

# AN APPLICATION OF PROPENSITY- SCORE MATCHING: THE EFFECT OF CHILDBEARING ON OBESITY RISK

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# Motivation

- To what extent does childbearing contribute to obesity prevalence in women?
- Importance:
  - **Societal:** Ethnic and SES-based disparities in obesity
  - **Public health burden:** To what extent do child-bearing patterns contribute to rising obesity prevalence?
  - **Individual:** Women's decision-making and expectations

# Methodological challenges in previous studies

- **Study design 1: In post-menopausal women, compare parous vs non-parous**
  - Minority of childless women different from other women
  - Timing: Did wt gain precede births – or occur long after?
  - Generalizability: elderly target population, births in 1940s-1970s
- **Study design 2: Compare post-pregnancy vs pre-pregnancy weight**
  - Does not account for existing trajectory of weight gain
  - Generalizability: births in 1970s, 1980s; not population-based
- **Study design 3: Compare weight gain or obesity incidence in parous versus non-parous women**
  - Parous women are different from non-parous women
  - Generalizability: Births in 1970s, 1980s; not population-based
  - Possible selection bias: lots of exclusions

# Methodological challenges in previous studies

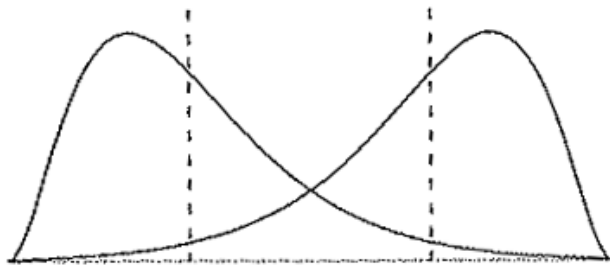
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# Why propensity scores?

## Address confounding

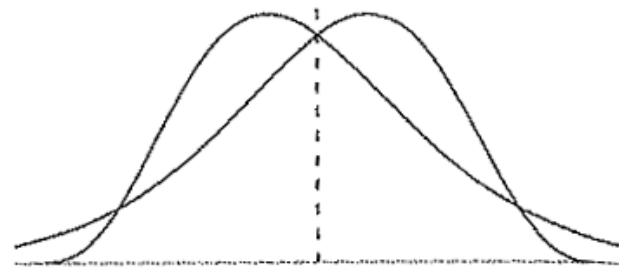
- Lack of comparability between exposed and unexposed, “exchangeability”
- Two aspects of lack of comparability
  - Imbalance – **uncontrolled confounding**
  - Lack of overlap (“positivity”) – **extrapolation**
    - indicates possible uncontrolled confounding, no data on which to evaluate

# How model-dependent are our inferences?



x

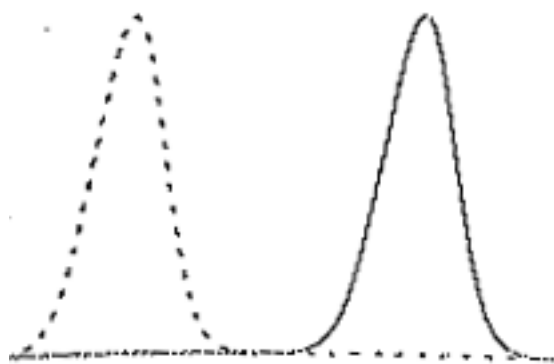
**Severe imbalance, good overlap**



x

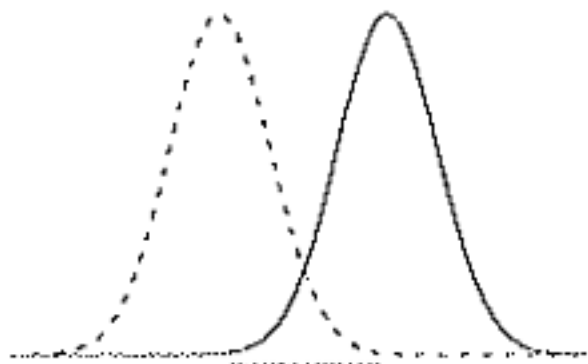
**Slight imbalance, good overlap**

**Severe imbalance, no overlap**

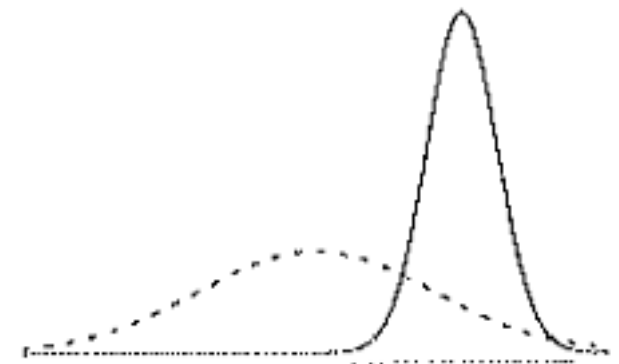


x

**Moderate imbalance, partial overlap**



x



x

From Gelman and Hill, 2003

# Study Design

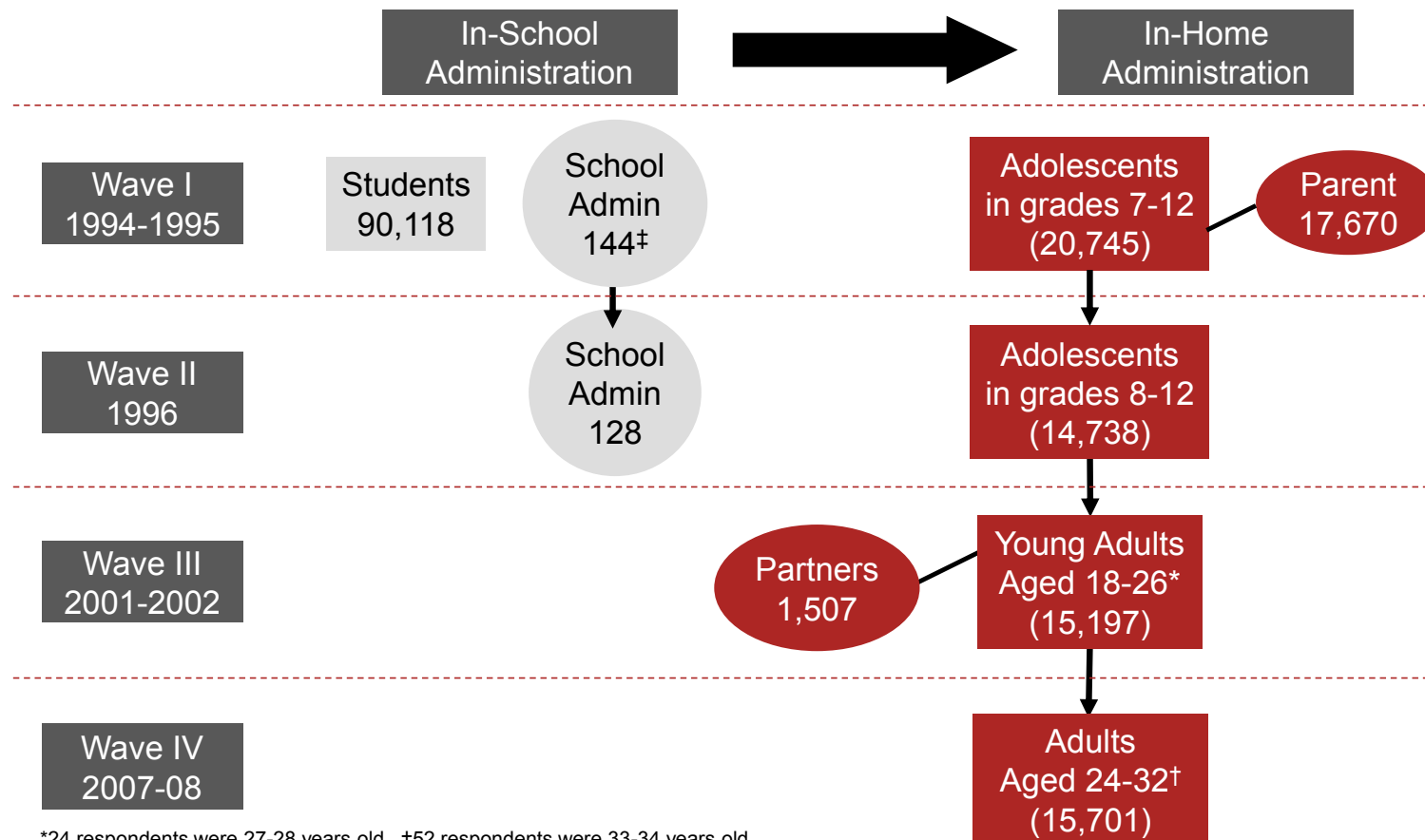
- Goals:
  - Achieve generalizability
  - Prevent selection bias
  - Improve exchangeability (achieve balance, test for non-overlap)
- Design: Prospective longitudinal cohort comparing parous and non-parous women
  - **Data:** Population-representative contemporary population of child-bearing women
  - **Restriction & variable selection:** Limit exclusions & induced bias
  - **Analysis:** Women in prime child-bearing years & propensity-score matching

# Data: Add Health



*Social, Behavioral, and Biological Linkages Across the Life Course*

## Longitudinal Design



\*24 respondents were 27-28 years old. †52 respondents were 33-34 years old.

‡ 144 schools participated in in school administration. School administration questionnaires from 143 of these schools



# Implementation of p-score matching

- **Exposure:** parous/non-parous, wave 4
- **Outcome:** obesity ( $\text{BMI} \geq 30.0 \text{ kg/m}^2$ ) at wave 4
- Step 1: Logistic regression to predict “propensity” to parity (exposure)
- Step 2: Assign each respondent a p-score,  $\text{Pr}(\text{parous})$
- Step 3: Match each parous women to 1+ non-parous women (ATT)
  - **Stata's psmatch2** (January 2012)
    - How good a match is necessary? calipers: 0.1 sd of p-score
    - Boot-strapping to get 95% CIs, 100 iterations

E. Leuven and B. Sianesi. (2003). "PSMATCH2: Stata module to perform full Mahalanobis and propensity score matching, common support graphing, and covariate imbalance testing". <http://ideas.repec.org/c/boc/bocode/s432001.html>.

```
-----  
help for psmatch2  
-----
```

### Mahalanobis and Propensity score Matching

```
psmatch2 depvar [indepvars] [if exp] [in range] [, outcome(varlist)  
pscore(varname) neighbor(integer) ai(integer) radius  
caliper(real) mahalanobis(varlist) kernel llr  
kerneltype(type) bwidth(real) spline nknots(integer)  
common trim(real) noreplacement descending odds index  
logit ties quietly w(matrix) ate]
```

### Description

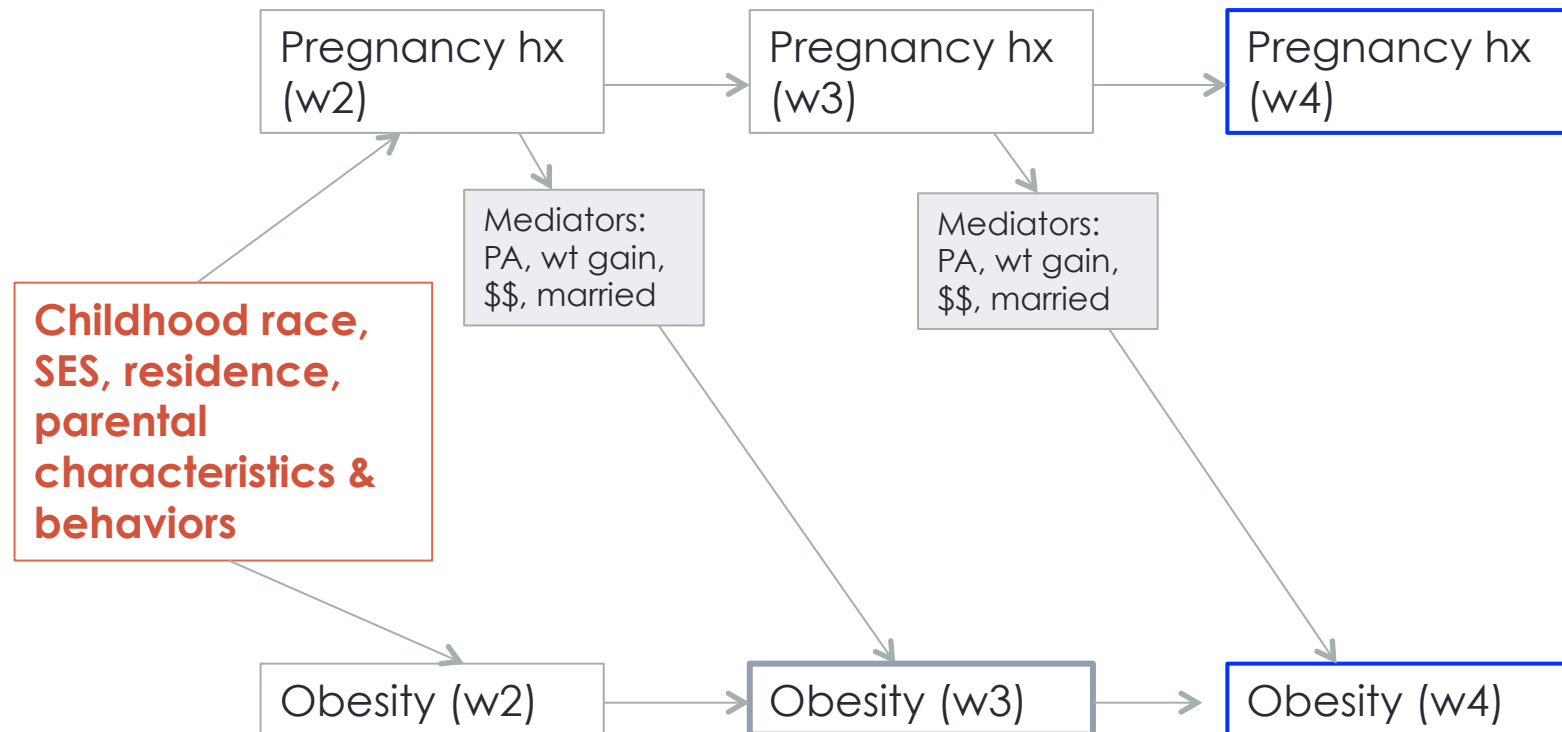
**psmatch2** implements full Mahalanobis matching and a variety of propensity score matching methods to adjust for pre-treatment observable differences between a group of treated and a group of untreated. Treatment status is identified by `depvar==1` for the treated and `depvar==0` for the untreated observations.

**psmatch2** is being continuously improved and developed. Make sure to keep your version up-to-date as follows

```
. ssc install psmatch2, replace
```

By default **psmatch2** calculates approximate standard errors on the

# Choosing matching algorithm



**Predictor variables:** Unique ID of school attended at the first survey, age, age<sup>2</sup>, US region (S, NE, MW, W), urbanicity, regionxurbanicity, parental education, Black race, Blackxparental education, immigrant, immigrant black, mexican, cuban, Puerto Rican, Central Am, Other Hispanic, Hispanicximmigrant, chinese, filipino, Vietnamese, etc.,

# Creating matching variable, logit1a

qui logistic parous

```
age_yr_w4 ageyrsq_w4 region1 region2 region4
rural suburb
ruralxreg1 ruralxreg2 ruralxreg4
suburbxreg1 suburbxreg2 suburbxreg4
highedcat1 highedcat2 highedcat3 highedcat4 highedcat6
black nonbwrace
blackxhighed1 blackxhighed2 blackxhighed3 blackxhighed4 blackxhighed6
usborn gennonblwh genblack
mexam cuban puertorican centrsoutham otherhisp hispmix
genmexam gencuban genpuertorican gencentrsoutham genotherhisp
chinese filipino japan asiaindn korean vietnam asianoth asianmix
```

**i.scid\_n**

```
if w4_selectf==1, coef;
```

```
*****
```

```
* CONSTRUCTING FINAL SAMPLE
```

```
• childhood to w4
```

```
*****.
```

```
predict pr1a_w4 if w4_selectf==1;
```

```
gen logit1a=log(pr1a_w4/(1-pr1a_w4));
```

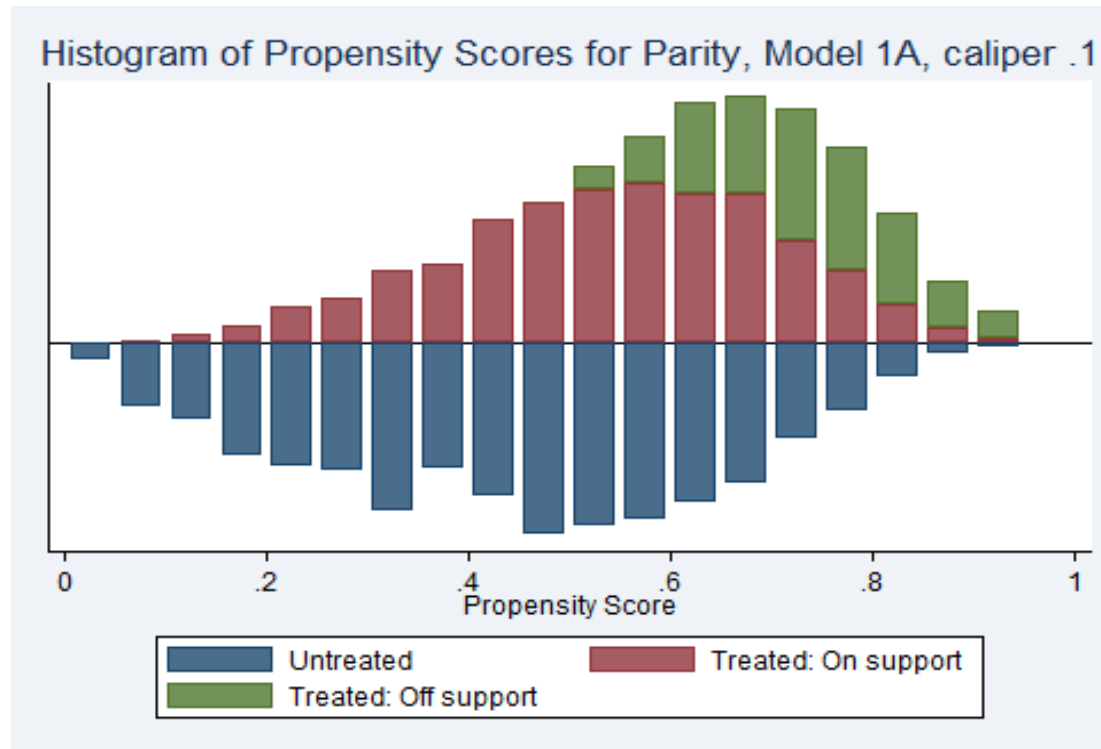
# Lack of comparability

Descriptive characteristics by parity in unmatched sample

	Parous	Non-Parous	Total
% (N)	52.3 (3593)	47.7 (3186)	100 (6779)
<b>Region (adolescent residence)</b>			
South	44.1%	32.0%	38.3%
Midwest	32.1%	30.9%	31.5%
West	14.2%	18.9%	16.5%
Northeast	9.7%	18.2%	13.7%
<b>Mother's education</b>			
< HS	23.6%	12.8%	18.5%
> college	3.6%	10.1%	6.7%

% are weighted for complex survey sampling and non-response

# Distribution of p-scores by parity



Slight imbalance, good overlap

```
psgraph, p(pr1a_w4) saving (hist_1log_1-1.gph,replace)  
title("Histogram: Model 1A, caliper .1logit, no replace");
```

# Covariate balance before & after matching

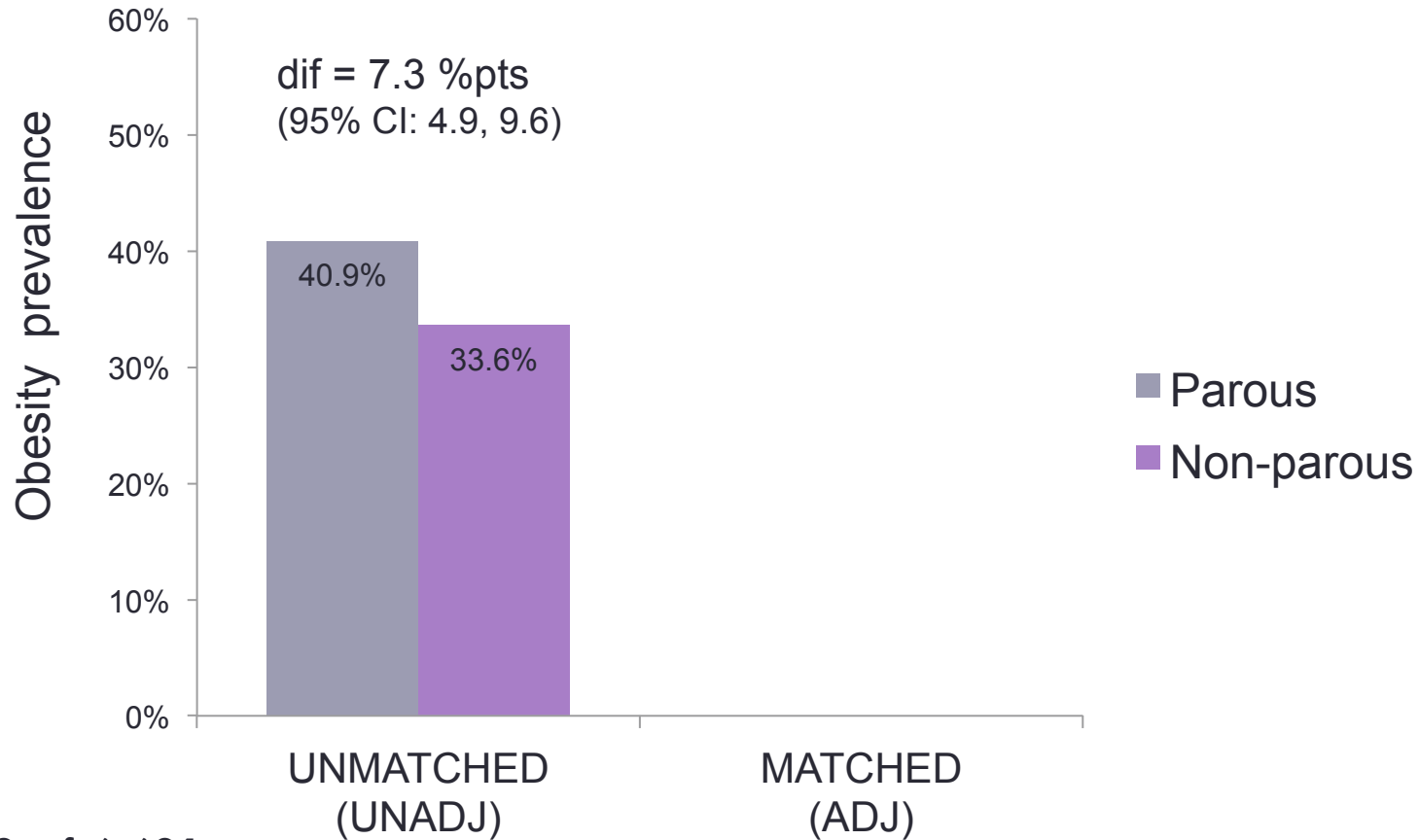
Variable	Sample	Parous	Non-parous	% bias	% reduction  bias
<b>Age (years), w4</b>					
	Unmatched	28.7	28.1	34	
	Matched	28.5	28.5	-2	94.2
<b>US REGION, w1 (ref=SOUTH)</b>					
West	Unmatched	20.8%	24.8%	-10	
	Matched	22.8%	25.3%	-6	40
Midwest	Unmatched	24.8%	26.1%	-3	
	Matched	23.5%	26.1%	-1	84
Northeast	Unmatched	10.8%	16.2%	-16	
	Matched	13.2%	11.6%	4.5	72
<b>URBANICITY, w1 (ref=urban)</b>					
Rural	Unmatched	19.7%	14.2%	15	
	Matched	16.5%	18.4%	-5	66
suburb	Unmatched	52.8%	53.7%	-2	
	Matched	52.9%	53.7%	-2	16.6
<b>PARENTAL EDUCATION</b>					
<high school	Unmatched	14.6%	7.7%	22	
	Matched	10.6%	10.8%	-1	97.2
vocational degree	Unmatched	8.8%	8.4%	1	
	Matched	10.3%	9.9%	1	-4
<b>RACE &amp; ETHNICITY</b>					
non-H BLACK	Unmatched	25.7%	21.8%	9	
	Matched	24.2%	24.5%	-1	93.1
non-H CHINESE	Unmatched	0.3%	2.3%	-18	
	Matched	0.4%	0.1%	3	81.4

pstest <varlist>

% do not account for complex survey design or non-response

# % Obesity in Parous vs Non-parous

Demographic match, 1:1 neighbor, no replacement



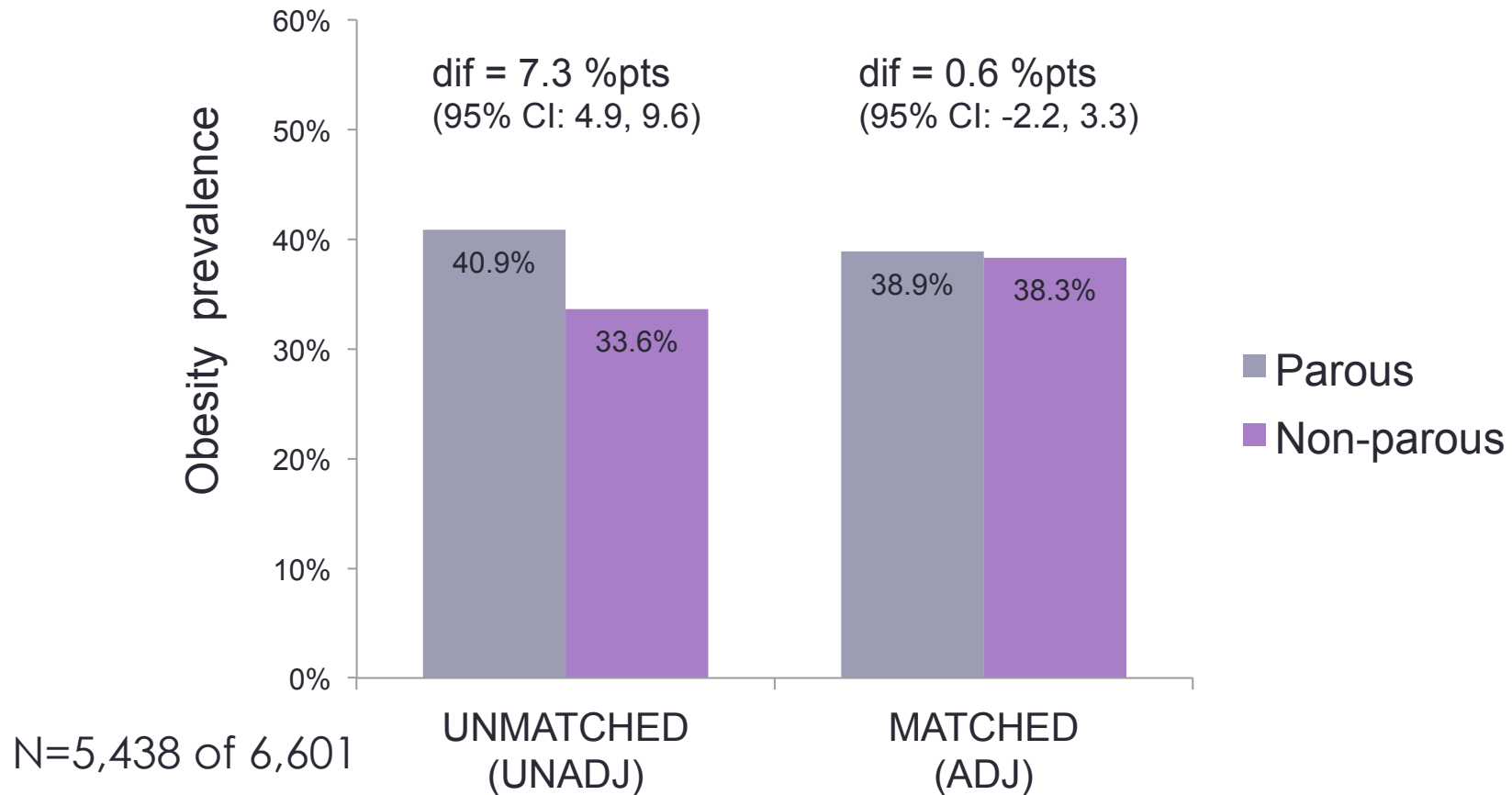
N=5,438 of 6,601

```
psmatch2 parous, outcome(ob_constr_w4) pscore(logit1a) caliper(.1)  
com noreplacement descending;
```



# % Obesity in Parous vs Non-parous

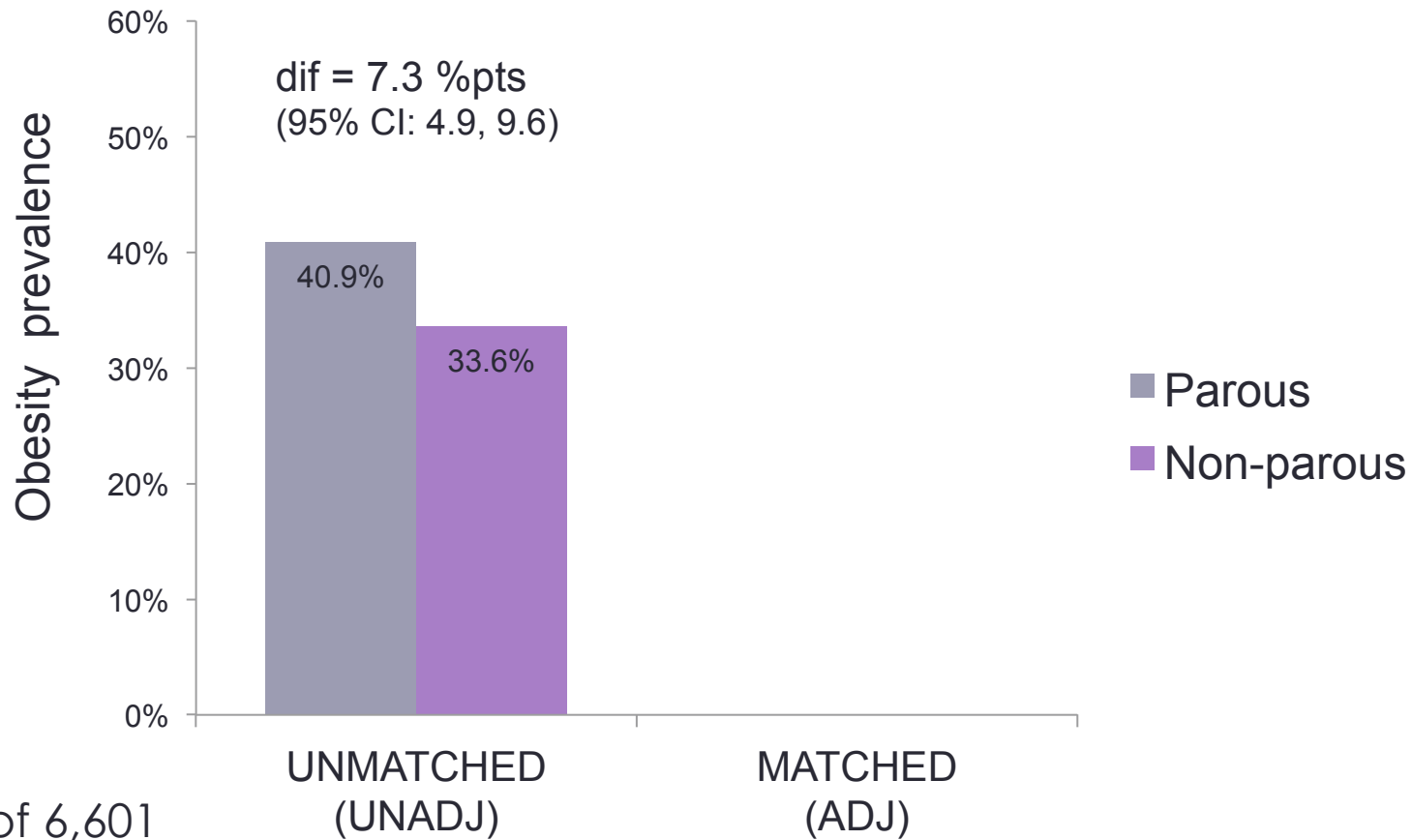
Demographic match, 1:1 neighbor, no replacement



```
psmatch2 parous, outcome(ob_constr_w4) pscore(logit1a) caliper(.1)  
com noreplacement descending;
```

# % Obesity in Parous vs Non-parous

Demographic match, 1:1 neighbor, with replacement

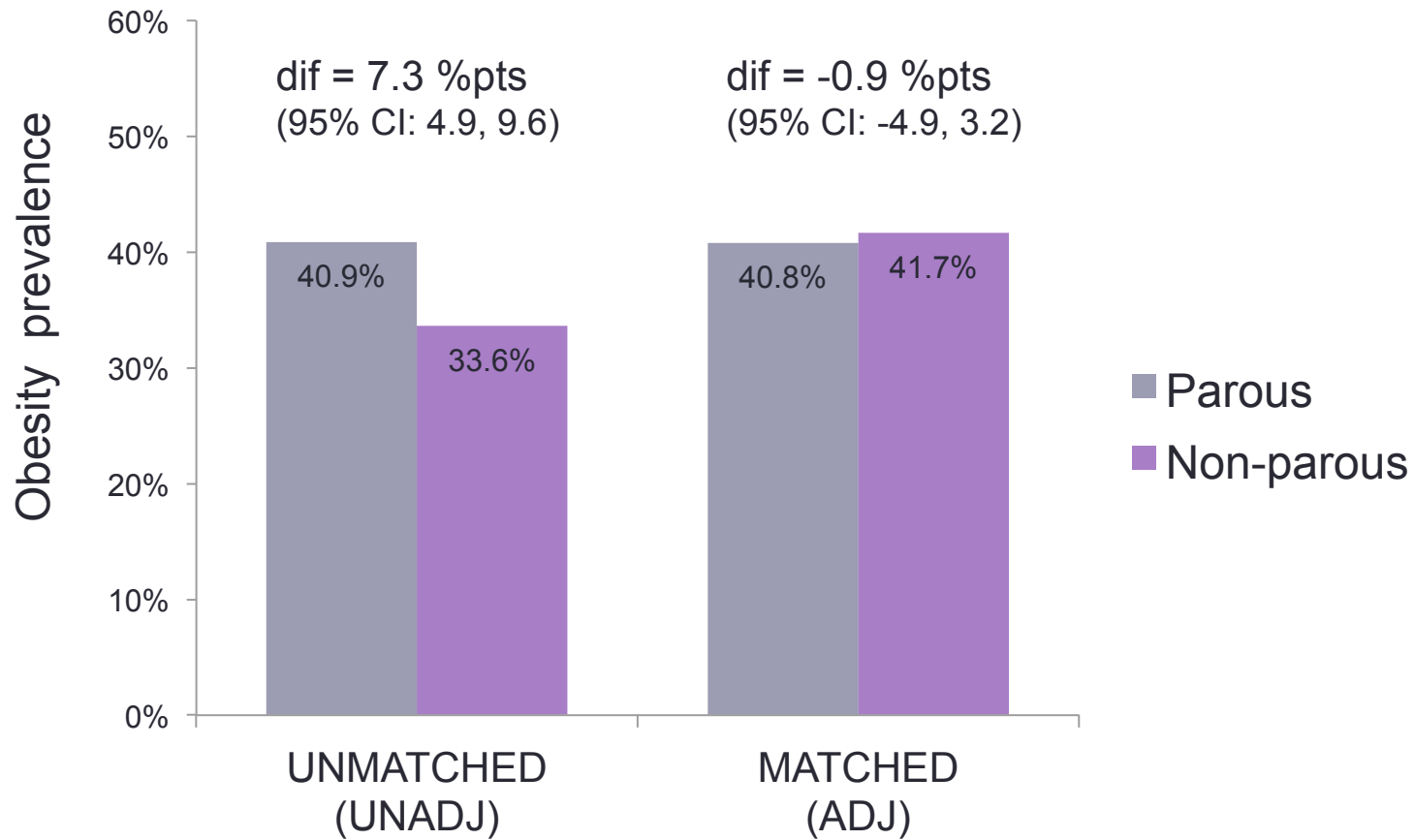


N=6,592 of 6,601

```
psmatch2 parous, outcome(ob_constr_w4) pscore(logit1 a) caliper(.1) com;
```

# % Obesity in Parous vs Non-parous

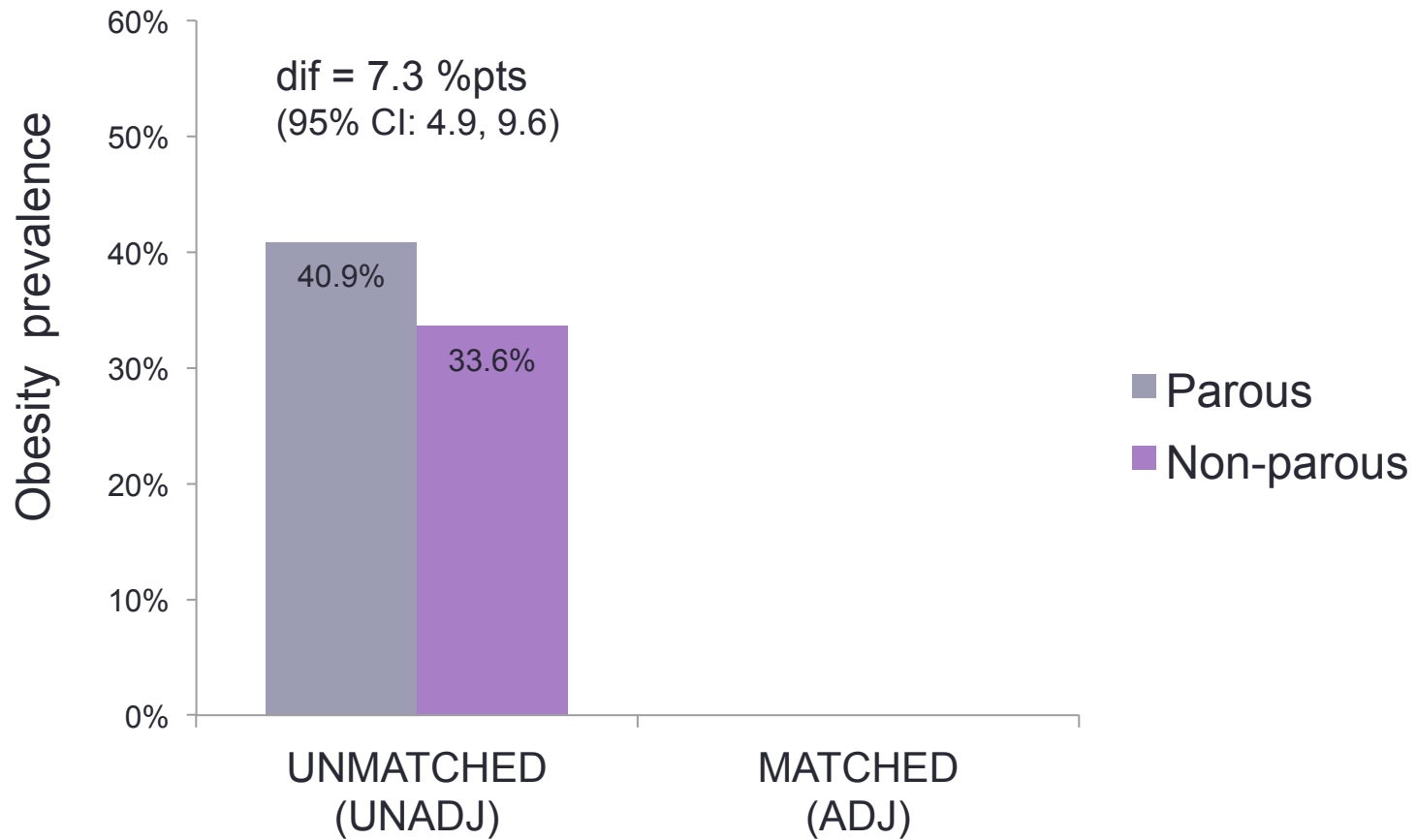
Demographic match, 1:1 neighbor, with replacement



N=6,592 of 6,601

# % Obesity in Parous vs Non-parous

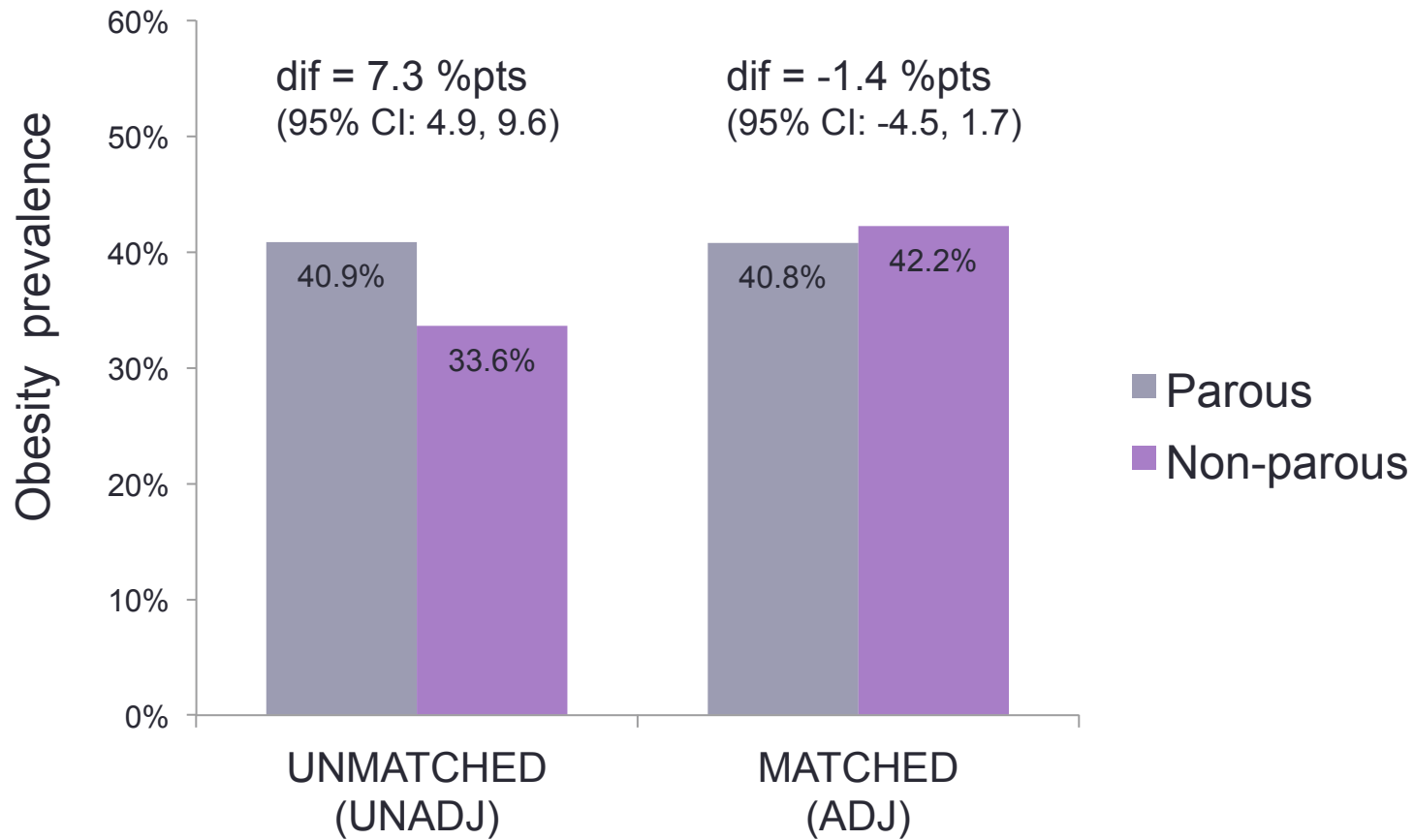
Demographic match, 4:1 neighbor, with replacement



N=6,592 of 6,601

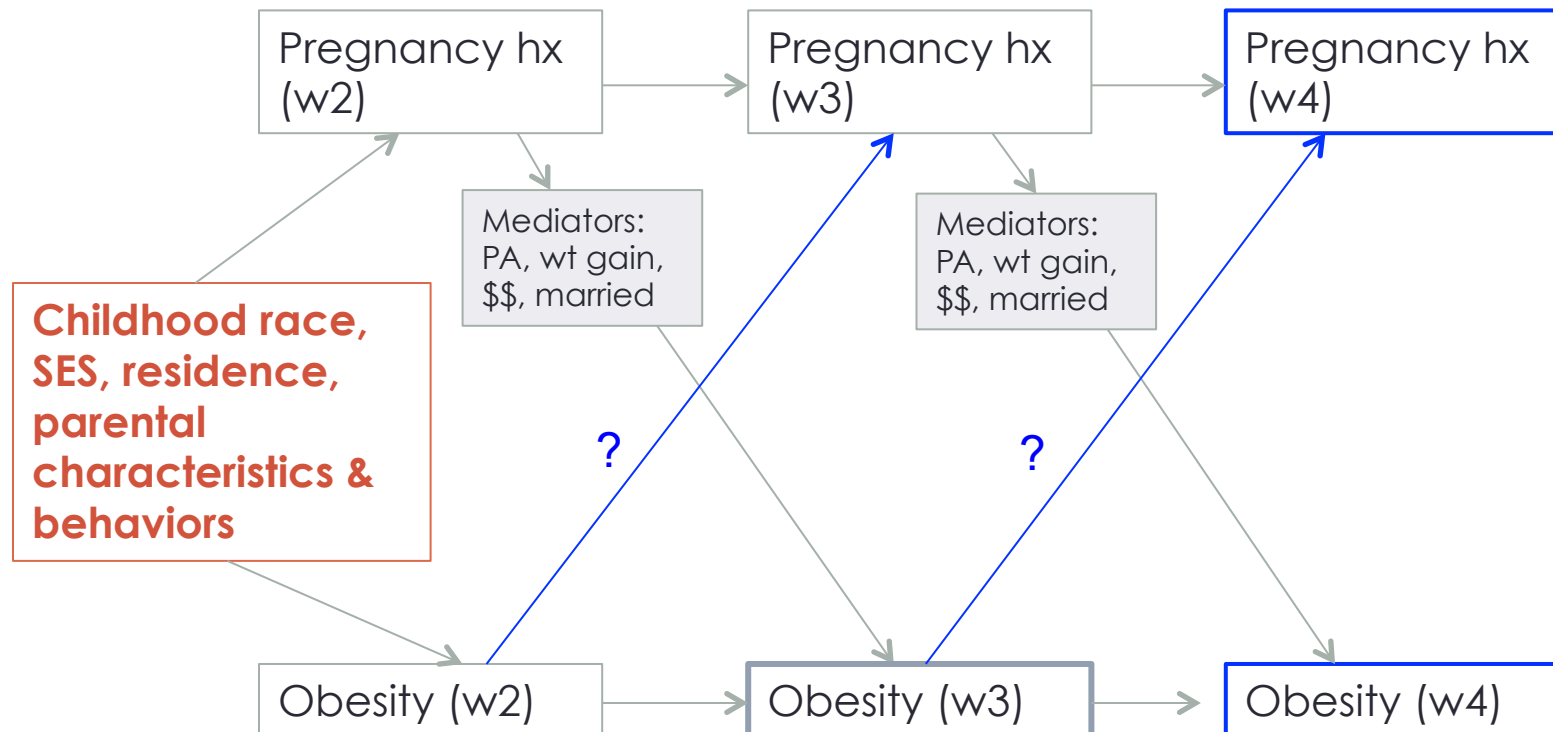
# % Obesity in Parous vs Non-parous

Demographic match, 4:1 neighbor, with replacement



N=6,592 of 6,601

# Motivation for w3-w4 incidence analysis



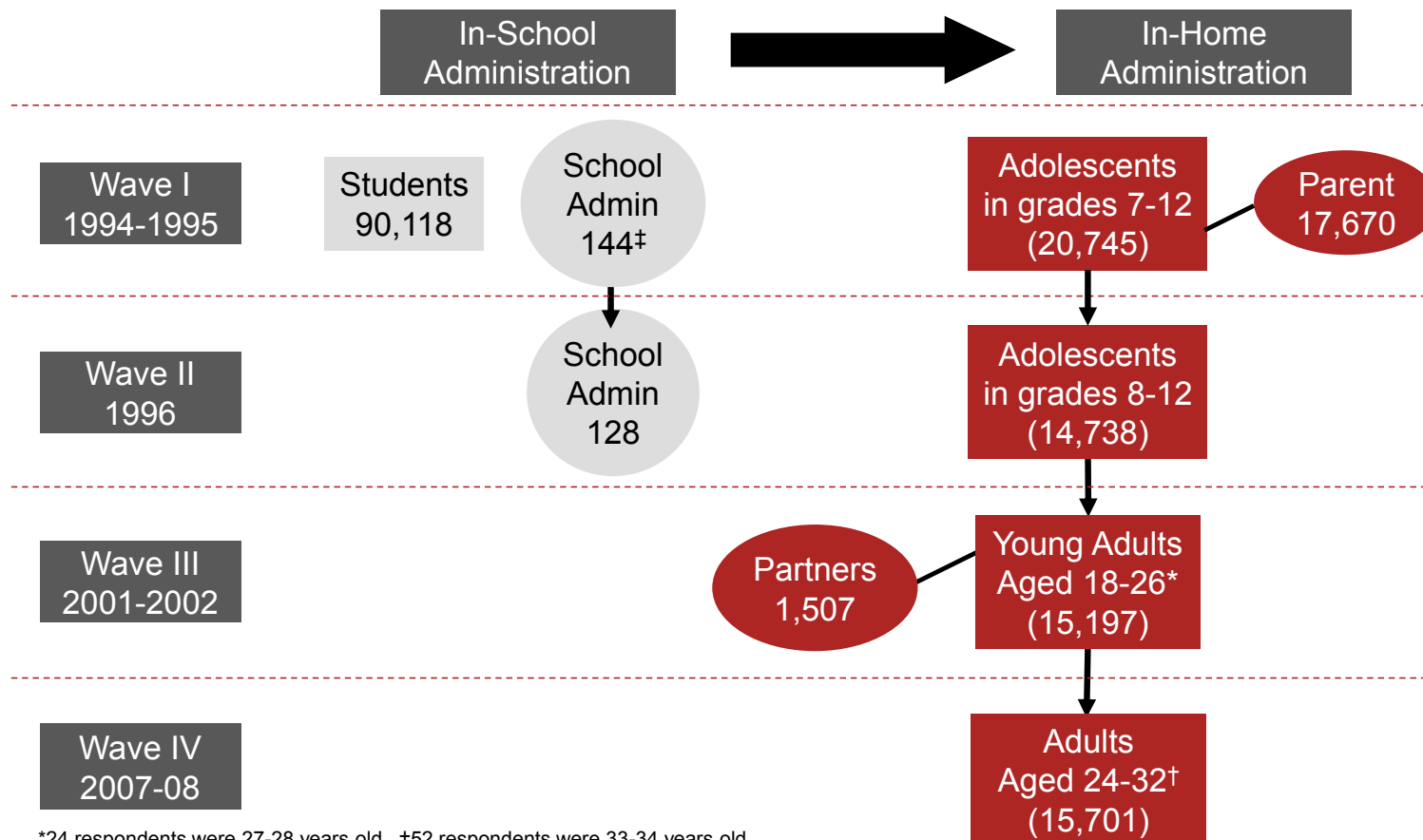
Frisco, M. L., M. M. Weden, A. M. Lippert and K. D. Burnett (2012). "The multidimensional relationship between early adult body weight and women's childbearing experiences." Soc Sci Med **74**(11): 1703-1711.

# Add Health: w3-w4 incidence



*Social, Behavioral, and Biological Linkages Across the Life Course*

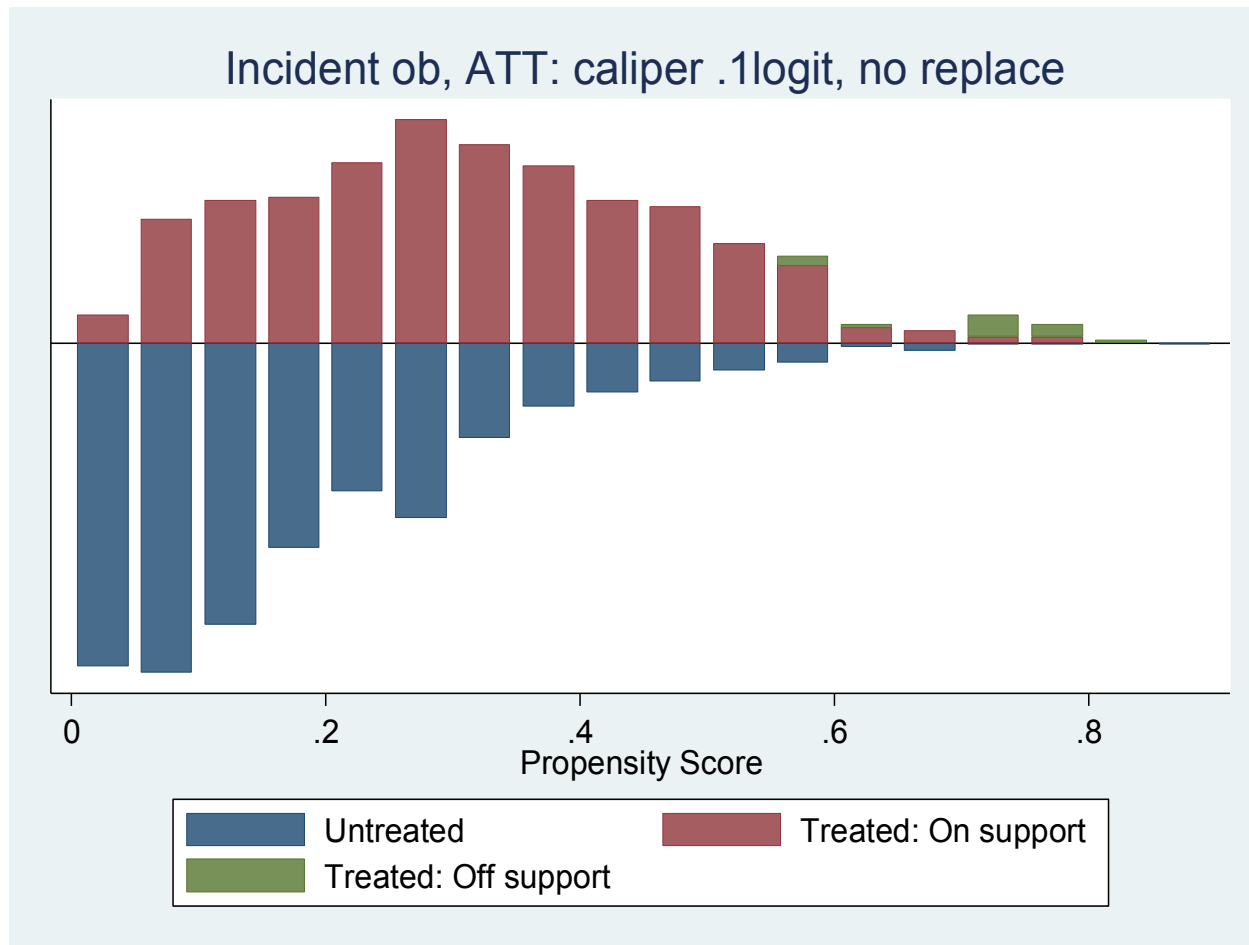
## Longitudinal Design



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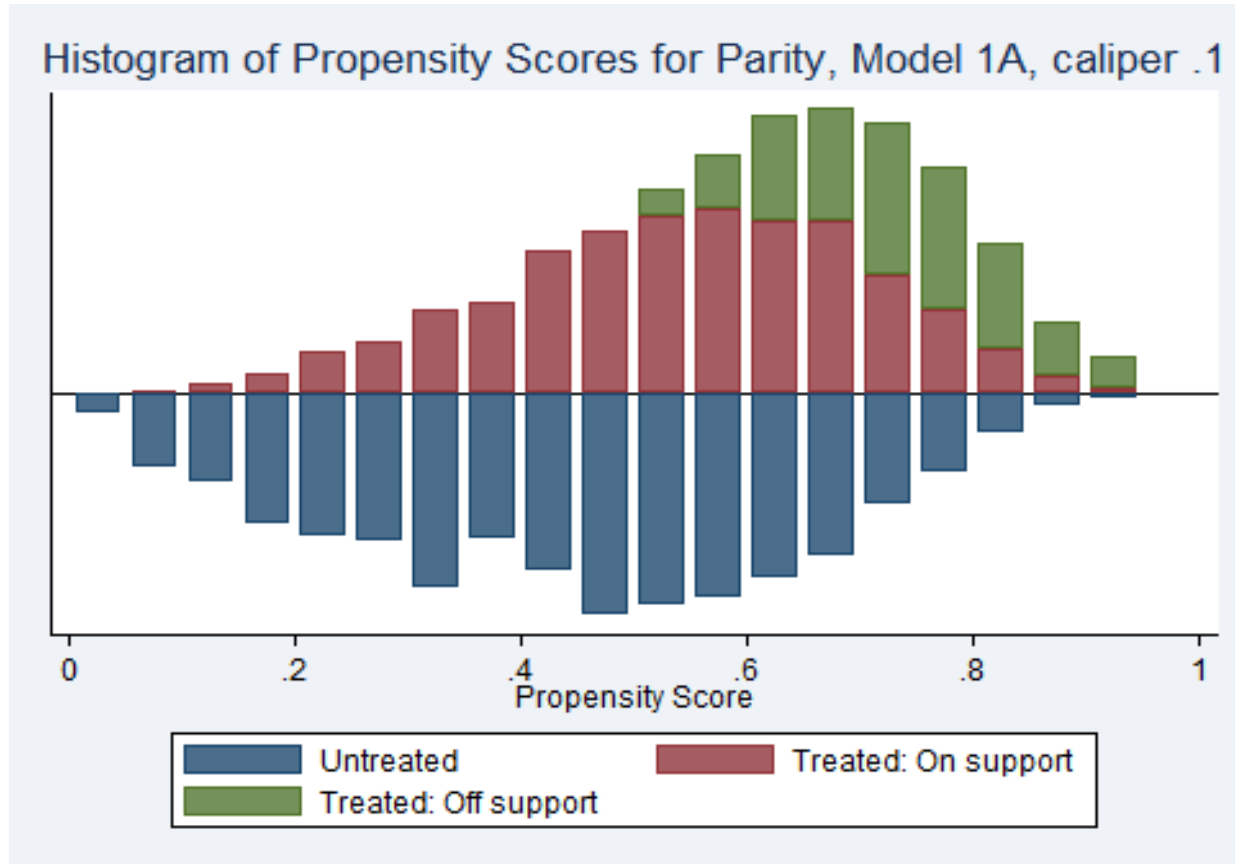
# Distribution of p-scores by parity: incidence analysis



More (moderate) imbalance, good overlap



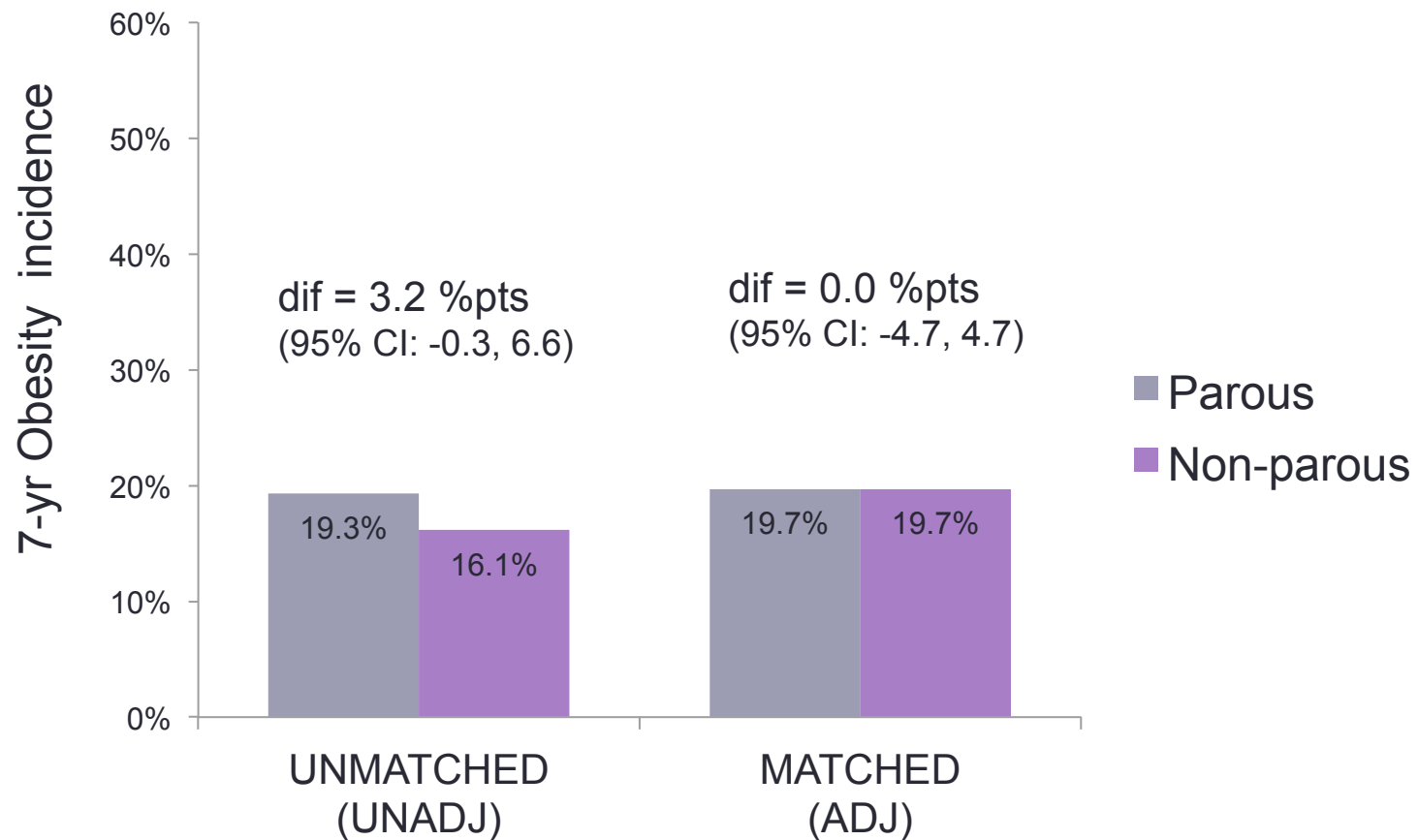
Compare this prevalence graph to incidence graph above



Slight imbalance, good overlap

# % Incident Obesity in Parous vs Non-parous

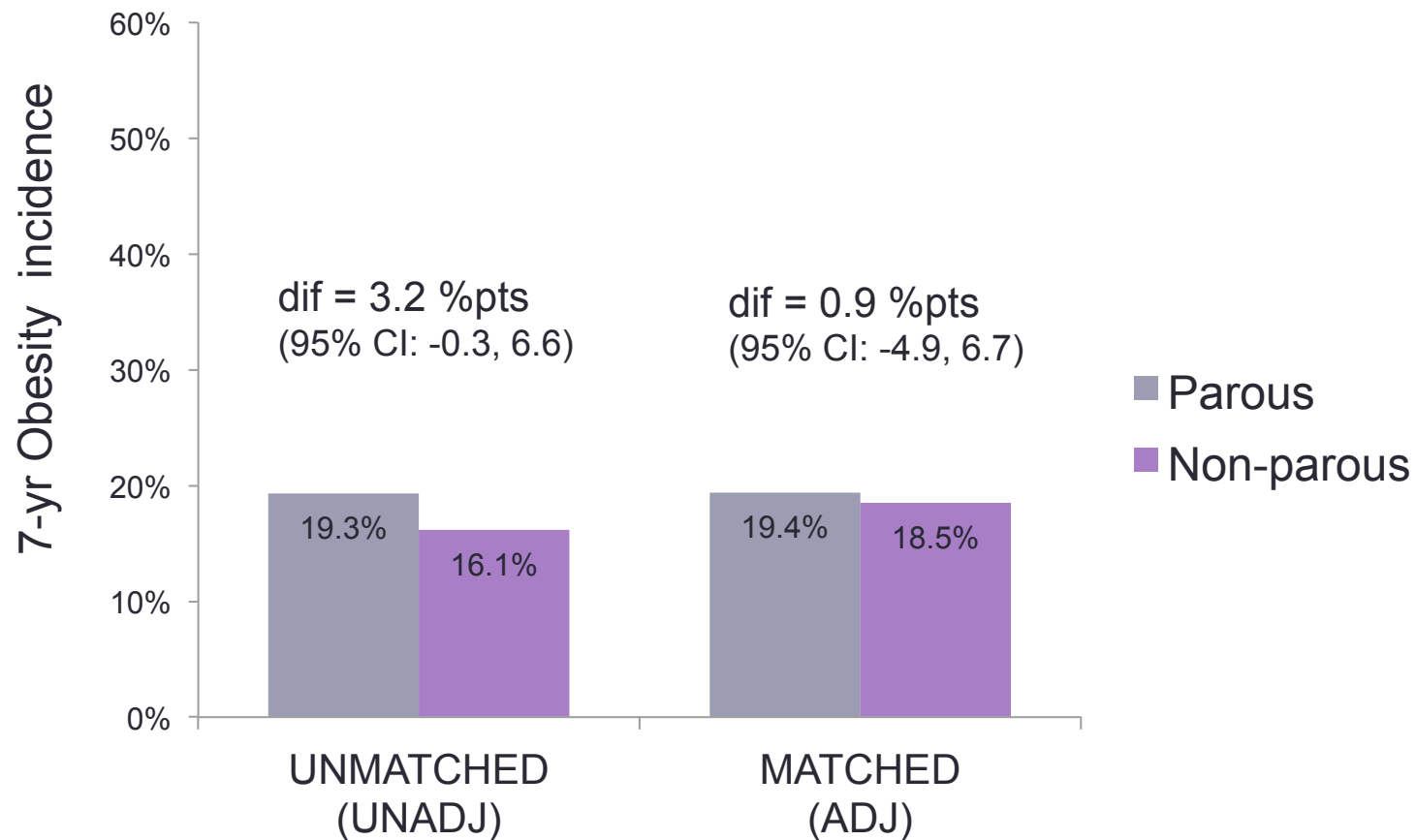
Demographic match, 1:1 neighbor, no replacement



N=2,715 of 2,731

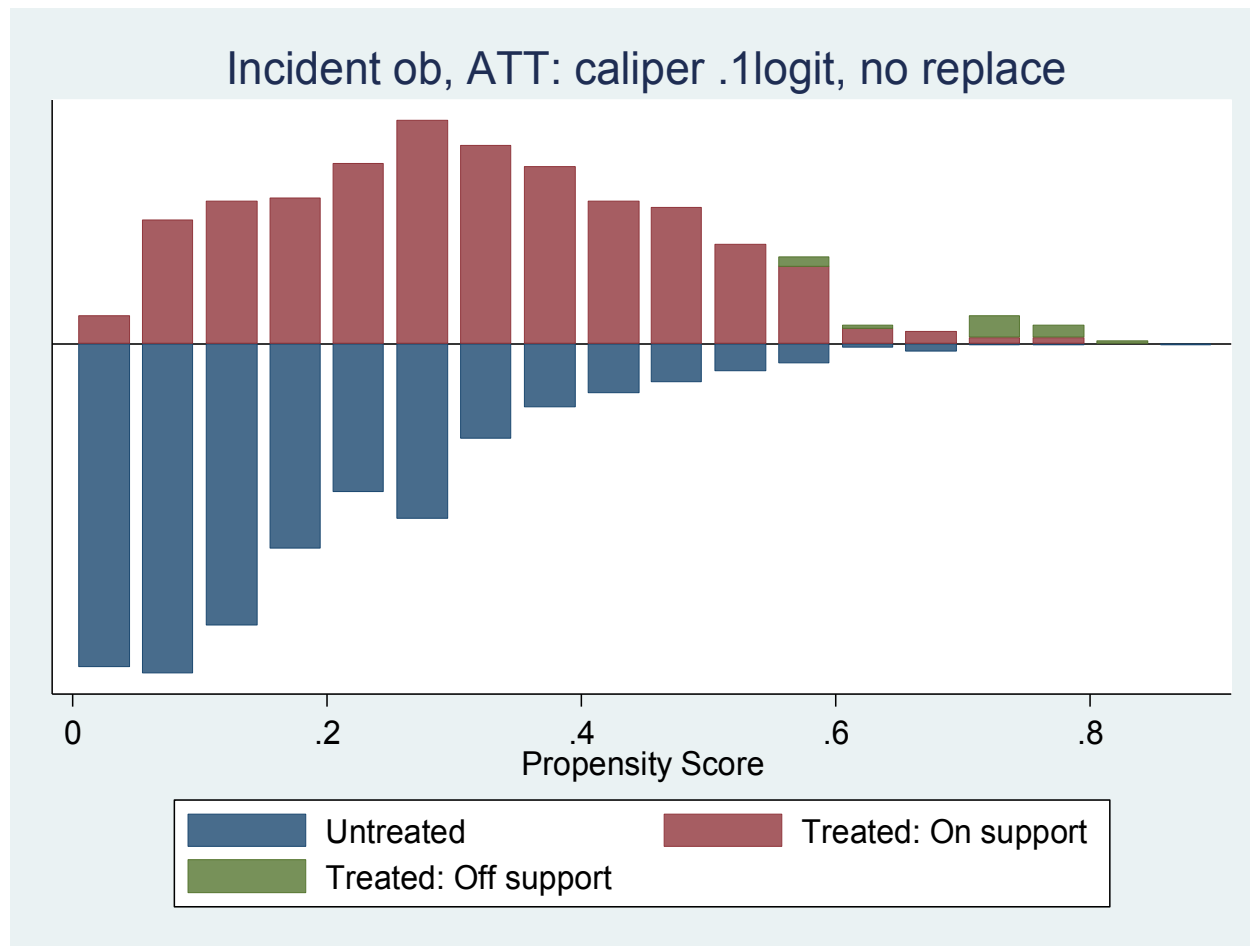
# % Incident Obesity in Parous vs Non-parous

Demographic match, 1:1 neighbor, with replacement



N=2,730 of 2,731

# Distribution of p-scores by parity: incidence analysis



Moderate imbalance, good overlap

# Interpreting results

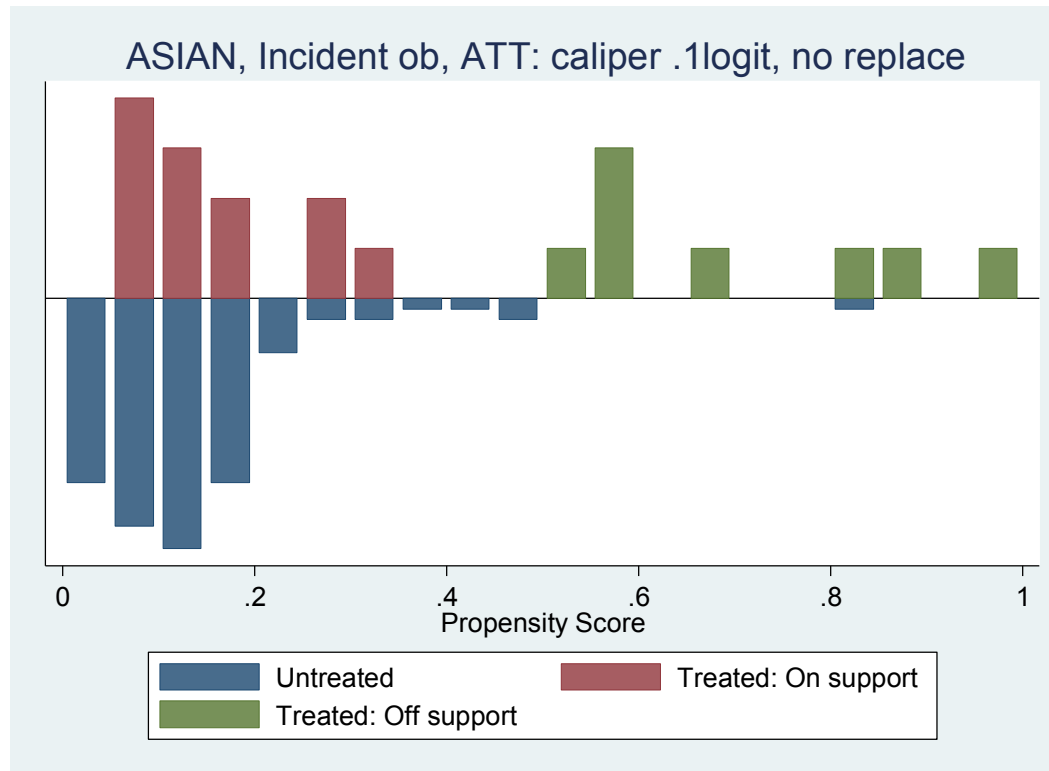
- **Summary:** No evidence that child-bearing contributes to obesity incidence or prevalence in young U.S. women
- **Target population:** women who gave birth by early 20s (incidence) or age ~30 years (prevalence),
  - Population: young mothers of mostly pre-school aged children
  - Causal action: postponing or abstaining from childbearing
- **Heterogeneity**
  - Marginal effect could mask risk/protectiveness of child-bearing
  - Could attempt to identify at-risk women, e.g., race, pre-preg BMI
- **Mechanisms**
  - Child-bearing: biological and social phenomenon

# Acknowledgements

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  - RWJF H&SS small grant, University of Michigan
  - Carolina Population Center Summer-in-Residence Grant Writing Fellowship (2012)
  - NCI K01CA172717
- Collaborators:
  - Mariah Cheng, Katherine Hoggatt, Anna-Maria Siega Riz

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# Example with overlap problem



# THE END

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