Treatment Effects in Experimental Data: False Discovery Risks and Correction Procedures**

## **Abstract**

We review the statistical models applied to test for heterogeneous treatment effects in the recent empirical literature, with a particular focus on data from randomized field experiments. We show that testing for heterogeneous treatment effects is highly common, and likely to result in a large number of false discoveries when conventional standard errors are applied. We demonstrate that applying correction procedures developed in the statistics literature can fully address this issue, and discuss the implications of multiple testing adjustments for power calculations and experimental design.

## I. INTRODUCTION

Multiple testing refers to any instance that involves the simultaneous testing of more than one hypothesis (Joseph P. Romano et al., 2010b). Even though economists have long been aware of the basic problem of conducting inference in the presence of multiple hypotheses, relatively little attention has been given to this issue in the empirical literature. In this paper we investigate one particularly common case of multiple testing in applied econometrics, the search for heterogeneous treatment effects within experimental data.

Few things have shaped empirical work on economics in the last decade as much as the arrival and establishment of randomized controlled trials (RCTs). The number of randomized experiments has grown rapidly across all continents over the past years. The projects listed on the Innovations for Poverty Actions (IPA) and the Abdul Latif Jameel Poverty Action Lab (JPAL) websites suggests that currently more than 100 RCT studies are either planned or in the field (JPAL, 2011); John List's "Field Experiments" website currently lists more than 250 field experiments (John A. List, 2011). One of the main advantages of experiments is the relative simplicity of the statistical analysis required to conduct causal inference. With properly done randomization, estimating causal effects corresponds to a simple conditional or unconditional mean comparison between treatment and control groups, with limited need or scope for more sophisticated empirical models.

Most experiments are designed to estimate the average effect of a given treatment of interest. However, researchers may become interested in the interactions of a treatment with some baseline characteristics of interest during or after data collection, and wish to test for heterogeneous treatment effects ex-post. In some cases, researchers may learn during fieldwork that the magnitude of the treatment effect hinges on a variable measured at baseline. In other cases, researchers may find that the theoretical framework applied has clear predictions regarding the expected behavioral changes across different subgroups of interest.

In order to provide some sense of how frequently papers test for heterogeneous treatment effects in the experimental, we review all articles using field experiment data published in the top 10

journals according to the 2009 Engemann and Wall ranking (2009) as well as the Journal of Development Economics (the top field journal) from 2005 and 2010. Out of 34 articles we classified as field-experiment-based, 26 (76%) estimate separate treatment effects for subgroups, and 10 articles (29%) estimate treatment effects for ten or more subgroups.

While testing for heterogeneous treatment effects through interaction terms or subgroup analyses is clearly desirable, applying traditional standard errors and p-values is not appropriate. Given that each interaction term represents a separate hypothesis beyond the original experimental design, trying out multiple interaction terms corresponds to multiple hypothesis testing, and results in a substantially increased false discovery risk in the empirical analysis.

To illustrate the severity of this issue, we use data from the Programa de Educación, Salud y Alimentación (PROGRESA) in Mexico, and run Monte-Carlo simulations to estimate a large number of heterogeneous treatment effects within the experimental data. We show that any researcher randomly choosing 10 baseline variables as proxy for an underlying characteristic of interest has a 62% chance of finding at least one variable significant at the 5% level. Given that the joint (Bernoulli) distribution for 10 independent binary variables implies a cumulative probabilities of finding at least one irrelevant factor significant with  $\alpha = 0.05$  is 40%, this implies that a majority of significant interactions uncovered in our PROGRESA regressions likely represent false discoveries. To provide a better sense of the study design implications of multiple testing, we compute ex-ante adjustments needed to sample size if researchers plan to investigate one or multiple interaction effects ex-post. We show that the required sample size adjustments are relatively small as long as the number of tested heterogeneous treatment effects is reasonably small.

The multiple hypothesis testing highlighted in this paper issue is not new, and has been faced by researchers in several other disciplines such as genomics or brain imaging. Possibly motivated by applications in these quickly evolving fields, recent statistical research has produced a number of powerful methods to correct for multiple hypothesis testing as summarized in Farcomeni (2008). While most conclusions of the paper apply to almost all empirical work, we have chosen this particular focus for two main reasons. First, heterogeneous treatment effects, even though they

are very common in empirical work, have not received much attention in the literature on multiple testing and are not even mentioned in the most recent review by Romano, Shaikh and Wolf (Joseph P. Romano et al., 2010a). Second, the design of field experiments allows researchers to consider heterogeneous treatment effects in the power calculations and thus to fully solve the problem of multiple testing prior to the collection of data.

The paper is related to the broader literature on heterogeneous treatment effects discussed in Angrist (2004), Green and Kern (2010) as well as Imai and Strauss (2011). While these papers primarily focus on optimal model specification in the presence of heterogeneous treatment effects, we mostly focus on the multiple-testing issue associated with sequential subgroup testing in this paper.

The rest of the paper is organized as follows: We start with a review of the methods used in recently published papers in the field in section 2. We discuss the theoretical and empirical distribution of heterogeneous treatment effects using PROGRESA data in Section 3. We introduce the corrections for multiple testing proposed by the statistics literature in Section 4, and analyze the practical implications of the various correction models in Section 5. In Section 6, we discuss the implications of multiple testing for study design and power calculations; section 7 concludes.

## **II. LITERATURE REVIEW**

Based on the Engemann and Wall (2009) ranking of Journals, we surveyed articles in the top 10 ranked journals as well as the Journal of Development Economics as the most commonly cited field journal. We use the classification proposed by Harrison and List (2004) as a guide to determine how to classify field experiments. We focus exclusively on “natural field experiments,” which Harrison and List describe as experiments where a “non-standard<sup>1</sup> subject pool” makes decisions where there is a “field context in either the commodity, task, or information set that the subjects can use” and “the environment is one where the subjects

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<sup>1</sup> In the context of experiments in Harrison and List (2004), a standard subject pool would be undergraduates recruited to perform an experiment in a laboratory setting

naturally undertake these tasks and where the subjects do not know that they are in an experiment.” (Glenn W. Harrison and John A. List, 2004, p. 1014)

In order to be considered for our literature review, a paper must present evidence from a study where a treatment intervention is randomly assigned by design of the study. Therefore, natural experiments and field experiments where treatments are not randomly assigned are excluded from our review.<sup>2</sup> Furthermore, we exclude papers that focus on econometric methods using data from natural field experiments. We refer to the papers that satisfy this definition as “strict” natural field experiments. While this classification may appear restrictive, it yields a clear decision rule allowing for a consistent review of the existing literature of interest. The main point made in this paper clearly applies to a larger set of empirical papers.

Table 1 shows the journal list, as well as the total number of articles, the number of “strict” natural field experiments.

TABLE 1  
“STRICT” NATURAL FIELD EXPERIMENTS PUBLISHED 2005-2009

Journal Rank	Journal Name	Total Articles	"Strict Natural Field Experiments"
1	Quarterly Journal of Economics	283	11
2	Journal of Political Economy	296	1
3	Econometrica	420	5
4	American Economic Review	644	8
5	Review of Economic Studies	292	0
6	Journal of Labor Economics	201	0
7	Journal of Economic Growth	87	0
8	Review of Economics and Statistics	456	1
9	Economic Journal	498	1
10	American Economic Review Papers and Proceedings	592	5
30	Journal of Development Economics	461	2
	Total	4230	34

<sup>2</sup> Examples include the analysis of data where subjects were quasi-randomly matched as in studies of choices in speed dating (Fisman et al, 2006), the impact of random roommate assignment and random assignment by lottery (Angrist, Bettenger and Kremer 2006).

Of 4,230 articles surveyed in these 11 journals over the period 2005-2009, 34 articles feature evidence from strict natural field experiments. Some of the reviewed journals did not publish any study based on a field experiment (Journal of Labor Economics and Journal of Economic Growth), while the Quarterly Journal of Economics published more than 10 studies based on experiments fitting our “strict” natural field experiment description over the same period.

In Table 2 we take a closer look at the econometric strategy employed by articles featuring evidence from strict natural field experiments. We do not show the full title of each paper, but rather show the paper’s index number, and provide the full listing as well as references in the Appendix. We report two measures of heterogeneous treatment effect testing: the number of subgroups for which treatment effects are estimated, and the number of interaction effects between the treatment variable and baseline characteristics tested.

In order to classify a reported result as a “heterogeneous treatment effect test” we followed a series of rules avoiding double-counting of treatment effects as well as incorrect classifications of regressions reflecting a particular study design. First, we only consider tests reported in the main tables of a paper, and exclude all results either shown in an appendix or only mentioned in the text. Second, we do not count reported tests reflecting the original research design. In many instances, researchers test for increasing effects over time, and also for heterogeneous treatment effects across different geographic sites. While one could argue that different time periods and sites reflect distinct sub-groups, we consider them as separate experiments, and thus do not count them as instances of heterogeneous treatment tests.<sup>3</sup> We also do not count heterogeneous treatment effect tests based on baseline characteristics that are measured after the experimental randomization, which are very uncommon but raise a different set of statistical concerns.

In some cases, papers combine subgroup analysis with estimates of interactions with treatments. We count the number of subgroups and interactions separately, and simply report the

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<sup>3</sup> While experiments may vary by time and location, most researchers do not consider differences across sites and over time to be a primary theoretical hypothesis generated by an economic experiment.

total number of subgroups and the total number of interactions. Furthermore, we only count each interaction or subgroup analysis once, even if they are considered for more than one dependent variable. Multiple dependent variable testing is very common in the literature as well, and the associated statistical problem similar to the heterogeneous treatment effects analyzed in this paper (Duflo, Glennester, et al 2008). The complications from considering dependent variables measured across time are discussed in recent work by McKenzie (2010).

Out of the 34 papers analyzed, 21 articles (62%) estimate separate treatment effects for subgroups, while 16 articles (47%) estimate interaction effects between the treatment and baseline characteristics. Only 8 articles (24%) neither estimate interaction effects nor consider the effect of treatment on subgroups. In some cases, testing for heterogeneous treatment effects is extensive: As Table 2 shows, 10 articles (29%) estimated 10 or more subgroup or interaction effects. Some examples of common interactions or subgroup analyses are sex, age, wealth and education. None of the article corrects (or mentions) multiple hypothesis testing in the empirical analysis.



TABLE 2  
STRUCTURE OF EMPIRICAL MODEL IN FIELD EXPERIMENTS

Article Index	Type(s) of Dependent Variable(s): Binary, Continuous or boTh	Number of Subsamples	Number of Interactions Estimated
1	T	2	0
2	B	7	0
3	C	0	4
4	T	11	0
5	T	16	11
6	T	4	0
7	T	0	0
8	C	0	0
9	C	0	0
10	T	0	1
11	T	2	4
12	C	21	10
13	B	0	0
14	B	5	3
15	C	0	3
16	T	15	0
17	B	0	1
18	B	0	0
19	T	6	1
20	C	2	0
21	C	0	0
22	C	7	0
23	C	10	0
24	C	60	2
25	C	3	11
26	C	0	0
27	T	1	1
28	C	7	0
29	B	4	14
30	B	23	0
31	T	2	2
32	C	0	2
33	C	9	6
34	B	0	0

### III. TESTING FOR HETEROGENEOUS TREATMENT EFFECTS

While most experiments are designed to investigate the average effects of a specific treatment of interest on a given outcome, researchers often may wish to investigate differences in the impact of the treatment by sex, ethnicity, income level, or other individual or household characteristics. In the simplest case, one may want to simply investigate one particular interaction of interest (possibly reflecting a particular model prediction or anecdotal evidence from the program rollout); in other instances, the researcher may simply be curious to see which factors modify an intervention's impact. Testing for heterogeneous treatment effects without adjusting the estimated standard errors for multiple testing after the fact, is highly likely to result in incorrect statistical inference. Given that 95% confidence intervals are constructed to allow for a false discovery probability of 0.05 on each interaction term, the probability of getting  $k$  significant  $p$ -values with zero true effects is given by the following binomial distribution:

$$(1) \quad f(k, m, \alpha) = \binom{m}{k} \alpha^k (1 - \alpha)^{(m-k)}.$$

Plugging in for  $\alpha = 0.05$  and  $m = 10$ , the probabilities of one, two, and more false discoveries with 10 different interaction terms are given by 31.5%, 7.5% and 1.2% as described in in Table 3 below.

TABLE 3  
THEORETICAL (BINOMIAL) DISTRIBUTION OF FALSE DISCOVERIES WITH 10 RANDOM INTERACTION  
TERMS AND ALPHA=0.05.

Event	Probability
No hypothesis significant	0.598
One hypothesis significant	0.315
Two hypothesis significant	0.075
Three or more hypotheses significant	0.012

The results shown in Table 3 are valid under the assumption that the interaction terms chosen for empirical analysis are orthogonal to each other. While this may be a reasonable assumption in the case where researchers simply wish to explore interactions with a series of baseline covariates such as sex, urban residence, ethnicity or religious affiliation, the assumption of independent interaction terms appears implausible in cases where researchers pursue a specific model prediction. Hypotheses regarding interaction effects are often generated by structural models developed by the researcher. Since the baseline characteristic of interest (say poverty) may not be easy to measure or directly available in the data, researchers may resort to trying out a series of proxies for the variable of interest, which can be presumed to be correlated. With correlated independent variables tested sequentially, the likelihood of finding one proxy significant will be positively associated with the likelihood of find other proxies used in the regressions significant, so that the basic binomial distribution underlying Table 3 can no longer be used to derive expected distributions of false discoveries.

While generalized false discovery distributions are hard to derive theoretically, it is relatively easy to generate false discovery distributions numerically. In order to illustrate this, we run a series of Monte-Carlo simulations under a range of dependence assumptions. Following the setup in Table 3, we assume that researchers conduct 10 independent tests within each experiment. In our baseline scenario, we assume that variables (interaction terms) are independent. As the first row of Table 4 shows, this yields exactly the distribution of false discoveries predicted by the binomial distribution (Table 3), with on average 40% of regressions showing at least one significant result.

To operationalize correlations among interaction terms, we assume that all included variables contain a common random variable component. One may think of the common random variable as representing the true variable of interest, and of the 10 independent variables as the proxy for this variable; alternatively, one may view the common component as a reflection of the more general dynamics between the variables included.

In order to provide a general sense of the induced changes in the distribution of false discoveries, we simulate a wide range of correlations, starting from a correlation of across proxies of 0.06

(Table 4, row 2) to a correlation of 0.8. As Table 4 shows, removing the independence assumption does indeed lower the false discovery risk. The reductions are, however, rather minor under most scenarios. Under the assumption that the 10 proxies display a correlation of 0.25, the false discovery risk drops from 40 to 36 percent. Even if one is willing to make the assumption that the average correlation is over 0.5, the false discovery risk remains at 26%, meaning that choosing interactions at random will lead to a more than one in four chance of finding at least one interaction significant at the 95% level. What is more concerning is that the risk of finding multiple false positives increases substantially with correlated interaction terms. While there is only a 1 in 100 chance of finding 3 or more false positives with 10 randomly chosen interaction terms, the risk of finding multiple false positives exceeds 5% with correlations over 0.5. Given that a large number of significant (and close to significant) interaction terms will likely convince even more skeptical researchers and reviewers, the chance of false discoveries being positively reviewed and published may actually be higher than lower with correlated interaction terms.

TABLE 4

EMPIRICAL DISTRIBUTION OF FALSE DISCOVERIES WITH CORRELATED INTERACTION TERMS AND ALPHA=0.05

Correlation with unobservable	Correlation of proxies	Probability of at least one false discovery	Probability of two false discoveries	Probability of three or more false discoveries
0.000	0.000	40.2	8.0	1.2
0.250	0.063	39.1	7.3	1.5
0.500	0.250	36.1	7.1	2.6
0.750	0.563	26.5	5.3	5.6
0.900	0.810	17.1	5.3	6.5

### Heterogeneous Treatment Effects in Practice: PROGRESA

In order to illustrate how the distribution of estimated heterogeneous treatment effects looks in practice, we randomly test for such effects within the experimental data collected as part of the the Programa de Educación, Salud y Alimentación (PROGRESA). PROGRESA, now called OPORTUNIDADES , is a community-level randomized experiment designed to increase school attendance among the poor through a conditional cash transfer program. By providing a cash

transfer to poor families large enough to compensate for lost wages from child labor<sup>4</sup> (Emmanuel Skoufias, 2005), the conditional cash transfer program was aiming at changing parental schooling decisions.

While PROGRESA's impact on schooling has been well documented (Paul Schultz, 2004), it seems natural to ask whether the program impact was contingent on, or mediated by, specific household characteristics of interest at baseline. One may, for example, conjecture that the program impact increases with measures of household poverty or vulnerability. The PROGRESA baseline data from the 1997 includes a large array of measures one could use as potential markers for poverty: size of the household, access to piped water and electricity, asset holdings, characteristics of the dwelling, household size and many more. Given the difficulties associated with correctly measuring the income and wealth level of the household, it appears plausible that an interested researcher would consider a larger set of measures, and we shall for simplicity assume that each researcher uses 10 proxies in his analysis. While this may appear high at first sight, 10 interaction terms appear fairly common in the literature: as our review shows, the average paper analyzes 6.4 subgroups and tests for 2.2 interaction effects. The researcher then estimates the following model:

$$(2) \quad y_i = \alpha + \beta T_i + \delta W_i + \lambda(T_i x W_i) + \varepsilon_i,$$

where  $y_i$  is the outcome of interest (in this case schooling),  $T$  is the treatment indicator (1 if the household was targeted by PROGRESA),  $W_i$  is one of the 10 poverty indicators coded, and  $T_i x W_i$  is the interaction between the poverty indicator and PROGRESA. While we focus on interaction-term-based empirical models in our simulations, it is easy to see that the results will look virtually the same if separate regressions were conducted for each subgroup of interest.

Given that virtually any baseline variable could be interpreted as a proxy for household poverty or vulnerability, we code all available baseline variables with non-zero variation within the treatment and control groups into binary variables. The total list of binary indicators (148 binary

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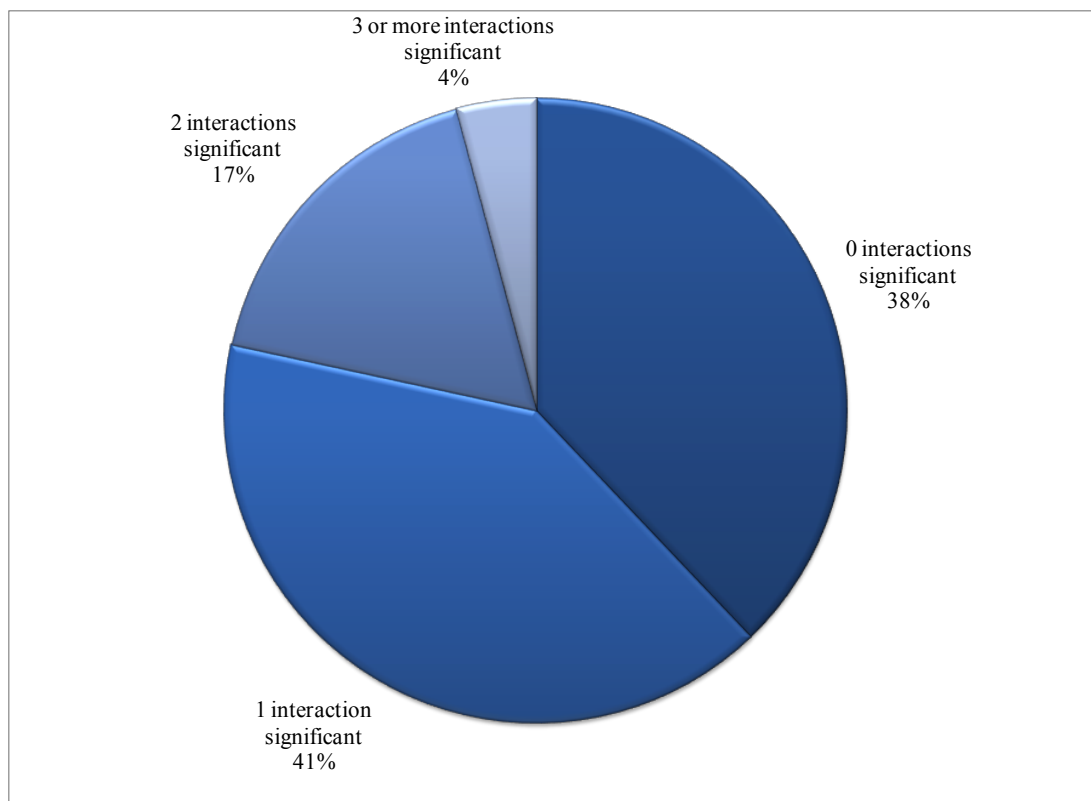
<sup>4</sup> A detailed description of the program as well as links to several evaluation studies are available at <http://www.ifpri.org/dataset/mexico-evaluation-progresa>.

variables) is shown in Appendix Table 2. We assume that each researcher randomly chooses 10 variables out of the set of 148, and runs 10 separate regressions as described in equation (1).<sup>5</sup> We run a Monte-Carlo simulation with 10,000 experiments, where we randomly chose a set of 10 out the 148 variables to interact with the treatment in each round. The Monte-Carlo simulation can thus be viewed as an approximation of a setting where a large number of independent researchers work on a given data set, and each of them subjectively chooses 10 variables as proxies for a specific variable of interest. We focus on binary indicators of heterogeneity which is common in the literature. The average correlation across variables is rather low, with a maximum correlation of 0.25, and an average correlation of 0.03 across the 10 proxies used in each simulation.

The results of the Monte-Carlo experiment are displayed in Figure 1. With 10 random binary regressors from the PROGRESA baseline data, more than 62% of cases (or independent researchers picking 10 interaction terms) find at least one interaction term significant at the 5% level; in 17% of all cases, 2 interaction terms are significant, and in 4% of cases, 3 or more interaction terms are significant.

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<sup>5</sup> For data collected to test a specific hypothesis, we might expect that the choices of our hypothetical “researchers” would be correlated instead of being independent. However, evaluating how groups respond differentially to social programs may not always be determined by specific theoretical hypotheses. Even when there are theoretical hypotheses, reasonable people might disagree about the best variables used to proxy for theoretically defined variables such as vulnerability.



*Notes:* Based on 10000 random block of size 11, consisting of the main treatment effect and ten randomly selected and independently tested interaction terms.

FIGURE 1

Empirical Distribution of Significant Coefficients at  $\alpha = 0.05$

Given that our proxies have relatively low correlation and that our Monte Carlo simulations of theoretical distributions suggest that we would expect at least one significant effect in about 40% of cases with zero correlation, we expect that over half of the statistically significant coefficients can be presumed to be false discoveries.

#### IV. STATISTICAL CORRECTIONS FOR MULTIPLE TESTING

Statisticians have been long aware of the problem of multiple hypothesis testing. For the purposes of this paper, we consider the frequentist approach to controlling for multiple testing.

The probability of the union of two events  $A_1$  and  $A_2$  is equal to the sum of the two probabilities  $P(A_1)$  and  $P(A_2)$  minus the probability of the intersect, i.e.

$$(3) \quad P(A_1 \cup A_2) = P(A_1) + P(A_2) - P(A_1 \cap A_2)$$

This means that the sum of  $P(A_1)$  and  $P(A_2)$  constitutes an upper bound of  $P(A_1 \cup A_2)$ . If  $A_1$  and  $A_2$  describe very similar events, this upper bound may be distant from the true probability; if  $A_1$  and  $A_2$  are nearly independent ( $P(A_1 \cap A_2) \approx 0$ ), this upper bound will be very close to the true probability. In the case of multiple events, equation (1) becomes a bit more complex, but the intuition remains exactly the same as in the two events case.

Based on this basic notion, Boole's inequality states that for a finite set of events  $A_1, \dots, A_m$  the probability of one event happening can never be greater than the sum of the probabilities of each individual event, i.e.

$$(4) \quad P\left(\bigcup_{i=1}^m A_i\right) \leq \sum_{i=1}^m P(A_i).$$

Building on this inequality, the Italian mathematician Carlo Emilio Bonferroni proposed a solution to the multiple testing problem. Assume we want to test  $m$  (dependent or independent) hypotheses at level  $\alpha$ . Boole's inequality<sup>6</sup> implies that at least one of the hypotheses comes out significant with probability less or equal to  $m\alpha$ . However, in order to keep the chance of false discoveries (Type I errors) at the desired level, we would like this upper bound to be  $\alpha$  and not  $m\alpha$ . Bonferroni showed that this can be achieved by testing each single hypothesis at the level  $\alpha' = \alpha / m$ . This is called the Bonferroni correction, designed to control the so-called familywise error rate (FWER).

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<sup>6</sup> Boole's inequality is sometimes also referred to as Bonferroni's inequality.



Duflo et al. (2008a) argue that Bonferroni type corrections may not be very useful in the context of economic field experiments, because the control of Type I errors might come at the cost of high Type II errors (less power). At standard 95% confidence intervals, testing for 10 effects simultaneously would require marginal p-values of 0.005 for *each* individual variable. Recent statistical and econometric research has produced a number of alternative methods to correct for multiple hypothesis testing, which are much more powerful than the simple Bonferroni method (Joseph P. Romano and Michael Wolf, 2005). We focus on frequentist methods here but recent work by Gelman, Hill and Yajima (2010) proposes Bayesian multilevel models which can address the problem of multiple comparisons and increase efficiency. Another possibility would be to consider bootstrap approaches to estimating uniform test critical values. Westfall and Young (1994) provide a re-sampling based procedure for correcting for multiple hypothesis testing.

Well known among natural scientists, but still not on the radar of most economists is the multiple testing approach introduced in a seminal paper by Benjamini and Hochberg (Y. Benjamini and Y. Hochberg, 1995a). Rather than focusing on the FWER, the authors define the false discovery rate (FDR) as the expectation of the false discovery proportion (FDP), i.e, the proportion of the rejected null hypothesis that are erroneously rejected. If all null hypotheses are true, the FDR is equivalent to the FWER. Further, if not all null hypotheses are true, it can be shown that any procedure that controls the FWER also controls the FDR. If a procedure controls the FDR only, a gain in power may be expected. The potential for increase is larger when more of the hypotheses are non-true.

Consider testing a set of hypotheses  $H_1, H_2, \dots, H_m$  based on the corresponding p-values  $p_1, p_2, \dots, p_m$ . Let  $p_{(1)} \leq p_{(2)} \leq \dots \leq p_{(m)}$  be the ordered p-values, and denote by  $H_{(i)}$  the null hypothesis corresponding to  $p_{(i)}$ . Let  $k$  be the largest  $i$  for which

$$(5) \quad p_{(i)} \leq \frac{i}{m} \alpha$$

Then reject all  $H_{(i)}$  with  $i = 1, 2, \dots, k$ . Benjamini and Hochberg (1995a) show that this procedure controls the FDR at  $\alpha$ ; independence of the test statistics is not needed for the proof.

It is easiest to illustrate the differences between the FWER and the FDR approaches with an example. Consider an experiment with one treatment, but 10 different dependent variables. The ordered p-values on each of the 10 estimated coefficients look as follows:

$$p_{(1)} = 0.001, p_{(2)} = 0.004, p_{(3)} = 0.006, p_{(4)} = 0.008, p_{(5)} = 0.010$$

$$p_{(6)} = 0.040, p_{(7)} = 0.050, p_{(8)} = 0.060, p_{(9)} = 0.100, p_{(10)} = 0.400$$

Without any adjustment for multiple testing we reject seven of the ten hypotheses at a 5 percent level of significance. The Bonferroni-adjustment requires a p-value of  $\alpha' = \frac{0.05}{10} = 0.005$ , which means that only two out of the ten hypotheses get rejected. With the Benjamini and Hochberg method we check the condition

$$(6) \quad p_{(i)} \leq \frac{i}{10} \alpha.$$

sequentially starting with  $i = 10$ . The first p-value to satisfy the condition is  $p_{(5)}$  with  $0.01 < 0.025$ ; it is straightforward to see that the condition is also satisfied for any  $i < 5$ , so that the FDR adjustment leads to a rejection of 5 out of the 10 tested hypotheses.

Genovese and Wasserman (2006) show that the Benjamini and Hochberg method is optimal in the sense that it minimizes the false non-discovery rate (FNR) subject to the constraint that the FDR is controlled at level  $\alpha$ , where the FNR is defined as the expectation of the proportion of non-rejections that are incorrect. In other words, the Benjamini and Hochberg method keeps the number of type II errors as small as possible, i.e. the chance of not rejecting a hypothesis when it is false. While this correction is generally less conservative than the methods based on FWER

(which continue to be most frequently used), it is also subject to the criticism of the independent test assumptions. While correction models under more general or arbitrary dependence structures would clearly be desirable, such approaches have not yet been developed (Joseph P. Romano and Michael Wolf, 2005).

## V. CORRECTING FOR MULTIPLE HYPOTHESES TESTS IN PRACTICE

There are two important aspects to consider when it comes to the applicability of these correction procedures: 1) the technical knowledge required for implementation; 2) the statistical and empirical consequences in terms of type I and type II errors. The first aspect is fortunately straightforward. Thanks to the *multproc* package available as a user-written add-on in Stata<sup>®</sup> (Roger Newson, 2003), both the FDR and FWER methods are easily implemented in practice. The *multproc* package takes p-values from a set of variables (from single or multiple regressions) as inputs, and calculates corrected critical p-values for a range of correction procedures. The FWER correction is intuitive, as critical p-values are simply divided by the number of hypotheses tested ( $m$ ). In the case of single control variable and one interaction term, this implies that the critical p-value for significance at the 95% level shifts from 0.05 to  $0.05/2 = 0.025$ . It is easy to see that this adjustment keeps the likelihood of a false discovery at the desired low level independent of the number of hypotheses tested.

The FDR adjustment is slightly more complex since it is taking true discoveries into account, and, as a result, adjusts the p-values to a lesser extent than the FWER method. To see how well these adjustments work, we show the implications of the adjustment with truly independent variables (theoretical binomial) in a first step, and then revisit the PROGRESA results presented in Section 3.

Given that the FDR deviates from the FWER correction only if at least one hypothesis is false, it is easy to see that the two corrections have virtually the same effect if we assume 10 independent

interactions without true effect. With  $\alpha = 0.05$  and  $m = 10$  the adjusted p-value under both correction models can be approximated by<sup>7</sup>

$$(7) \quad p_{adj} = \frac{\alpha}{m} = \frac{0.05}{10} = 0.005$$

With the adjusted probability of 0.005, we can get the joint distribution within blocks of 10 by plugging the adjusted p-value into the corresponding binomial distribution. As the results displayed in Table 4 show, the chance of false discoveries is indeed reduced to just below 5%.

TABLE 4  
THEORETICAL (BINOMIAL) DISTRIBUTION OF FALSE DISCOVERIES WITH 10 RANDOM INTERACTION  
TERMS AND ALPHA=0.005

Event	Probability
No hypothesis significant	0.951
One hypothesis significant	0.048
Two or more hypothesis significant	0.010

The interpretation of the FDR differs from the typical statistical testing done by economists. The FDR correction is probabilistic, meaning that with a critical value of 0.05, we expect 5% of our tested hypotheses to be significant by chance. However, since the number of hypotheses generated by most experiments is relatively small, we will end up concluding that we expect some fraction of a hypothesis to be rejected erroneously, which can be difficult to interpret conceptually. Nonetheless, we would argue that the FDR correction allows for an appropriate level of caution in interpreting the results of testing many hypotheses, without becoming so conservative that we can no longer draw important conclusions about heterogeneity in responses to treatment.

<sup>7</sup> Technically, the FDR calculates separate p-values for each hypothesis. The adjusted FWER p-value corresponds to the p-value for the variable with the lowest p-value. The p-value of the second variable would be 0.01, the third 0.015 and so on. In practice the first one will already rule out 95% such that the result is very similar to FWER, but it is not the same.

To provide a better sense of how powerful these corrections are in practice, we show the PROGRESA results displayed in Figure 1 with corrected p-values in Figure 2 below. The assumption underlying the correction is that within each experiment we test 11 hypotheses, the main treatment variable plus 10 randomly selected interaction terms. As Figure 2 shows, neither correction affects the significance of the main treatment effect, which is significant in all cases with both corrections. Large differences emerge, however, with respect to the interaction terms. While we see at least one significant result in 62% of the specifications if no correction is applied, the likelihood of finding a statistically significant results drops by 60% (FDR) and 70% (FWER), respectively, after the correction is applied. This, however, does not mean that researchers applying either correction will never find significant results – as Figure 2 clearly illustrates, the chance of finding one or more significant results in a Table showing 10 interaction terms in the PROGRESA sample is 18% with the Bonferroni FWER correction, and 24% with the Benjamini-Hochberg FDR correction.

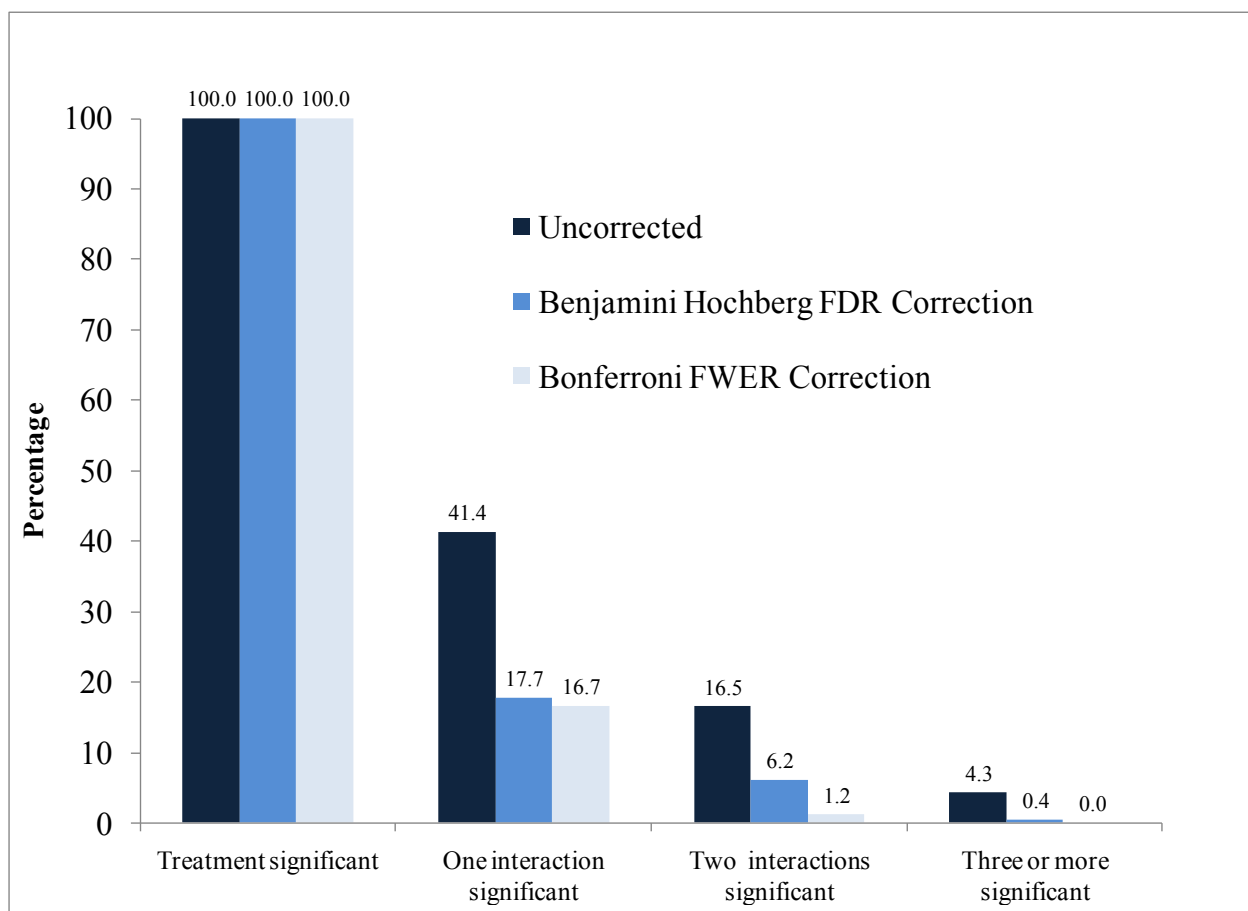


FIGURE 2

## VI. IMPLICATIONS FOR STUDY DESIGN AND POWER CALCULATIONS

Given that interaction terms may often be of major importance of researchers designing an experiment, one of the key questions is how much of an adjustment is needed to sample size ex-ante if the researcher plans to test for interaction effects ex-post. To understand what the two corrections imply in terms of power, we show a set of numerical simulations in this section. As a first step, we assume a sample size of 2000, with a corresponding (unadjusted) power of 0.5 and investigate how much power is lost in expected terms once we adjust for multiple hypotheses testing. As Figure 3 illustrates, the power drops in a non-linear fashion from 0.35 to about 0.2 for the FDR approach; as expected the drop is larger for the FWER approach, where the power drops to 0.1 if 10 hypotheses are tested simultaneously.

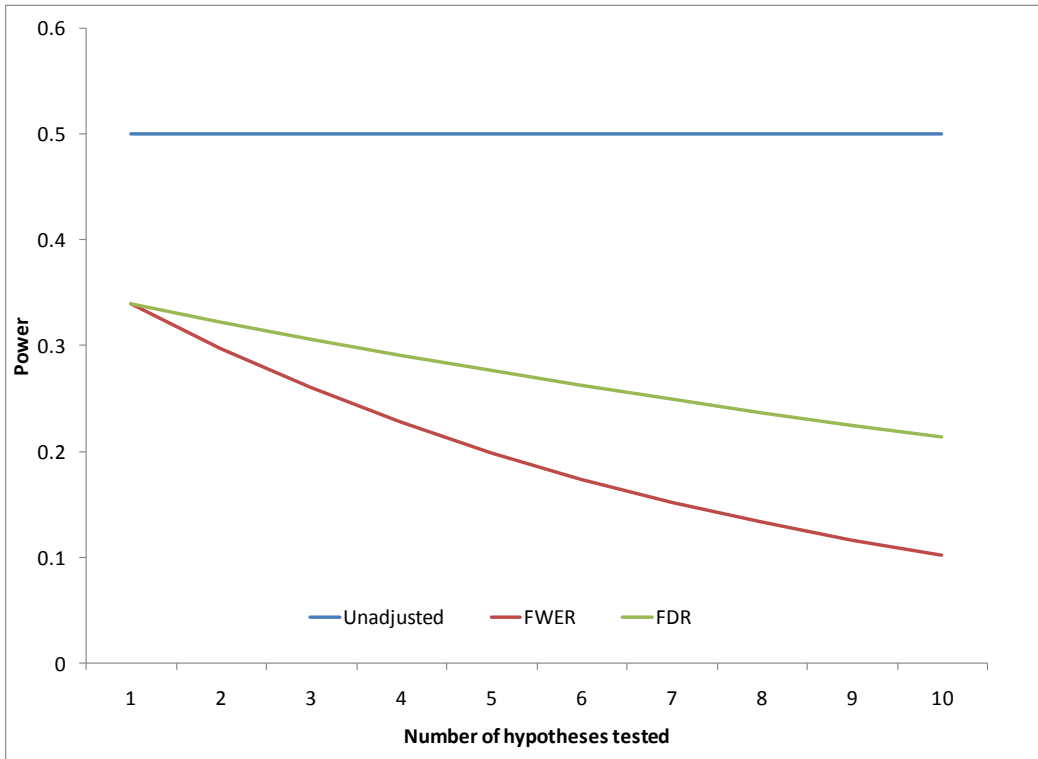


FIGURE 3  
Power with Corrections

While these losses in terms of power may dissuade researchers from applying these corrections in practice, the necessary sample adjustment may not be as large as one may think (or fear) as long as the number of interactions the researcher is interested in is reasonably small.

Figures 4 illustrate this point for one and two control variables and their interaction terms, respectively. As pointed out before, standard power calculations do not apply here as two separate treatment groups lower effective group sizes, and increase estimated standard errors. As Figure 6 shows, the power of the study with sample size 5000 and a treatment effect of 0.05 is 0.9 if no interaction term is included. With the interaction term, the power drops to about 0.7. The drop in power due to the multiple testing corrections is comparable in magnitude. With a sample of 5000, the power with one interaction term drops to about 0.5, while the power with two interaction terms is about 0.45.

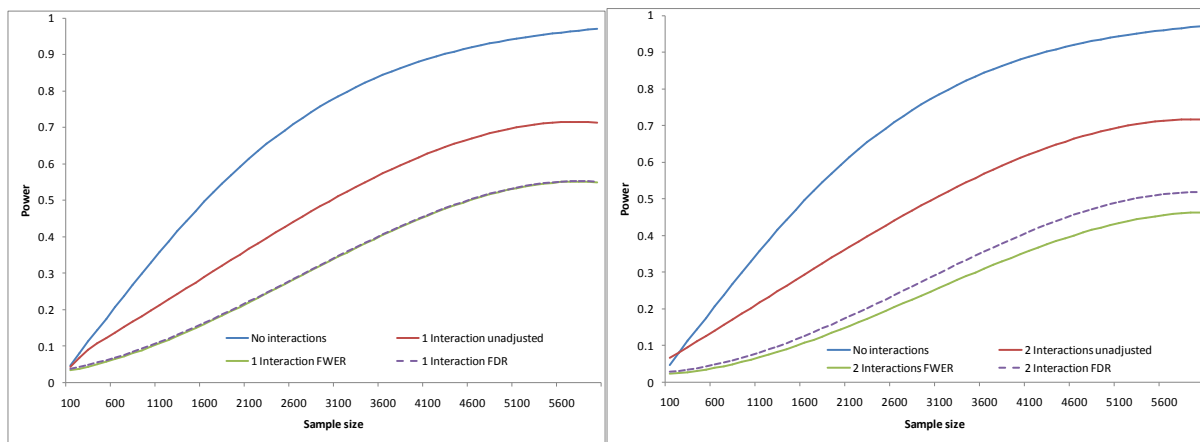


FIGURE 4

Power with one and two Interaction Terms

To see the implications for study design, we show necessary samples with and without corrections for multiple testing in Figure 5 below. As the figure shows, the absolute differences in sample size are rather small for large effect sizes; in relative terms, doing the FWER adjustment implies an average increase in necessary sample size (assumed power is 0.9) of about 28 percent with one interaction, 55% with two interactions, and about 67% with three

interactions.<sup>8</sup> These adjustments are not trivial and may appear overly conservative. As discussed earlier, the FWER adjustment reduces the risk of false discoveries under the most conservative assumption of independence across events. Smaller sample size adjustments could in theory be generated by using the FDR approach and by relaxing the assumption regarding event independence; however, this would require imposing a large set of additional distributional assumptions researchers will struggle to make during early stages of field experiments. From a pragmatic perspective, it seems best to base initial sample size calculations on FWER adjusted standard errors. The FWER adjustment will keep the risk of false discoveries at the desired low levels and guarantee sufficient power for either FWER or FDR standard error adjustments ex-post.

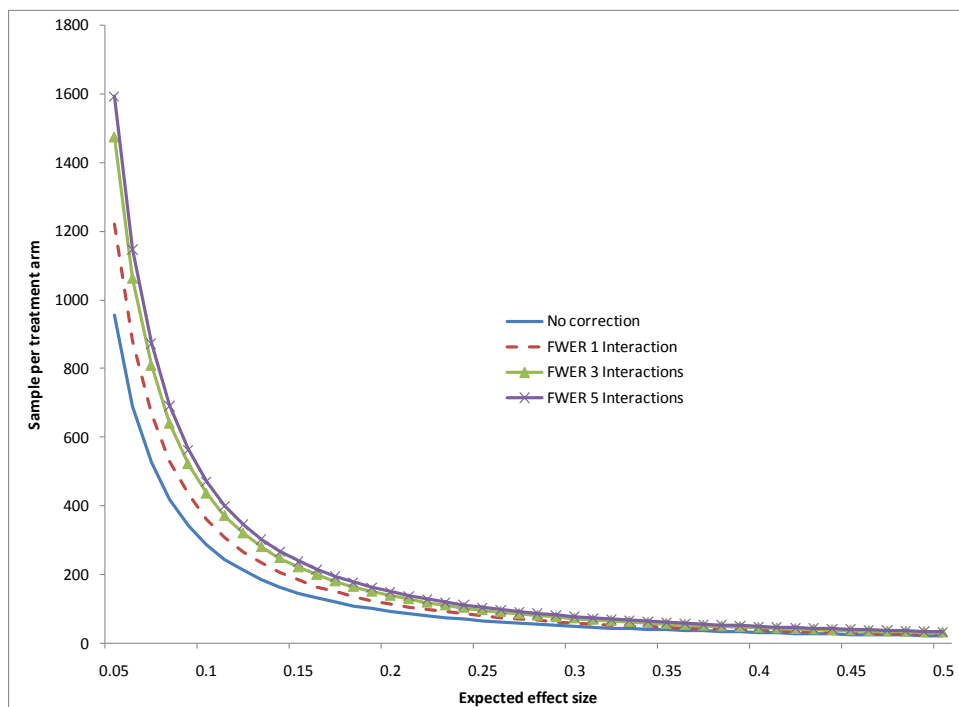


FIGURE 5  
Absolute Sample Size with and without FWER Correction

<sup>8</sup> We assume that each interaction term is tested separately, so that each regression yields 3 coefficients (covariate, treatment and covariate\*treatment) and 3 p-values that need to be adjusted.



## VII. DISCUSSION AND CONCLUSIONS

Even though the testing of heterogeneous treatment effects is very common in current empirical work based on experimental data, multiple testing corrections are generally not applied. In this paper we demonstrate that standard statistical inference is not valid when multiple heterogeneous treatment effects are tested, and that ignoring this issue is likely to generate a large number of false discoveries. Without any true effect, the likelihood of finding at least one result significant at the 5% level is 40% with 10 uncorrelated interaction effects. In the PROGRESA example, we find that more than half of the results significant at the 5% level can be assumed to reflect false discoveries. This risk appears high and, more importantly, unnecessary given the readily available correction models developed in the statistics literature.

There are three main critiques of the correction procedures discussed here. The first concern regards the actual reporting of statistical tests conducted. Even if proper adjustments are applied to the final set of interaction terms reported, the underlying variable selection is unobservable ex-post, and may itself be the result of pre-testing. One possible approach to address this issue might be a central registration system similar to the ones used in medical trials. In fact, Duflo et al (2008a) suggest that granting agencies create such a database of projects and their ex-ante designs.<sup>9</sup>

The second concern relates to the definition of what counts as a distinct hypothesis. Given that many variables may be used as proxies for a specific factor of interest such as income or human capital, it may be tempting to argue that all interacted variables are related, and thus reflect one single hypothesis. However, given that the correlation between any two proxies of interest is small in most cases empirically,<sup>10</sup> treating multiple measures of a specific factor of interest as single hypothesis appears not advisable from a statistical perspective. Furthermore, our analysis shows that correlation of interaction terms does not solve the problem of false discoveries and may in fact make it worse in some cases.

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<sup>9</sup> Some researchers have begun to publish their analysis prior to conducting experiments. Another potential alternative is the use of interdisciplinary system where researchers in any field can currently post their designs prior to conducting experiments, such as <http://clinicaltrials.gov/>.

<sup>10</sup> In the PROGRESA data, the highest correlation between any two indicator variables is 0.27.

Finally and maybe most importantly, there is the concern that applying more stringent standard errors increases the chance of type-II errors, i.e. the chance of not rejecting a hypothesis when it is false. While adjusting p-values clearly comes with some loss of power, we have shown in this paper that the cost in terms of additional sample size required for researchers planning to test for heterogeneous treatment effects ex-post appears well worth the benefit in terms of reduced false discovery risk.

Overall, a wider application of multiple testing procedures in the economics literature appears highly desirable. Testing for heterogeneous treatment effects is of obvious interest to researchers, and neither can, nor should, be avoided in practice. The resulting risk of false discoveries is high, but can be reduced to a minimum if correction procedures are applied.

## Appendix Table 1: Index of Strict Natural Experiments

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**Appendix Table 2: Binary Variables Used in PROGRESA Regressions**

1 bathroom	38 head_nospanish	75 lost_limb	112 rooftype4
2 bathroom_water	39 head_o60	76 mental	113 rooftype5
3 bedrooms1	40 head_perm_unable	77 migrant	114 rooftype6
4 bedrooms2	41 head_preparatoria	78 needs_help_to_move	115 rooftype7
5 bedrooms3	42 head_primary	79 no_waterelectric	116 rooftype8
6 bedrooms4	43 head_primary_income	80 noincome	117 rooftype9
7 bedrooms5	44 head_profesional	81 not_childofhead	118 second_income
8 bedrooms6plus	45 head_retired	82 one_child	119 shared_building
9 blender	46 head_secondary	83 owns_agri_land	120 spouse_away
10 blind	47 head_single	84 owns_animals	121 stove
11 budget_control_head	48 head_single_female	85 owns_cattle	122 three_children
12 budget_control_other	49 head_socialsecurity	86 owns_chicken	123 treatment_dif
13 budget_control_shared	50 head_temp_unable	87 owns_donkey	124 treatment_hospital
14 budget_control_spouse	51 head_u20	88 owns_goatsorsheeps	125 treatment_imss
15 car	52 head_widowed	89 owns_horse	126 treatment_issste
16 cd	53 head_working	90 owns_land	127 treatment_othergov
17 deaf	54 hhsizel0plus	91 owns_multiple_pieces	128 treatment_ssa
18 decision_head	55 hhsizel2	92 owns_ox	129 truck
19 dumb	56 hhsizel3	93 owns_pigs	130 tv
20 electric_lights	57 hhsizel4	94 owns_rabbits	131 two_children
21 fan	58 hhsizel5	95 piped_inside	132 video
22 father_athome	59 hhsizel6	96 piped_water	133 walltype1
23 female	60 hhsizel7	97 radio	134 walltype10
24 five_or_more_children	61 hhsizel8	98 receives_apoyoINI	135 walltype11
25 floortype1	62 hhsizel9	99 receives_becapapicacion	136 walltype12
26 floortype2	63 house_paid	100 receives_desayuno_escolar	137 walltype13
27 floortype3	64 house_paying	101 receives_despensa_DIF	138 walltype14
28 floortype4	65 house_provided	102 receives_empleotemporal	139 walltype15
29 four_children	66 house_rented	103 receives_leche	140 walltype2
30 fridge	67 inshool_97	104 receives_ninosdesolid	141 walltype3
31 head_2060	68 kids_medical_head	105 receives_tortilla	142 walltype4
32 head_basica	69 kids_medical_other	106 rooftype1	143 walltype5
33 head_dialect	70 kids_medical_shared	107 rooftype10	144 walltype6
34 head_female	71 kids_medical_spouse	108 rooftype11	145 walltype7
35 head_literate	72 Laundry	109 rooftype12	146 walltype8
36 head_married	73 light_meter	110 rooftype2	147 walltype9
37 head_noschool	74 Literate	111 rooftype3	148 water_heater

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