

[1] "Mixed – treated"

Exact binomial test

data: `sum(df.good$Is.Latent)` and `length(df.good$Is.Latent)`

number of successes = 187, number of trials = 415, p-value = 0.9804

alternative hypothesis: true probability of success is greater than 0.5

95 percent confidence interval:

0.4095823 1.0000000

sample estimates:

probability of success

0.4506024

Generalized linear mixed model fit by maximum likelihood (Laplace
Approximation) [glmerMod]
Family: binomial (logit)
Formula: ls.Latent ~ (1 | Patient)
Data: df.good

AIC	BIC	logLik	deviance	df.resid
494.3	502.3	-245.1	490.3	413

Scaled residuals:

Min	1Q	Median	3Q	Max
-1.9719	-0.8059	-0.4718	0.9473	2.1195

Random effects:

Groups	Name	Variance	Std.Dev.
Patient	(Intercept)	1.213	1.101

Number of obs: 415, groups: Patient, 5

Fixed effects:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-0.3812	0.5066	-0.752	0.452

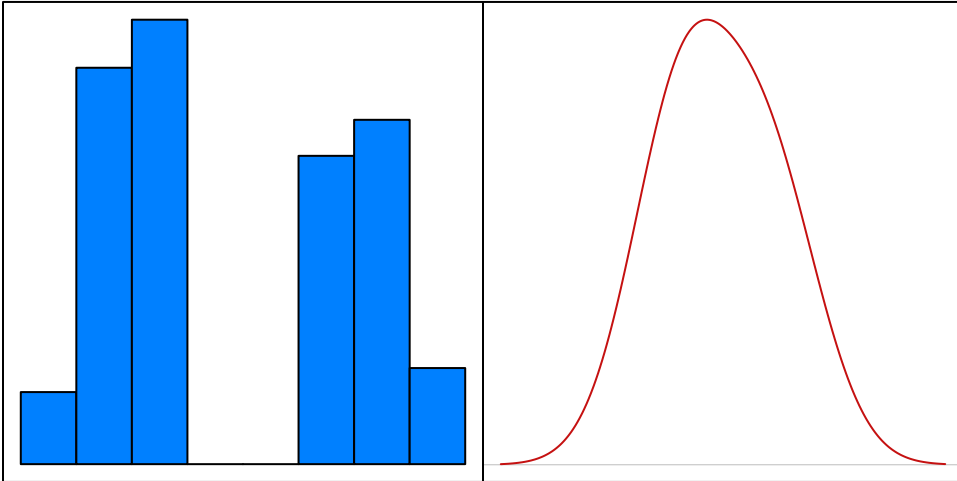
[1] "AIC: 494.275102208077, null AIC: 573.254942284639"

[2] "p-value: 0"

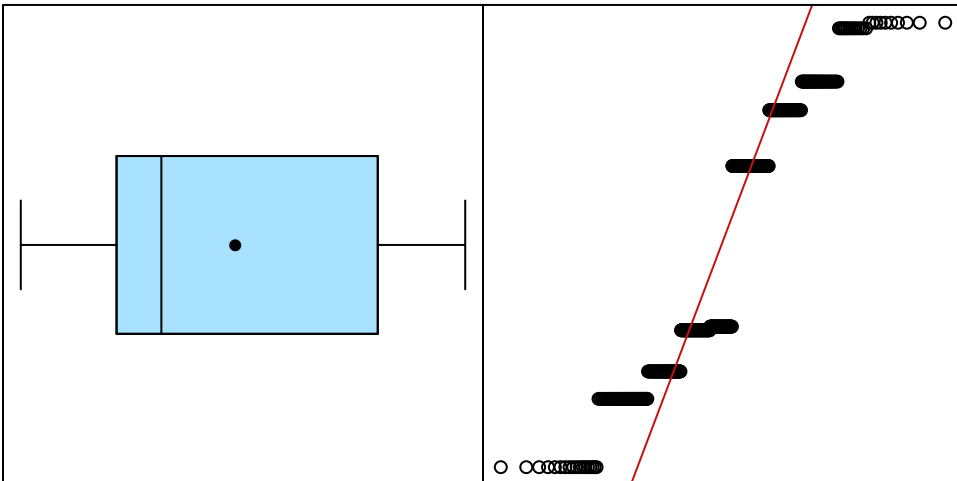
[3] "mean: 0.405846524867213"

EXPLORATORY DATA ANALYSIS

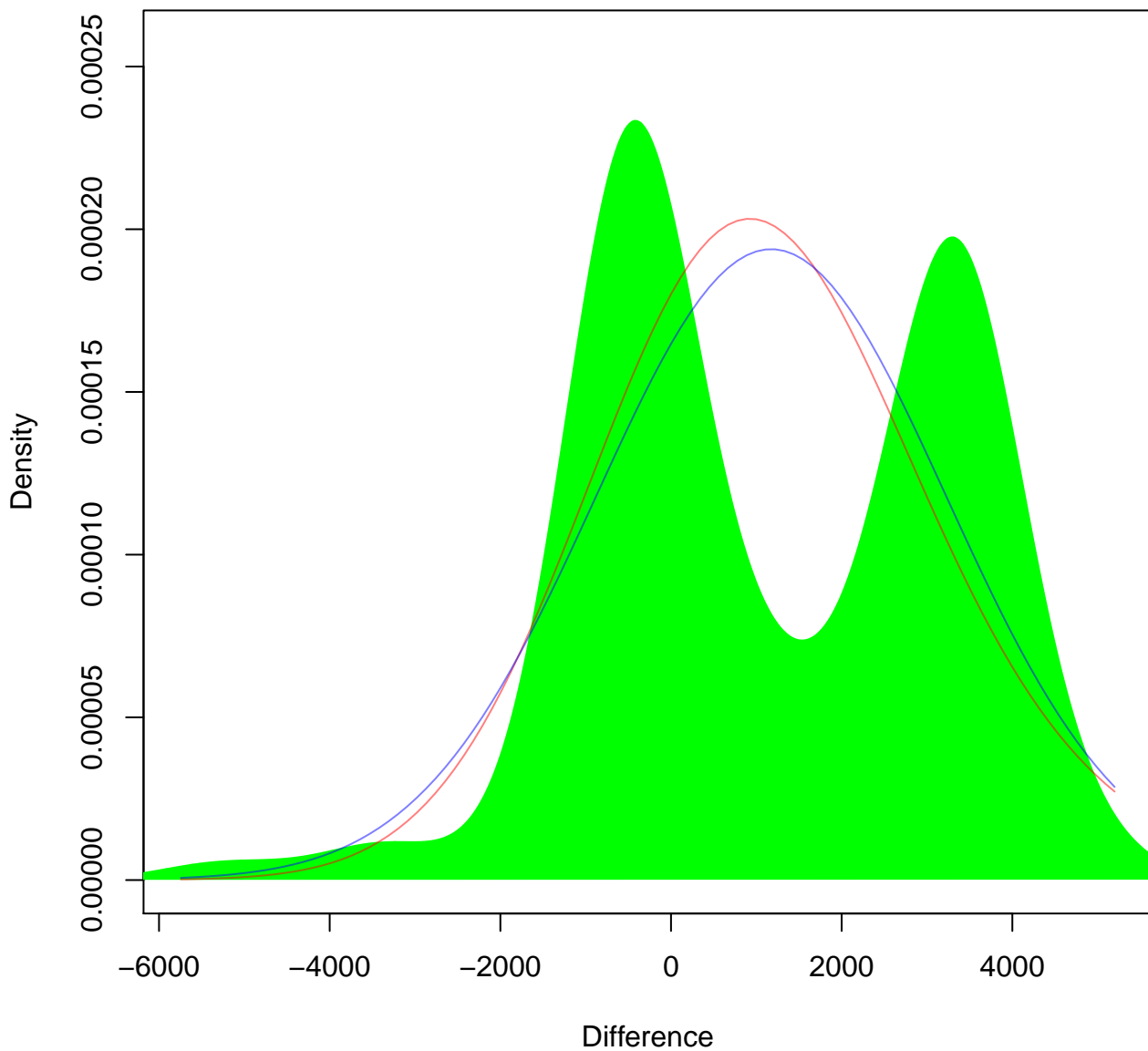
histogram of resid(bin.glme.test) Density of resid(bin.glme.test)



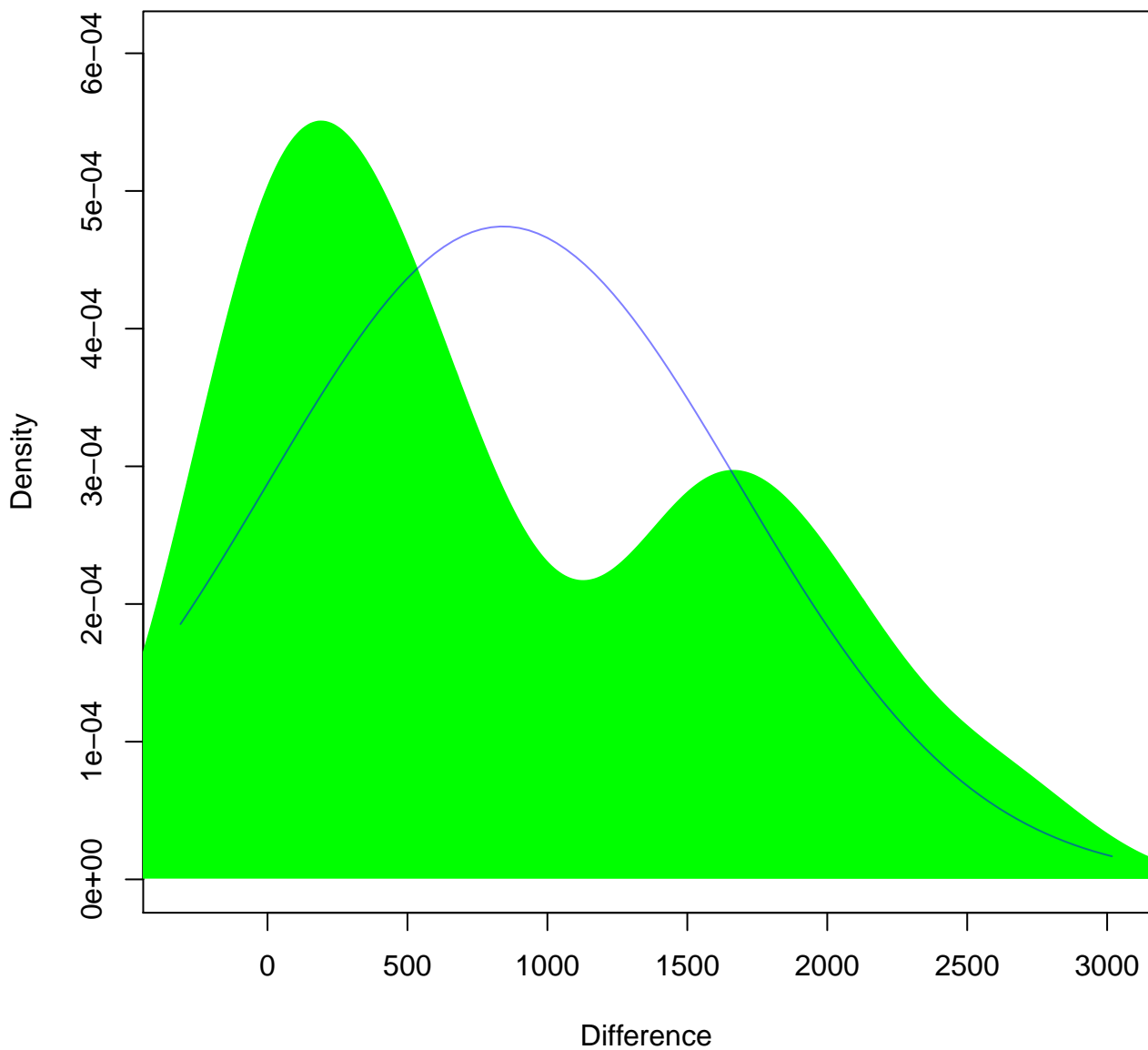
Boxplot of resid(bin.glme.test) Q-Q Plot of resid(bin.glme.test)



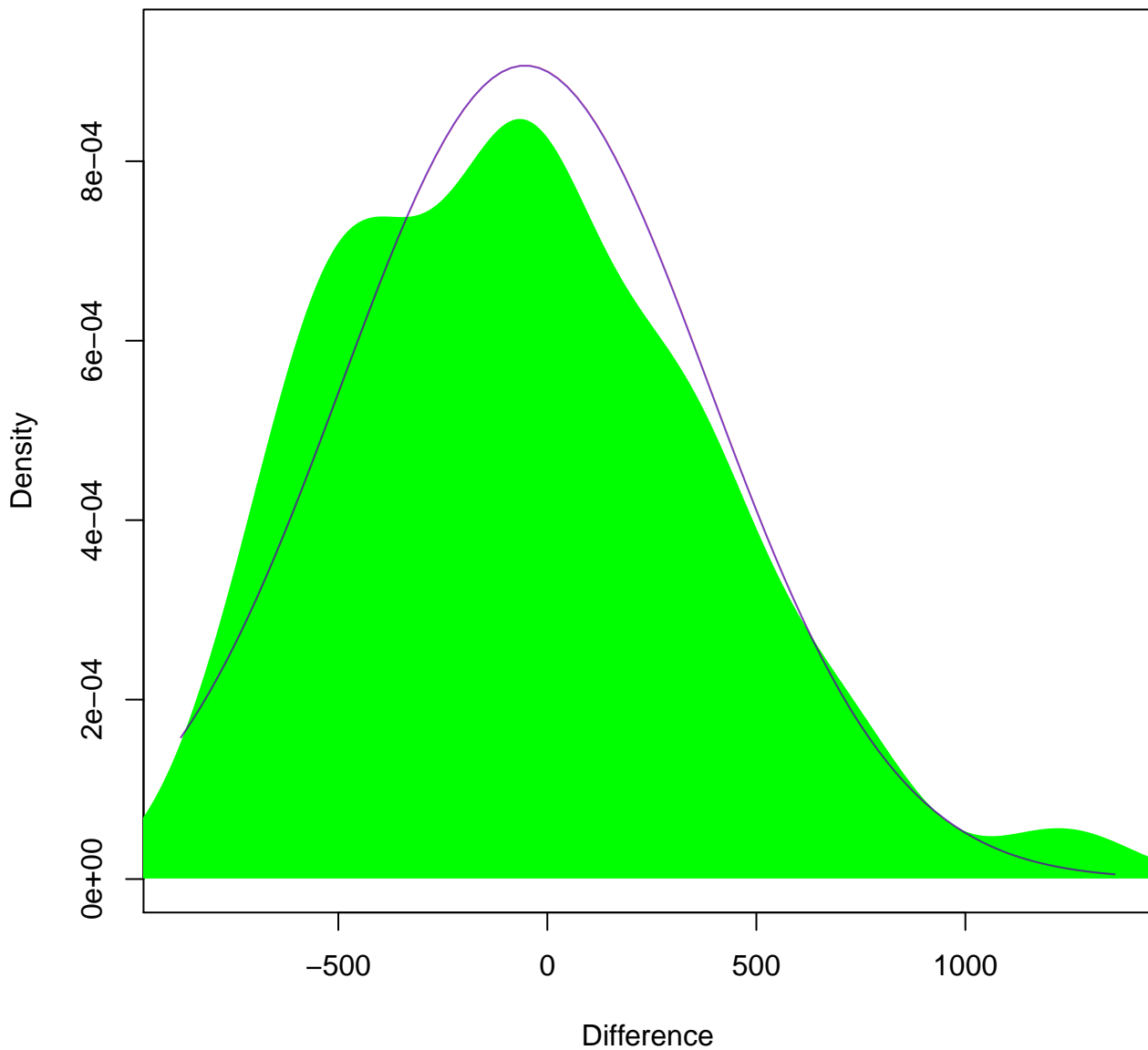
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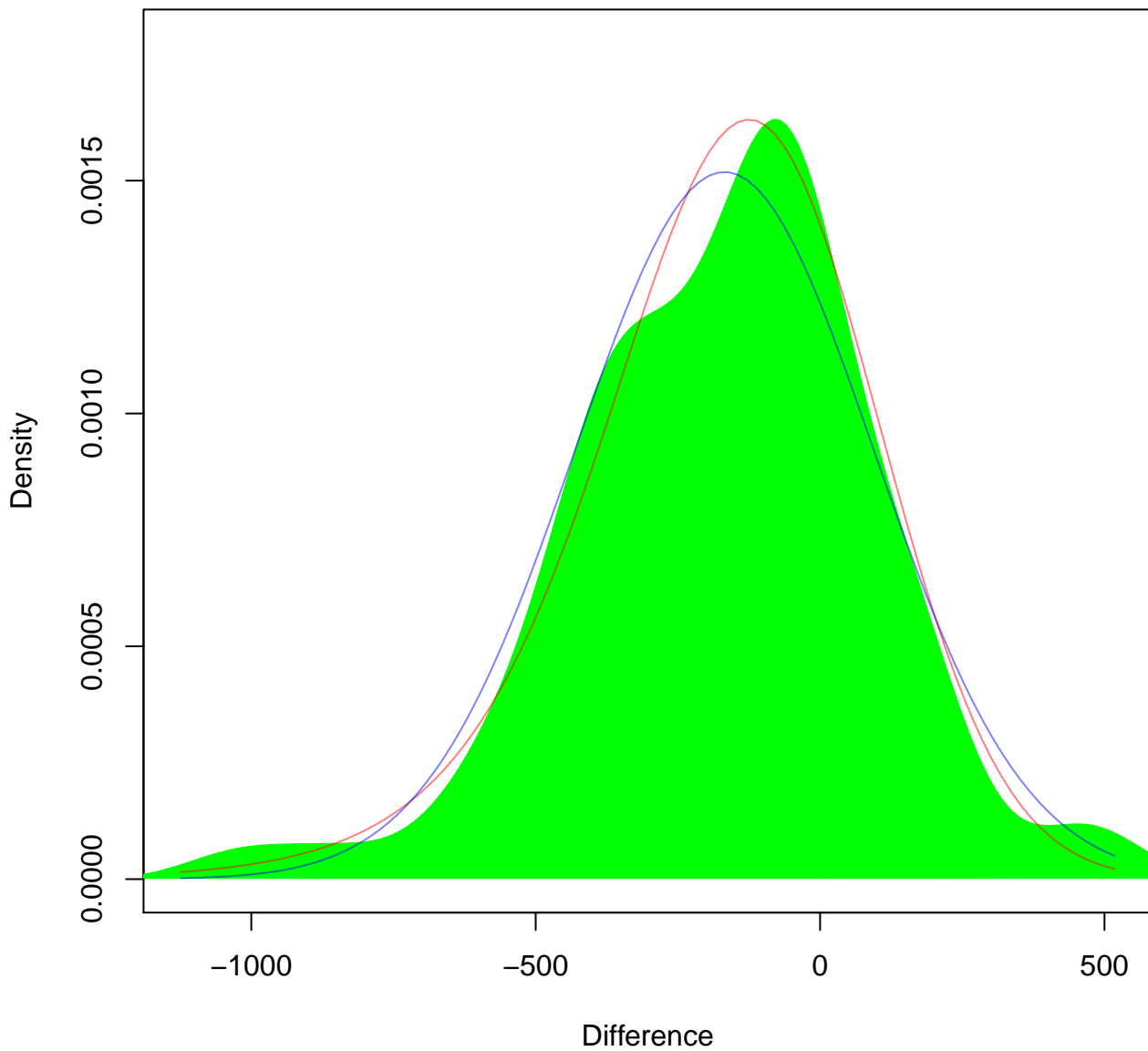
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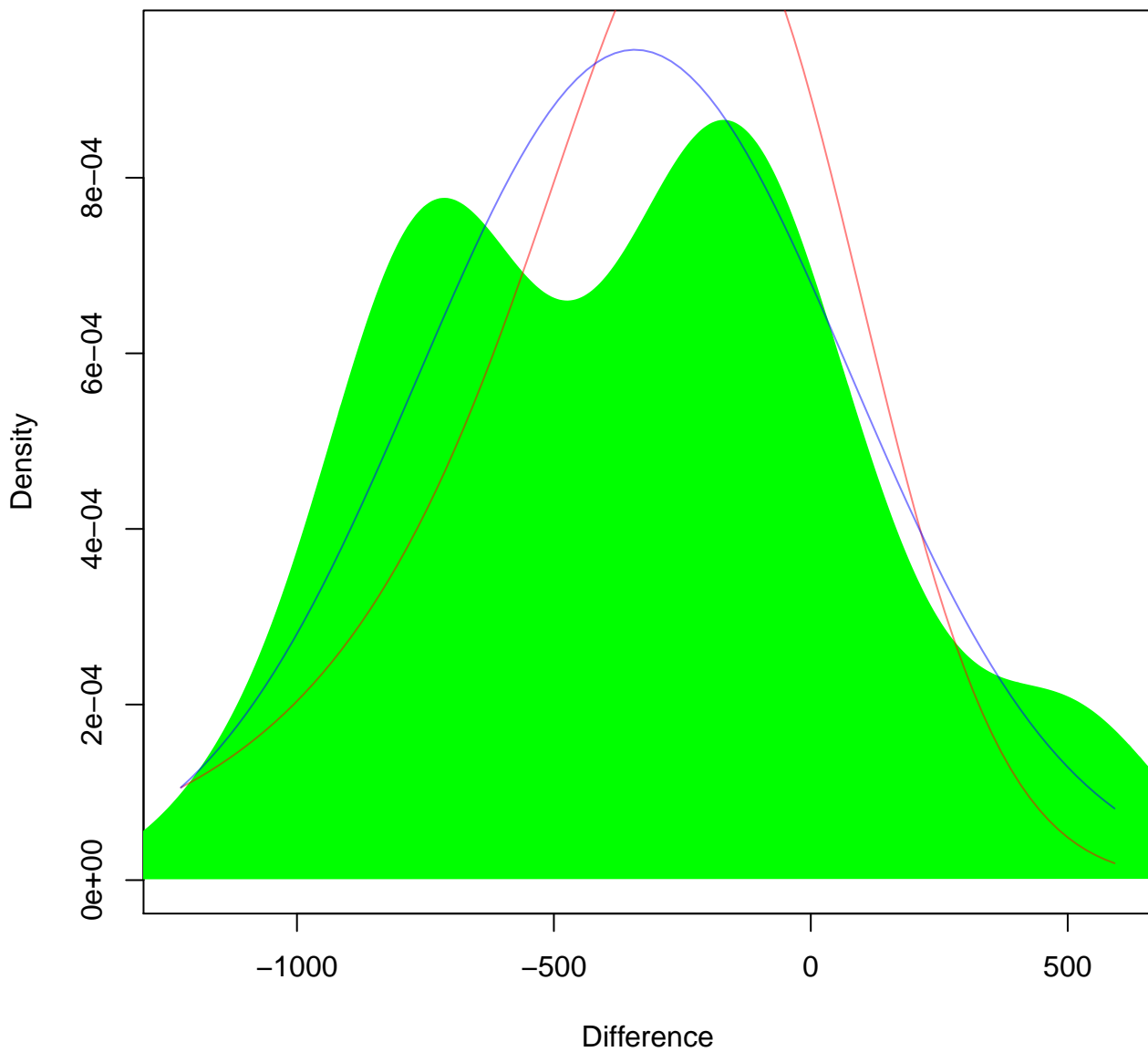
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Overall

