

# Interactions

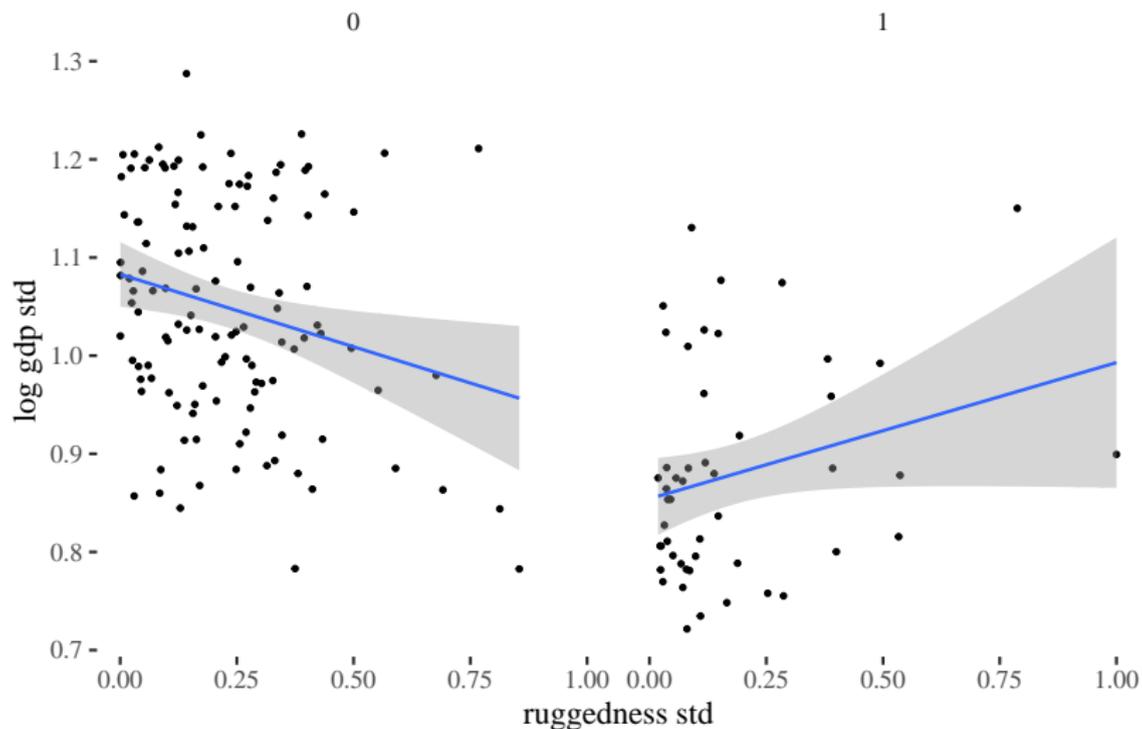
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## What is interaction/moderation?

The importance of one predictor depends upon another predictor. Interaction is a kind of conditioning, a way of allowing posterior distributions of parameters to be conditional on further aspects of the data.

# African economy and bad geography

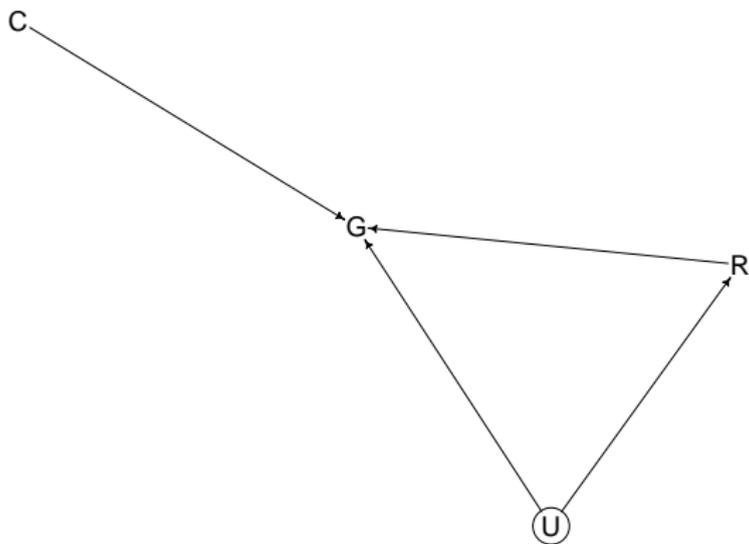
## Africa and impact of ruggedness



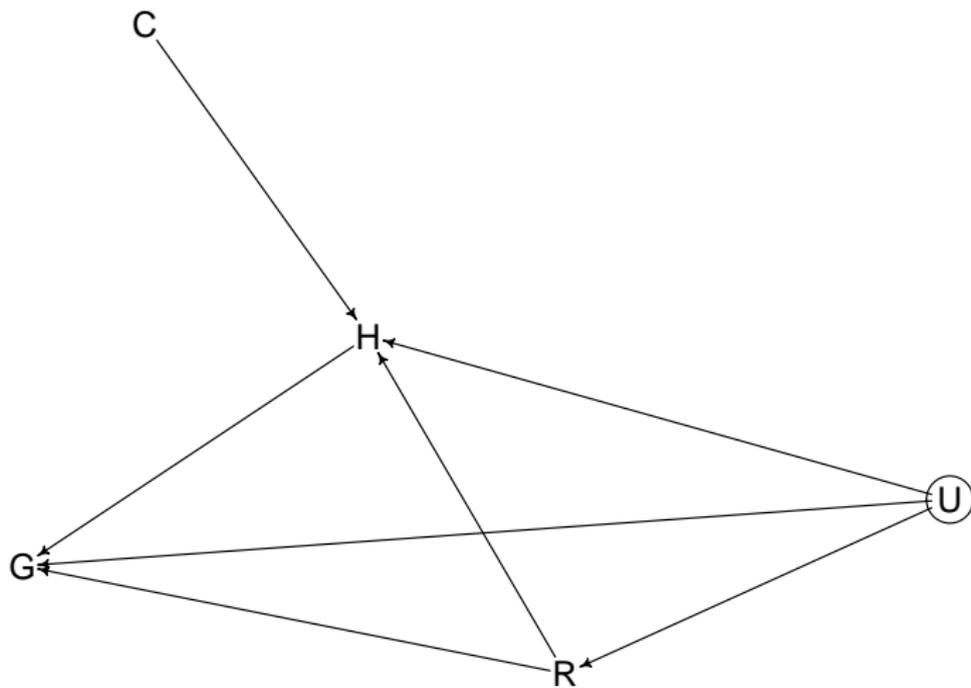
# African economy and bad geography

DAG fails to represent interaction

$$G = f(R, C)$$



## African economy and bad geography



## African economy and bad geography

```
africaNoInteraction <- quap(  
  alist(  
    log_gdp_std ~ dnorm( mu , sigma ) ,  
    mu <- a + b*( rugged_std - 0.215 ) ,  
    a ~ dnorm( 1 , .1 ) ,  
    b ~ dnorm( 0 , .3 ) ,  
    sigma ~ dexp( 1 )  
  ) , data=dd )
```

# African economy and bad geography

```
precis(africaNoInteraction)
```

```
##           mean          sd          5.5%          94.5%  
## a      0.999998585 0.010412558  0.98335731 1.01663986  
## b      0.001987997 0.054796473 -0.08558735 0.08956334  
## sigma 0.136505158 0.007397203  0.12468300 0.14832732
```

## African economy and bad geography

```
dd$cid <- ifelse( dd$cont_africa==1 , 1 , 2 )

africaJustIndicator <- quap(
  alist(
    log_gdp_std ~ dnorm( mu , sigma ) ,
    mu <- a[cid] + b*( rugged_std - 0.215 ) ,
    a[cid] ~ dnorm( 1 , 0.1 ) ,
    b ~ dnorm( 0 , 0.3 ) ,
    sigma ~ dexp( 1 )
  ) , data=dd )
```

# African economy and bad geography

```
compare(africaNoInteraction, africaJustIndicator)
```

```
##                WAIC      SE    dWAIC      dSE    pWAIC      weight
## africaJustIndicator -252.1934 15.30424  0.00000      NA 4.272339 1.000000e+00
## africaNoInteraction -188.8633 13.24558 63.33009 15.08302 2.642283 1.770299e-14
```

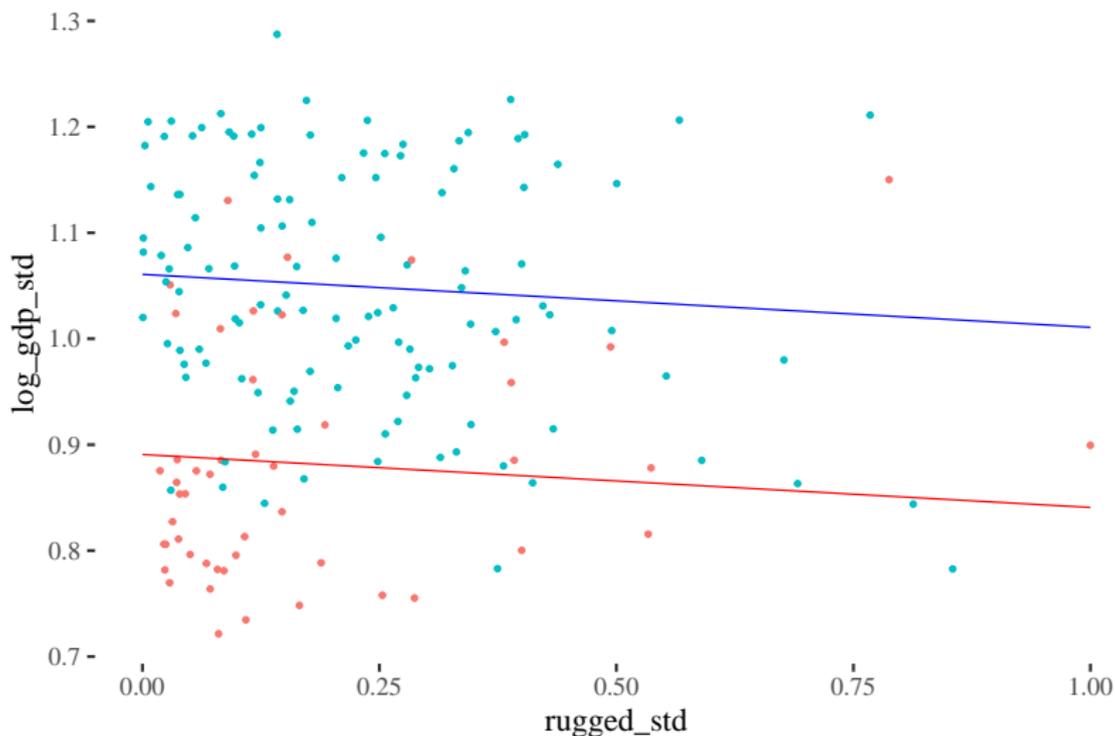
## African economy and bad geography

```
precis( africaJustIndicator , depth=2 )
```

##		mean	sd	5.5%	94.5%
##	a[1]	0.88040635	0.015937612	0.8549350	0.90587773
##	a[2]	1.04916045	0.010185961	1.0328813	1.06543958
##	b	-0.04649695	0.045688524	-0.1195160	0.02652214
##	sigma	0.11239189	0.006091687	0.1026562	0.12212758

# African economy and bad geography

Model with indicator



## African economy and bad geography

```
africaInteraction <- quap(  
  alist(  
    log_gdp_std ~ dnorm( mu , sigma ) ,  
    mu <- a[cid] + b[cid]*( rugged_std - 0.215 ) ,  
    a[cid] ~ dnorm( 1 , 0.1 ) ,  
    b[cid] ~ dnorm( 0 , 0.3 ) ,  
    sigma ~ dexp( 1 )  
  ) , data=dd )
```

# African economy and bad geography

```
precis( africaInteraction , depth=2 )
```

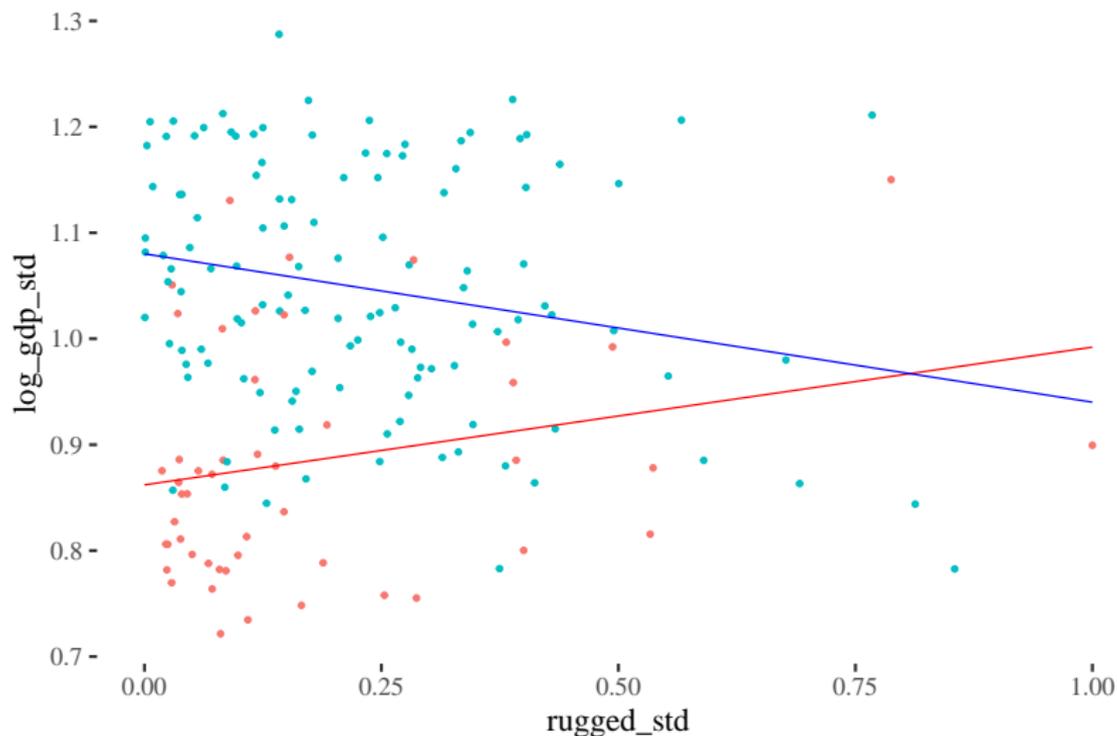
```
##           mean          sd      5.5%      94.5%
## a[1]  0.8865638 0.015675774  0.86151085  0.91161668
## a[2]  1.0505694 0.009936658  1.03468870  1.06645009
## b[1]  0.1325055 0.074204803  0.01391188  0.25109910
## b[2] -0.1425764 0.054749677 -0.23007697 -0.05507585
## sigma 0.1094947 0.005935376  0.10000881  0.11898056
```

```
compare(africaNoInteraction, africaJustIndicator, africaInteraction)
```

```
##           WAIC          SE      dWAIC      dSE      pWAIC
## africaInteraction -259.0097 15.21834  0.000000      NA 5.251338
## africaJustIndicator -252.3183 15.41667  6.691372  6.798868 4.254228
## africaNoInteraction -188.6429 13.26207 70.366785 15.457349 2.725360
##           weight
## africaInteraction 9.659633e-01
## africaJustIndicator 3.403671e-02
## africaNoInteraction 5.069995e-16
```

# African economy and bad geography

Model with interaction



# Tulips

```
## 'data.frame':    27 obs. of  4 variables:  
## $ bed      : Factor w/ 3 levels "a","b","c": 1 1 1 1 1 1 1 1 1 1 2 ...  
## $ water    : int   1 1 1 2 2 2 3 3 3 1 ...  
## $ shade    : int   1 2 3 1 2 3 1 2 3 1 ...  
## $ blooms: num   0 0 111 183.5 59.2 ...
```

In the absence of light, no amount of water will help

# Tulips

```
d$blooms_std <- d$blooms / max(d$blooms)
d$water_cent <- d$water - mean(d$water)
d$shade_cent <- d$shade - mean(d$shade)
```

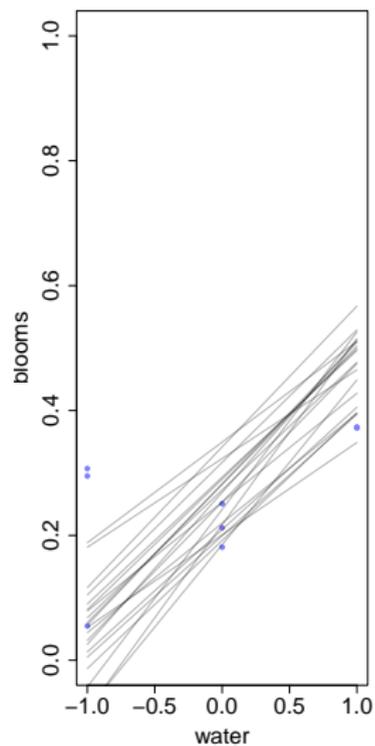
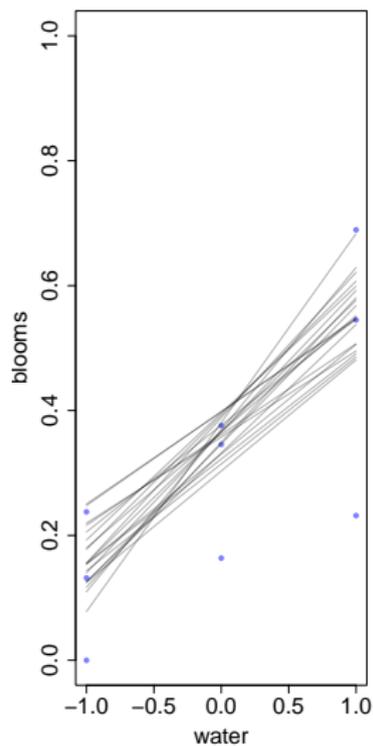
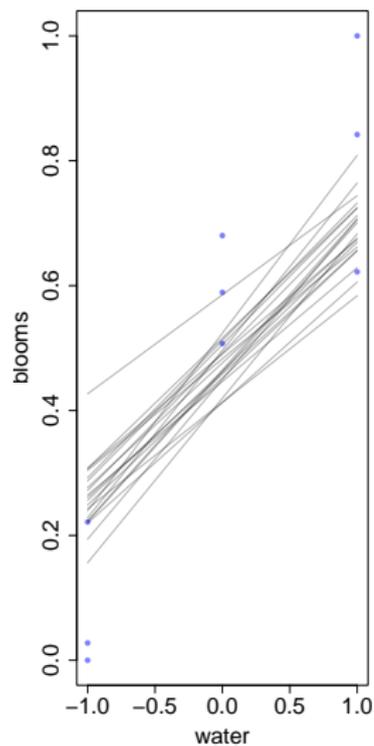
# Tulips

```
tulipsNoInteraction <- quap(  
  alist(  
    blooms_std ~ dnorm( mu , sigma ) ,  
    mu <- a + bw*water_cent + bs*shade_cent ,  
    a ~ dnorm( 0.5 , 0.25 ) ,  
    bw ~ dnorm( 0 , 0.25 ) ,  
    bs ~ dnorm( 0 , 0.25 ) ,  
    sigma ~ dexp( 1 )  
  ) , data=d )
```

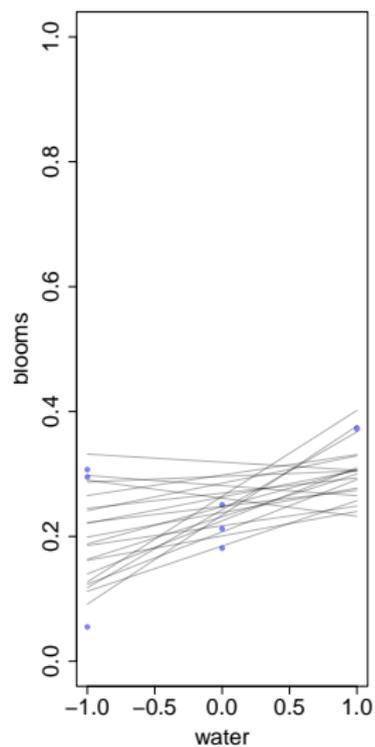
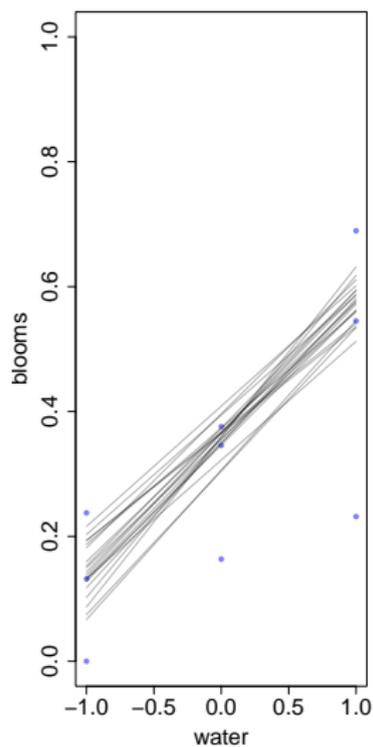
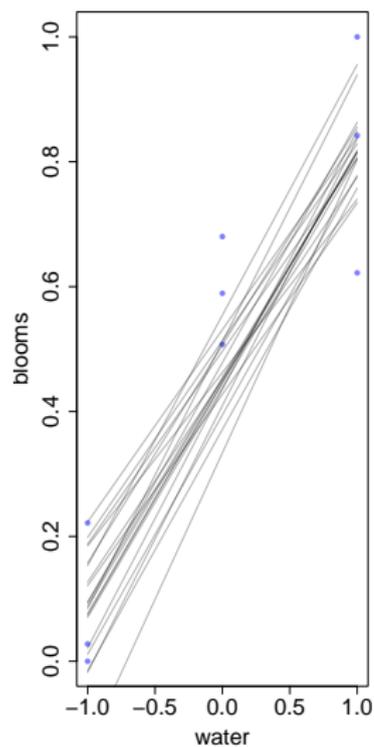
# Tulips

```
tulipsInteraction <- quap(  
  alist(  
    blooms_std ~ dnorm( mu , sigma ) ,  
    mu <- a + bw*water_cent + bs*shade_cent +  
      bws*water_cent*shade_cent ,  
    a ~ dnorm( 0.5 , 0.25 ) ,  
    bw ~ dnorm( 0 , 0.25 ) ,  
    bs ~ dnorm( 0 , 0.25 ) ,  
    bws ~ dnorm( 0 , 0.25 ) ,  
    sigma ~ dexp( 1 )  
  ) , data=d )
```

# Tulips: no interaction



# Tulips: with interaction



# Tulips

```
compare(tulipsInteraction, tulipsNoInteraction)
```

```
##                WAIC          SE    dWAIC      dSE    pWAIC      weight
## tulipsInteraction  -20.44760 11.309416 0.000000      NA 7.390437 0.98700314
## tulipsNoInteraction -11.78767  9.052902 8.659931 8.831736 5.491489 0.01299686
```