## Comparing the significance of prot\_000000001830 and prot\_000000000877 to OTU\_11

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## The significance of prot\_00000000877 to OTU\_11

```
log_transform = function(x) {
   constant = min(x[x > 0])/2.
   x = x + constant
   x = x/sum(x)
   log(x)
}

otu_log_mat = apply(preprocess_mat(infiles[4]), 1, log_transform)
genes_log_mat = apply(preprocess_mat(infiles[1]), 1, log_transform)

prot_000000000877 = genes_log_mat['prot_000000000877',]
otu_11 = otu_log_mat['OTU_11',]

mapping_maria = load_data_table('mapping_maria_BL.txt')

dependent = as.numeric(mapping_maria$Persistence)
lm_prot_000000000877 = lm(dependent~prot_000000000877)
lm_otu_11 = lm(dependent~otu_11)

summary(lm_prot_000000000877)
```

```
##
## Call:
## lm(formula = dependent ~ prot_00000000877)
##
## Residuals:
               1Q Median
                               3Q
## -0.4919 -0.1997 -0.0073 0.1252 0.7298
##
## Coefficients:
##
                    Estimate Std. Error t value Pr(>|t|)
                                0.2680 -1.121
## (Intercept)
                     -0.3003
                                                   0.276
## prot_00000000877 -0.1870
                                 0.0293 -6.381 3.17e-06 ***
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.2896 on 20 degrees of freedom
## Multiple R-squared: 0.6706, Adjusted R-squared: 0.6542
## F-statistic: 40.72 on 1 and 20 DF, p-value: 3.165e-06
```

```
##
## Call:
## lm(formula = dependent ~ otu_11)
##
## Residuals:
##
      Min
             1Q Median
                           ЗQ
                                 Max
## -0.4344 -0.3039 -0.1407 0.3911 1.0604
##
## Coefficients:
##
            Estimate Std. Error t value Pr(>|t|)
## (Intercept) 0.70757
                     0.27592 2.564
                                       0.0185 *
                       0.05171 -2.528 0.0200 *
## otu_11
           -0.13071
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.4392 on 20 degrees of freedom
## Multiple R-squared: 0.2421, Adjusted R-squared: 0.2042
## F-statistic: 6.389 on 1 and 20 DF, p-value: 0.02001
coxtest(lm_prot_000000000877, lm_otu_11)
## Cox test
##
## Model 1: dependent ~ prot_00000000877
## Model 2: dependent ~ otu 11
                Estimate Std. Error z value Pr(>|z|)
##
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
jtest(lm_prot_000000000877, lm_otu_11)
## J test
##
## Model 1: dependent ~ prot_00000000877
## Model 2: dependent ~ otu_11
                Estimate Std. Error t value Pr(>|t|)
## M1 + fitted(M2) 0.48557 0.26224 1.8516 0.07969 .
## M2 + fitted(M1) 0.89977
                          0.15757 5.7103 1.669e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
The significance of prot 00000001830 to OTU 11
```

summary(lm\_otu\_11)

prot\_00000001830 = genes\_log\_mat['prot\_000000001830',]

```
lm_prot_00000001830 = lm(dependent~prot_00000001830)
summary(prot_00000001830)
     Min. 1st Qu. Median
                          Mean 3rd Qu.
## -13.830 -9.577 -7.393 -8.261 -6.772 -3.853
coxtest(lm_prot_00000001830, lm_otu_11)
## Cox test
##
## Model 1: dependent ~ prot_00000001830
## Model 2: dependent ~ otu_11
                Estimate Std. Error z value Pr(>|z|)
## fitted(M2) ~ M1 -7.8107 0.57436 -13.5991 < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
jtest(lm_prot_00000001830, lm_otu_11)
## J test
##
## Model 1: dependent ~ prot_00000001830
## Model 2: dependent ~ otu_11
                Estimate Std. Error t value Pr(>|t|)
## M1 + fitted(M2) 0.62513 0.30641 2.0401 0.0554847 .
## M2 + fitted(M1) 0.87495 0.20937 4.1790 0.0005092 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```