

Washc2—Q6PGL7

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

R2m	R2c
0.9408193	0.9408193
R2m	R2c
0.7627664	0.9331342
R2m	R2c
0.7003719	0.8813102

Table 1: Abundance (1—Mixture) + Condition

		<i>Dependent variable:</i>
		Abundance
ConditionControl.F4		−0.977*** (0.123)
ConditionControl.F5		−0.570*** (0.123)
ConditionControl.F6		−0.216* (0.123)
ConditionControl.F7		−0.164 (0.123)
ConditionControl.F8		−0.264** (0.123)
ConditionControl.F9		−0.503*** (0.123)
ConditionMutant.F10		−1.398*** (0.123)
ConditionMutant.F4		−1.858*** (0.123)
ConditionMutant.F5		−1.528*** (0.123)
ConditionMutant.F6		−1.285*** (0.123)
ConditionMutant.F7		−1.258*** (0.123)
ConditionMutant.F8		−1.292*** (0.123)
ConditionMutant.F9	2	−1.546*** (0.123)
Constant		7.644*** (0.087)
Observations		42
Log Likelihood		5.661
Akaike Inf. Crit.		20.677
Bayesian Inf. Crit.		48.480

Note: * = 0.1, ** = 0.05, *** = 0.01

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

	<i>Dependent variable:</i>
	Abundance
GenotypeControl	7.260*** (0.103)
GenotypeMutant	6.192*** (0.103)
Observations	42
Log Likelihood	5.142
Akaike Inf. Crit.	-2.284
Bayesian Inf. Crit.	4.667
<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