

Model draft

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model 1

```
mod1 <- lmer(c_weight~as.factor(fuel_bin)+w_age+bmi+as.factor(gender)+as.factor(residence)+ as.factor(w

c(summary(mod1)$coefficient[2,1:2], confint(mod1, method="Wald")[4,1:2],nobs(mod1),"birthweight",
  paste0(round(summary(mod1)$coefficient[2,1:2][1],2)," (",
  round(confint(mod1, method="Wald")[4,1:2][1],2)," ",
  round(confint(mod1, method="Wald")[4,1:2][2],2),")")
```

```
##           Estimate           Std. Error           2.5 %
##      "22.9431895860758"      "39.7648796496601"      "-54.9945423768277"
##           97.5 %
##      "100.880921548979"           "2960"           "birthweight"
##
## "22.94 (-54.99, 100.88)"
```

with anemic

```
mod_ane <- lmer(c_weight~as.factor(fuel_bin)+w_age+bmi+as.factor(gender)+as.factor(residence)+ as.factor

paste0(round(summary(mod_ane)$coefficient[2,1:2][1],2)," (",
  round(confint(mod_ane, method="Wald")[4,1:2][1],2)," ",
  round(confint(mod_ane, method="Wald")[4,1:2][2],2),")")
```

```
## [1] "32.38 (-46.84, 111.59)"
```

using only documented weights

```
doc_weight <- filter(newdata, document == 1)
mod_doc <- lmer(c_weight~as.factor(fuel_bin)+w_age+bmi+as.factor(gender)+as.factor(residence)+
  as.factor(wealth)+as.factor(education)+as.factor(marital_s)+(1|region),na.action=na.omit, data =

paste0(round(summary(mod_doc)$coefficient[2,1:2][1],2)," (",
  round(confint(mod_doc, method="Wald")[4,1:2][1],2)," ",
  round(confint(mod_doc, method="Wald")[4,1:2][2],2),")")
```

```
## [1] "67.58 (-19.62, 154.77)"
```

```
##anemia
```

```
mod_doc_a <- lmer(c_weight~as.factor(fuel_bin)+w_age+bmi+as.factor(gender)+as.factor(residence)+as.factor(wealth)+as.factor(education)+as.factor(marital_s)+(1|region),na.action=na.omit, data = doc_a)
```

```
paste0(round(summary(mod_doc_a)$coefficient[2,1:2][1],2)," (",  
round(confint(mod_doc_a, method="Wald")[4,1:2][1],2)," ",  
round(confint(mod_doc_a, method="Wald")[4,1:2][2],2),")")
```

```
## [1] "78.47 (-9.76, 166.71)"
```

```
##GLM
```

```
modglm <- glmer(bin_weight~as.factor(fuel_bin)+w_age+bmi+as.factor(gender)+as.factor(residence)+  
as.factor(wealth)+as.factor(education)+as.factor(marital_s)+(1|region),na.action=na.omit, data =  
control = glmerControl(optimizer = "bobyqa",nAGQ=9))
```

```
paste0(round(exp(summary(modglm)$coefficient[2,1:2][1]),2)," (",  
round(exp(confint(modglm, method="Wald")[4,1:2][1]),2)," ",  
round(exp(confint(modglm, method="Wald")[4,1:2][2]),2),")")
```

```
## [1] "0.68 (0.97, 1.01)"
```

Not significant

```
##documented
```

```
dglm <- glmer(bin_weight~as.factor(fuel_bin)+w_age+bmi+as.factor(gender)+as.factor(residence)+  
as.factor(wealth)+as.factor(education)+as.factor(marital_s)+(1|region),na.action=na.omit, data =  
control = glmerControl(optimizer = "bobyqa",nAGQ=9))
```

```
##bmi categorised, take out smoke, married be binary, model per residence status, poorest vs richest
```

```
paste0(round(exp(summary(dglm)$coefficient[2,1:2][1]),2)," (",  
round(exp(confint(dglm, method="Wald")[4,1:2][1]),2)," ",  
round(exp(confint(dglm, method="Wald")[4,1:2][2]),2),")")
```

```
## [1] "0.55 (0.96, 1.01)"
```

Not significant