Simulation of Simpson's Paradox With Palmer Penguin Data

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Background

Simpson's paradox occurs when a bivariate association is reversed in a multivariate model. This example using the Palmer Penguins Data was inspired by a tweet by Andrew Heiss.



Figure 1: Palmer Penguins Illustration from @allison_horst

To begin, a little definition of penguin terminology is in order. Note the diagram defining culmen depth below.



Figure 2: Culmen Depth from @allison_horst

Setup

- . clear all
- . cd "/Users/agrogan/Desktop/newstuff/simpsonsparadox"/Users/agrogan/Desktop/newstuff/simpsonsparadox
- . use "penguins.dta"

Bivariate

. twoway (scatter culmen_depth_mm body_mass_g) ///

```
> (lfit culmen_depth_mm body_mass_g), ///
> title("Culmen Depth by Body Mass") ///
> caption("Palmer Penguin Data") ///
> scheme(michigan)
. graph export mygraph1.png, width(1000) replace
(file /Users/agrogan/Desktop/newstuff/simpsonsparadox/mygraph1.png written in PNG format)
```

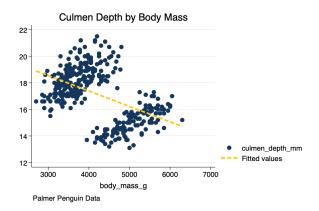


Figure 3: Scatterplot and Linear Fit

. regress cul	men_depth_mm b	ody_mass_g					
Source	SS df		MS	Number of obs		=	342
				F(1,	340)	=	97.41
Model	296.15994	1	296.15994	Prob	> F	=	0.0000
Residual	1033.67459	340	3.04021939	R-sq	R-squared		0.2227
				Adj	R-squared	=	0.2204
Total	1329.83453	341	3.89980801	Root	MSE	=	1.7436
culmen_dep_m	Coef.	Std. Err.	t	P> t	[95% Cd	onf.	Interval]
body_mass_g _cons	0011621 22.03395	.0001177		0.000 0.000	001393 21.0433		0009305 23.02455

Multivariate

```
. twoway (scatter culmen_depth_mm body_mass_g) ///
> (lfit culmen_depth_mm body_mass_g), ///
> by(species, title("Culmen Depth by Body Mass") caption("Palmer Penguin Data")) ///
> scheme(michigan)
. graph export mygraph2.png, width(1000) replace
(file /Users/agrogan/Desktop/newstuff/simpsonsparadox/mygraph2.png written in PNG format)
```

The association is reversed when we take into account multiple variables.

. regress	culme	en_depth_mm body	y_mass_g	species			
Sou	rce	SS	df	MS	Number of obs	=	342
					F(2, 339)	=	225.41
Mo	del	759.047284	2	379.523642	Prob > F	=	0.0000
Resid	ual	570.787248	339	1.6837382	R-squared	=	0.5708
					Adj R-squared	=	0.5683
То	tal	1329.83453	341	3.89980801	Root MSE	=	1.2976
	- '						

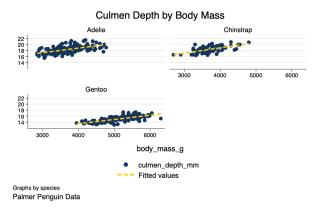


Figure 4: Scatterplot and Linear Fit

culmen_dep_m	Coef.	Std. Err.	t	P> t	[95% Conf.	Interval]
body_mass_g	.0004877	.0001326	3.68	0.000	.0002269	.0007485
species	-1.974985	.1191142	-16.58	0.000	-2.209281	-1.740689
_cons	18.89014	.4200224	44.97	0.000	18.06396	19.71631