I am basically evaluating Mendel's claim on hybrid production on the basis of his observed counts, using significance level 0.1 by first reporting a table of observed values, expected values, and discrepancies between them and finding which values appear most discrepant. I am also interested in the null distribution for the test statistic here (including its degrees of freedom) to see I would reject the null hypothesis here.

Found at https://library.si.edu/digital-library/book/versucheberpflan00mend (image is also on canvas)

Hypothesized Ratios

Warning: package 'pander' was built under R version 4.2.1

Table 1: Relative ratios under the Mendel's proposal (continued below)

Category NullRatio	ABC 1	ABc 1	AbC 1	Abc 1	aBC	aBc 1	abC 1	abc 1	ABCc 2	AbCc 2
Category NullRatio	aBCc	abCc 2	ABbC		3bc 2	aBbC	aBbc	AaBC 2	AaBc 2	AabC 2
Category NullRatio	Aabc 2	ABbC	c aBb		AaBCc	e Aab(Cc Aa	BbC 4	AaBbc 4	AbBbCc 8

Observed, Expected, and Discrepancies

Table 4: Observed vs. Expected values for Experiment 1 (continued below)

	ABC	ABc	AbC	Abc	aBC	aBc
NullProportion	0.0156	0.0156	0.0156	0.0156	0.0156	0.0156
Expected	9.98	9.98	9.98	9.98	9.98	9.98
Observed	8	14	9	11	8	10
Discrepancy	0.394	1.62	0.0971	0.103	0.394	2.45e-05

	abC	abc	ABCc	AbCc	aBCc	abCc
NullProportion	0.0156	0.0156	0.0312	0.0312	0.0312	0.0312
Expected	9.98	9.98	20	20	20	20
Observed	10	7	22	17	25	20
Discrepancy	2.45 e-05	0.892	0.207	0.441	1.27	4.89 e - 05

	ABbC	ABbc	aBbC	aBbc	AaBC	AaBc
$egin{aligned} \mathbf{NullProportion} \\ \mathbf{Expected} \end{aligned}$	0.0312 20	0.0312 20	0.0312 20	0.0312 20	0.0312 20	0.0312 20

	ABbC	ABbc	aBbC	aBbc	AaBC	AaBc
Observed	15	18	19	24	14	18
Discrepancy	1.24	0.194	0.047	0.814	1.78	0.194

	AabC	Aabc	ABbCc	aBbCc	AaBCc	AabCc
NullProportion	0.0312	0.0312	0.0625	0.0625	0.0625	0.0625
Expected	20	20	39.9	39.9	39.9	39.9
Observed	20	16	45	36	38	40
Discrepancy	4.89e-05	0.789	0.642	0.388	0.094	9.78e-05

	AaBbC	AaBbc	AbBbCc
NullProportion	0.0625	0.0625	0.125
Expected	39.9	39.9	79.9
Observed	49	48	78
Discrepancy	2.06	1.63	0.044

Test Statistic and p-value

[1] 0.9511498

Discussion

The categories with the most discrepancies are AaBbC, AaBC, AaBbc, ABc, aBCc, and ABbC. These all have discrepancies of over 1, meaning the numbers observed for these categories deviate significantly from the expected values. Categories with discrepancies of over 0.5 include abc, aBbc, Aabc, and ABbCc, these values also have abnormally large observed numbers compared to the expected. With 27 categories, there are 26 degrees of freedom. The chi-squared test results in a p-value of 0.9511498, which is very large, so there is a lack of evidence against the null hypothesis, and that the discrepancies are a result of random chance.

Appendix

```
knitr::opts_chunk$set(echo = FALSE)
library(pander)
Category <- c("ABC", "ABc", "AbC", "Abc",</pre>
               "aBC", "aBc", "abC", "abc",
               "ABCc", "AbCc", "aBCc", "abCc",
               "ABbC", "ABbc", "aBbC", "aBbc",
               "AaBC", "AaBc", "AabC", "Aabc",
               "ABbCc", "aBbCc",
               "AaBCc", "AabCc",
               "AaBbC", "AaBbc",
               "AbBbCc")
NullRatio <- c(1, 1, 1, 1,
               1, 1, 1, 1,
               2, 2, 2, 2,
               2, 2, 2, 2,
               2, 2, 2, 2,
               4, 4,
               4, 4,
               4, 4,
               8)
ATable <- rbind(Category, NullRatio)
panderOptions('table.continues', '')
pander(ATable, caption="Relative ratios under the Mendel's proposal")
NullProportion <- NullRatio/sum(NullRatio) # converting ratios to proportions
Observed <- c(8,14,9,11,8,10,10,7,22,17,25,20,15,18,19,24,14,18,20,16,45,36,38,40,49,48,78) # from Mend
Expected <-sum(Observed)*NullProportion # converting proportions to Expected counts
Discrepancy <- (Expected-Observed)^2/Expected</pre>
Table <- rbind(NullProportion,Expected,Observed,Discrepancy)</pre>
pander(Table,
      col.names=Category,
      digits=3.
      caption="Observed vs. Expected values for Experiment 1")
TestStat <- sum(Discrepancy) # chi-squared test statistic</pre>
p_value <- 1-pchisq(TestStat,26) # don't forget the degrees of freedom!
p_value
```