

# Interaction

February 7, 2017

Biostatistics 208

# Announcement about homework assignments:

- Please include relevant Stata code and output for every question
- File name prefixes for submitted assignments: "LastName HW#" (e.g. "Shiboski HW2")

# Interaction in Linear Regression Models

- Defining interaction
- Regression models including interaction terms
- Estimating effects in subgroups using `lincom` in Stata
- Interpretation of coefficients

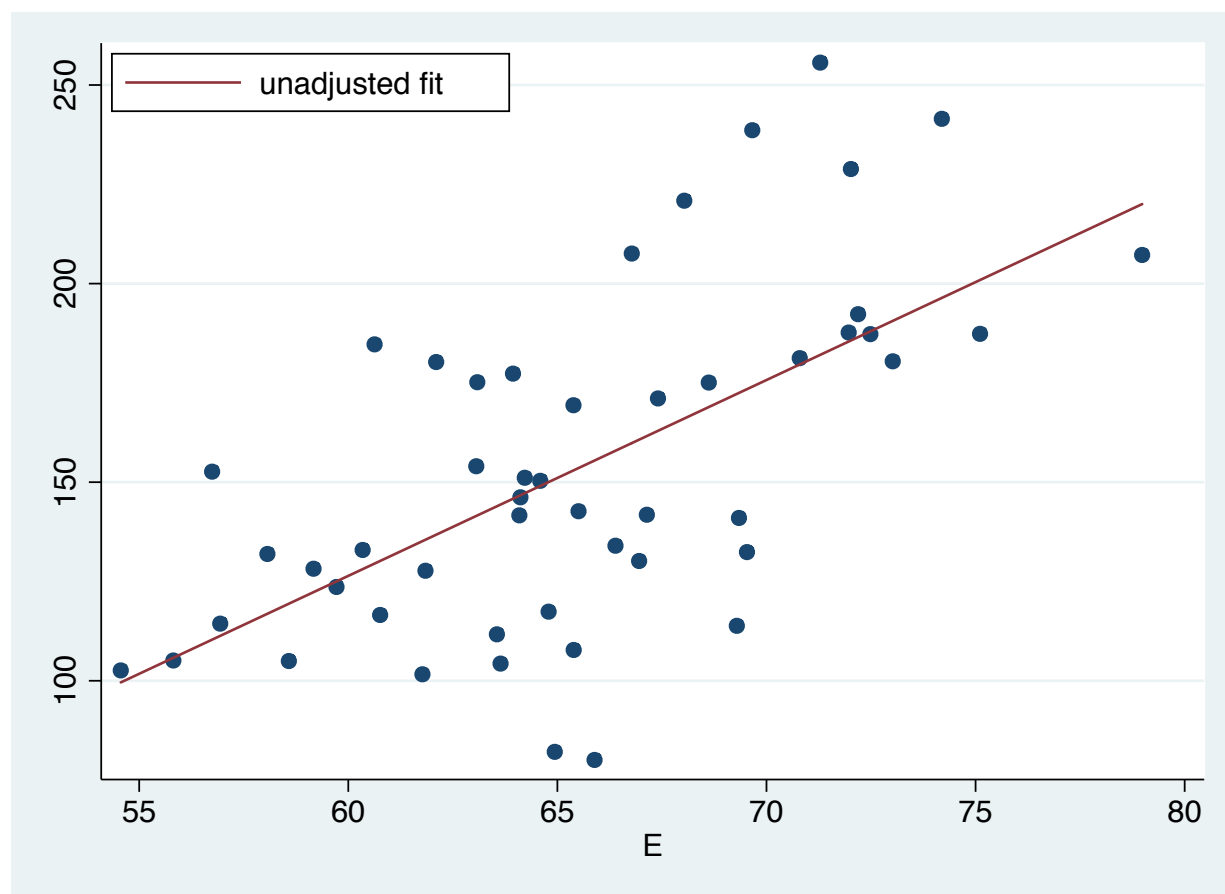
# Interaction

- **Confounding and mediation:** association of primary predictor with outcome *differs after adjustment*
  - adjustment variable *explains* or *masks* unadjusted effect of predictor
- **Interaction:** association of primary predictor with the outcome *differs across levels of the interaction variable*
  - interaction variable *modifies/moderates* estimated effect of primary predictor

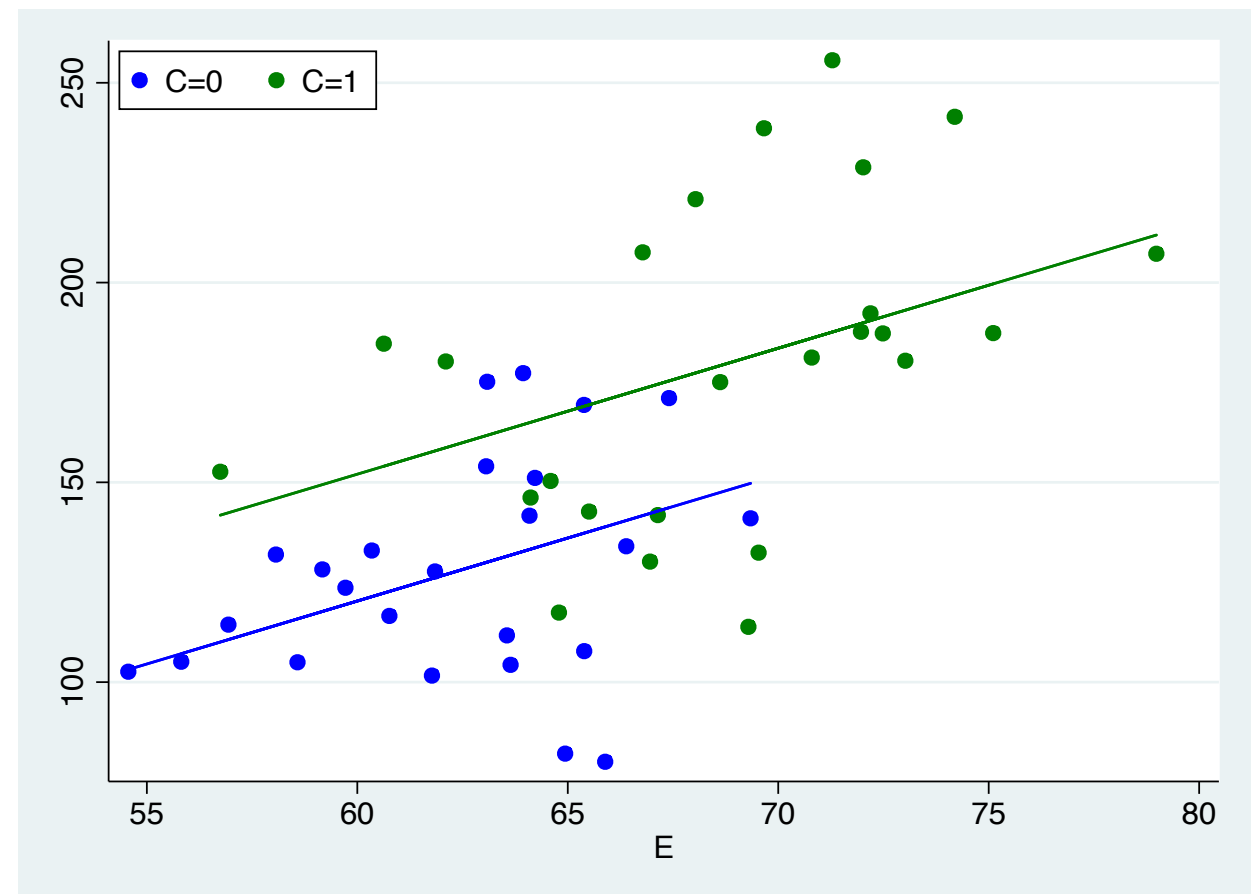
# Confounding example based on simulated data (slide #9 - last lecture)

- continuous exposure  $E$
- binary confounder  $C$  - no interaction with  $E$
- continuous outcome  $y = \beta_0 + \beta_1 E + \beta_2 C + \varepsilon$

Unadjusted



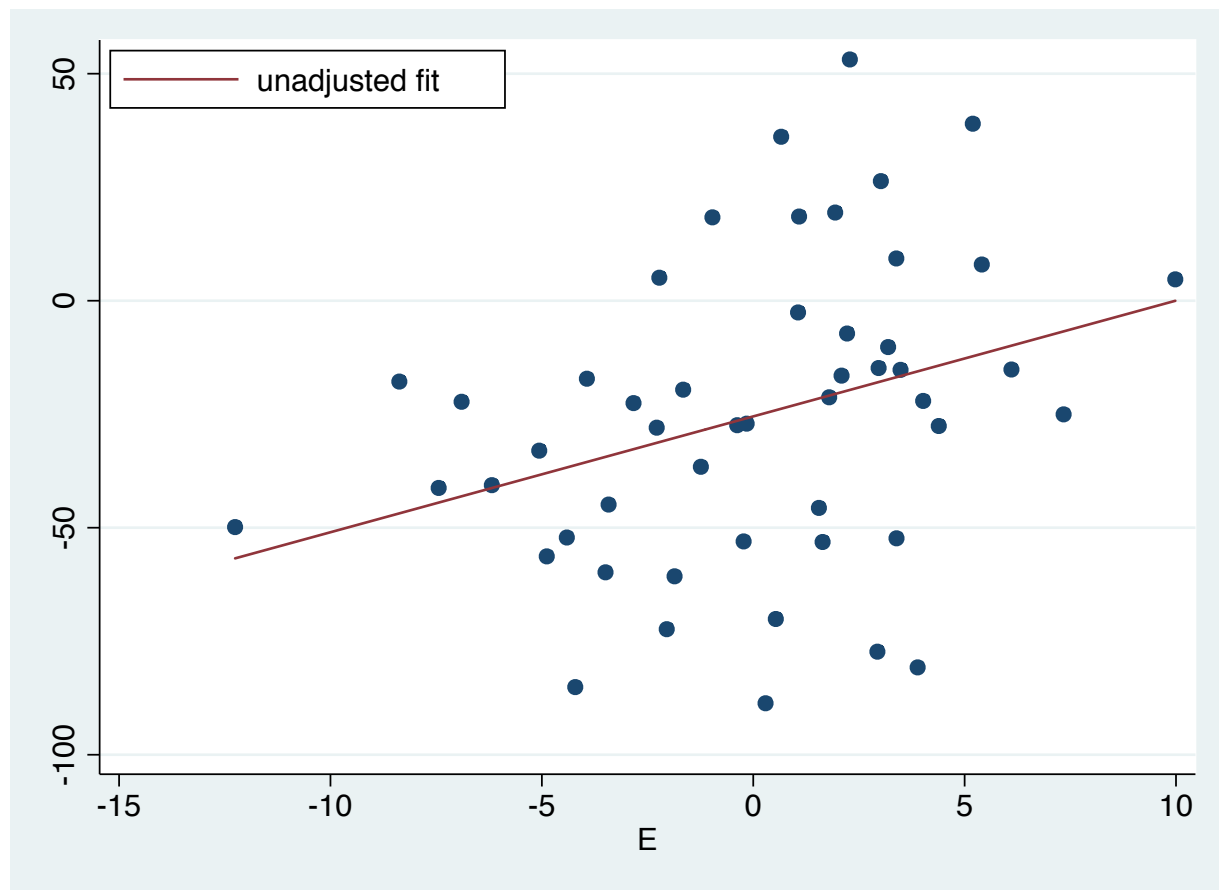
Adjusted



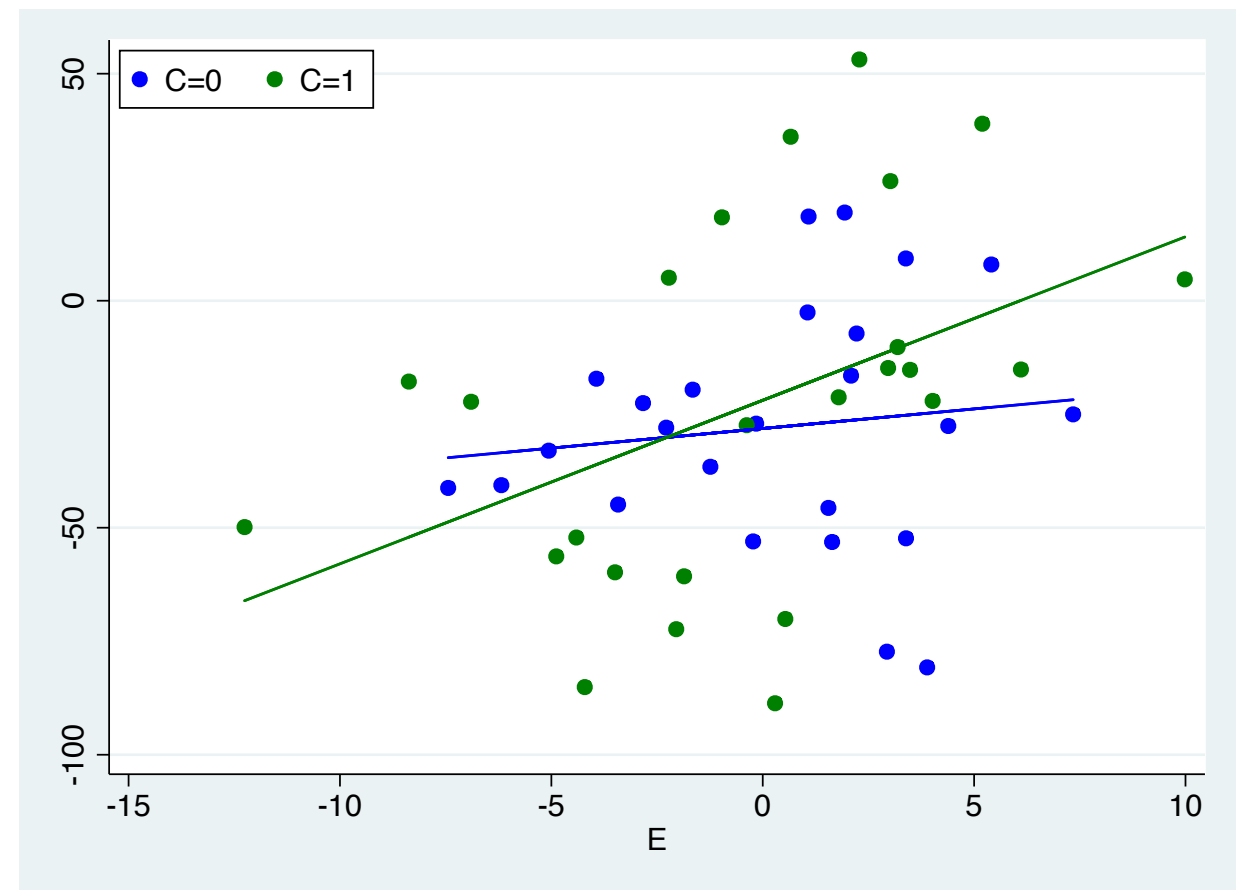
# Example based on simulated data including interaction

- continuous exposure  $E$
- binary predictor  $C$  - interaction with  $E$
- continuous outcome  $y = \beta_0 + \beta_1 E + \beta_2 C + \beta_3 CE + \varepsilon$

Unadjusted



Adjusted



# Linear regression models including interaction

To allow interaction between two predictors  $x_1$  &  $x_2$  in a regression model for a continuous outcome  $y$ , include both predictors individually, as *main effects* along with their product (for *interaction*):

$$E[y|x_1, x_2] = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_3 x_1 \times x_2$$

Resulting model allows separate linear effects for  $x_1$  &  $x_2$  :

- Outcome mean as a function of  $x_1$ , with  $x_2$  fixed at a constant value  $c$ :

$$\begin{aligned} E[y|x_1, x_2 = c] &= \beta_0 + \beta_1 x_1 + \beta_2 c + \beta_3 x_1 \times c \\ &= (\beta_0 + \beta_2 c) + (\beta_1 + \beta_3 c) x_1 \end{aligned}$$

effect of  $x_1$ : 

effect of  $x_2$ : 

intercept: 

- Outcome mean as a function  $x_2$ , with  $x_1$  fixed at a constant value  $c$ :

$$\begin{aligned} E[y|x_1 = c, x_2] &= \beta_0 + \beta_1 c + \beta_2 x_2 + \beta_3 c \times x_2 \\ &= (\beta_0 + \beta_1 c) + (\beta_2 + \beta_3 c) x_2 \end{aligned}$$

- Allows different intercepts & slopes as  $c$  varies
- Reduces to standard linear model with no interaction (  $\beta_3 = 0$  )
- $\beta_3 \neq 0$  represents *interaction* between  $x_1$  &  $x_2$

# Example: interaction of BMI and physical activity (PA) effects on SBP in HERS

Motivating questions:

- Excess weight increases SBP, but PA may blunt this effect
- Does higher BMI lead to higher SBP only among inactive?
- Does protective effect of PA on SBP increase with BMI?

- Predictors:

```
egen mb = mean(BMI)
```

```
gen BMIC = BMI - mb
```

```
recode physact (1/2=0) (3/5=1), g(active)
```

- Centered BMI:  $BMIC = BMI - 28.6$  (centered on sample mean for interpretability)

- Physically active (indicator for activity generated by including `i.active` in model):

`1.active` (0 = inactive, 1 = active)

- Interaction (product) term `1.active#c.BMIC` (# denotes product term in model):

$$1.active\#c.BMIC = 1.active \times c.BMIC$$
$$= 0 \text{ if } active = 0$$
$$= BMIC \text{ if } active = 1$$

*Note:* “`c.`” syntax denotes a continuous predictor in a regression model - used for including continuous variables in interaction terms

# Effects of PA - dependence on BMI

- Model for mean SBP:

$$E[\text{SBP} | \text{active}, \text{BMI}c] = \beta_0 + \beta_1 1.\text{active} + \beta_2 c.\text{BMI}c + \beta_3 1.\text{active} \# c.\text{BMI}c$$

- Estimated mean SBP for each level of PA:

$$E[\text{SBP} | \text{active}=0, \text{BMI}c] = \beta_0 + \beta_2 c.\text{BMI}c$$

$$E[\text{SBP} | \text{active}=1, \text{BMI}c] = \beta_0 + \beta_1 + (\beta_2 + \beta_3) c.\text{BMI}c$$

- Difference is  $\beta_1 + \beta_3 c.\text{BMI}c$
- Effect of PA depends on BMI

# Effects of BMI among active & inactive participants

$$E[\text{SBP} | \text{active}, \text{BMI}c] = \beta_0 + \beta_1 1.\text{active} + \beta_2 c.\text{BMI}c + \beta_3 1.\text{active} \# c.\text{BMI}c$$

For inactive:

$$E[\text{SBP} | \text{active}=0, \text{BMI}c] = \beta_0 + \beta_2 c.\text{BMI}c$$

- $\beta_2$  gives change in mean SBP (*mmHg*) for unit (*kg/m<sup>2</sup>*) increase in BMI

For active:

$$E[\text{SBP} | \text{active}=1, \text{BMI}c] = (\beta_0 + \beta_1) + (\beta_2 + \beta_3) c.\text{BMI}c$$

- $\beta_2 + \beta_3$  gives change in mean SBP for unit increase in BMI
- Effect of BMI depends on PA

# Interaction model for SBP in baseline HERS dataset

```
. regress SBP i.active##c.BMIc
```

| Source   | SS         | df   | MS         | Number of obs | = | 2758   |
|----------|------------|------|------------|---------------|---|--------|
| Model    | 4579.65484 | 3    | 1526.55161 | F( 3, 2754)   | = | 4.24   |
| Residual | 992671.256 | 2754 | 360.447079 | Prob > F      | = | 0.0054 |
|          |            |      |            | R-squared     | = | 0.0046 |
|          |            |      |            | Adj R-squared | = | 0.0035 |
| Total    | 997250.911 | 2757 | 361.715963 | Root MSE      | = | 18.985 |

| SBP           | Coef.               | Std. Err. | t      | P> t  | [95% Conf. Interval] |
|---------------|---------------------|-----------|--------|-------|----------------------|
| 1.active      | $-.5468102 \beta_1$ | .8611566  | -0.63  | 0.525 | -2.235388 1.141768   |
| BMIc          | $.0512451 \beta_2$  | .1138269  | 0.45   | 0.653 | -.1719496 .2744397   |
| active#c.BMIc |                     |           |        |       |                      |
| 1             | $.2293768 \beta_3$  | .1406966  | 1.63   | 0.103 | -.0465047 .5052584   |
| _cons         | $135.5883 \beta_0$  | .7508309  | 180.58 | 0.000 | 134.116 137.0605     |

Note: syntax "i.active##c.BMIc" is shorthand for including:  
i.active, c.BMIc & i.active#c.BMIc.

Equivalent command from earlier Stata versions ( $\leq 10$ ):

```
xi: regress sbp i.active*BMIc
```

# Equivalent specification of previous model

```
. regress SBP i.active c.BMIc i.active#c.BMIc
```

| Source   | SS         | df   | MS         | Number of obs = 2758   |  |  |  |
|----------|------------|------|------------|------------------------|--|--|--|
| Model    | 4579.65484 | 3    | 1526.55161 | F( 3, 2754) = 4.24     |  |  |  |
| Residual | 992671.256 | 2754 | 360.447079 | Prob > F = 0.0054      |  |  |  |
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| SBP           | Coef.     | Std. Err. | t      | P> t  | [95% Conf. Interval] |          |
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| 1.active      | -.5468102 | .8611566  | -0.63  | 0.525 | -2.235388            | 1.141768 |
| BMIc          | .0512451  | .1138269  | 0.45   | 0.653 | -.1719496            | .2744397 |
| active#c.BMIc |           |           |        |       |                      |          |
| 1             | .2293768  | .1406966  | 1.63   | 0.103 | -.0465047            | .5052584 |
| _cons         | 135.5883  | .7508309  | 180.58 | 0.000 | 134.116              | 137.0605 |

## Interpretation of coefficients: `_cons` ( $\beta_0$ )

$$E[\text{SBP} | \text{active}, \text{BMIC}] = \beta_0 + \beta_1 1.\text{active} + \beta_2 \text{c.BMIC} + \beta_3 1.\text{active} \# \text{c.BMIC}$$

- Interpretation: average SBP in inactive participants with `BMIC = 0`  
(i.e., BMI set at mean value of  $28.6 \text{ kg/m}^2$ ):

$$E[\text{SBP} | \text{active}=0, \text{BMIC}=0] = \beta_0$$

- Average SBP among inactive with BMI =  $28.6 \text{ kg/m}^2$  is  $135.6 \text{ mmHg}$  (95% CI: 134.1, 137.1)

# Interpretation of coefficients: 1.active ( $\beta_1$ )

$$E[\text{SBP} | \text{active}, \text{BMIC}] = \beta_0 + \beta_1 1.\text{active} + \beta_2 c.\text{BMIC} + \beta_3 1.\text{active} \# c.\text{BMIC}$$

- Interpretation: effect of being active for participants with  $\text{BMIC} = 0$  :

$$E[\text{SBP} | \text{active} = 1, \text{BMIC} = 0] = \beta_0 + \beta_1$$

$$E[\text{SBP} | \text{active} = 0, \text{BMIC} = 0] = \beta_0$$

- Difference is  $\beta_1$  (i.e. effect of being active for those with average BMI)
  - At BMI of  $28.6 \text{ kg/m}^2$  (mean), PA reduces SBP by  $0.55 \text{ mmHg}$  (95% CI: -2.2, 1.1)
- For other values of BMI the effect of PA on SBP involves the coefficient  $\beta_3$

# Interpretation of coefficients: BMIc ( $\beta_2$ )

$$E[\text{SBP} | \text{active}, \text{BMIc}] = \beta_0 + \beta_1 1.\text{active} + \beta_2 \text{BMIc} + \beta_3 1.\text{active} \# c.\text{BMIc}$$

- BMIc ( $\beta_2$ ): effect of a 1-kg/m<sup>2</sup> increase in BMI, if inactive

$$E[\text{SBP} | \text{active} = 0, \text{BMIc} = k+1] = \beta_0 + \beta_2 (k+1)$$

$$E[\text{SBP} | \text{active} = 0, \text{BMIc} = k] = \beta_0 + \beta_2 k$$

- Difference is  $\beta_2$  for any value of BMI ( $k$ )
- Among inactive, a 1-kg/m<sup>2</sup> increase in BMI increases SBP by 0.05 mmHg (95% CI: -0.2, 0.3)

## Interpretation of coefficients: $1.active\#c.BMIc$ ( $\beta_3$ )

$$E[SBP|active, BMIc] = \beta_0 + \beta_1 1.active + \beta_2 BMIc + \beta_3 1.active\#c.BMIc$$

### 1. Difference between BMI effects in active and inactive

$$E[SBP|active = 1, BMIc = k+1] - E[SBP|active = 1, BMIc = k] -$$

$$E[SBP|active = 0, BMIc = k+1] - E[SBP|active = 0, BMIc = k]$$

$$= \{ ( \beta_0 + \beta_1 + ( \beta_2 + \beta_3 ) ( k+1 ) ) - ( \beta_0 + \beta_1 + ( \beta_2 + \beta_3 ) k ) \} -$$

$$\{ ( \beta_0 + \beta_2 ( k+1 ) ) - ( \beta_0 + \beta_2 k ) \} = \beta_3$$

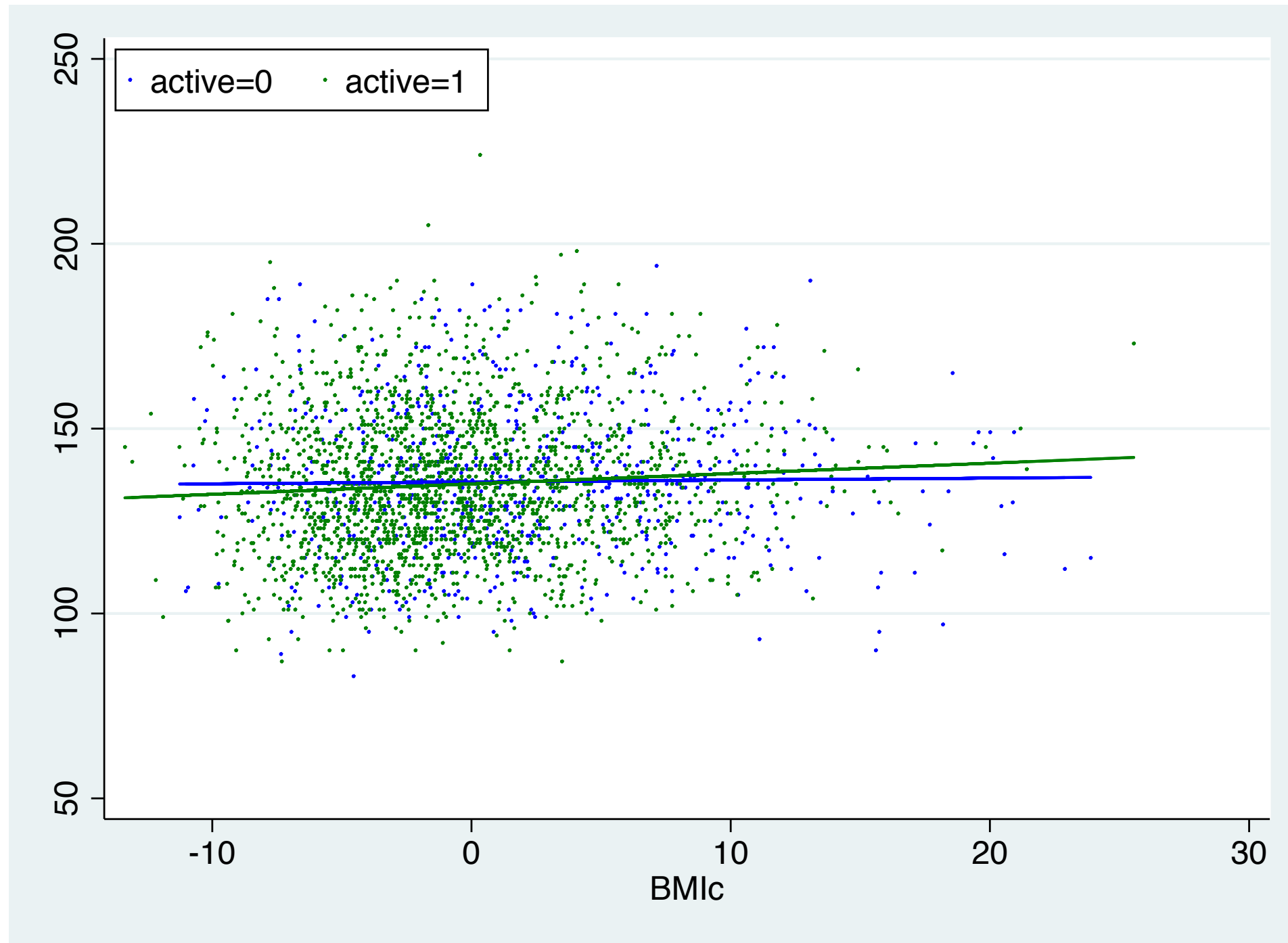
- among active, a unit  $kg/m^2$  increase in BMI predicts a *0.23 mmHg greater* increase in SBP than among inactive (95% CI: -0.05, 0.51)

### 2. Change in effect of PA for every $kg/m^2$ increase in BMI

- for every  $kg/m^2$  increase in BMI, the beneficial effect of being active on SBP *decreases by 0.23 mmHg* (95% CI: -0.05, 0.51)

### 3. $p$ -value of 0.10 for $1.active\#c.BMIc$ shows that evidence for interaction is weak

# Illustrating BMI - activity interaction in HERS



```
. regress SBP i.active##c.BM1c
. predict p
. graph twoway (scatter SBP BM1c if active==0, mcolor(blue) msize(vtiny))
(scatter SBP BM1c if active==1, mcolor(green) msize(vtiny)) (line p BM1c if
active==0, clcolor(blue)) (line p BM1c if active==1, clcolor(green)),
legend( order(1 "active=0" 2 "active=1") ring(0) pos(10) )
```

# Use of `lincom` for summarizing interaction effects

- Effects of PA & BMI depend on interaction coefficient `1.active#c.BMIc` ( $\beta_3$ )
  - `lincom` statements can be used to illustrate effects (preferred to presenting raw coefficients)
  - Example - two `lincom` statements:

```
. lincom 1.active + 1.4*1.active#c.BMIc ( $\beta_1 + 1.4\beta_3$ )
```

```
( 1) 1.active + 1.4*1.active#c.BMIc = 0
```

```
-----+-----
      sbp |      Coef.   Std. Err.      t    P>|t|     [95% Conf. Interval]
-----+-----
      (1) |   -.2256828   .8505544    -0.27   0.791   -1.893472    1.442106
-----+-----
```

```
. lincom BMIc + 1.active#c.BMIc ( $\beta_2 + \beta_3$ )
```

```
( 1) BMIc + 1.active#c.BMIc = 0
```

```
-----+-----
      sbp |      Coef.   Std. Err.      t    P>|t|     [95% Conf. Interval]
-----+-----
      (1) |   .2806219   .0826981     3.39   0.001    .1184653    .4427785
-----+-----
```

# Interpretation of first `lincom` result:

`lincom 1.active + 1.4*1.active#c.BM Ic` ( $\beta_1 + 1.4\beta_3$ )

- Interpretation: effect of PA for an individual with BMI of 30  $kg/m^2$ 
  - for BMI=30,  $BMIc=(30-28.6) = 1.4 \text{ kg/m}^2$

$$E[SBP|x] = \beta_0 + \beta_1 1.active + \beta_2 BMIc + \beta_3 1.active \# c.BM Ic$$

*Active:*  $E[SBP_{active} = 1, BMIc = 1.4] = \beta_0 + \beta_1 + 1.4(\beta_2 + \beta_3)$

*Inactive:*  $E[SBP_{active} = 0, BMIc = 1.4] = \beta_0 + 1.4\beta_2$

- Difference is  $\beta_1 + 1.4\beta_3 = -0.23$
- At BMI of 30  $kg/m^2$ , PA reduces mean SBP by 0.23  $mmHg$  (95% CI: -1.9, 1.4)

# Interpretation of second `lincom` result:

`lincom c.BMIc + 1.active#c.BMIc` ( $\beta_2 + \beta_3$ )

- Interpretation: effect of a  $1\text{-kg/m}^2$  increase in BMI, for an active individual

$$E[\text{SBP}|x] = \beta_0 + \beta_1 1.\text{active} + \beta_2 \text{BMIc} + \beta_3 1.\text{active}\#\text{c.BMIc}$$

$$E[\text{SBP}|\text{active} = 1, \text{BMIc} = k+1] = \beta_0 + \beta_1 + (k+1)(\beta_2 + \beta_3)$$

$$E[\text{SBP}|\text{active} = 1, \text{BMIc} = k] = \beta_0 + \beta_1 + k(\beta_2 + \beta_3)$$

- Difference is  $\beta_2 + \beta_3 = 0.28$ , for any value of  $k$
- Among active, a  $1\text{-kg/m}^2$  increase in BMI increases SBP by  $0.28\text{ mmHg}$   
(95% CI: -0.12, 0.44)

# Figuring out what `lincom` statement to use:

- **Example 1:** Estimate effect of PA at BMI of 35 (i.e.  $BMI_c = 35 - 28.6 = 6.4 \text{ kg/m}^2$ ):

$$E[SBP|active, BMI_c] = \beta_0 + \beta_1 1.active + \beta_2 BMI_c + \beta_3 1.active \# c.BMI_c$$

| Active | BMI <sub>c</sub> | _cons | 1.active | BMI <sub>c</sub> | 1.active#BMI <sub>c</sub> |
|--------|------------------|-------|----------|------------------|---------------------------|
| yes    | 6.4              | 1     | 1        | 6.4              | 6.4                       |
| no     | 6.4              | 1     | 0        | 6.4              | 0                         |
|        | Diff.            | 0     | 1        | 0                | 6.4                       |

- Use `lincom 1.active + 6.4 * 1.active#c.BMIc` (output on next slide)

# Figuring out what `lincom` statement to use:

## \* Example 1

```
. lincom 1.active + 6.4 * 1.active#c.BMIc
```

```
( 1) 1.active + 6.4*1.active#c.BMIc = 0
```

| SBP | Coef.    | Std. Err. | t    | P> t  | [95% Conf. Interval] |          |
|-----|----------|-----------|------|-------|----------------------|----------|
| (1) | .9212014 | 1.1367    | 0.81 | 0.418 | -1.307668            | 3.150071 |

*Interpretation:* The estimated difference in average SBP between active and inactive individuals with a BMI of 35  $\text{kg/m}^2$  is 0.92 mmHg (95% CI: -1.37, 3.15).

Using margins to get the group means:

```
. margins active, at(BMIc=6.4)
```

Adjusted predictions

Number of obs = 2758

Model VCE : OLS

Expression : Linear prediction, `predict()`

at : BMIc = 6.4

|        | Delta-method |           |        |       |                      |          |
|--------|--------------|-----------|--------|-------|----------------------|----------|
|        | Margin       | Std. Err. | t      | P> t  | [95% Conf. Interval] |          |
| active |              |           |        |       |                      |          |
| 0      | 135.9163     | .8817122  | 154.15 | 0.000 | 134.1874             | 137.6451 |
| 1      | 136.8375     | .7174047  | 190.74 | 0.000 | 135.4307             | 138.2442 |

## Figuring out what `lincom` statement to use:

- **Example 2:** Estimate effect of additional 5  $\text{kg/m}^2$  of BMI among active individuals:

$$E[\text{SBP}|\text{active}, \text{BMIC}] = \beta_0 + \beta_1 1.\text{active} + \beta_2 \text{BMIC} + \beta_3 1.\text{active}\#c.\text{BMIC}$$

| Active | BMIC  | _cons | 1.active | BMIC  | 1.active#BMIC |
|--------|-------|-------|----------|-------|---------------|
| yes    | $k+5$ | 1     | 1        | $k+5$ | $k+5$         |
| yes    | $k$   | 1     | 1        | $k$   | $k$           |
|        | Diff. | 0     | 0        | 5     | 5             |

- Use `lincom 5 * (BMIC + 1.active#c.BMIC)`

# Figuring out what `lincom` statement to use:

## \* Example 2

```
. lincom 5 * (BMIC + 1.active#c.BMIC)
```

```
( 1) 5*BMIC + 5*1.active#c.BMIC = 0
```

| -----       |  |         |           |      |       |                      |
|-------------|--|---------|-----------|------|-------|----------------------|
| SBP         |  | Coef.   | Std. Err. | t    | P> t  | [95% Conf. Interval] |
| -----+----- |  |         |           |      |       |                      |
| (1)         |  | 1.40311 | .4134907  | 3.39 | 0.001 | .5923264 2.213893    |
| -----       |  |         |           |      |       |                      |

*Interpretation:* The estimated effect on average SBP of an increase of 5  $\text{kg/m}^2$  in BMI among active individuals is 1.40  $\text{mmHg}$  (95% CI: 0.59, 2.21)

## Figuring out what `lincom` statement to use:

- **Example 3:** Estimate effect of additional 5  $\text{kg/m}^2$  among the *inactive*:

$$E[\text{SBP}|\text{active}, \text{BMIC}] = \beta_0 + \beta_1 1.\text{active} + \beta_2 \text{BMIC} + \beta_3 1.\text{active}\#c.\text{BMIC}$$

| Active | BMIC  | _cons | 1.active | BMIC  | 1.active#BMIC |
|--------|-------|-------|----------|-------|---------------|
| no     | $k+5$ | 1     | 0        | $k+5$ | 0             |
| no     | $k$   | 1     | 0        | $k$   | 0             |
|        | Diff. | 0     | 0        | 5     | 0             |

- Use `lincom 5*BMIC`

# Figuring out what `lincom` statement to use:

## \* Example 3

```
. lincom 5*BMIc
```

```
( 1) 5*BMIc = 0
```

| -----       |  |          |           |      |       |                      |
|-------------|--|----------|-----------|------|-------|----------------------|
| SBP         |  | Coef.    | Std. Err. | t    | P> t  | [95% Conf. Interval] |
| -----+----- |  |          |           |      |       |                      |
| (1)         |  | .2562254 | .5691343  | 0.45 | 0.653 | -.8597478 1.372199   |
| -----       |  |          |           |      |       |                      |

*Interpretation:* The estimated effect on average SBP of an increase of 5  $\text{kg/m}^2$  in BMI among inactive individuals is 0.26 *mmHg* (95% CI: -0.86, 1.37)

# Interactions between continuous predictors

- Be careful with interactions between continuous variables
  - non-linearity, influential points can be a problem
  - easier to interpret if one or both dichotomized, but information gets lost
- Restricted cubic splines work (next week), but results only interpretable by plotting
- An example will be covered in this week's lab

# Interactions between multi-category predictors

- Hard to detect and interpret
- Example - predictors with 3 and 4 categories:
  - $(3 - 1) \times (4 - 1) = 6$  interaction parameters
- Manually-constructed product terms often don't work; best to use Stata syntax
  - newer versions ( $\geq 11$ ): `regress y i.catvar1##i.catvar2`
  - earlier versions ( $\leq 10$ ): `xi: regress y i.catvar1*i.catvar2`

# Additive models, interactions and outcome scale

- Outcome scale:
  - as measured vs log-transformed; absolute vs percent change
  - logistic, Poisson, negative binomial models:  $E[y | x]$  linked to predictors via logit or log transformations
- Additive models excluding interaction terms hold exactly on at most one outcome scale (may hold approximately on  $> 1$ )
- Changes of scale can eliminate need for interaction term

# Effects of hormone therapy (ht) on change in LDL (ldlch)

. tab ht

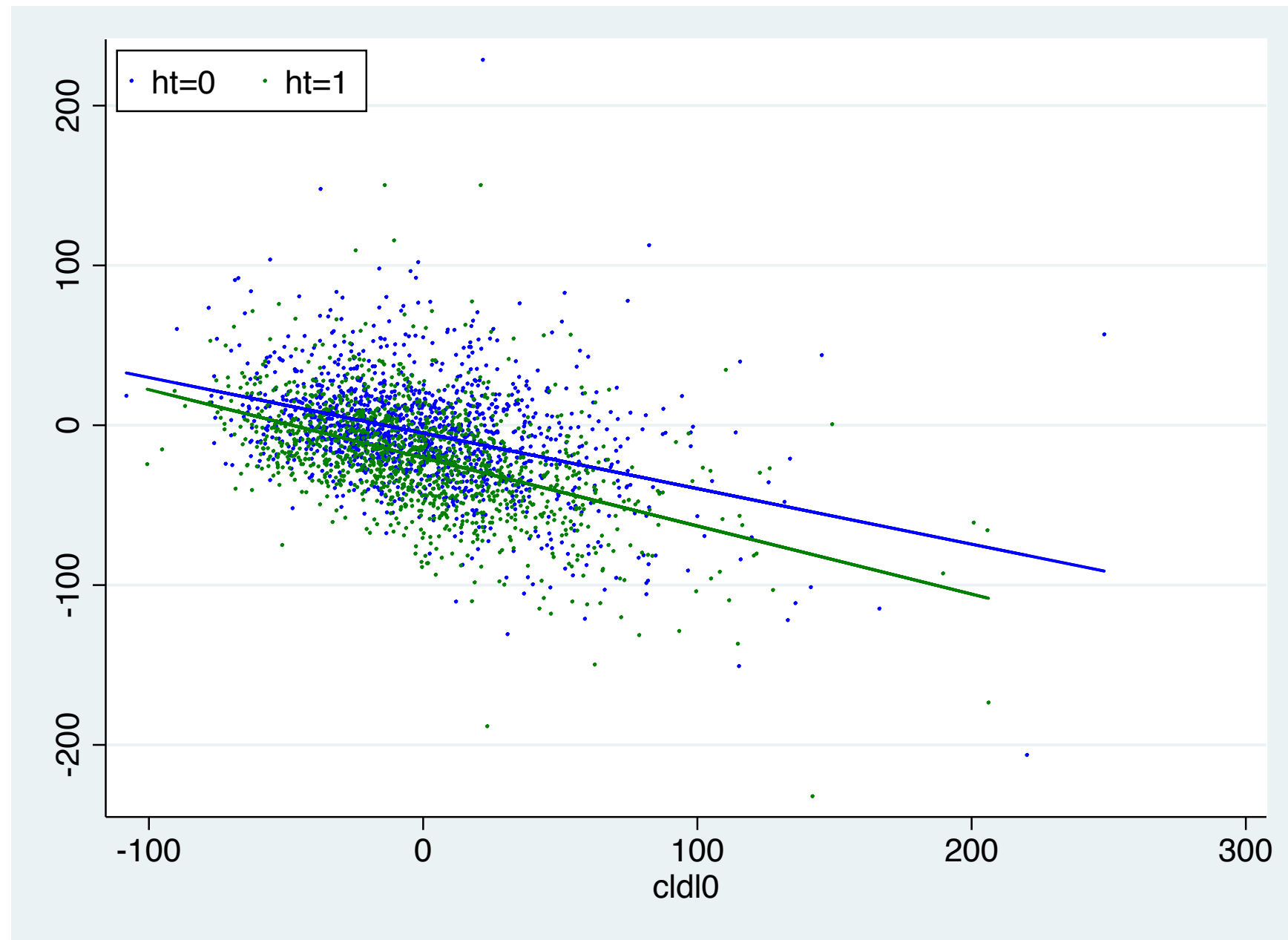
| assignment to<br>hormone therapy | Freq. | Percent | Cum.   |
|----------------------------------|-------|---------|--------|
| placebo                          | 1,383 | 50.05   | 50.05  |
| hormone therapy                  | 1,380 | 49.95   | 100.00 |
| Total                            | 2,763 | 100.00  |        |

. regress ldlch ht##c.cldl0 (ldlch is change from baseline in LDL level, cldl0 is centered baseline level)

| Source          | SS         | df        | MS         | Number of obs = 2597   |                      |           |  |
|-----------------|------------|-----------|------------|------------------------|----------------------|-----------|--|
| Model           | 721218.969 | 3         | 240406.323 | F( 3, 2593) = 258.81   |                      |           |  |
| Residual        | 2408575.51 | 2593      | 928.876015 | Prob > F = 0.0000      |                      |           |  |
| Total           | 3129794.48 | 2596      | 1205.62191 | R-squared = 0.2304     |                      |           |  |
|                 |            |           |            | Adj R-squared = 0.2295 |                      |           |  |
|                 |            |           |            | Root MSE = 30.477      |                      |           |  |
| ldlch           | Coef.      | Std. Err. | t          | P> t                   | [95% Conf. Interval] |           |  |
| ht              |            |           |            |                        |                      |           |  |
| hormone therapy | -15.47703  | 1.196246  | -12.94     | 0.000                  | -17.82273            | -13.13134 |  |
| cldl0           | -.3477064  | .0225169  | -15.44     | 0.000                  | -.3918593            | -.3035534 |  |
| ht#c.cldl0      |            |           |            |                        |                      |           |  |
| hormone therapy | -.0786871  | .0316365  | -2.49      | 0.013                  | -.1407226            | -.0166517 |  |
| _cons           | -4.888739  | .8408392  | -5.81      | 0.000                  | -6.537523            | -3.239955 |  |

*Interpretation:* among post-menopausal women with CHD, HT induces greater absolute reductions in LDL among women with higher baseline levels

# Effects of hormone therapy (ht) on change in LDL (ldlch)



# Effects of hormone therapy (ht) on change in LDL (ldlch)

\* Use of margins to estimate effect of HT at diff. levels of (centered) baseline LDL  
\* Effect among indiv. at mean, and at 25 and 50 units more than the mean base. level  
. margins ht, at(cldl0=(0 25 50))

Adjusted predictions

Number of obs = 2597

Model VCE : OLS

Expression : Linear prediction, predict()

1.\_at : cldl0 = 0  
2.\_at : cldl0 = 25  
3.\_at : cldl0 = 50

|                   |  | Delta-method |           |        |       |                      |
|-------------------|--|--------------|-----------|--------|-------|----------------------|
|                   |  | Margin       | Std. Err. | t      | P> t  | [95% Conf. Interval] |
| _at#ht            |  |              |           |        |       |                      |
| 1#placebo         |  | -4.888739    | .8408392  | -5.81  | 0.000 | -6.537523 -3.239955  |
| 1#hormone therapy |  | -20.36577    | .8508778  | -23.94 | 0.000 | -22.03424 -18.6973   |
| 2#placebo         |  | -13.5814     | 1.017519  | -13.35 | 0.000 | -15.57663 -11.58617  |
| 2#hormone therapy |  | -31.02561    | 1.017472  | -30.49 | 0.000 | -33.02075 -29.03047  |
| 3#placebo         |  | -22.27406    | 1.41331   | -15.76 | 0.000 | -25.04539 -19.50273  |
| 3#hormone therapy |  | -41.68545    | 1.401366  | -29.75 | 0.000 | -44.43336 -38.93754  |

*Interpretation:* estimated effect of HT increases with higher baseline levels

# HT and percent change LDL

```
. gen ldlpctch = (ldlch / ldl0) * 100
. regress ldlpctch ht##c.cldl0 (ldlpctch is % change from baseline in LDL level)
```

| Source   | SS         | df   | MS         | Number of obs = 2597 |   |        |  |
|----------|------------|------|------------|----------------------|---|--------|--|
| Model    | 233394.163 | 3    | 77798.0545 | F( 3, 2593)          | = | 165.33 |  |
| Residual | 1220171.82 | 2593 | 470.563756 | Prob > F             | = | 0.0000 |  |
|          |            |      |            | R-squared            | = | 0.1606 |  |
|          |            |      |            | Adj R-squared        | = | 0.1596 |  |
| Total    | 1453565.98 | 2596 | 559.925263 | Root MSE             | = | 21.692 |  |

| ldlpctch        | Coef.     | Std. Err. | t      | P> t  | [95% Conf. Interval] |           |
|-----------------|-----------|-----------|--------|-------|----------------------|-----------|
| ht              |           |           |        |       |                      |           |
| hormone therapy | -10.79035 | .8514335  | -12.67 | 0.000 | -12.45991            | -9.120789 |
| cldl0           | -.2162437 | .0160265  | -13.49 | 0.000 | -.2476697            | -.1848176 |
| ht#c.cldl0      |           |           |        |       |                      |           |
| hormone therapy | .0218767  | .0225175  | 0.97   | 0.331 | -.0222773            | .0660307  |
| _cons           | -1.284977 | .5984713  | -2.15  | 0.032 | -2.458507            | -.1114465 |

*Interpretation: percent reductions in LDL induced by HT do not clearly vary by baseline level*

# Data dredging for interaction in multi-predictor models

- With 10 predictors
  - 45 possible 2-way interactions – 120 possible 3-way interactions
- Considering them all inflates type-I error, uncovers trivial interactions in big datasets
- If goal is outcome prediction, criteria for considering interactions differs (lecture 7)

# Focused data dredging

- Focus on plausible interactions (Harrell et al., Stat Med, 1996;15:361-87)
  - treatment and severity of disease
  - treatment and calendar time
  - age and risk factors
  - measurement and state during measurement (e.g. l. ventric. function and stress level)
  - *with primary predictor*

# If we hadn't centered BMI ...

- In BMI and PA example, if we did not center BMI
  - $\beta_0$ : average SBP if inactive, BMI = 0
  - $\beta_1$ : effect of PA if BMI = 0
  - estimated effect of PA at any plausible BMI requires `lincom`
- With centered BMI:
  - $\beta_0$  and  $\beta_1$  refer to participants with average BMI

# If we hadn't centered BMI ...

```
. regress SBP i.active##c.BMI
```

| Source   | SS         | df   | MS         | Number of obs = 2758   |  |  |  |
|----------|------------|------|------------|------------------------|--|--|--|
| Model    | 4579.65484 | 3    | 1526.55161 | F( 3, 2754) = 4.24     |  |  |  |
| Residual | 992671.256 | 2754 | 360.447079 | Prob > F = 0.0054      |  |  |  |
| Total    | 997250.911 | 2757 | 361.715963 | R-squared = 0.0046     |  |  |  |
|          |            |      |            | Adj R-squared = 0.0035 |  |  |  |
|          |            |      |            | Root MSE = 18.985      |  |  |  |

| SBP          | Coef.     | Std. Err. | t     | P> t  | [95% Conf. Interval] |          |
|--------------|-----------|-----------|-------|-------|----------------------|----------|
| 1.active     | -7.102228 | 4.251184  | -1.67 | 0.095 | -15.43806            | 1.233603 |
| BMI          | .0512451  | .1138269  | 0.45  | 0.653 | -.1719496            | .2744397 |
| active#c.BMI |           |           |       |       |                      |          |
| 1            | .2293768  | .1406966  | 1.63  | 0.103 | -.0465047            | .5052584 |
| _cons        | 134.1237  | 3.544294  | 37.84 | 0.000 | 127.174              | 141.0735 |

- Average SBP among inactive with BMI = 0  $kg/m^2$  is 134.1 *mmHg*
- At BMI of 0  $kg/m^2$ , PA reduces SBP by 7.1 *mmHg*

# Jeff's (Epi 203) criteria for reporting interaction

- *Substantively trivial* (e.g., RRs of 3.2 and 3.5 for two levels of a binary interacting factor): ignore, regardless of  $p$ -value for interaction
- *Substantively important* (e.g., RRs of 2.0 and 3.5): report if interaction  $p$ -value  $< 0.10$
- *Qualitative interaction* (e.g., RRs of -2.0 and 2.0): report if interaction  $p$ -value  $< 0.15$

# Possible additional criteria for reporting

- Trying to prove something (e.g., show that BiDil should be approved for heart failure in African American patients):
  - need strong evidence, especially for ad hoc finding, possibly penalized for multiple comparisons
- Weak evidence for interaction that undermines your controversial hypothesis:
  - good to report as a sensitivity analysis

# Further recommendations<sup>1</sup>

- Not enough to show a within-subgroup effect; need to show that effects significantly differ by subgroup
- Focus on a priori interactions, those with primary predictor
- Evaluate interactions for strength, biological plausibility
- Identify post hoc interactions
- In ambiguous cases, present both overall and stratified results

<sup>1</sup> Wang et al., *NEJM*, 2008;357(21):2189-94

# Summary: Interaction

- Summary: Interaction
- Interaction: association of one predictor with outcome modified by another predictor
- Modeled using product terms, made automatically by Stata `##` operator
- Subgroup-specific regression lines for continuous predictor not parallel
- Centering makes “main effects” easier to interpret
- Power not always low, but can be

# Confounding, mediation, and interaction

- Confounders are a cause of both outcome and primary predictor, or surrogates for such a cause
- Potential confounders must be associated with predictor and independently with the outcome
- Confounding can account for the some or all of the unadjusted association between a predictor and an outcome

# Mediation vs confounding

- Mediators are on causal pathway from predictor to outcome
- In multi-predictor models, mediation and confounding behave alike, must be distinguished on substantive grounds
- Some problems too complex for this paradigm:
  - LDL both confounds, mediates statin effects on CHD
- You have the substantive expertise to draw the DAG

# Interaction vs confounding/mediation

- Interaction vs confounding/mediation
- Important to distinguish between:
  - change in the coefficient for a primary predictor in two models *with and without adjustment for a confounder or mediator*, which we sometimes use to assess confounding and/or mediation
  - differences in the estimated effect of a primary predictor *across strata defined by an effect modifier* in a single model that includes interaction term(s)

# Confounding, mediation, or interaction?

1. Is alendronate effect on BMD reduced in vitamin-D deficient patients?
2. Is alendronate effect on vertebral fracture explained by changes in BMD?
3. Is HT effect on CHD explained by a healthier lifestyle?
4. Are beta-blockers less effective among African Americans?
5. Is diabetes a stronger risk factor for CHD events in women than in men?