

Washc1—Q8VDD8

Tyler W. A. Bradshaw

October 15, 2020

marginal (fixed effects) and conditional (total variance) R2:

R2m	R2c
0.9232196	0.9298347
R2m	R2c
0.8844438	0.9266716
R2m	R2c
0.7003719	0.8813102

Table 1: Abundance (1—Mixture) + Condition

		<i>Dependent variable:</i>	
		Abundance	
ConditionControl.F4		−0.925***	(0.220)
ConditionControl.F5		−0.605***	(0.220)
ConditionControl.F6		−0.307	(0.220)
ConditionControl.F7		−0.260	(0.220)
ConditionControl.F8		−0.473**	(0.220)
ConditionControl.F9		−0.692***	(0.220)
ConditionMutant.F10		−2.232***	(0.220)
ConditionMutant.F4		−2.707***	(0.220)
ConditionMutant.F5		−2.345***	(0.220)
ConditionMutant.F6		−2.274***	(0.220)
ConditionMutant.F7		−2.117***	(0.220)
ConditionMutant.F8		−2.343***	(0.220)
ConditionMutant.F9	2	−2.358***	(0.220)
Constant		7.862***	(0.163)
Observations		42	
Log Likelihood		−11.538	
Akaike Inf. Crit.		55.077	
Bayesian Inf. Crit.		82.880	

Notes: \* = 0.1, \*\* = 0.05, \*\*\* = 0.01

Table 2: Abundance  $\sim 0 + (1 - \text{BioFraction}) + \text{Genotype}$ 

	<i>Dependent variable:</i>
	Abundance
GenotypeControl	7.396*** (0.098)
GenotypeMutant	5.523*** (0.098)
Observations	42
Log Likelihood	-12.350
Akaike Inf. Crit.	32.700
Bayesian Inf. Crit.	39.651
<i>Note:</i>	*p<0.1; **p<0.05; ***p<0.01

Table 3: Abundance  $\sim 0 + (1 - \text{BioFraction}) + (1 - \text{Protein}) + \text{Genotype}$ 

	<i>Dependent variable:</i>
	Abundance
GenotypeControl	7.319*** (0.151)
GenotypeMutant	5.953*** (0.151)
Observations	210
Log Likelihood	-52.214
Akaike Inf. Crit.	114.427
Bayesian Inf. Crit.	131.163
<i>Note:</i>	*p<0.1; **p<0.05; ***p<0.01