



NSW NATIONAL PARKS & WILDLIFE SERVICE

# River Red Gum Ecological Thinning Trial

Monitoring report 2020 – Appendices





© 2020 State of NSW and Department of Planning, Industry and Environment

With the exception of photographs, the State of NSW and Department of Planning, Industry and Environment are pleased to allow this material to be reproduced in whole or in part for educational and non-commercial use, provided the meaning is unchanged and its source, publisher and authorship are acknowledged. Specific permission is required for the reproduction of photographs.

The Department of Planning, Industry and Environment (DPIE) has compiled this report in good faith, exercising all due care and attention. No representation is made about the accuracy, completeness or suitability of the information in this publication for any particular purpose. DPIE shall not be liable for any damage which may occur to any person or organisation taking action or not on the basis of this publication. Readers should seek appropriate advice when applying the information to their specific needs.

All content in this publication is owned by DPIE and is protected by Crown Copyright, unless credited otherwise. It is licensed under the [Creative Commons Attribution 4.0 International \(CC BY 4.0\)](#), subject to the exemptions contained in the licence. The legal code for the licence is available at [Creative Commons](#).

DPIE asserts the right to be attributed as author of the original material in the following manner: © State of New South Wales and Department of Planning, Industry and Environment 2020.

Cover photo: Botanists conducting floristic monitoring in thinned river red gum forest. Evan Curtis.

This report should be cited as: Gorrod E, Travers S, Ellis M, Hope B, McAllister D, Val J, O'Kelly T and Curtis E 2020, *River red gum ecological thinning trial: Monitoring report 2020 – Appendices*, NSW National Parks and Wildlife Service, Hurstville.

Published by:

Environment, Energy and Science  
Department of Planning, Industry and Environment  
Locked Bag 5022, Parramatta NSW 2124  
Phone: +61 2 9995 5000 (switchboard)  
Phone: 1300 361 967 (Environment, Energy and Science enquiries)  
TTY users: phone 133 677, then ask for 1300 361 967  
Speak and listen users: phone 1300 555 727, then ask for 1300 361 967  
Email: [info@environment.nsw.gov.au](mailto:info@environment.nsw.gov.au)  
Website: [www.environment.nsw.gov.au](http://www.environment.nsw.gov.au)

Report pollution and environmental incidents  
Environment Line: 131 555 (NSW only) or [info@environment.nsw.gov.au](mailto:info@environment.nsw.gov.au)  
See also [www.environment.nsw.gov.au](http://www.environment.nsw.gov.au)

ISBN 978-1-922431-41-7  
EES 2020/0268  
June 2020

Find out more about your environment at:

**[www.environment.nsw.gov.au](http://www.environment.nsw.gov.au)**













# Contents

Appendix A: Photo points	1
Appendix B: Model summaries – Tree parameters	6
B.1 Tree growth rates	6
B.2 Tree mortality	9
Appendix C: Model summaries – Recruitment	12
C.1 Germinants	12
C.2 Seedlings	15
C.3 Saplings	19
Appendix D: Model summaries – Canopy condition	23
D.1 Tree crown extent	23
D.2 Visually assessed canopy cover	32
D.3 Remotely sensed canopy cover	36
D.4 Plant area index	39
Appendix E: Model summaries – Fuel hazard	42
E.1 Overall fuel hazard	42
E.2 Surface fuel hazard	45
E.3 Near surface fuel hazard	54
E.4 Combined surface and near surface fuel hazard	63
E.5 Elevated fuel hazard	66
Appendix F: Model summaries – Floristic composition	75
F.1 Plant species richness	75
F.2 Plant cover	81
F.3 Floristic community composition 2019–20	87
F.4 Threatened plant species	87
Appendix G: Model summaries – Birds	88
G.1 Bird species richness	88
G.2 Bird community composition	90
G.3 Threatened bird species	90
Appendix H: Model summaries – Bats	91
H.1 Overall bat activity	91
H.2 Bat guild activity	93
H.3 Individual bat species	96
Appendix I: Plant species list	99
I.1 Native plant species list	99
I.2 Exotic plant species list	103
Appendix J: Bird species list	107

# Appendix A: Photo points

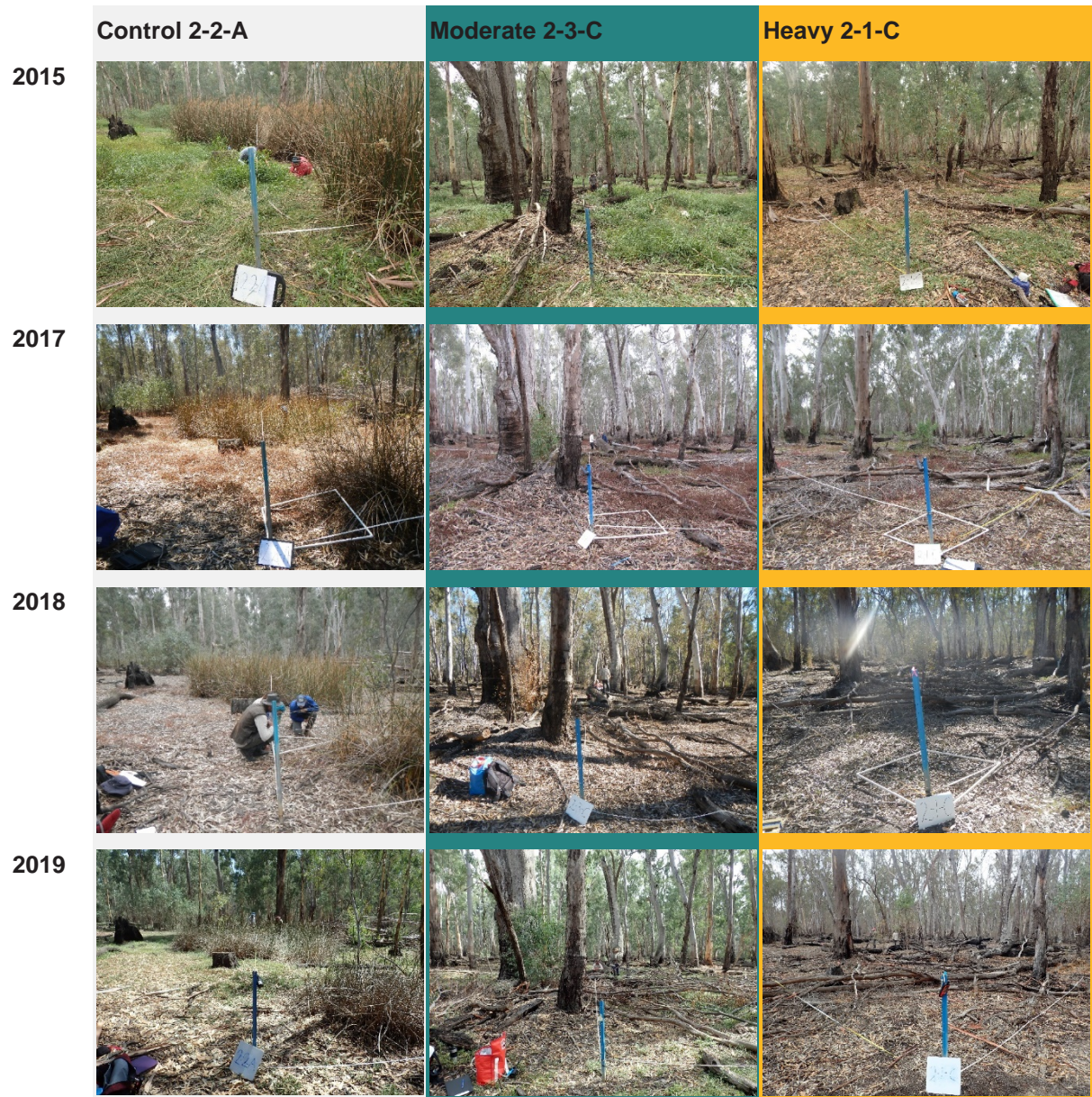
Photos from the floristic plots of a selection of sites for each survey year.

## SITE 1 – Site Quality 2

	Control 1-2-C	Moderate 1-1-A	Heavy 1-3-B
2015			
2017			
2018			
2019			

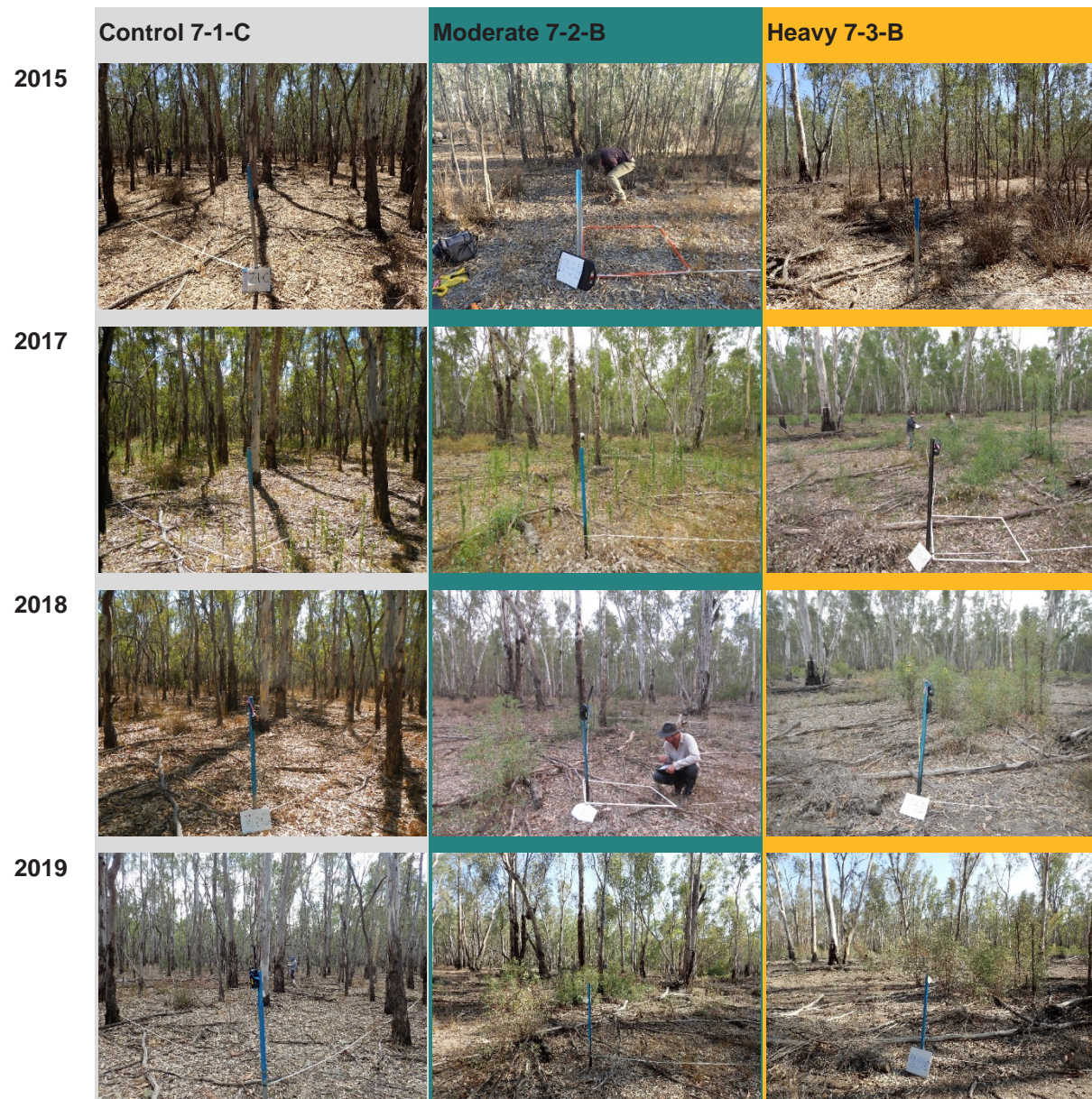


## SITE 2 – Site Quality 1





## SITE 7 – Site Quality 2

















## SITE 10 – Site Quality 2





## SITE 22 – Site Quality 1

	Control 22-2-B	Moderate 22-1-A	Heavy 22-3-C
2015			
2017			
2018			
2019			



## Appendix B: Model summaries – Tree parameters

### B.1 Tree growth rates

**Table 1 Model fitting summary: tree growth rate by tree size**

<b>Response</b>	Growth.per.year – change in tree diameter at breast height in millimetres between initial and most recent surveys, divided by the number of decimal years passed between survey dates Continuous positive variable
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	growth.per.year <-5 (13 values) growth.per.year >15 (5 values) diameter.first >1100 (26 values)
<b>Fixed factors</b>	Diameter.first.m – tree diameter at breast height in metres at initial survey (metres used to make the scale of the response and predictors similar) Ecological thinning treatment (control, moderate, heavy) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two interactions were included: diameter first * site quality and diameter first * treatment. No other interactions were trialled
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Marginal = 11.73% Conditional = 32.77%
<b>Confidence comments</b>	Moderate. Dunn Smyth residual tests indicated significant deviance from uniformity, with residuals slightly smaller than expected for a central portion of the range. Outlier test significant identified approximately 20 outliers, most of which were for large values

## Model results summary 1: tree growth rate by tree size

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula:
## growth.per.year ~ diameter.first.m * Site quality + diameter.first.m *
##   Treatment + (1 | fsiteplot) + (1 | fsite)
##   Data: dat
##
## REML criterion at convergence: 9164.4
##
## Scaled residuals:
##   Min      1Q  Median      3Q      Max
## -4.1369 -0.6207 -0.0283  0.5739  4.2752
##
## Random effects:
##   Groups      Name          Variance Std.Dev.
##   fsiteplot (Intercept) 1.103     1.050
##   fsite      (Intercept) 1.029     1.015
##   Residual                6.813     2.610
## Number of obs: 1901, groups: fsiteplot, 66; fsite, 22
##
## Fixed effects:
##
##              Estimate Std. Error   df t value
## (Intercept)    4.5068    0.5221  77.0797   8.632
## diameter.first.m    0.2946    0.5940 1854.5258   0.496
## Site qualitySQ2   -1.7385    0.6083  37.2975  -2.858
## TreatmentMod - Flood    0.7338    0.6546 197.1809   1.121
## TreatmentFlood - Mod    2.4234    0.6557 194.1694   3.696
## TreatmentHeavy - Flood    1.5386    0.6632 201.8874   2.320
## TreatmentFlood - Heavy    3.0036    0.6555 192.0522   4.582
## diameter.first.m:Site qualitySQ2   -0.3855    0.6205 1860.8221  -0.621
## diameter.first.m:TreatmentMod - Flood   -0.5823    0.8942 1857.1444  -0.651
## diameter.first.m:TreatmentFlood - Mod   -3.4499    0.9191 1860.0505  -3.754
## diameter.first.m:TreatmentHeavy - Flood  -2.8380    0.9128 1875.3159  -3.109
## diameter.first.m:TreatmentFlood - Heavy  -4.8433    0.9534 1871.4947  -5.080
##
##              Pr(>|t|)
## (Intercept)    6.10e-13 ***
## diameter.first.m    0.619968
## Site qualitySQ2    0.006936 **
## TreatmentMod - Flood    0.263633
## TreatmentFlood - Mod    0.000285 ***
## TreatmentHeavy - Flood    0.021340 *
## TreatmentFlood - Heavy    8.27e-06 ***
## diameter.first.m:Site qualitySQ2    0.534445
## diameter.first.m:TreatmentMod - Flood    0.515037
## diameter.first.m:TreatmentFlood - Mod    0.000180 ***
## diameter.first.m:TreatmentHeavy - Flood    0.001906 **
## diameter.first.m:TreatmentFlood - Heavy    4.15e-07 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```



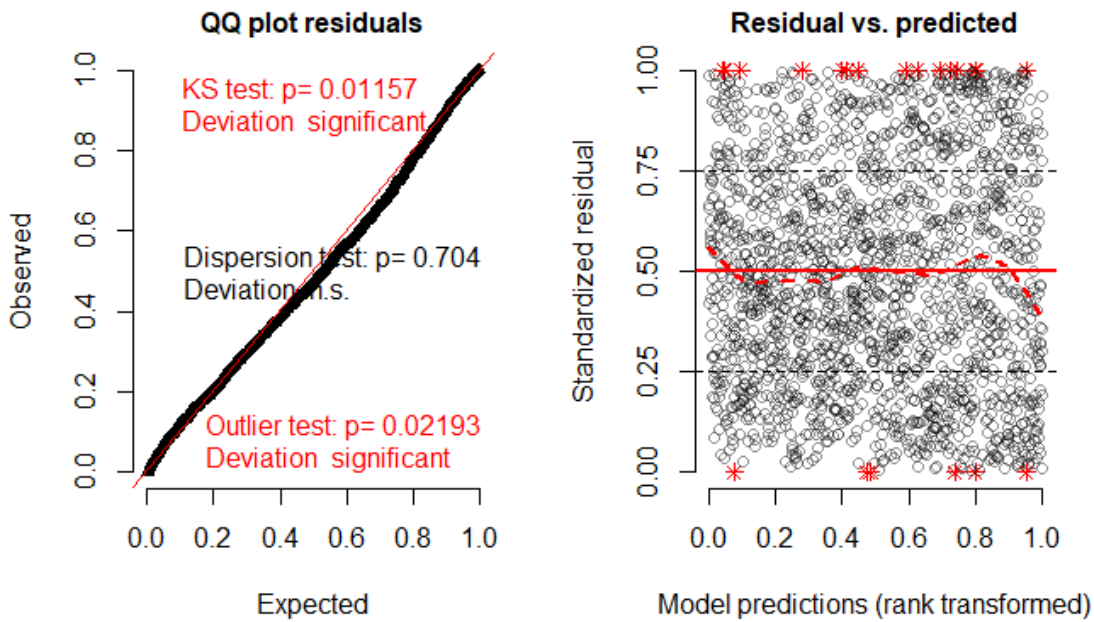


Figure 1 Dunn Smyth simulated residuals: tree growth rate by tree size

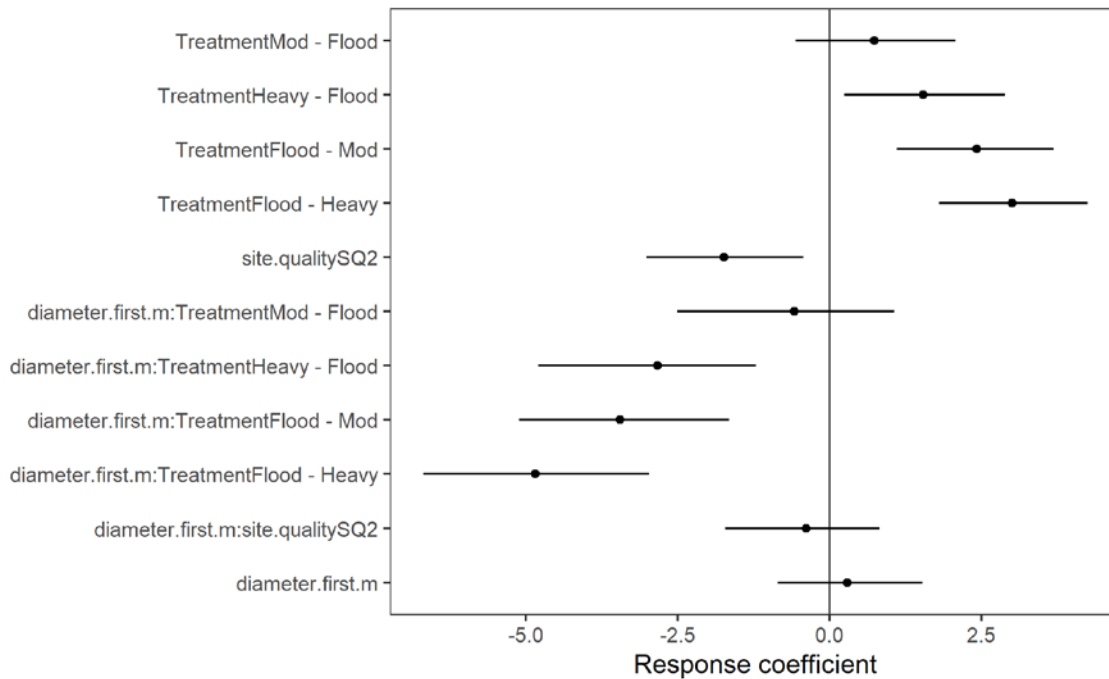


Figure 2 Bootstrapped confidence intervals for fixed effects: tree growth rate by tree size

Table 2 Bootstrapped model fitted values for average conditions: tree growth rate by tree size

Initial diameter	Site quality	Control	Moderately thinned then flooded	Heavily thinned then flooded	Flooded then moderately thinned	Flooded then heavily thinned
20 cm	SQ1	4.7435	5.3395	5.5360	6.1929	6.7656
80 cm	SQ1	4.6997	5.0888	4.0926	4.4939	3.9193
20 cm	SQ2	2.5929	3.3391	3.6847	4.5690	4.6901
80 cm	SQ2	2.6395	2.7991	1.9365	2.3967	1.8777

## B.2 Tree mortality

**Table 3 Model fitting summary: tree mortality**

<b>Response</b>	Proportion of 50 trees per 9 hectare plot that were dead Modelled as the ratio of dead to live trees
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	glmer from lme4
<b>Distribution used</b>	Binomial
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2) Also ran a model with the order of flooding and thinning included, which was not significant
<b>Fixed factor interactions</b>	Interaction between thinning treatment and survey year Also trialled a three-way interaction between thinning treatment, survey year and site quality, which was less parsimonious
<b>Random factors</b>	site (a factor over sites), and siteplot (a factor over 9 ha treatment plots)
<b>R<sup>2</sup></b>	Marginal = 1.16% Conditional = 9.15%
<b>Confidence comments</b>	Moderate, no deviations detected but R <sup>2</sup> values low

### Model results summary 2: tree mortality

```

y <- cbind(success = DAT.livedead$dead, failure = DAT.livedead$live)
model2 <- glmer(y ~ year * Treatment + Site quality + (1|siteplot) + (1|site),
               data = DAT.livedead, family = binomial,
               control=glmerControl(optimizer="bobyqa", optCtrl = list(maxfun = 2e5
)))

summary(model2)

## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula: y ~ year * Treatment + Site quality + (1 | siteplot) + (1 | site)
## Data: DAT.livedead
## Control: glmerControl(optimizer = "bobyqa", optCtrl = list(maxfun = 2e+05))
##

```



```
##      AIC      BIC  logLik deviance df.resid
##  1163.7  1217.3  -566.8  1133.7    249
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.7608 -0.5635 -0.0569  0.4143  2.7455
##
## Random effects:
## Groups   Name      Variance Std.Dev.
## siteplot (Intercept) 0.19629  0.4430
## site     (Intercept) 0.09314  0.3052
## Number of obs: 264, groups: siteplot, 66; site, 22
##
## Fixed effects:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -2.45088    0.18285  -13.404  <2e-16 ***
## year2017      -0.17829    0.15868   -1.124   0.2612
## year2018      -0.03614    0.15406   -0.235   0.8145
## year2019      -0.11167    0.15645   -0.714   0.4754
## TreatmentModerate  0.06409    0.20358    0.315   0.7529
## TreatmentHeavy   0.05958    0.20382    0.292   0.7701
## Site qualitySQ2 -0.06881    0.18118   -0.380   0.7041
## year2017:TreatmentModerate  0.53195    0.21183    2.511   0.0120 *
## year2018:TreatmentModerate  0.30047    0.20974    1.433   0.1520
## year2019:TreatmentModerate  0.34803    0.21195    1.642   0.1006
## year2017:TreatmentHeavy   0.33764    0.21535    1.568   0.1169
## year2018:TreatmentHeavy   0.34763    0.20948    1.660   0.0970 .
## year2019:TreatmentHeavy   0.44105    0.21098    2.091   0.0366 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

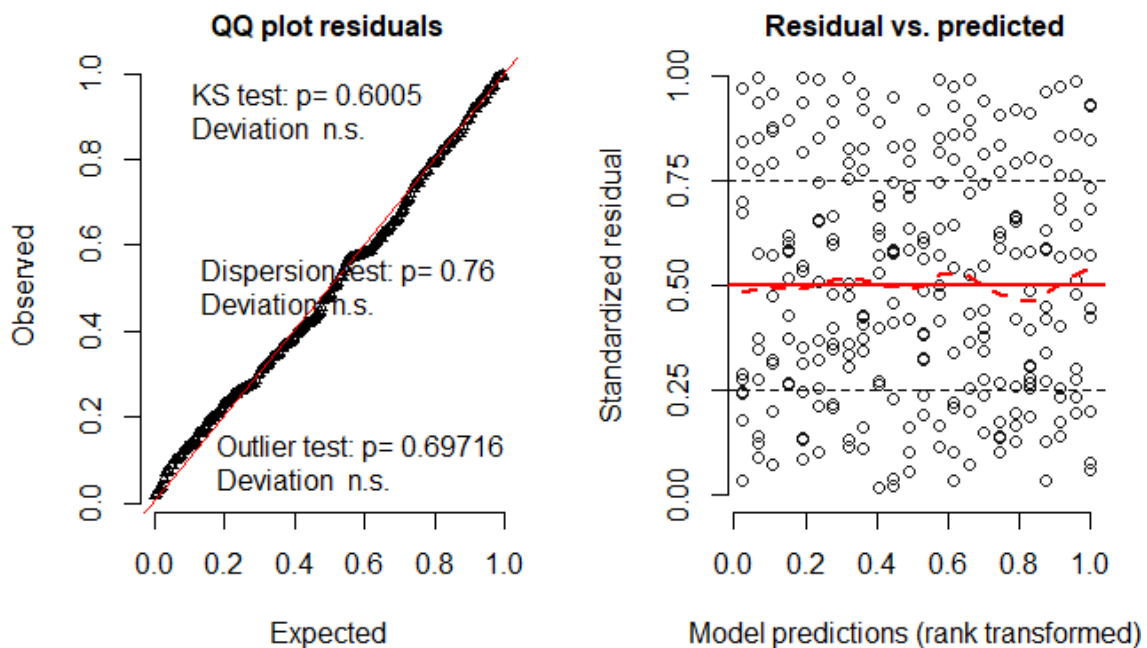
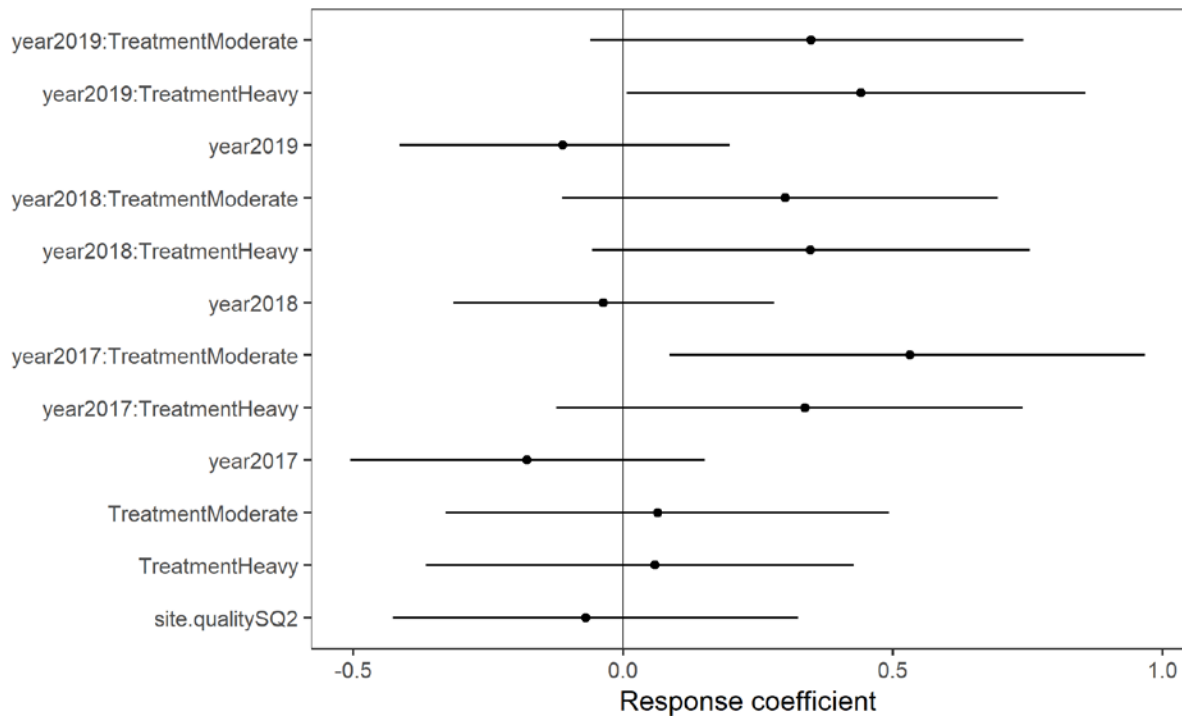


Figure 3 Dunn Smyth simulated residuals: tree mortality



**Figure 4** Bootstrapped confidence intervals for fixed effects: tree mortality

**Table 4** Bootstrapped model fitted values for average conditions: tree mortality (proportion of trees that were dead)

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	0.0793	0.0809	0.0841
2017	SQ1	0.0652	0.1122	0.1017
2018	SQ1	0.0764	0.1079	0.1074
2019	SQ1	0.0718	0.1082	0.1135
2015	SQ2	0.0786	0.0748	0.0744
2017	SQ2	0.0650	0.1041	0.0868
2018	SQ2	0.0751	0.1039	0.1027
2019	SQ2	0.0651	0.0961	0.1098



# Appendix C: Model summaries – Recruitment

## C.1 Germinants

### C.1.1 Germinant presence–absence

**Table 5 Model fitting summary: germinant presence–absence**

<b>Response</b>	Probability of germinants occurring on 9 hectare plot
<b>Response transformation</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	glmer function from lme4 package
<b>Distribution used</b>	Binomial
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Interaction between thinning treatment and survey year Also trialled a model with three-way interaction between thinning treatment, survey year and site quality, which failed to complete due to insufficient data
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Marginal = 27.2% Conditional = 60.1%
<b>Confidence comments</b>	Moderate to high, with some mid values having larger residuals than expected; and some low values having lower residuals than expected. No assumptions violated

### Model results summary 3: germinant presence–absence

```
## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula: germ.pres ~ Survey year * Treatment + Site quality + (1 | fsite)
## Data: dat
## Control: glmerControl(optimizer = "bobyqa", optCtrl = list(maxfun = 1e+05))

##      AIC      BIC  logLik deviance df.resid
##  239.2   289.3  -105.6   211.2     250

## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.8465 -0.3688 -0.1894 -0.0837  4.5716

## Random effects:
## Groups Name          Variance Std.Dev.
## fsite (Intercept)  2.712     1.647
## Number of obs: 264, groups: fsite, 22
## Fixed effects:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -1.757e+00  9.047e-01  -1.942  0.05213 .
## Survey year2017  1.281e+00  9.523e-01   1.345  0.17850
## Survey year2018  9.031e-01  9.657e-01   0.935  0.34970
## Survey year2019  1.633e+00  9.456e-01   1.727  0.08416 .
## TreatmentModerate -2.748e-07  1.031e+00   0.000  1.00000
## TreatmentHeavy    7.966e-07  1.031e+00   0.000  1.00000
## Site qualitySQ2  -2.501e+00  9.217e-01  -2.713  0.00666 **
## Survey year2017:TreatmentModerate  1.011e+00  1.326e+00   0.762  0.44578
## Survey year2018:TreatmentModerate  3.781e-01  1.350e+00   0.280  0.77940
## Survey year2019:TreatmentModerate -1.149e+00  1.364e+00  -0.842  0.39987
## Survey year2017:TreatmentHeavy    6.868e-01  1.326e+00   0.518  0.60459
## Survey year2018:TreatmentHeavy   -9.031e-01  1.413e+00  -0.639  0.52261
## Survey year2019:TreatmentHeavy   -1.633e+00  1.399e+00  -1.167  0.24303
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

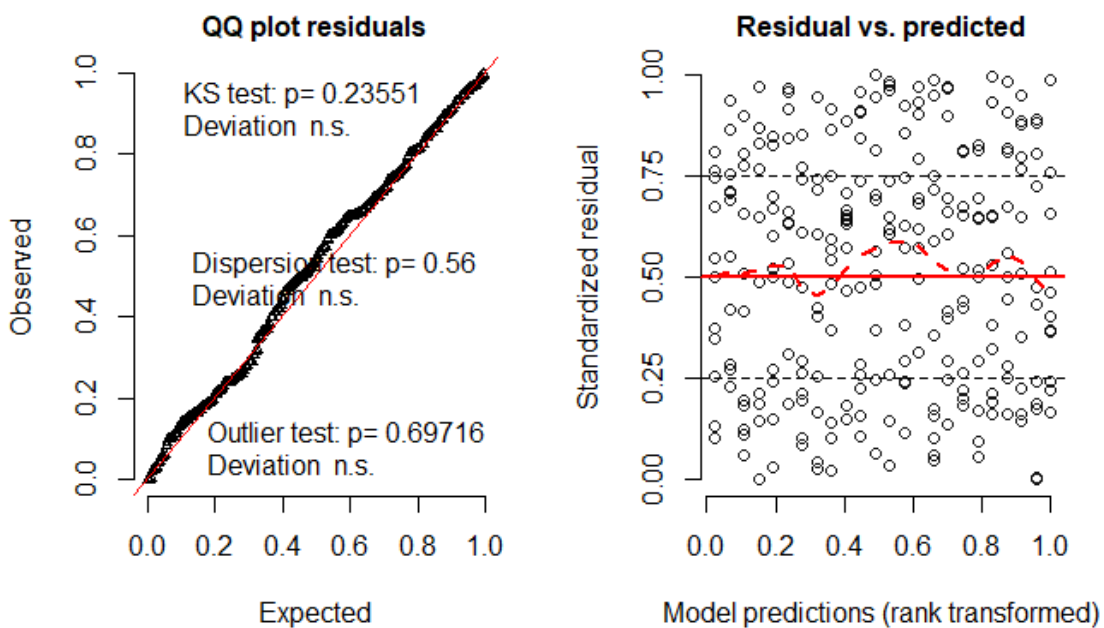
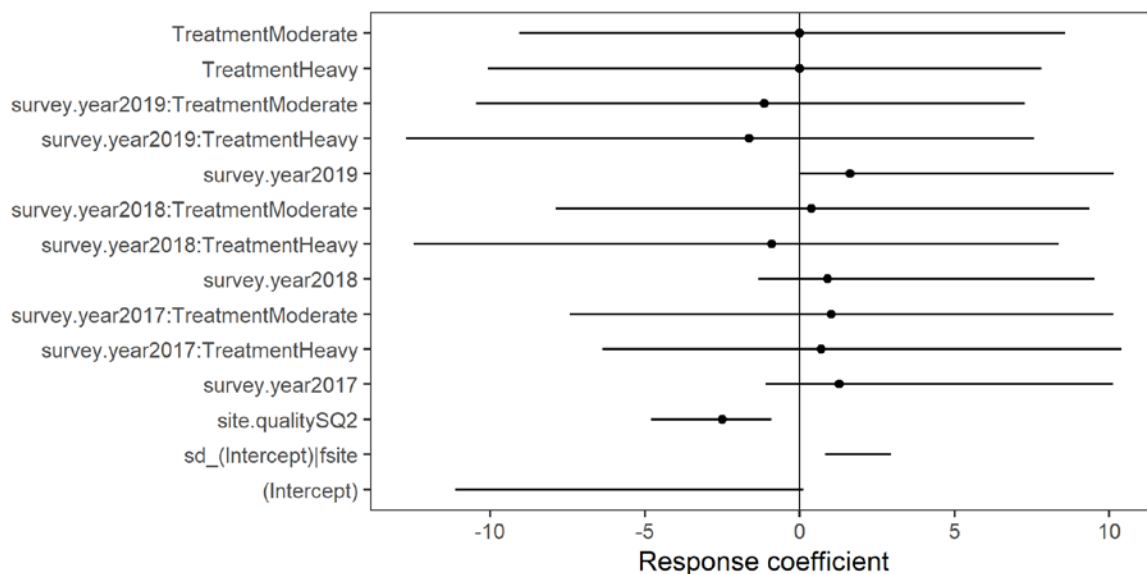


Figure 5 Dunn Smyth simulated residuals: germinant presence–absence



**Figure 6** Bootstrapped confidence intervals for fixed effects: germinant presence–absence

**Table 6** Bootstrapped fitted values for average conditions: germinant presence–absence (probability of germinants occurring per 9 hectare plot)

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	0.1358	0.1413	0.1522
2017	SQ1	0.3933	0.6420	0.5564
2018	SQ1	0.2894	0.3837	0.1574
2019	SQ1	0.4643	0.2191	0.1430
2015	SQ2	0.0129	0.0143	0.0132
2017	SQ2	0.0483	0.1199	0.0915
2018	SQ2	0.0350	0.0451	0.0148
2019	SQ2	0.0696	0.0219	0.0134

## C.1.2 Germinant abundance

Not modelled.



## C.2 Seedlings

### C.2.1 Seedling presence–absence

Not modelled.

### C.2.2 Seedling abundance

**Table 7 Model fitting summary: seedling abundance**

<b>Response</b>	Number of seedlings present (total of three 0.04 ha subplots)
<b>Response transformation used</b>	Transformed using $\log_e(x + 1)$
<b>Other transformations compared</b>	Square root
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	Negative binomial
<b>Outliers removed</b>	No outliers removed A gaussian model was trialled with values greater than 600 removed
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Trialled interaction between thinning treatment and survey year Trialled a model with no interactions A model with three-way interaction between thinning treatment, survey year and site quality was more parsimonious and detected significant effects
<b>Random factors</b>	fsite (a factor over sites)
<b>R<sup>2</sup></b>	Marginal = 10.9% Conditional = 54.9%
<b>Confidence comments</b>	High, no assumptions violated

**Model results summary 4: seedling abundance**

```

dat$Lseedlings<-log(dat$seedlings+1)
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Lseedlings ~ Survey year * Treatment * Site quality + (1 | fsite)
## Data: dat

## REML criterion at convergence: 817.2

## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.6128 -0.5228  0.0226  0.5302  4.0757

## Random effects:
## Groups   Name                Variance Std.Dev.
## fsite    (Intercept)  1.096    1.047
## Residual                    1.123    1.059
## Number of obs: 264, groups: fsite, 22

## Fixed effects:
##                                     Estimate Std. Error
## (Intercept)                        2.93201   0.44905
## Survey year2017                     0.34959   0.45177
## Survey year2018                     0.88134   0.45177
## Survey year2019                     0.11629   0.45177
## TreatmentModerate                   0.56219   0.45177
## TreatmentHeavy                      0.44946   0.45177
## Site qualitySQ2                     0.05899   0.63505
## Survey year2017:TreatmentModerate  -0.85172   0.63890
## Survey year2018:TreatmentModerate  -1.01485   0.63890
## Survey year2019:TreatmentModerate   0.19694   0.63890
## Survey year2017:TreatmentHeavy     -0.11403   0.63890
## Survey year2018:TreatmentHeavy     -0.52872   0.63890
## Survey year2019:TreatmentHeavy     -0.32814   0.63890
## Survey year2017:Site qualitySQ2    -0.47418   0.63890
## Survey year2018:Site qualitySQ2    -1.47462   0.63890
## Survey year2019:Site qualitySQ2    -1.31478   0.63890
## TreatmentModerate:Site qualitySQ2  -1.07609   0.63890
## TreatmentHeavy:Site qualitySQ2     -1.31491   0.63890
## Survey year2017:TreatmentModerate:Site qualitySQ2  1.71851   0.90354
## Survey year2018:TreatmentModerate:Site qualitySQ2  1.87452   0.90354
## Survey year2019:TreatmentModerate:Site qualitySQ2  0.80194   0.90354
## Survey year2017:TreatmentHeavy:Site qualitySQ2    1.31591   0.90354
## Survey year2018:TreatmentHeavy:Site qualitySQ2    1.86054   0.90354
## Survey year2019:TreatmentHeavy:Site qualitySQ2    2.14715   0.90354
##                                     df t value Pr(>|t|)
## (Intercept)                        65.15335  6.529 1.16e-08
## Survey year2017                    220.00001  0.774  0.4399
## Survey year2018                    220.00001  1.951  0.0523
## Survey year2019                    220.00001  0.257  0.7971
## TreatmentModerate                  220.00001  1.244  0.2147
## TreatmentHeavy                     220.00001  0.995  0.3209
## Site qualitySQ2                    65.15335  0.093  0.9263
## Survey year2017:TreatmentModerate  220.00001 -1.333  0.1839
## Survey year2018:TreatmentModerate  220.00001 -1.588  0.1136
## Survey year2019:TreatmentModerate  220.00001  0.308  0.7582
## Survey year2017:TreatmentHeavy     220.00001 -0.178  0.8585
## Survey year2018:TreatmentHeavy     220.00001 -0.828  0.4088
## Survey year2019:TreatmentHeavy     220.00001 -0.514  0.6080
## Survey year2017:Site qualitySQ2    220.00001 -0.742  0.4588

```



```

## Survey year2018:Site qualitySQ2          220.00001  -2.308  0.0219
## Survey year2019:Site qualitySQ2          220.00001  -2.058  0.0408
## TreatmentModerate:Site qualitySQ2        220.00001  -1.684  0.0935
## TreatmentHeavy:Site qualitySQ2           220.00001  -2.058  0.0408
## Survey year2017:TreatmentModerate:Site qualitySQ2 220.00001  1.902  0.0585
## Survey year2018:TreatmentModerate:Site qualitySQ2 220.00001  2.075  0.0392
## Survey year2019:TreatmentModerate:Site qualitySQ2 220.00001  0.888  0.3758
## Survey year2017:TreatmentHeavy:Site qualitySQ2   220.00001  1.456  0.1467
## Survey year2018:TreatmentHeavy:Site qualitySQ2   220.00001  2.059  0.0407
## Survey year2019:TreatmentHeavy:Site qualitySQ2   220.00001  2.376  0.0183
##
## (Intercept)                                ***
## Survey year2017
## Survey year2018                             .
## Survey year2019
## TreatmentModerate
## TreatmentHeavy
## Site qualitySQ2
## Survey year2017:TreatmentModerate
## Survey year2018:TreatmentModerate
## Survey year2019:TreatmentModerate
## Survey year2017:TreatmentHeavy
## Survey year2018:TreatmentHeavy
## Survey year2019:TreatmentHeavy
## Survey year2017:Site qualitySQ2
## Survey year2018:Site qualitySQ2             *
## Survey year2019:Site qualitySQ2             *
## TreatmentModerate:Site qualitySQ2          .
## TreatmentHeavy:Site qualitySQ2             *
## Survey year2017:TreatmentModerate:Site qualitySQ2 .
## Survey year2018:TreatmentModerate:Site qualitySQ2 *
## Survey year2019:TreatmentModerate:Site qualitySQ2
## Survey year2017:TreatmentHeavy:Site qualitySQ2
## Survey year2018:TreatmentHeavy:Site qualitySQ2 *
## Survey year2019:TreatmentHeavy:Site qualitySQ2 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

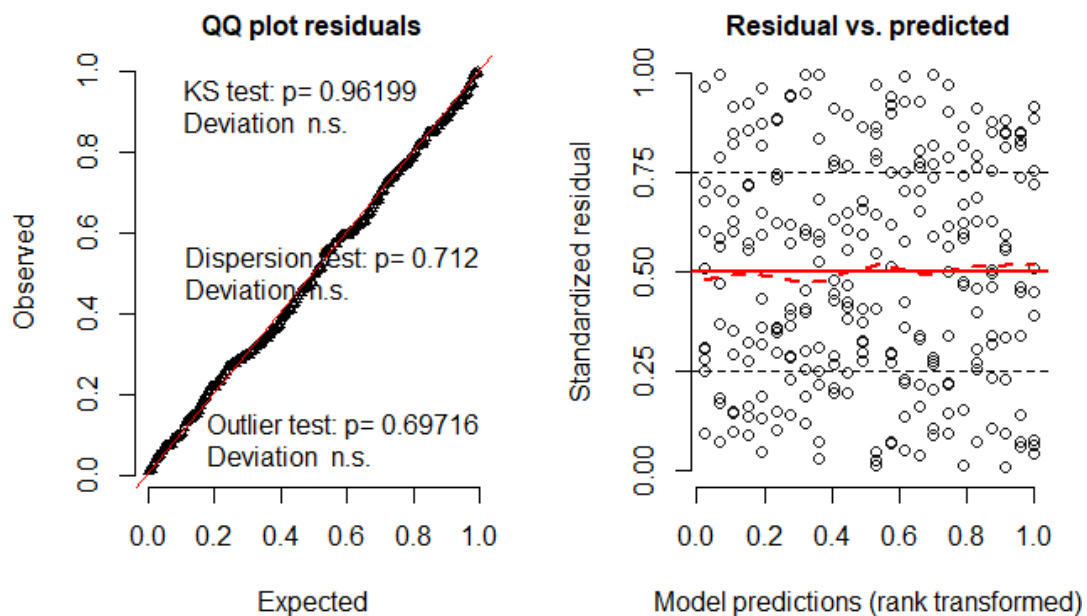
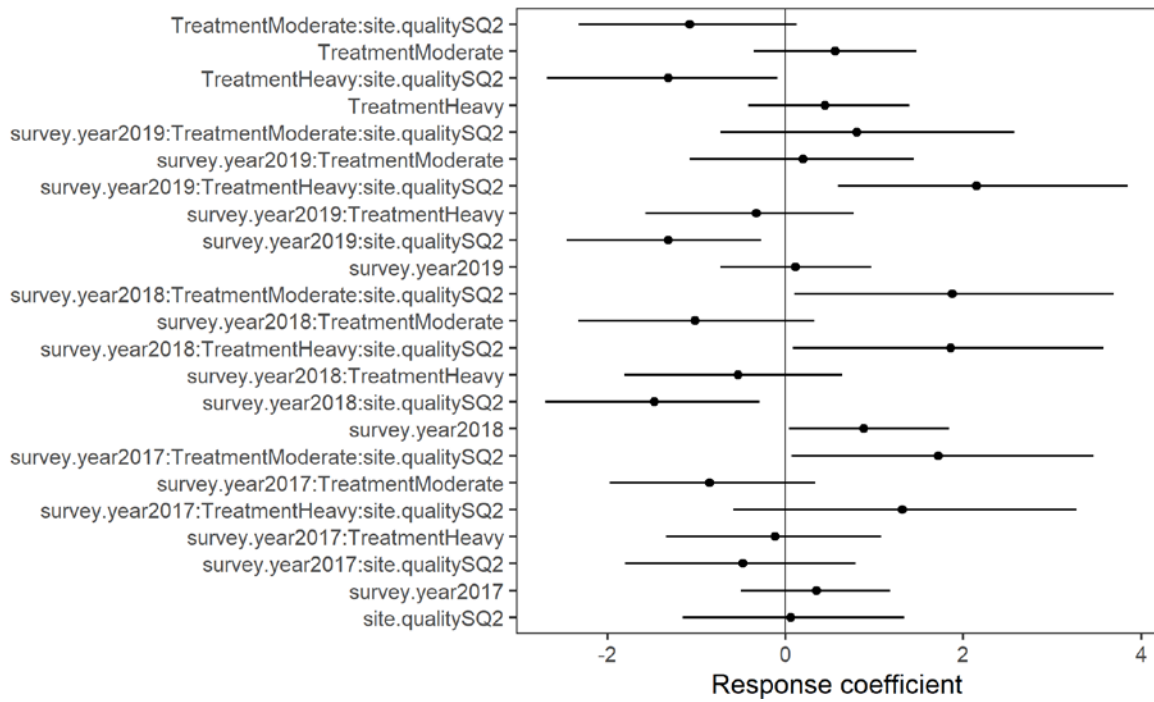


Figure 7 Dunn Smyth simulated residuals: seedling abundance



**Figure 8** Bootstrapped confidence intervals for fixed effects: seedling abundance

**Table 8** Bootstrapped fitted values for average conditions: seedling abundance

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	17.2718	32.9701	28.4781
2017	SQ1	26.9601	19.9246	37.0526
2018	SQ1	46.3531	28.8572	42.3082
2019	SQ1	20.4866	45.4674	23.5706
2015	SQ2	17.8609	10.6769	7.7861
2017	SQ2	16.3969	25.8775	23.3185
2018	SQ2	10.3936	13.8200	17.3604
2019	SQ2	4.7678	8.4469	14.3858



## C.3 Saplings

### C.3.1 Sapling presence–absence

Not modelled.

### C.3.2 Sapling abundance

**Table 9 Model fitting summary: sapling abundance**

<b>Response</b>	Total number of saplings recorded in three 0.04 ha subplots in each 9 ha plot
<b>Response transformation used</b>	$\text{Log}_e(x + 1)$
<b>Other transformations compared</b>	Square root was considered (histogram of square root of sapling abundance was inspected) but model was not run
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	Negative binomial
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year Also trialled a model with three-way interaction between thinning treatment, survey year and site quality, which was less parsimonious
<b>Random factors</b>	f <sub>site</sub> (a factor over sites)
<b>R<sup>2</sup></b>	Marginal = 31.9% Conditional = 66.1%
<b>Confidence comments</b>	Moderate. The normal QQ plot (not the Dunn Smyth simulated residuals shown below) indicates larger than expected residuals for large values

**Model results summary 5: sapling abundance**

```

dat$Lsaplings<-log(dat$saplings +1)

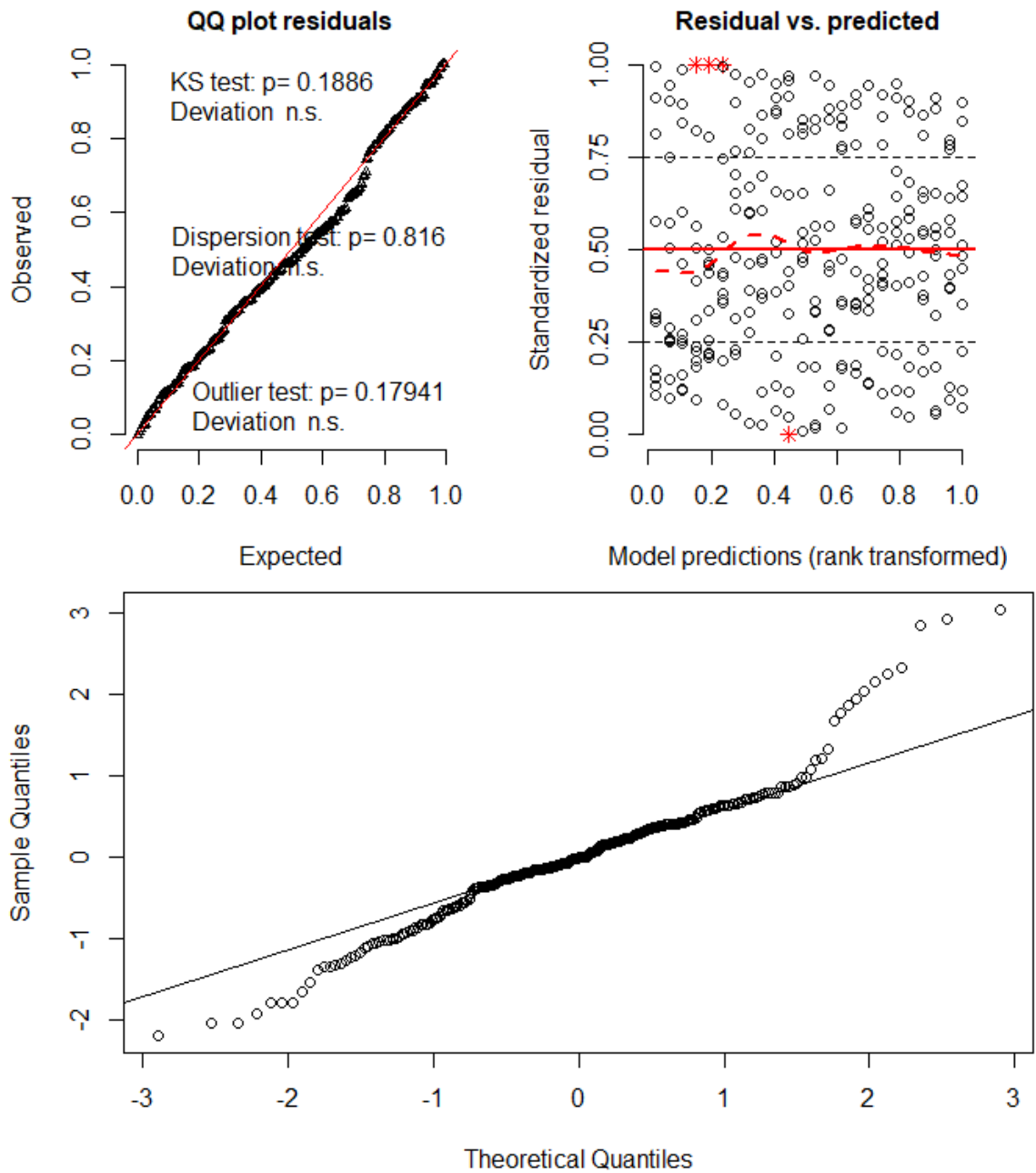
model6_log <- lmer(Lsaplings ~ Survey year*Treatment + Site quality + (1 | fsite),
                  data = dat)

summary(model6_log)

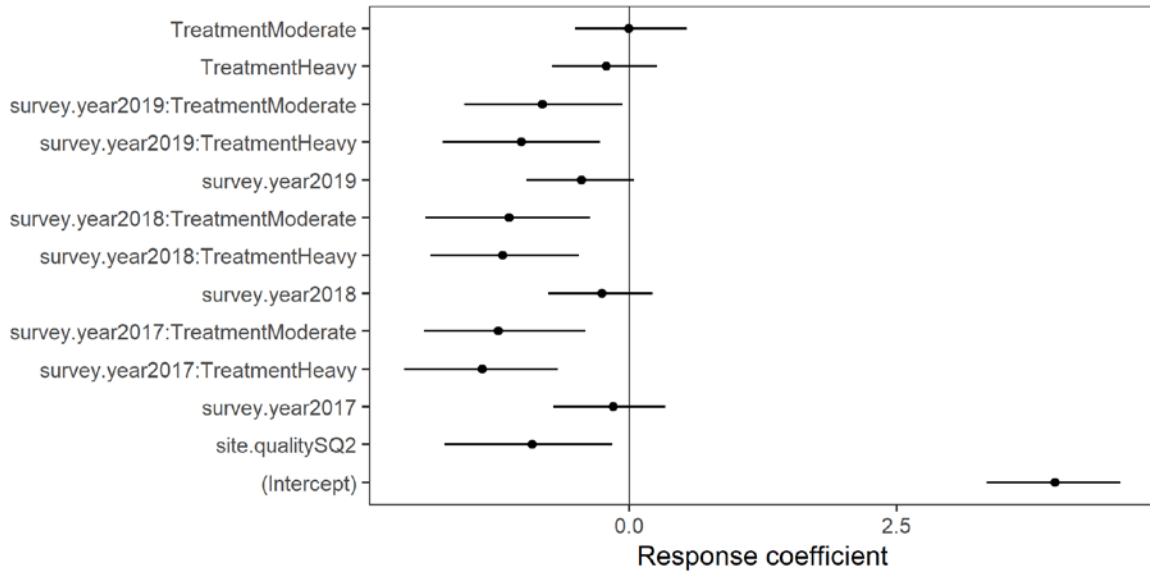
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Lsaplings ~ Survey year * Treatment + Site quality + (1 | fsite)
## Data: dat
##
## REML criterion at convergence: 727.6
##
## Scaled residuals:
##   Min       1Q   Median       3Q      Max
## -2.5576 -0.4353 -0.0119  0.4712  3.5569
##
## Random effects:
## Groups Name          Variance Std.Dev.
## fsite  (Intercept) 0.7403  0.8604
## Residual          0.7345  0.8570
## Number of obs: 264, groups: fsite, 22
##
## Fixed effects:
##              Estimate Std. Error      df t value
## (Intercept)    3.988383   0.321667   39  12.399
## Survey year2017 -0.152453   0.258397  231  -0.590
## Survey year2018 -0.257070   0.258397  231  -0.995
## Survey year2019 -0.448996   0.258397  231  -1.738
## TreatmentModerate -0.008306   0.258397  231  -0.032
## TreatmentHeavy  -0.219193   0.258397  231  -0.848
## Site qualitySQ2 -0.912877   0.381751   19  -2.391
## Survey year2017:TreatmentModerate -1.228845   0.365428  231  -3.363
## Survey year2018:TreatmentModerate -1.129331   0.365428  231  -3.090
## Survey year2019:TreatmentModerate -0.814448   0.365428  231  -2.229
## Survey year2017:TreatmentHeavy -1.378018   0.365428  231  -3.771
## Survey year2018:TreatmentHeavy -1.189243   0.365428  231  -3.254
## Survey year2019:TreatmentHeavy -1.012669   0.365428  231  -2.771
##
##              Pr(>|t|)
## (Intercept)    3.1e-15 ***
## Survey year2017    0.555769
## Survey year2018    0.320843
## Survey year2019    0.083610 .
## TreatmentModerate 0.974385
## TreatmentHeavy    0.397159
## Site qualitySQ2    0.026734 *
## Survey year2017:TreatmentModerate 0.000903 ***
## Survey year2018:TreatmentModerate 0.002244 **
## Survey year2019:TreatmentModerate 0.026794 *
## Survey year2017:TreatmentHeavy 0.000207 ***
## Survey year2018:TreatmentHeavy 0.001307 **
## Survey year2019:TreatmentHeavy 0.006040 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```





**Figure 9** Dunn Smyth simulated residuals (top panels) and normal QQ plot (bottom pane): sapling abundance



**Figure 10** Bootstrapped confidence intervals for fixed effects: sapling abundance

**Table 10** Bootstrapped fitted values for average conditions: sapling abundance (total of three 0.04 hectare subplots)

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	52.9491	54.4580	40.6011
2017	SQ1	43.9568	13.1281	8.5304
2018	SQ1	43.6266	12.4474	9.3628
2019	SQ1	33.2792	14.2173	9.4484
2015	SQ2	21.1552	20.9354	16.0500
2017	SQ2	17.1027	4.9699	2.7762
2018	SQ2	15.7366	4.4380	3.3320
2019	SQ2	13.1764	5.1329	3.0101



# Appendix D: Model summaries – Canopy condition

## D.1 Tree crown extent

### D.1.1 Average crown extent

Table 11 Model fitting summary: average crown extent

<b>Response</b>	Average crown extent. Average of crown extent (5% categories) assessed for 30 trees per 9 hectare plot A continuous, positive variable
<b>Response transformations used</b>	None
<b>Response transformations compared</b>	None
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Trialled a model with a two-way interaction between thinning treatment and survey year and an additive effect of site quality A model with three-way interaction between thinning treatment, survey year and site quality was more parsimonious and detected significant effects
<b>Random effects</b>	fsite (a factor over sites)
<b>R<sup>2</sup></b>	Marginal = 38.1% Conditional = 61.0%
<b>Confidence comments</b>	High. Met all assumptions

**Model results summary 6: average crown extent**

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: mean_extent ~ year * Treatment * Site quality + (1 | site)
## Data: dat.time

## REML criterion at convergence: 1279.7

## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.8943 -0.5673  0.0228  0.5061  2.3398

## Random effects:
## Groups   Name                Variance Std.Dev.
## site     (Intercept)  4.714    2.171
## Residual                    8.009    2.830
## Number of obs: 264, groups: site, 22

## Fixed effects:
##                                     Estimate Std. Error  df
## (Intercept)                        76.93156    1.07548  95
## year2017                            4.82602    1.20676 220
## year2018                            3.90178    1.20676 220
## year2019                            5.20481    1.20676 220
## TreatmentModerate                   2.16353    1.20676 220
## TreatmentHeavy                      -1.49142    1.20676 220
## Site qualitySQ2                     -2.74974    1.52096  95
## year2017:TreatmentModerate          -1.11808    1.70662 220
## year2018:TreatmentModerate          -0.07262    1.70662 220
## year2019:TreatmentModerate          -0.28474    1.70662 220
## year2017:TreatmentHeavy              0.87073    1.70662 220
## year2018:TreatmentHeavy              1.40051    1.70662 220
## year2019:TreatmentHeavy              0.94596    1.70662 220
## year2017:Site qualitySQ2             0.29519    1.70662 220
## year2018:Site qualitySQ2             0.43156    1.70662 220
## year2019:Site qualitySQ2            -3.47753    1.70662 220
## TreatmentModerate:Site qualitySQ2   -1.69331    1.70662 220
## TreatmentHeavy:Site qualitySQ2       1.13671    1.70662 220
## year2017:TreatmentModerate:Site qualitySQ2  1.33856    2.41352 220
## year2018:TreatmentModerate:Site qualitySQ2  1.16040    2.41352 220
## year2019:TreatmentModerate:Site qualitySQ2  4.02665    2.41352 220
## year2017:TreatmentHeavy:Site qualitySQ2  -0.48833    2.41352 220
## year2018:TreatmentHeavy:Site qualitySQ2    0.22692    2.41352 220
## year2019:TreatmentHeavy:Site qualitySQ2    2.84814    2.41352 220
##                                     t value Pr(>|t|)
## (Intercept)                        71.532 < 2e-16 ***
## year2017                            3.999 8.68e-05 ***
## year2018                            3.233 0.00141 **
## year2019                            4.313 2.43e-05 ***
## TreatmentModerate                   1.793 0.07437 .
## TreatmentHeavy                      -1.236 0.21782
## Site qualitySQ2                     -1.808 0.07377 .
## year2017:TreatmentModerate          -0.655 0.51306
## year2018:TreatmentModerate          -0.043 0.96610
## year2019:TreatmentModerate          -0.167 0.86764
## year2017:TreatmentHeavy              0.510 0.61042
## year2018:TreatmentHeavy              0.821 0.41274
## year2019:TreatmentHeavy              0.554 0.57994
## year2017:Site qualitySQ2             0.173 0.86283
## year2018:Site qualitySQ2             0.253 0.80060
```

```

## year2019:Site qualitySQ2                -2.038  0.04278 *
## TreatmentModerate:Site qualitySQ2       -0.992  0.32219
## TreatmentHeavy:Site qualitySQ2          0.666  0.50607
## year2017:TreatmentModerate:Site qualitySQ2 0.555  0.57973
## year2018:TreatmentModerate:Site qualitySQ2 0.481  0.63114
## year2019:TreatmentModerate:Site qualitySQ2 1.668  0.09667 .
## year2017:TreatmentHeavy:Site qualitySQ2  -0.202  0.83984
## year2018:TreatmentHeavy:Site qualitySQ2   0.094  0.92518
## year2019:TreatmentHeavy:Site qualitySQ2   1.180  0.23924
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
    
```

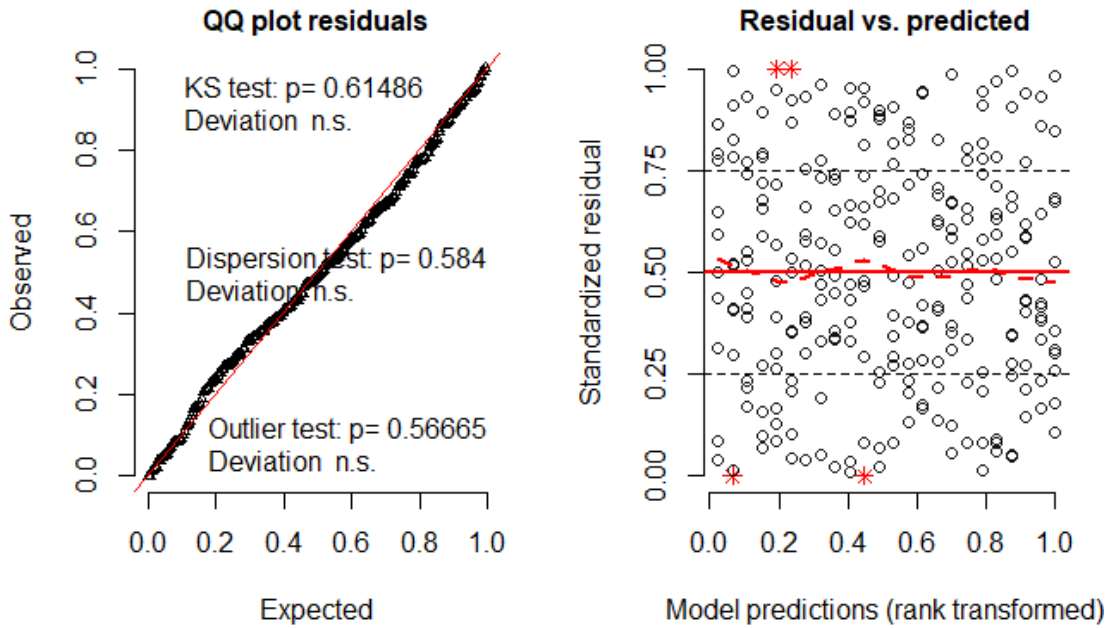


Figure 11 Dunn Smyth simulated residuals: average crown extent

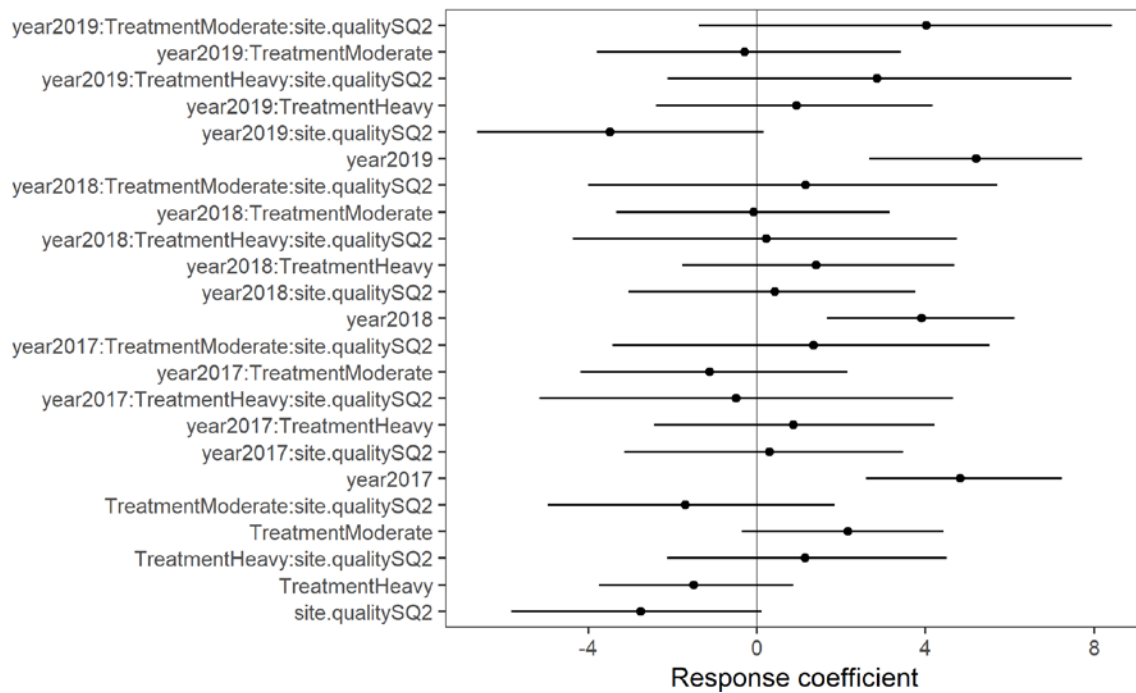


Figure 12 Bootstrapped confidence intervals for fixed effects: average crown extent



**Table 12 Bootstrapped fitted values for average conditions: average crown extent**

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	77.0392	79.0953	75.2230
2017	SQ1	81.6934	82.7277	80.9850
2018	SQ1	80.8175	82.7698	80.8600
2019	SQ1	81.9851	83.7398	81.5946
2015	SQ2	74.2049	74.5359	73.7595
2017	SQ2	79.4031	79.9347	79.2039
2018	SQ2	78.4629	80.1466	79.7105
2019	SQ2	75.7144	80.1640	79.2886

## D.1.2 Proportion of trees with recent crown extent decline

**Table 13 Model fitting summary: proportion of trees with recent crown extent decline**

<b>Response</b>	Proportion of 30 trees per 9 hectare plot with recent decline (between 2018 and 2019) in crown extent Ratio of trees with decline to trees with stable or increasing crown extent
<b>Response transformations used</b>	None
<b>Response transformations compared</b>	None
<b>R package and function</b>	glmer from the lme4 package
<b>Distribution used</b>	Binomial
<b>Other distributions compared</b>	Negative binomial
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and site quality
<b>Random factors</b>	fsite (a factor over sites)
<b>R<sup>2</sup></b>	Marginal = 8.7% Conditional = 27.0%
<b>Confidence comments</b>	Moderate. Assumptions met but some minor overdispersion apparent

## Model results summary 7: proportion of trees with recent decline in crown extent

```

y<-cbind(success = dat.num.trees2$Decline, failure = dat.num.trees2$Notdecline)

model2 <- glmer(y ~ Treatment * Site quality + (1|fsite), data=dat.num.trees2, family = binomial)

summary(model2)

## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula: y ~ Treatment * Site quality + (1 | fsite)
## Data: dat.num.trees2
##
##          AIC          BIC    logLik deviance df.resid
##    379.6      394.9   -182.8   365.6      59
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.5768 -0.8330 -0.1300  0.8484  3.4002
##
## Random effects:
## Groups Name          Variance Std.Dev.
## fsite (Intercept) 0.8248   0.9082
## Number of obs: 66, groups: fsite, 22
##
## Fixed effects:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -2.3983    0.3382  -7.092 1.32e-12 ***
## TreatmentModerate  0.3512    0.2408   1.458 0.14471
## TreatmentHeavy    0.2127    0.2446   0.870 0.38456
## Site qualitySQ2    1.7875    0.4510   3.964 7.38e-05 ***
## TreatmentModerate:Site qualitySQ2 -1.2517    0.3040  -4.117 3.83e-05 ***
## TreatmentHeavy:Site qualitySQ2  -0.8006    0.3020  -2.651 0.00803 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) TrtmnM TrtmnH st.SQ2 TM: .SQ
## TretmntMdrt -0.397
## TreatmentHvy -0.388  0.538
## sit.qltySQ2 -0.749  0.298  0.291
## TrtmnM: .SQ2  0.316 -0.792 -0.426 -0.339
## TrtmnH: .SQ2  0.315 -0.436 -0.810 -0.339  0.500

```

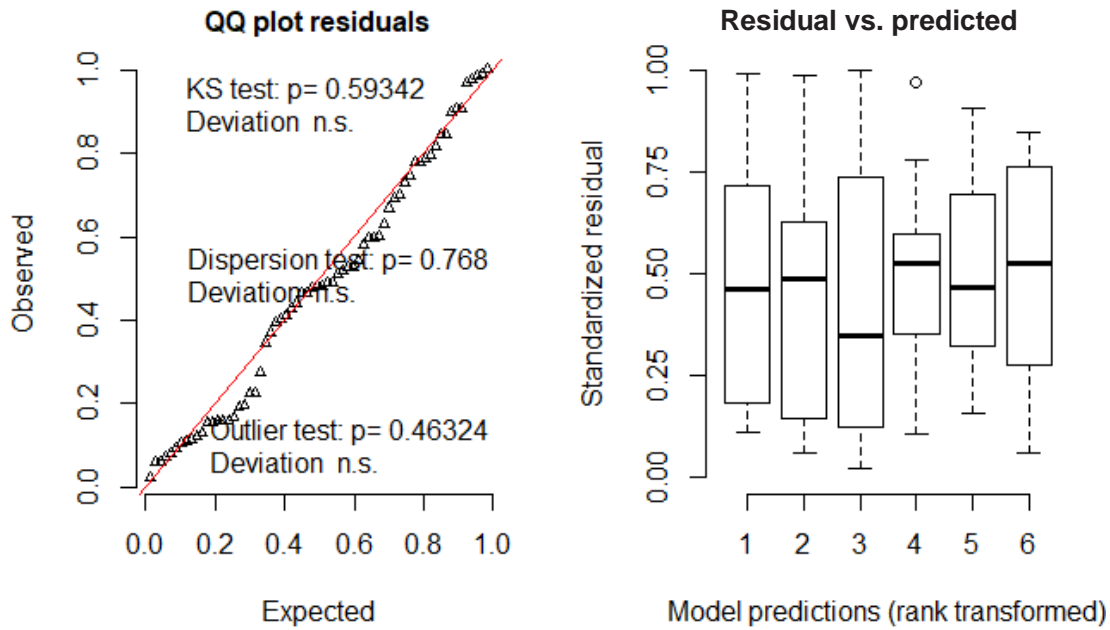


Figure 13 Dunn Smyth simulated residuals: proportion of trees with recent decline in crown extent

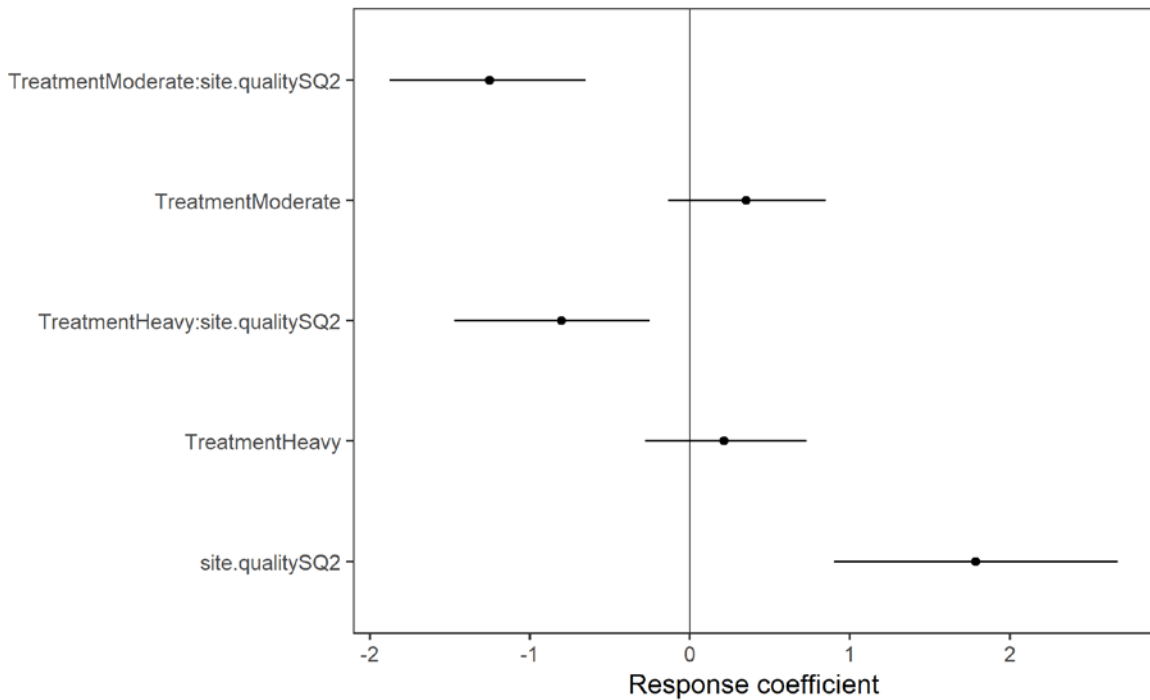


Figure 14 Bootstrapped confidence intervals for fixed effects: proportion of trees with recent decline in crown extent

Table 14 Bootstrapped model fitted values for average conditions: proportion of trees with recent decline in crown extent

Survey year	Site quality	Control	Moderate	Heavy
2019	SQ1	0.0784	0.1146	0.1021
2019	SQ2	0.3331	0.1833	0.2354



### D.1.3 Magnitude of recent crown extent decline

**Table 15 Model fitting summary: magnitude of recent crown extent decline**

<b>Response</b>	Probability of being in each 5% category of crown extent decline (e.g. category 1 is –5%; category 2 is –10%) Ordinal categorical variable
<b>Response transformations used</b>	None
<b>Response transformations compared</b>	None
<b>R package and function</b>	brm from brms package Note that cumulative ordinal regressions using the clm, clmm, and clmm2 from the ordinal package were attempted but the package was not well-developed enough to be useful
<b>Distribution used</b>	Cumulative with equidistant thresholds
<b>Other distributions compared</b>	Negative binomial, which violated many assumptions
<b>Outliers removed</b>	None
<b>Fixed factor</b>	Ecological thinning treatment (control, moderate, heavy) Site quality
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and site quality
<b>Random factor</b>	fsite (a factor over sites) and fsiteplot (a factor over plots nested within sites). Note that most plots had more than one tree with recent decline.
<b>Confidence comments</b>	Moderate

### Model results summary 8: magnitude of recent crown extent decline

```

model4_brms_cm <- brm(formula = change_recent_abs_or ~ Site quality * Treatment +
(1|fsite) + (1|fsiteplot),
  data = dat.decline.recent, family = cumulative(threshold = "equidistant
"),
  prior = set_prior("normal(0,1)"), control = list(adapt_delta = 0.99))

summary(model4_brms_cm)

## Family: cumulative
## Links: mu = logit; disc = identity
## Formula: change_recent_abs_or ~ Site quality * Treatment + (1 | fsite) + (1 | f
siteplot)
## Data: dat.decline.recent (Number of observations: 2045)
## Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup samples = 4000
##
## Group-Level Effects:
## ~fsite (Number of levels: 21)

```

```

##           Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)      1.12      0.37      0.51      1.98 1.00      1292      1281
##
## ~fsiteplot (Number of levels: 61)
##           Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)      0.83      0.20      0.50      1.27 1.00      1032      1708
##
## Population-Level Effects:
##
##           Estimate Est.Error 1-95% CI u-95% CI Rhat
## Intercept[1]          1.92      0.47      1.05      2.87 1.00
## Intercept[2]          2.48      0.47      1.62      3.43 1.00
## Intercept[3]          3.04      0.47      2.17      4.00 1.00
## Intercept[4]          3.60      0.48      2.73      4.57 1.00
## Intercept[5]          4.16      0.48      3.28      5.14 1.00
## Intercept[6]          4.72      0.48      3.83      5.71 1.00
## Intercept[7]          5.29      0.49      4.38      6.27 1.00
## Intercept[8]          5.85      0.50      4.92      6.84 1.00
## Intercept[9]          6.41      0.50      5.47      7.41 1.00
## Intercept[10]         6.97      0.51      6.01      7.97 1.00
## Intercept[11]         7.53      0.52      6.56      8.55 1.00
## Intercept[12]         8.09      0.53      7.10      9.11 1.00
## Site qualitySQ2        0.56      0.55     -0.52      1.64 1.00
## TreatmentModerate     -0.12      0.42     -0.94      0.69 1.00
## TreatmentHeavy         0.21      0.41     -0.61      0.98 1.00
## Site qualitySQ2:TreatmentModerate -0.51      0.50     -1.51      0.52 1.00
## Site qualitySQ2:TreatmentHeavy   -0.73      0.52     -1.73      0.29 1.00
##
##           Bulk_ESS Tail_ESS
## Intercept[1]          2927      3115
## Intercept[2]          2932      3080
## Intercept[3]          2945      3064
## Intercept[4]          2969      3219
## Intercept[5]          2997      3114
## Intercept[6]          3036      3038
## Intercept[7]          3085      3197
## Intercept[8]          3139      3144
## Intercept[9]          3198      3279
## Intercept[10]         3269      3266
## Intercept[11]         3341      3085
## Intercept[12]         3420      2999
## Site qualitySQ2        2737      2774
## TreatmentModerate      3444      3018
## TreatmentHeavy         3068      2924
## Site qualitySQ2:TreatmentModerate  3306      3259
## Site qualitySQ2:TreatmentHeavy    2826      3013
##
## Family Specific Parameters:
##           Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## delta      0.56      0.02      0.52      0.60 1.00      9487      2865
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).

```

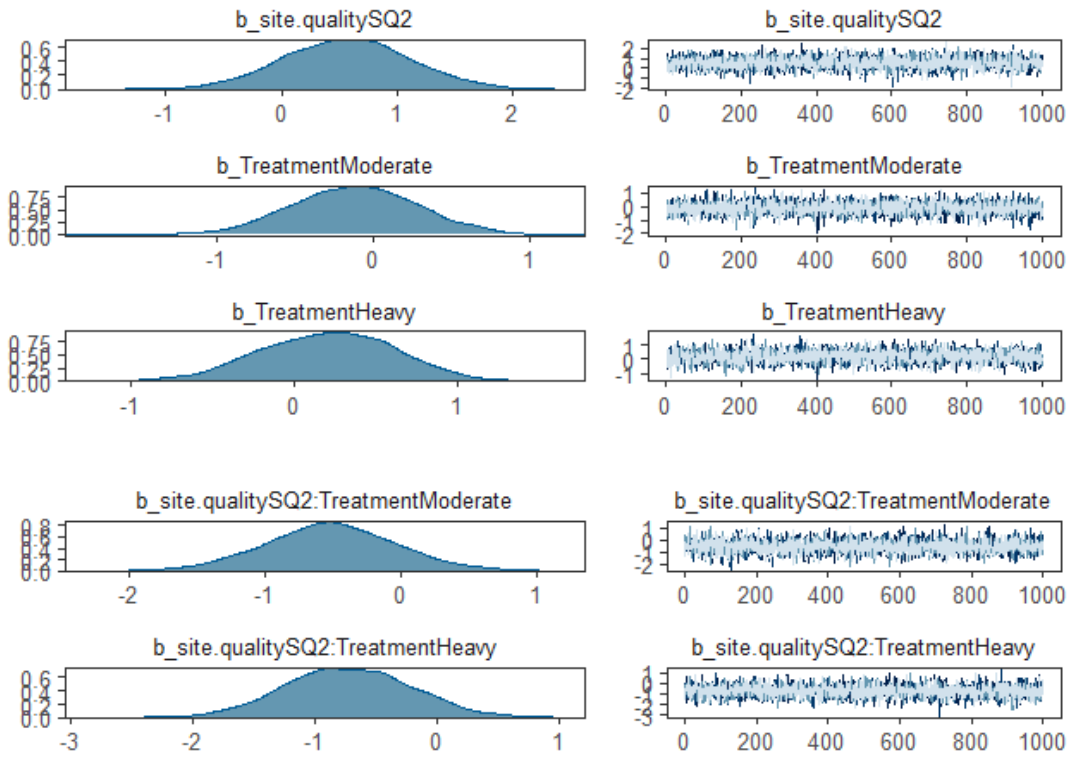


Figure 15 Posterior distributions (left) and chain mixing (right): magnitude of recent crown extent decline



## D.2 Visually assessed canopy cover

### D.2.1 Live canopy cover

**Table 16 Model fitting summary: visually assessed live canopy cover**

<b>Response</b>	Visually assessed canopy cover A continuous positive variable
<b>Response transformations used</b>	Divided by 100
<b>Response transformations compared</b>	Log <sub>e</sub> was trialled with a gaussian family (see distributions compared below)
<b>R package and function</b>	glmmTMB from the glmmTMB package
<b>Distribution used</b>	Beta
<b>Other distributions compared</b>	Gaussian (with a log transformation of the response) – failed Dunn Smyth tests Binomial – failed Dunn Smyth tests
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Also trialled a model with a two-way interaction between thinning treatment and survey year A model with a three-way interaction between thinning treatment, survey year and site quality was the most parsimonious
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Function not available to calculate R <sup>2</sup> for beta family
<b>Confidence comments</b>	Moderate to high. Standard QQ plot indicated some larger residuals than expected for higher values of the response, but passed all Dunn Smyth tests

## Model results summary 9: visually assessed live canopy cover

Family: beta ( logit )  
 Formula: cover.live/100 ~ Survey year \* Site quality \* Treatment + (1 | fsite) + (1 | fsiteplot)  
 Data: DAT.treecover

AIC	BIC	logLik	deviance	df.resid
-2842.8	-2716.7	1448.4	-2896.8	761

Random effects:

Conditional model:

Groups	Name	Variance	Std.Dev.
fsite	(Intercept)	0.006142	0.07837
fsiteplot	(Intercept)	0.053380	0.23104

Number of obs: 788, groups: fsite, 22; fsiteplot, 66  
 Overdispersion parameter for beta family (): 47.3

Conditional model:

	Estimate	Std. Error	z value	Pr(> z )	
(Intercept)	-1.93554	0.10659	-18.158	< 2e-16	***
Survey year2017	-0.12353	0.10893	-1.134	0.2568	
Survey year2018	-0.22946	0.11023	-2.082	0.0374	*
Survey year2019	-0.67622	0.12197	-5.544	2.96e-08	***
Site qualitySQ2	-0.27864	0.15333	-1.817	0.0692	.
TreatmentModerate	0.14749	0.14360	1.027	0.3044	
TreatmentHeavy	-0.12159	0.14583	-0.834	0.4044	
Survey year2017:Site qualitySQ2	0.32574	0.15607	2.087	0.0369	*
Survey year2018:Site qualitySQ2	0.14811	0.16081	0.921	0.3571	
Survey year2019:Site qualitySQ2	-0.22088	0.18415	-1.199	0.2304	
Survey year2017:TreatmentModerate	-0.35419	0.15445	-2.293	0.0218	*
Survey year2018:TreatmentModerate	-0.26230	0.15492	-1.693	0.0904	.
Survey year2019:TreatmentModerate	-0.10272	0.16758	-0.613	0.5399	
Survey year2017:TreatmentHeavy	-0.21603	0.15704	-1.376	0.1689	
Survey year2018:TreatmentHeavy	-0.17596	0.15893	-1.107	0.2682	
Survey year2019:TreatmentHeavy	-0.30503	0.17934	-1.701	0.0890	.
Site qualitySQ2:TreatmentModerate	-0.16035	0.20897	-0.767	0.4429	
Site qualitySQ2:TreatmentHeavy	-0.08018	0.21300	-0.376	0.7066	
Survey year2017:Site qualitySQ2:TreatmentModerate	0.02747	0.22397	0.123	0.9024	
Survey year2018:Site qualitySQ2:TreatmentModerate	0.15225	0.22775	0.668	0.5038	
Survey year2019:Site qualitySQ2:TreatmentModerate	0.32864	0.25361	1.296	0.1950	
Survey year2017:Site qualitySQ2:TreatmentHeavy	-0.12658	0.23073	-0.549	0.5833	
Survey year2018:Site qualitySQ2:TreatmentHeavy	0.04586	0.23533	0.195	0.8455	
Survey year2019:Site qualitySQ2:TreatmentHeavy	0.56076	0.26583	2.109	0.0349	*

---  
 Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

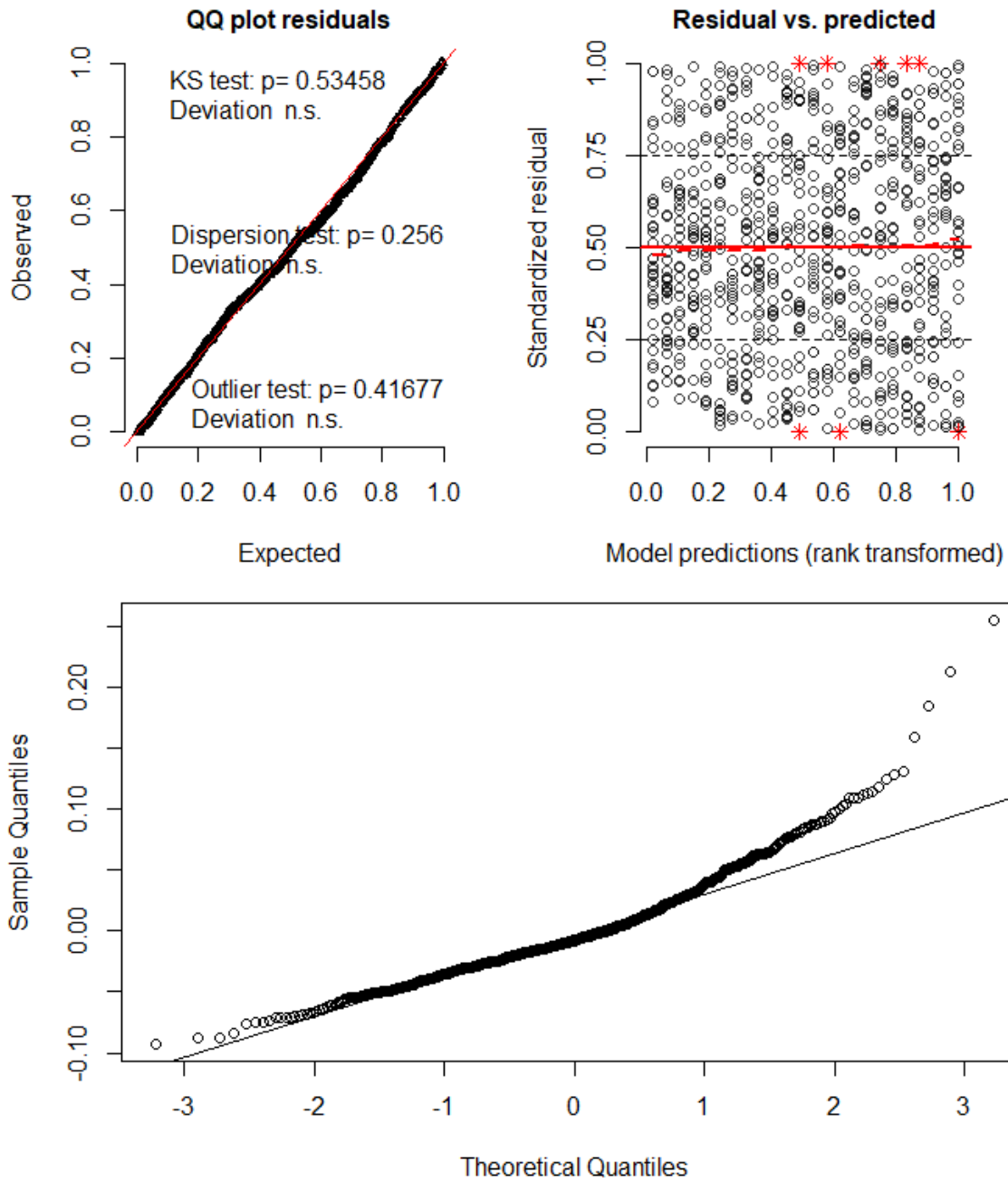


Figure 16 Dunn Smyth simulated residuals (top panels) and normal QQ plot (lower panel): visually assessed live canopy cover



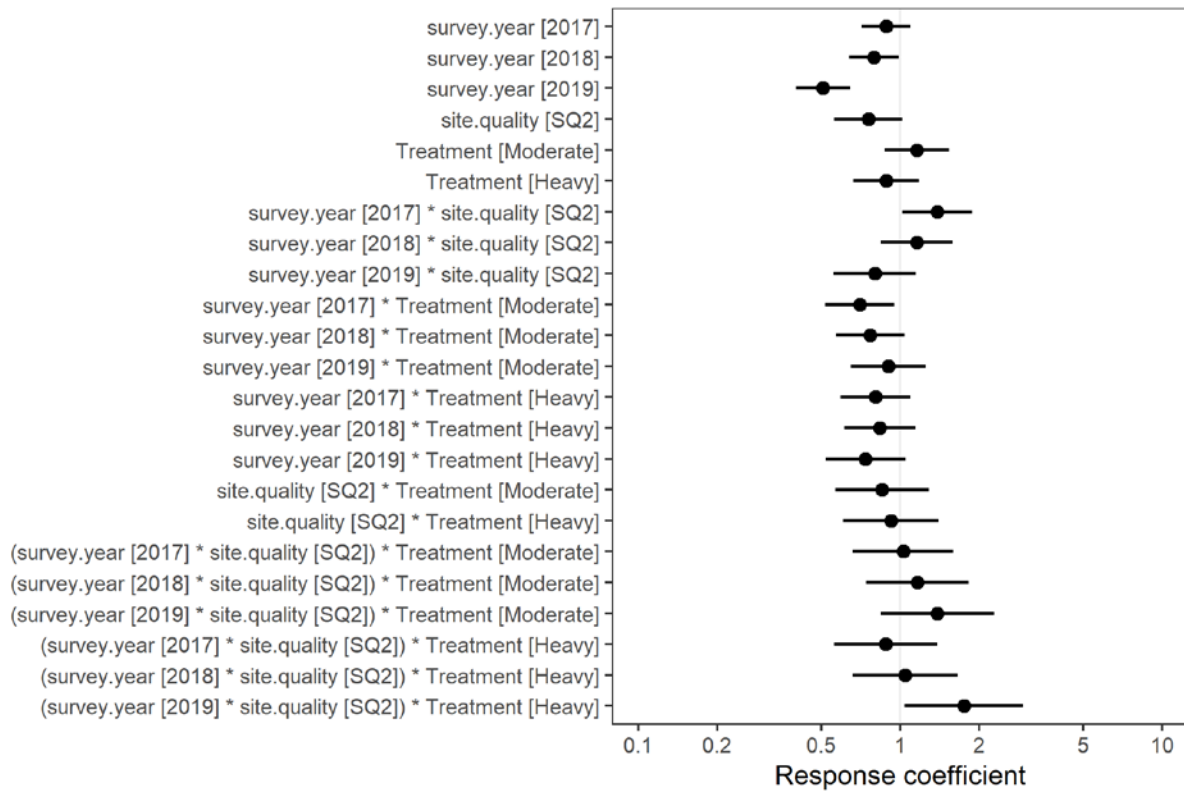


Figure 17 Wald confidence intervals for fixed effects: visually assessed live canopy cover

Table 17 Model fitted values for average conditions: visually assessed live canopy cover

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	12.6939	14.4203	11.4063
2017	SQ1	11.3868	9.4616	8.3980
2018	SQ1	10.3608	9.3420	7.9051
2019	SQ1	6.8848	7.1774	4.6038
2015	SQ2	9.9129	9.7986	8.2511
2017	SQ2	11.8707	8.7519	7.2486
2018	SQ2	9.2097	8.2321	6.7851
2019	SQ2	4.2941	5.2601	4.5214

## D.2.2 Dead canopy cover

No model was run because there is no ecological difference between 0.5% dead and 1% dead canopy cover.

## D.3 Remotely sensed canopy cover

**Table 18 Model fitting summary: remotely sensed foliage projective cover**

<b>Response</b>	Foliage projective cover (derived from Landsat images), median of up to 9 values per 9 hectare plot
<b>Response transformation used</b>	Logit
<b>Other transformations compared</b>	None
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning Site quality (SQ1, SQ2) Pass (of Landsat satellite, approximately every 16 days) CosTime (to represent seasonality) River flow maximum (megalitres per day) from Yarrawonga in the previous year
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and site quality
<b>Random factors</b>	A factor over 9 ha treatment plots A spherical correlation structure to account for autocorrelation among plots over time
<b>R<sup>2</sup></b>	Marginal = 52.7% Conditional = 72.6%
<b>Confidence comments</b>	Moderate to high. Dunn Smyth residual tests were not able to be run on a model with autocorrelation. Normal QQ plots and standardised residuals show some deviation from expected behaviour for highest and lowest values. No model violations detected

## Model results summary 10: remotely sensed foliage projective cover

```

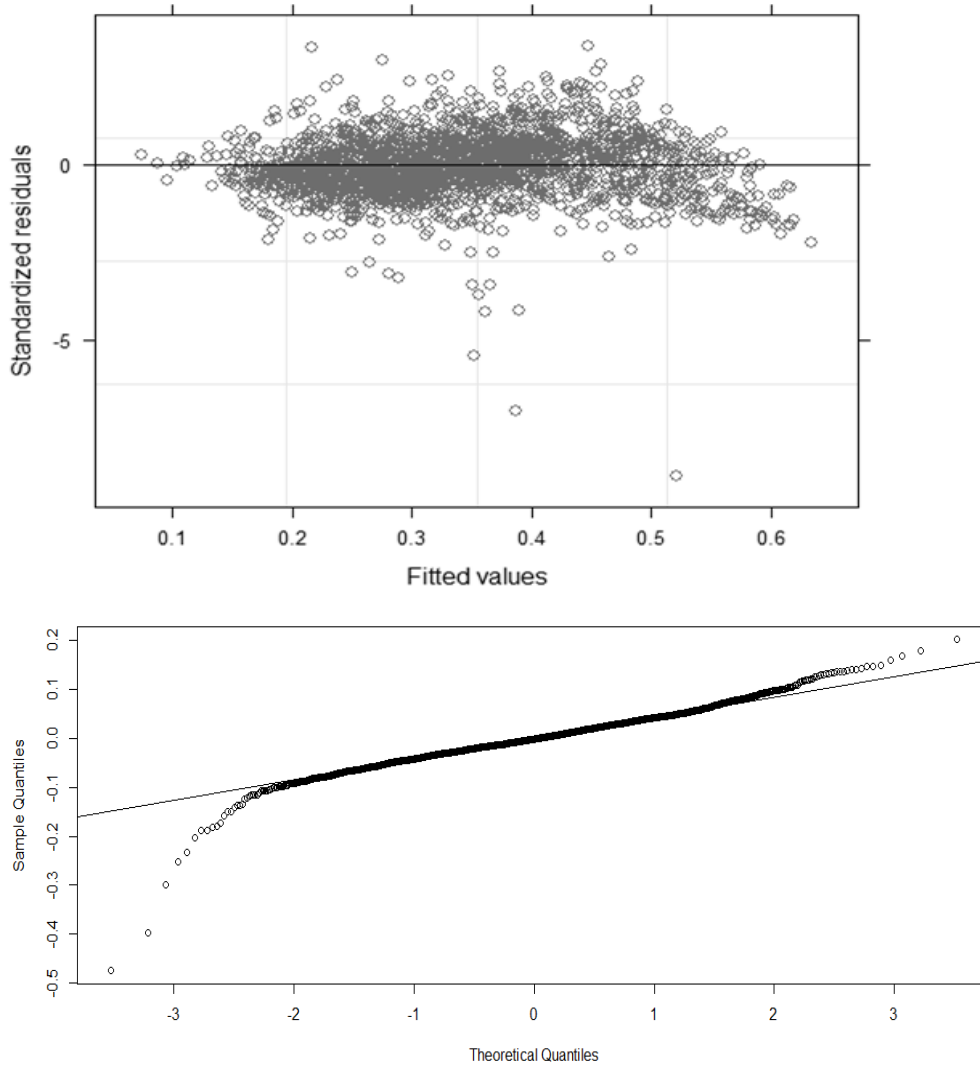
Linear mixed-effects model fit by REML
Data: p3x
      AIC      BIC    logLik
-8001.621 -7932.738 4012.811

Random effects:
Formula: ~1 | f
      (Intercept)  Residual
StdDev:  0.0523178 0.06134366

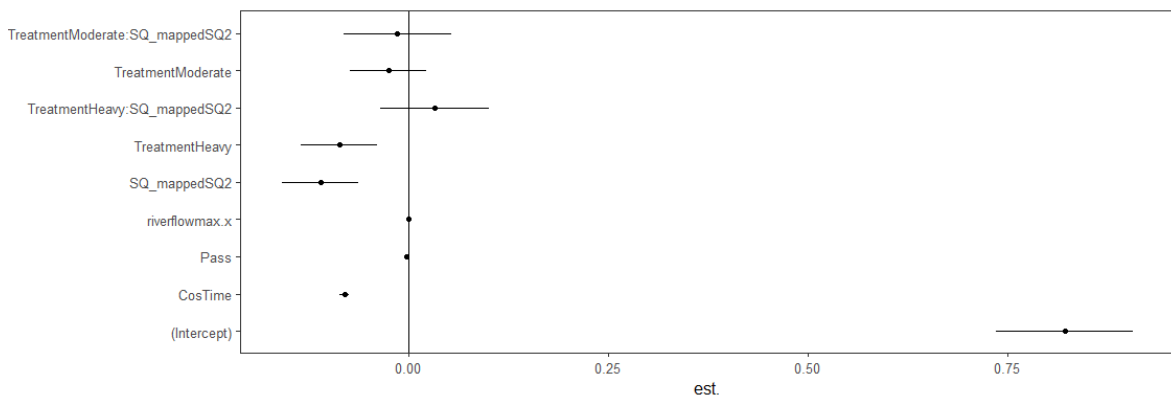
Correlation Structure: Spherical spatial correlation
Formula: ~Pass | f
Parameter estimate(s):
      range
8.758559
Fixed effects: FPC_med ~ Treatment * SQ_mapped + Pass + CosTime + riverflowmax.x
              Value Std.Error   DF   t-value p-value
(Intercept)    0.8211735 0.04355385 2239  18.854211 0.0000
TreatmentModerate -0.0247781 0.02386276   60  -1.038358 0.3033
TreatmentHeavy    -0.0861035 0.02386278   60  -3.608274 0.0006
SQ_mappedSQ2     -0.1096509 0.02386276   60  -4.595062 0.0000
Pass             -0.0015475 0.00015597 2239  -9.921636 0.0000
CosTime          -0.0798731 0.00318020 2239 -25.115775 0.0000
riverflowmax.x    0.0000003 0.00000006 2239   5.088963 0.0000
TreatmentModerate:SQ_mappedSQ2 -0.0137029 0.03374704   60  -0.406049 0.6862
TreatmentHeavy:SQ_mappedSQ2    0.0330160 0.03374706   60   0.978338 0.3318
Correlation:
              (Intr) TrtmnM TrtmnH SQ_SQ2 Pass   CosTim rvrfl. TM:SQ_
TreatmentModerate -0.274
TreatmentHeavy    -0.274 0.500
SQ_mappedSQ2     -0.274 0.500 0.500
Pass             -0.921 0.000 0.000 0.000
CosTime          -0.093 0.000 0.000 0.000 0.083
riverflowmax.x   -0.521 0.000 0.000 0.000 0.527 0.258
TreatmentModerate:SQ_mappedSQ2 0.194 -0.707 -0.354 -0.707 0.000 0.000 0.000
TreatmentHeavy:SQ_mappedSQ2    0.194 -0.354 -0.707 -0.707 0.000 0.000 0.000 0.500

Standardised Within-Group Residuals:
      Min      Q1      Med      Q3      Max
-7.72298832 -0.46197102 -0.02630681 0.46251712 3.28337711

Number of Observations: 2308
Number of Groups: 66
    
```



**Figure 18** Standardised residuals (top panel) and normal QQ plot (lower panel): remotely sensed foliage projective cover



**Figure 19** Wald confidence intervals for fixed effects: remotely sensed foliage projective cover



**Table 19 Model fitted values for average conditions: remotely sensed foliage projective cover**

Survey year	Site quality	Control	Moderate	Heavy
First pass: 2017	SQ1	0.49	0.43	0.40
Most recent pass: 2020	SQ1	0.36	0.35	0.31
First pass: 2017	SQ2	0.39	0.32	0.29
Most recent pass: 2020	SQ2	0.25	0.25	0.21

## D.4 Plant area index

**Table 20 Model fitting summary: plant area index**

<b>Response</b>	Plant area index is the amount of plant material relative to a unit of sky Five estimates were obtained per 9 hectare plot A positive continuous variable
<b>Response transformation used</b>	None
<b>Response transformations compared</b>	None
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Site quality (SQ1, SQ2)
<b>Fixed factors interactions</b>	Two-way interaction between thinning treatment and site quality
<b>Random factors</b>	site (a factor over sites), and siteplot (a factor over 9 ha treatment plots)
<b>R<sup>2</sup></b>	Marginal = 45.7% Conditional = 72.7%
<b>Confidence comments</b>	High. Passed all Dunn Smyth tests

### Model results summary II: plant area index

```
modelf <- lmer(PAI ~ Treatment * Site quality + (1|siteplot) + (1|site),
              data = DAT.PAI)
```

```
summary(modelf)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: PAI ~ Treatment * Site quality + (1 | siteplot) + (1 | site)
```

```
## Data: DAT.PAI
##
## REML criterion at convergence: -70
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.51841 -0.55997  0.03256  0.57244  3.16180
##
## Random effects:
## Groups   Name      Variance Std.Dev.
## siteplot (Intercept) 0.01418  0.1191
## site     (Intercept) 0.01819  0.1349
## Residual                0.03266  0.1807
## Number of obs: 330, groups: siteplot, 66; site, 22
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)      1.33564    0.05946 41.74981  22.461 < 2e-16
## TreatmentModerate -0.20636    0.06136 39.99822  -3.363  0.00171
## TreatmentHeavy    -0.45764    0.06136 39.99822  -7.458  4.35e-09
## Site qualitySQ2   -0.46836    0.08410 41.74981  -5.569  1.68e-06
## TreatmentModerate:Site qualitySQ2  0.08145    0.08678 39.99822   0.939  0.35355
## TreatmentHeavy:Site qualitySQ2    0.23564    0.08678 39.99822   2.715  0.00973
##
## (Intercept)          ***
## TreatmentModerate     **
## TreatmentHeavy        ***
## Site qualitySQ2       ***
## TreatmentModerate:Site qualitySQ2
## TreatmentHeavy:Site qualitySQ2    **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

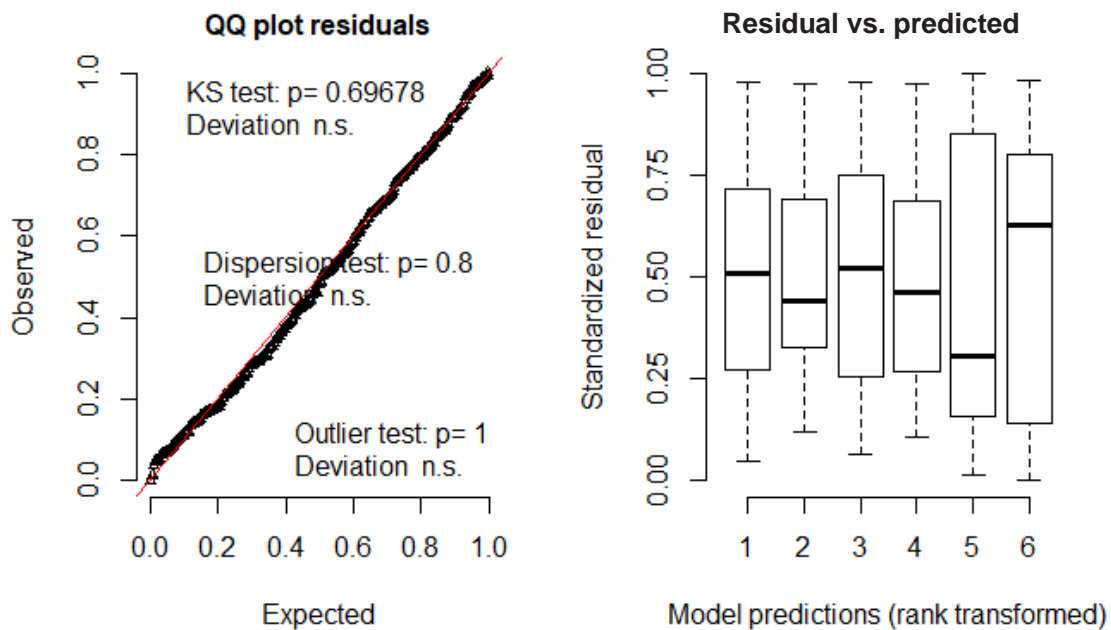
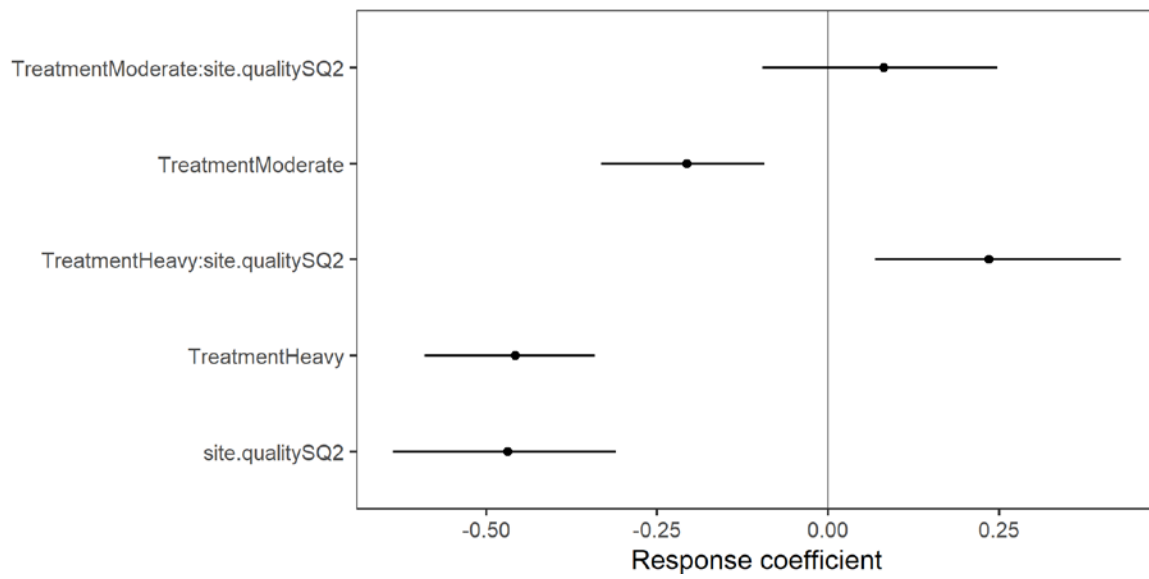


Figure 20 Dunn Smyth simulated residuals: plant area index



**Figure 21** Bootstrapped confidence intervals for fixed effects: plant area index

**Table 21** Bootstrapped model fitted values for average conditions: plant area index

Survey year	Site quality	Control	Moderate	Heavy
2019	SQ1	1.3280	1.1345	0.8794
2019	SQ2	0.8698	0.7266	0.6544

# Appendix E: Model summaries – Fuel hazard

## E.1 Overall fuel hazard

**Table 22 Model fitting summary: overall fuel hazard**

<b>Response</b>	Probability of being in each overall fuel hazard category
<b>Response transformation used</b>	None
<b>Response transformations compared</b>	None
<b>R package and function</b>	brms with family specified as cratio(threshold = flexible)
<b>Distribution used</b>	Continuation ratio with flexible category thresholds The cumulative model has an underlying distribution that spans all categories, which is appropriate for fuel hazard because hazard is an underlying continuous variable and responses can move up and down as well as jump categories. (In contrast, adjacent categories assumes each level of risk rating has its own distribution, and sequential models assume response data can only move sequentially through the levels). Flexible thresholds allow for the boundaries between hazard rating categories to have unequal sizes and account for different proportions of the data. This is supported by the input data where more sites report low or moderate ratings than very high or extreme ratings.
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>Confidence comments</b>	Low. A warning was issued that the Tail Effective Samples Size (ESS) was too low, indicating posterior variances and tail quantiles may be unreliable



## Model results summary 12: overall fuel hazard

```

## Family: cratio
## Links: mu = logit; disc = identity
## Formula: risk.total.or ~ Survey year * Treatment + Site quality + (1 | fsiteplot) + (1 | fsite)
## Data: DAT.fire (Number of observations: 789)
## Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
##           total post-warmup samples = 4000
##
## Group-Level Effects:
## ~fsite (Number of levels: 22)
##           Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)      0.39      0.18      0.04      0.75 1.00      553      787
##
## ~fsiteplot (Number of levels: 66)
##           Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)      0.50      0.16      0.13      0.80 1.00      532      302
##
## Population-Level Effects:
##           Estimate Est.Error 1-95% CI u-95% CI Rhat
## Intercept[1]      -2.65      0.34      -3.36      -1.98 1.00
## Intercept[2]       0.13      0.31      -0.49      0.75 1.00
## Intercept[3]       7.03      0.85       5.56      8.93 1.00
## Survey year2017     0.37      0.34      -0.27      1.05 1.00
## Survey year2018     1.35      0.37       0.61      2.08 1.00
## Survey year2019     1.42      0.38       0.68      2.19 1.00
## TreatmentModerate  -0.02      0.37      -0.74      0.68 1.00
## TreatmentHeavy     0.14      0.37      -0.59      0.87 1.00
## Site qualitySQ2     0.06      0.27      -0.48      0.59 1.00
## Survey year2017:TreatmentModerate -0.30      0.47      -1.23      0.63 1.00
## Survey year2018:TreatmentModerate -0.33      0.51      -1.30      0.67 1.00
## Survey year2019:TreatmentModerate -0.09      0.52      -1.11      0.91 1.00
## Survey year2017:TreatmentHeavy -0.23      0.49      -1.19      0.69 1.00
## Survey year2018:TreatmentHeavy -0.08      0.53      -1.07      0.96 1.00
## Survey year2019:TreatmentHeavy -0.39      0.52      -1.43      0.62 1.00
##
##           Bulk_ESS Tail_ESS
## Intercept[1]      2065      2590
## Intercept[2]      1772      2172
## Intercept[3]      3630      2445
## Survey year2017     1789      2332
## Survey year2018     1887      2815
## Survey year2019     1687      2090
## TreatmentModerate   1615      1965
## TreatmentHeavy     1583      2219
## Site qualitySQ2     2828      2882
## Survey year2017:TreatmentModerate  1987      2352
## Survey year2018:TreatmentModerate  2048      2812
## Survey year2019:TreatmentModerate  1920      2647
## Survey year2017:TreatmentHeavy    1899      2552
## Survey year2018:TreatmentHeavy    1967      2927
## Survey year2019:TreatmentHeavy    1864      2647
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).

```

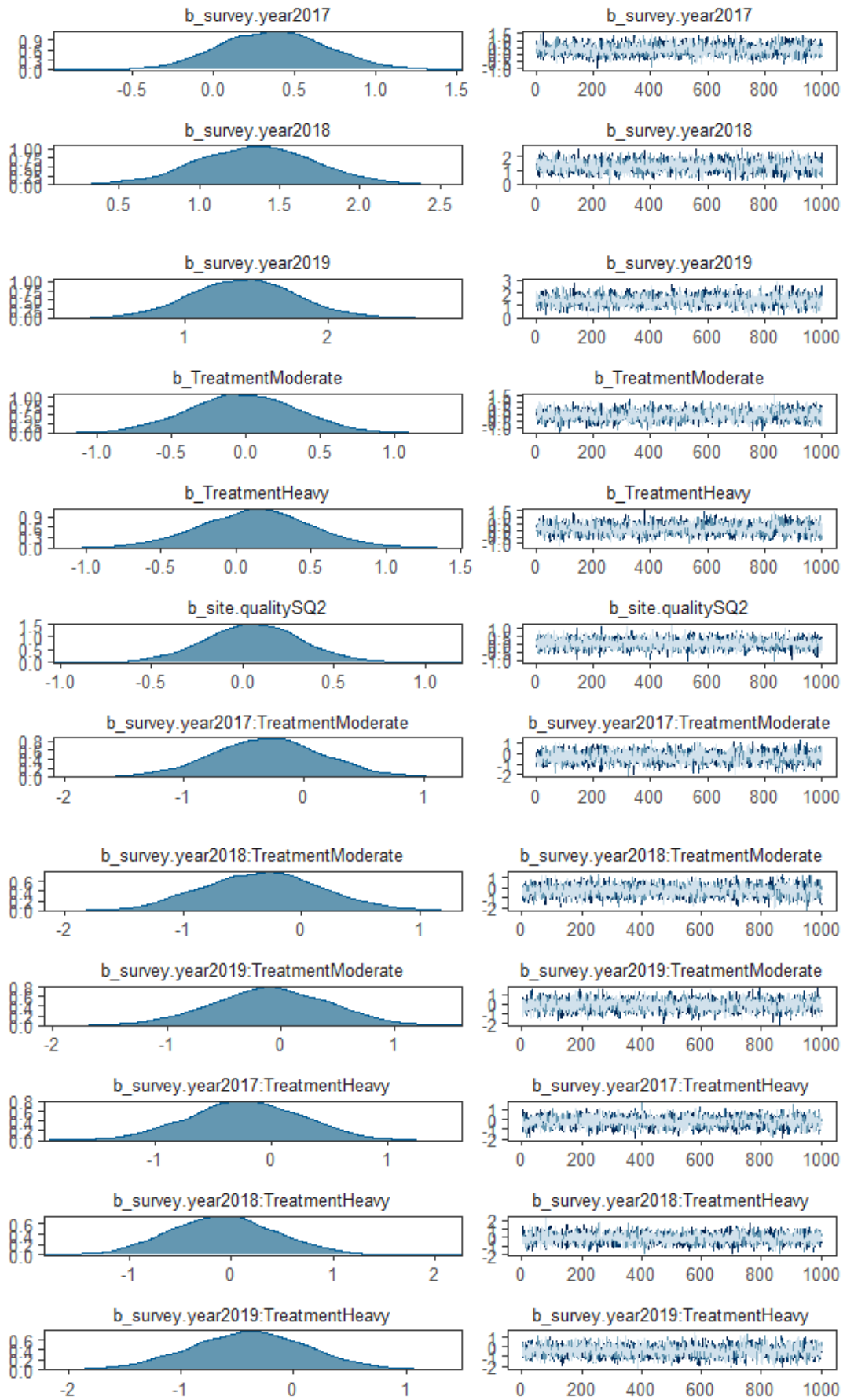


Figure 22 Posterior distributions (left) and chain mixing (right): overall fuel hazard

## E.2 Surface fuel hazard

### E.2.1 Litter depth

**Table 23 Model fitting summary: litter depth**

<b>Response</b>	Litter depth, measured at 30 locations in each 9 hectare plot, in millimetres A continuous positive variable Log transformed
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two two-way interactions between thinning treatment and survey year, and thinning treatment and site quality A three-way model between thinning treatment, survey year and site quality was trialled but was not as parsimonious
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Marginal = 3.5% Conditional = 18.7%
<b>Confidence comments</b>	Moderate. Failed the Dunn Smyth uniformity test, with minor deviations

**Model results summary 13: litter depth**

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: log(litter.depth) ~ Survey year * Treatment + Site quality *
##   Treatment + (1 | siteplot) + (1 | fsite)
##   Data: DAT.litter

## REML criterion at convergence: 1166
##
## Scaled residuals:
##   Min      1Q  Median      3Q      Max
## -4.2699 -0.5441  0.0628  0.6261  2.7392
##
## Random effects:
##   Groups   Name                Variance Std.Dev.
##   siteplot (Intercept) 0.02405  0.1551
##   fsite    (Intercept) 0.01787  0.1337
##   Residual                0.22452  0.4738
## Number of obs: 787, groups:  siteplot, 66; fsite, 22
##
## Fixed effects:
##
##              Estimate Std. Error      df t value
## (Intercept)      2.78216    0.08985 107.97966  30.963
## Survey year2017    0.26937    0.08316 713.00082   3.239
## Survey year2018    0.03320    0.08282 712.72343   0.401
## Survey year2019    0.05132    0.08248 712.39259   0.622
## TreatmentModerate -0.09638    0.11353 107.18907  -0.849
## TreatmentHeavy     0.12870    0.11325 106.14693   1.136
## Site qualitySQ2    -0.11687    0.10519  51.50363  -1.111
## Survey year2017:TreatmentModerate -0.10186    0.11713 712.69952  -0.870
## Survey year2018:TreatmentModerate  0.04128    0.11689 712.55878   0.353
## Survey year2019:TreatmentModerate  0.03144    0.11665 712.39259   0.270
## Survey year2017:TreatmentHeavy    -0.15802    0.11765 713.27191  -1.343
## Survey year2018:TreatmentHeavy    -0.02982    0.11645 712.56002  -0.256
## Survey year2019:TreatmentHeavy    -0.09142    0.11668 712.63909  -0.784
## TreatmentModerate:Site qualitySQ2  0.22032    0.12487  40.18186   1.764
## TreatmentHeavy:Site qualitySQ2     0.04607    0.12501  40.34749   0.369
##
##              Pr(>|t|)
## (Intercept)    < 2e-16 ***
## Survey year2017  0.00125 **
## Survey year2018  0.68866
## Survey year2019  0.53401
## TreatmentModerate  0.39778
## TreatmentHeavy    0.25834
## Site qualitySQ2    0.27172
## Survey year2017:TreatmentModerate  0.38477
## Survey year2018:TreatmentModerate  0.72407
## Survey year2019:TreatmentModerate  0.78759
## Survey year2017:TreatmentHeavy    0.17966
## Survey year2018:TreatmentHeavy    0.79797
## Survey year2019:TreatmentHeavy    0.43358
## TreatmentModerate:Site qualitySQ2  0.08526 .
## TreatmentHeavy:Site qualitySQ2     0.71440
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation matrix not shown by default, as p = 15 > 12.
## Use print(x, correlation=TRUE) or
##   vcov(x)      if you need it
```



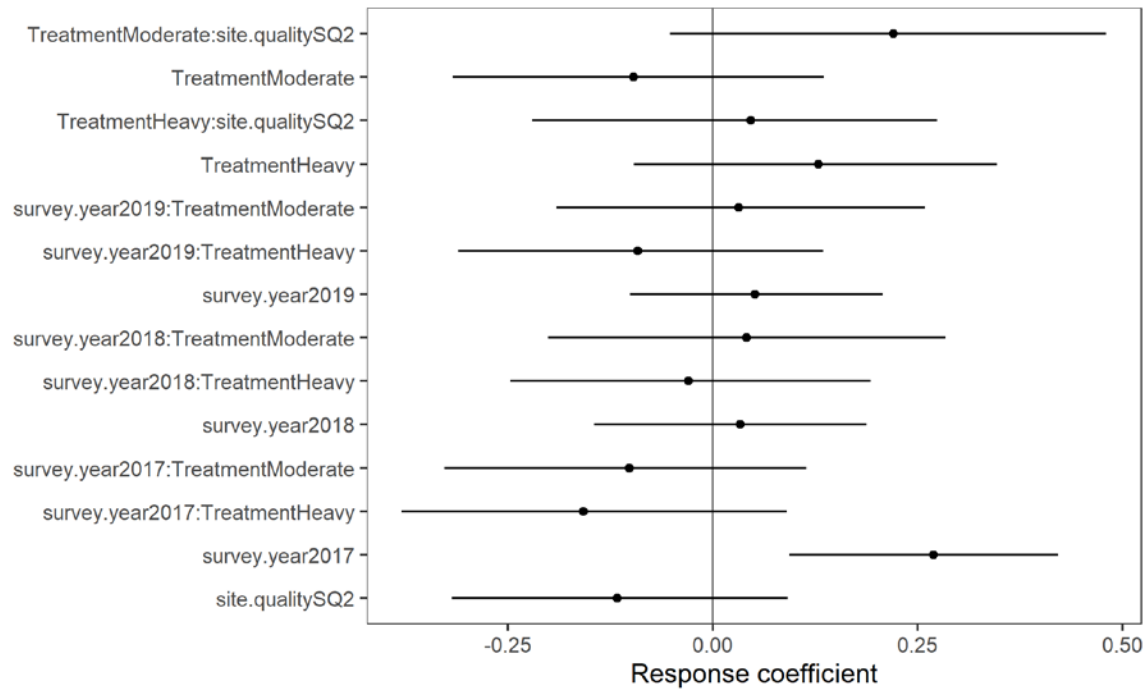
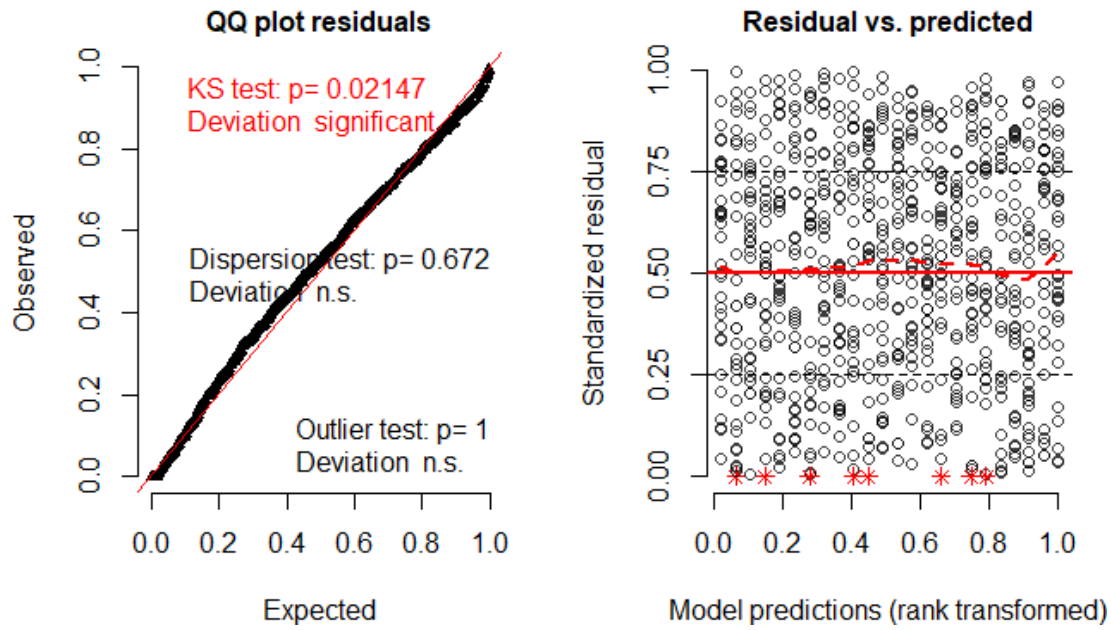


Figure 23 Bootstrapped confidence intervals for fixed effects: litter depth

Table 24 Bootstrapped model fitted values for average conditions: litter depth

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	16.0733	14.6850	18.4503
2017	SQ1	13.9698	16.4513	17.1007
2018	SQ1	21.8291	17.4387	20.6454
2019	SQ1	18.6883	19.0553	18.2791
2015	SQ2	16.7746	15.4586	18.4860
2017	SQ2	14.7309	17.9886	17.3682
2018	SQ2	16.5660	16.4356	17.5555
2019	SQ2	14.6351	17.0655	16.0848

## E.2.2 Litter cover

**Table 25 Model fitting summary: litter cover**

<b>Response</b>	Litter cover is visually estimated as a percentage of three 0.04 ha plots per 9 hectare plot Proportion variable
<b>Response transformation</b>	None
<b>Response transformations compared</b>	Log + 1 Square root Z transformed Logit 100 – x (where x is rounded to nearest whole value)
<b>R package and function</b>	glmmTMB from package glmmTMB
<b>Distribution used</b>	Beta
<b>Other distributions compared</b>	Binomial Negative binomial (using 100 – x transformation) Gaussian
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Interaction between thinning treatment and survey year Also trialled a model with three-way interaction between thinning treatment, survey year and site quality, which was less parsimonious
<b>Random factors</b>	site (a factor over sites), and siteplot (a factor over 9 ha treatment plots)
<b>R<sup>2</sup></b>	Function not available to calculate R <sup>2</sup> for beta family
<b>Confidence comments</b>	Low. Failed Dunn Smyth dispersion and uniformity tests

### Model results summary 14: litter cover

```
modelf.cover.beta <- glmmTMB(litter.prop ~ Survey year*Treatment + Site quality +
(1|siteplot) + (1|site),
  data = DAT.cover, family = beta_family())
```

```
summary(modelf.cover.beta)
```

```
## Family: beta ( logit )
## Formula:
## litter.prop ~ Survey year * Treatment + Site quality + (1 | siteplot) +
## (1 | site)
## Data: DAT.cover
##
##      AIC      BIC   logLik deviance df.resid
## -1301.2 -1226.5   666.6  -1333.2     771
```

```
##
## Random effects:
##
## Conditional model:
## Groups Name Variance Std.Dev.
## siteplot (Intercept) 0.01426 0.1194
## site (Intercept) 0.06226 0.2495
## Number of obs: 787, groups: siteplot, 66; site, 22
##
## Overdispersion parameter for beta family (:): 6.75
##
## Conditional model:
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) 1.321136 0.132926 9.939 < 2e-16 ***
## Survey year2017 -0.193306 0.142713 -1.355 0.17557
## Survey year2018 0.446458 0.150243 2.972 0.00296 **
## Survey year2019 0.600563 0.151181 3.972 7.11e-05 ***
## TreatmentModerate -0.005564 0.146063 -0.038 0.96961
## TreatmentHeavy 0.161643 0.148141 1.091 0.27521
## Site qualitySQ2 0.188078 0.126011 1.493 0.13555
## Survey year2017:TreatmentModerate -0.086018 0.199101 -0.432 0.66572
## Survey year2018:TreatmentModerate 0.118504 0.211171 0.561 0.57468
## Survey year2019:TreatmentModerate -0.158092 0.210896 -0.750 0.45348
## Survey year2017:TreatmentHeavy -0.295409 0.201194 -1.468 0.14203
## Survey year2018:TreatmentHeavy -0.148226 0.211401 -0.701 0.48320
## Survey year2019:TreatmentHeavy -0.188723 0.213477 -0.884 0.37667
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

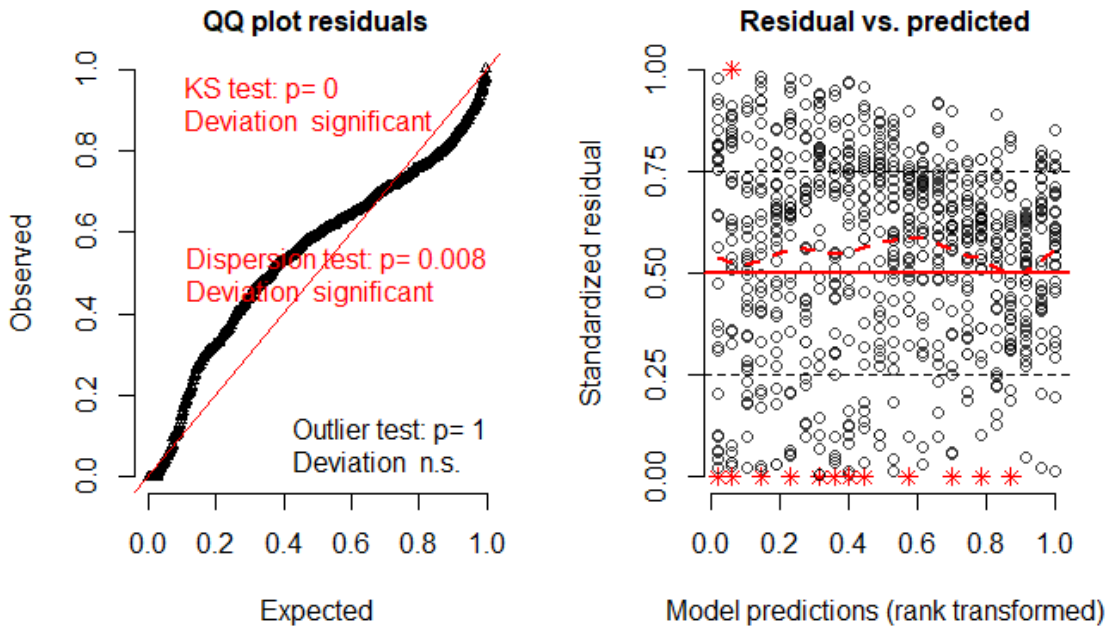


Figure 24 Dunn Smyth simulated residuals: litter cover

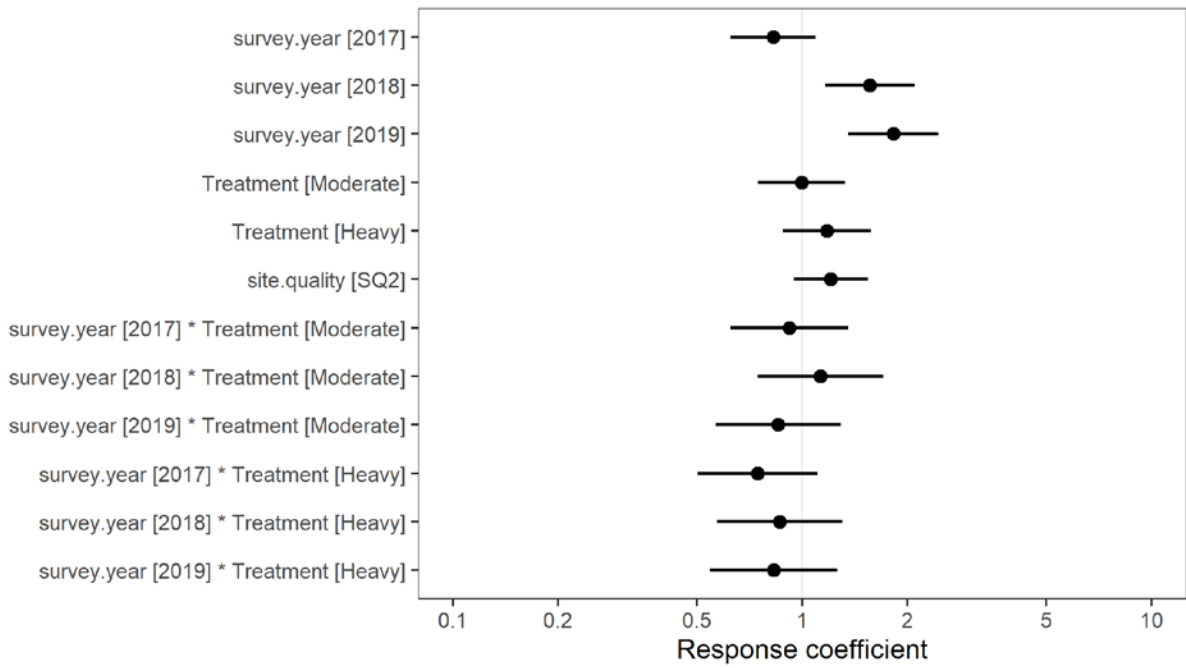


Figure 25 Wald confidence intervals for fixed effects: litter cover

Table 26 Model fitted values for average conditions: litter cover

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	83.4308	83.3537	85.5464
2017	SQ1	85.8702	85.8025	87.7201
2018	SQ1	80.5834	79.1102	78.4045
2019	SQ1	83.3584	82.0488	81.4191
2015	SQ2	88.7247	89.8061	88.8583
2017	SQ2	90.4737	91.4036	90.5887
2018	SQ2	90.1769	88.6288	89.9344
2019	SQ2	91.7216	90.3911	91.5137



## E.2.3 Surface fuel assessment

**Table 27 Model fitting summary: surface fuel hazard**

<b>Response</b>	Probability of being in each surface fuel hazard category
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	brm from the brms package
<b>Distribution used</b>	Continuation ratio with flexible category thresholds – see explanation for overall fuel hazard above
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Not applicable
<b>Confidence comments</b>	Moderate

**Model results summary 15: surface fuel hazard**

```

## Family: cratio
## Links: mu = logit; disc = identity
## Formula: risk.surface.or ~ Survey year * Treatment + Site quality + (1 | fsitep
lot) + (1 | fsite)
## Data: DAT.fire (Number of observations: 789)
## Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup samples = 4000
##
## Group-Level Effects:
## ~fsite (Number of levels: 22)
## Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 0.35 0.16 0.03 0.69 1.00 599 1076
##
## ~fsiteplot (Number of levels: 66)
## Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 0.53 0.11 0.31 0.75 1.00 956 1601
##
## Population-Level Effects:
## Estimate Est.Error l-95% CI u-95% CI Rhat
## Intercept[1] -2.67 0.30 -3.29 -2.07 1.00
## Intercept[2] -0.57 0.28 -1.10 -0.02 1.00
## Intercept[3] 1.17 0.29 0.61 1.74 1.00
## Intercept[4] 3.60 0.44 2.77 4.51 1.00
## Survey year2017 0.07 0.29 -0.50 0.63 1.00
## Survey year2018 0.66 0.29 0.11 1.23 1.00
## Survey year2019 0.51 0.29 -0.05 1.07 1.00
## TreatmentModerate -0.14 0.33 -0.77 0.52 1.00
## TreatmentHeavy 0.17 0.33 -0.46 0.84 1.00
## Site qualitySQ2 -0.06 0.24 -0.54 0.40 1.00
## Survey year2017:TreatmentModerate -0.11 0.41 -0.90 0.68 1.00
## Survey year2018:TreatmentModerate -0.09 0.40 -0.90 0.66 1.00
## Survey year2019:TreatmentModerate 0.22 0.41 -0.57 1.03 1.00
## Survey year2017:TreatmentHeavy -0.47 0.41 -1.28 0.33 1.00
## Survey year2018:TreatmentHeavy -0.20 0.41 -1.03 0.59 1.00
## Survey year2019:TreatmentHeavy -0.18 0.40 -0.96 0.58 1.00
## Bulk_ESS Tail_ESS
## Intercept[1] 2191 3068
## Intercept[2] 1980 2589
## Intercept[3] 2123 2858
## Intercept[4] 3287 3026
## Survey year2017 2290 2807
## Survey year2018 1889 2838
## Survey year2019 2212 3185
## TreatmentModerate 1963 2543
## TreatmentHeavy 2122 2443
## Site qualitySQ2 3094 2810
## Survey year2017:TreatmentModerate 2491 2868
## Survey year2018:TreatmentModerate 2202 2878
## Survey year2019:TreatmentModerate 2410 2552
## Survey year2017:TreatmentHeavy 2848 3132
## Survey year2018:TreatmentHeavy 2267 2737
## Survey year2019:TreatmentHeavy 2322 2948
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).

```

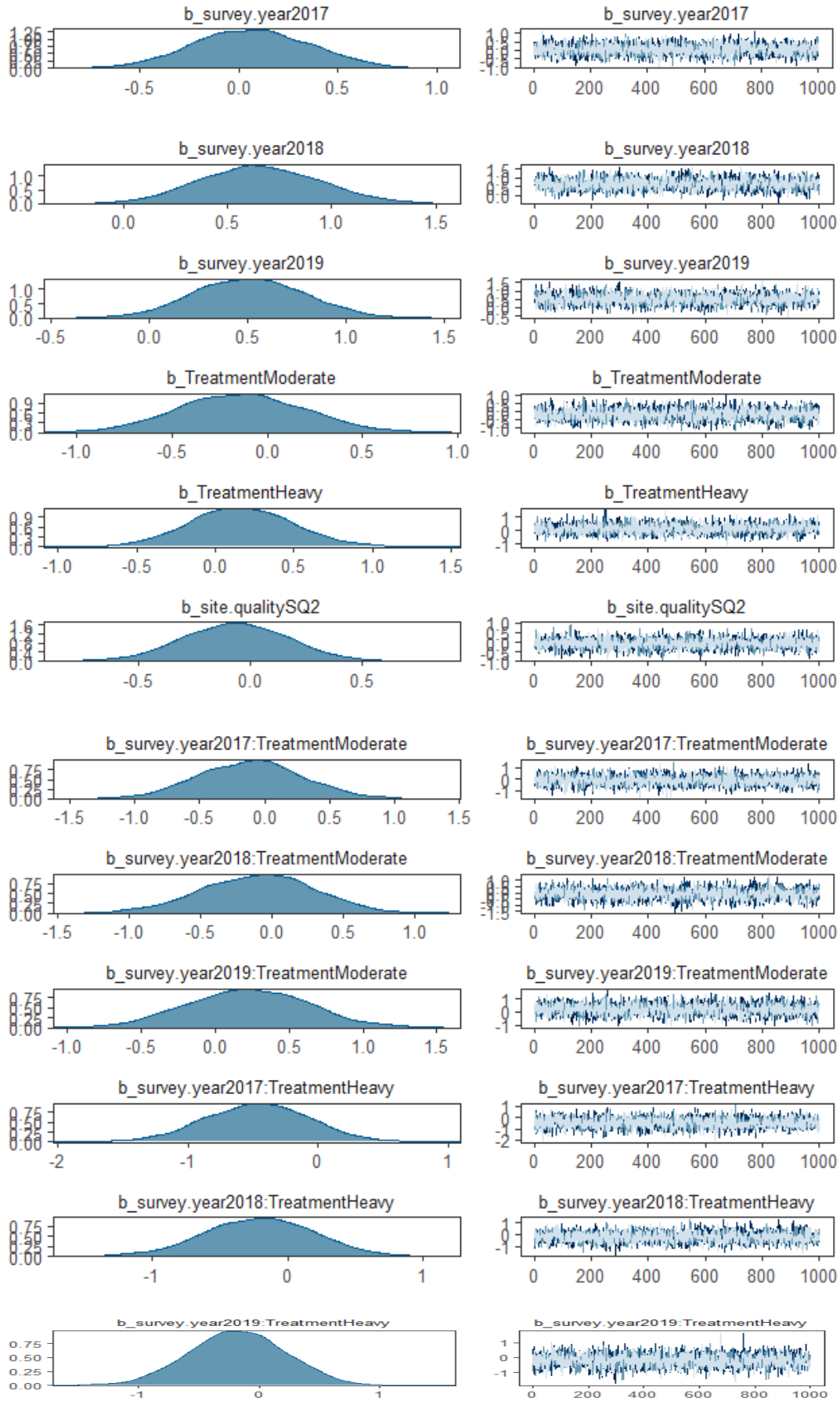


Figure 26 Posterior distributions (left) and chain mixing (right): surface fuel hazard

## E.3 Near surface fuel hazard

### E.3.1 Live near surface vegetation cover

**Table 28 Model fitting summary: live near surface vegetation cover**

<b>Response</b>	Percent live vegetation cover visually estimated for three 0.04 ha subplots in each 9 ha plot Percentage: positive continuous bounded variable
<b>Response transformation used</b>	$\text{Log}_e(x)$
<b>Other transformations compared</b>	None
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	Beta – failed Dunn Smyth tests
<b>Outliers removed</b>	One zero value recorded in 2015, likely an observer error.
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Marginal = 25.1% Conditional = 42.9%
<b>Confidence comments</b>	Moderate – High. Passed all Dunn Smyth tests

### Model results summary 16: live near surface vegetation cover

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: log(nearsurface.live) ~ Survey year * Treatment + Site quality +
## (1 | fsite) + (1 | fsiteplot)
## Data: DAT.vegcover1
##
## REML criterion at convergence: 2285
##
## Scaled residuals:
##   Min       1Q   Median       3Q      Max
## -3.7132 -0.5624  0.0045  0.6157  3.2115
##
## Random effects:
## Groups      Name                Variance Std.Dev.
## fsiteplot (Intercept) 0.06134  0.2477
```

```

## fsite (Intercept) 0.22814 0.4776
## Residual 0.92740 0.9630
## Number of obs: 794, groups: fsiteplot, 66; fsite, 22
##
## Fixed effects:
##
## Estimate Std. Error df t value
## (Intercept) -2.48937 0.19919 49.69669 -12.497
## Survey year2017 0.44989 0.16764 718.91630 2.684
## Survey year2018 -0.74758 0.16764 718.91630 -4.459
## Survey year2019 -1.01290 0.16764 718.91630 -6.042
## TreatmentModerate 0.16774 0.18352 257.93252 0.914
## TreatmentHeavy 0.01144 0.18358 257.78835 0.062
## Site qualitySQ2 -0.40441 0.22332 19.99104 -1.811
## Survey year2017:TreatmentModerate -0.46520 0.23708 718.91630 -1.962
## Survey year2018:TreatmentModerate -0.25429 0.23708 718.91630 -1.073
## Survey year2019:TreatmentModerate -0.26344 0.23708 718.91630 -1.111
## Survey year2017:TreatmentHeavy -0.14500 0.23666 719.10916 -0.613
## Survey year2018:TreatmentHeavy -0.16659 0.23666 719.10916 -0.704
## Survey year2019:TreatmentHeavy -0.26556 0.23712 719.25503 -1.120
##
## Pr(>|t|)
## (Intercept) < 2e-16 ***
## Survey year2017 0.00745 **
## Survey year2018 9.53e-06 ***
## Survey year2019 2.44e-09 ***
## TreatmentModerate 0.36157
## TreatmentHeavy 0.95038
## Site qualitySQ2 0.08522 .
## Survey year2017:TreatmentModerate 0.05012 .
## Survey year2018:TreatmentModerate 0.28382
## Survey year2019:TreatmentModerate 0.26685
## Survey year2017:TreatmentHeavy 0.54028
## Survey year2018:TreatmentHeavy 0.48172
## Survey year2019:TreatmentHeavy 0.26312
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

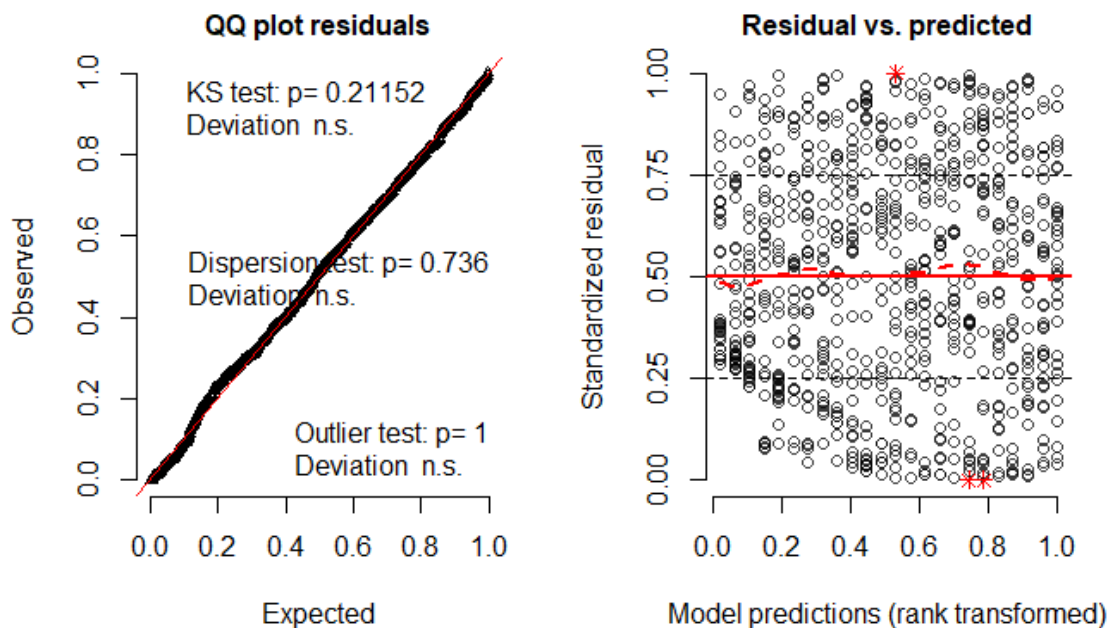
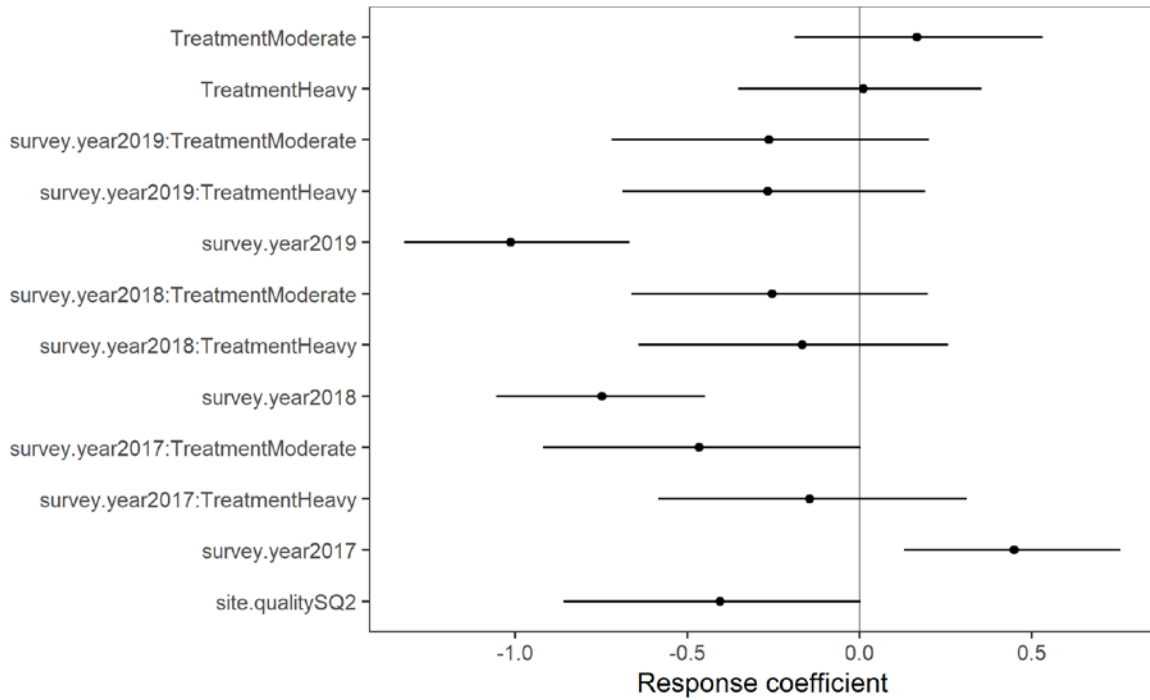


Figure 27 Dunn Smyth simulated residuals: live near surface vegetation cover





**Figure 28** Bootstrapped confidence intervals for fixed effects: live near surface vegetation cover

**Table 29** Bootstrapped model fitted values for average conditions: live near surface vegetation cover

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	8.6345	9.8558	8.4999
2017	SQ1	12.7172	9.7288	11.1641
2018	SQ1	3.9332	3.7513	3.3207
2019	SQ1	2.9972	2.6691	2.3984
2015	SQ2	5.5926	6.4783	5.9869
2017	SQ2	8.3257	6.8355	7.5032
2018	SQ2	2.7353	2.2973	2.1884
2019	SQ2	1.9620	1.8106	1.5690

### E.3.2 Dead near surface vegetation cover

**Table 30 Model fitting summary: dead near surface vegetation cover**

<b>Response</b>	Percent dead near surface visually estimated for three 0.04 ha subplots in each 9 ha plot Percentage: positive continuous bounded variable
<b>Response transformation used</b>	$\text{Log}_e(x + 0.001)$
<b>Other transformations compared</b>	$X + 0.001$ (a small constant to enable zero values to be modelled)
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	Beta
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year. Also trialled a model with a three-way interaction between thinning treatment, survey year and site quality, which had a higher AIC (that is, worse fit). The three-way model detected an additional significant effect of higher dead near surface cover on heavily thinned plots in SQ2 in 2018. However, the bootstrapped confidence interval included zero and therefore the two-way model was reported
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Function not available to calculate R <sup>2</sup> for beta distribution
<b>Confidence comments</b>	Moderate. Failed the Dunn Smyth uniformity test

### Model results summary 17: dead near surface vegetation cover

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula:
## log(nearsurface.dead + 1e-04) ~ Survey year * Treatment + Site quality +
## (1 | fsite) + (1 | fsiteplot)
## Data: DAT.vegcover
##
## REML criterion at convergence: 2393.1
##
## Scaled residuals:
##   Min       1Q   Median       3Q      Max
## -4.9234 -0.4903  0.0060  0.5382  3.1292
##
## Random effects:
```

```

## Groups      Name      Variance Std.Dev.
## fsiteplot (Intercept) 0.03827  0.1956
## fsite      (Intercept) 0.18653  0.4319
## Residual                1.08345  1.0409
## Number of obs: 795, groups: fsiteplot, 66; fsite, 22
##
## Fixed effects:
##
##              Estimate Std. Error      df t value
## (Intercept)   -4.50353    0.19250  61.52529 -23.395
## Survey year2017    1.18131    0.18120  719.83834   6.520
## Survey year2018    0.89815    0.18120  719.83834   4.957
## Survey year2019    0.84626    0.18120  719.83834   4.670
## TreatmentModerate -0.08854    0.19055  322.41217  -0.465
## TreatmentHeavy    0.26339    0.18993  319.20925   1.387
## Site qualitySQ2   -0.18097    0.20418  19.97766  -0.886
## Survey year2017:TreatmentModerate 0.06834    0.25625  719.83834   0.267
## Survey year2018:TreatmentModerate 0.07235    0.25625  719.83834   0.282
## Survey year2019:TreatmentModerate 0.09909    0.25625  719.83834   0.387
## Survey year2017:TreatmentHeavy -0.02459    0.25529  719.83834  -0.096
## Survey year2018:TreatmentHeavy -0.42554    0.25529  719.83834  -1.667
## Survey year2019:TreatmentHeavy -0.28747    0.25579  720.01159  -1.124
##
##              Pr(>|t|)
## (Intercept)   < 2e-16 ***
## Survey year2017 1.33e-10 ***
## Survey year2018 8.94e-07 ***
## Survey year2019 3.59e-06 ***
## TreatmentModerate    0.642
## TreatmentHeavy      0.166
## Site qualitySQ2     0.386
## Survey year2017:TreatmentModerate 0.790
## Survey year2018:TreatmentModerate 0.778
## Survey year2019:TreatmentModerate 0.699
## Survey year2017:TreatmentHeavy 0.923
## Survey year2018:TreatmentHeavy 0.096 .
## Survey year2019:TreatmentHeavy 0.261
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

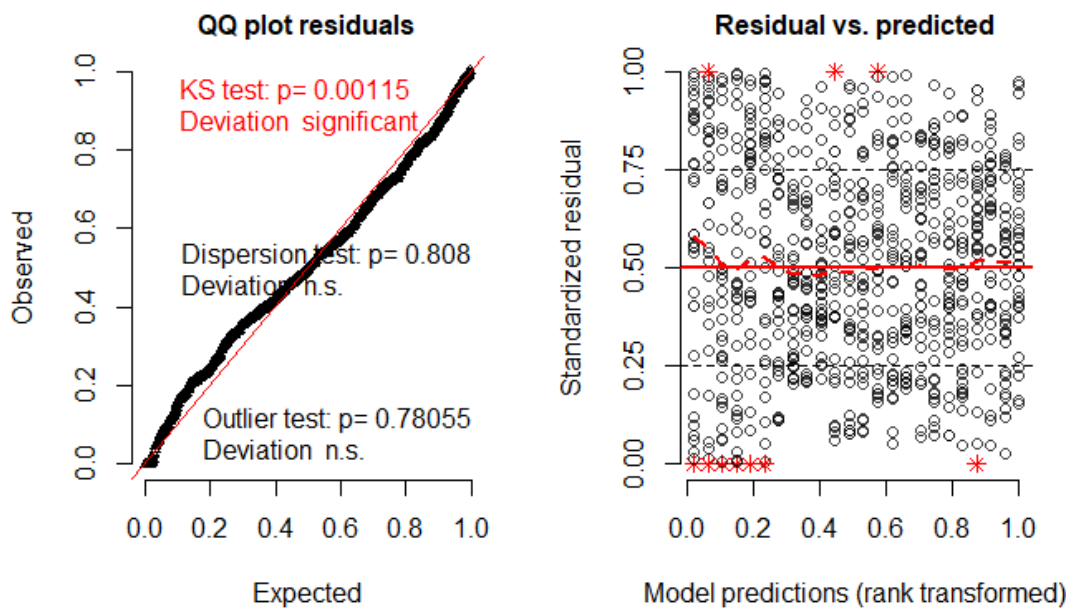
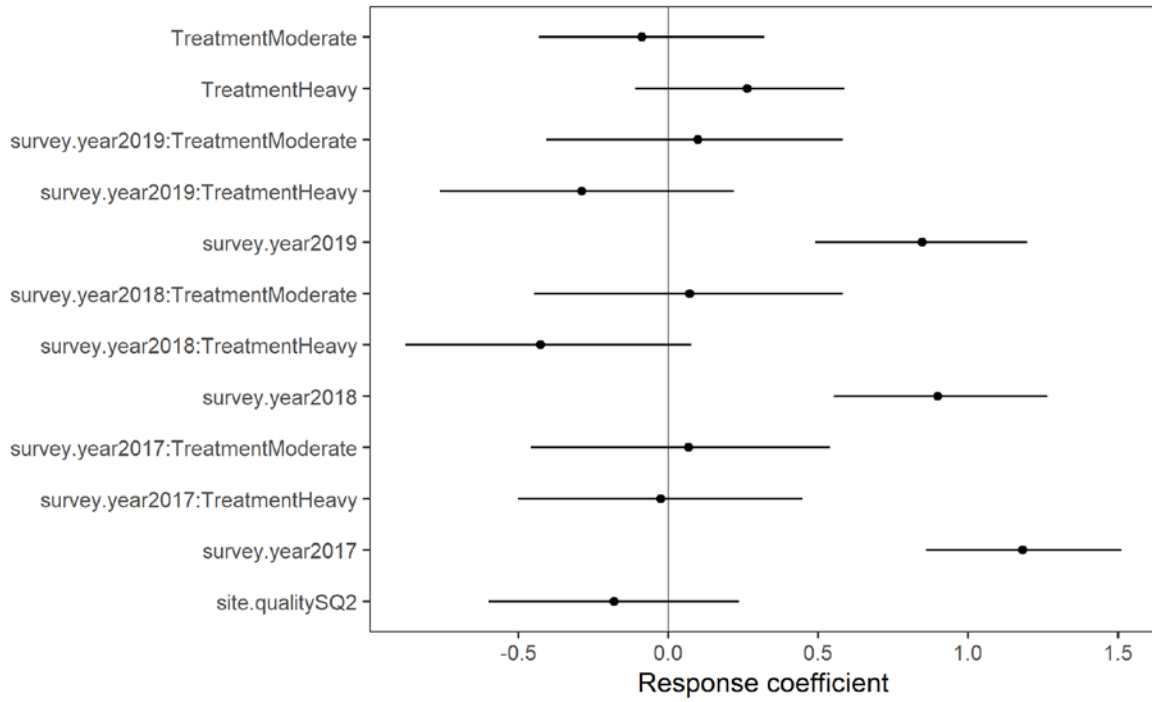


Figure 29 Dunn Smyth simulated residuals: dead near surface vegetation cover



**Figure 30** Bootstrapped confidence intervals for fixed effects: dead near surface vegetation cover

**Table 31** Bootstrapped model fitted values for average conditions: dead near surface vegetation cover

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	1.1014	0.9853	1.3681
2017	SQ1	3.6757	3.7828	4.3027
2018	SQ1	2.6153	2.9059	2.4583
2019	SQ1	2.7029	2.5790	2.3671
2015	SQ2	0.9144	0.8630	1.2340
2017	SQ2	3.0263	2.7434	3.7986
2018	SQ2	2.2868	2.0926	1.8039
2019	SQ2	2.2119	2.1365	2.1045

### E.3.3 Near surface fuel assessment

**Table 32 Model fitting summary: near surface fuel hazard**

<b>Response</b>	Probability of being in each near surface fuel hazard category
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	brm from the brms package
<b>Distribution used</b>	Continuation ratio with flexible category thresholds – see explanation for overall fuel hazard above
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	f <sub>site</sub> (a factor over sites) and f <sub>siteplot</sub> (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Not applicable
<b>Confidence comments</b>	Moderate



## Model results summary 18: near surface fuel hazard

```
## Family: cratio
## Links: mu = logit; disc = identity
## Formula: risk.nearsurface.or ~ Survey year * Treatment + Site quality + (1 | fs
iteplot) + (1 | fsite)
## Data: DAT.fire (Number of observations: 795)
## Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup samples = 4000
##
## Group-Level Effects:
## ~fsite (Number of levels: 22)
## Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 0.64 0.19 0.30 1.05 1.00 1005 859
##
## ~fsiteplot (Number of levels: 66)
## Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 0.56 0.14 0.30 0.84 1.01 1018 1831
##
## Population-Level Effects:
## Estimate Est.Error l-95% CI u-95% CI Rhat
## Intercept[1] -1.63 0.37 -2.36 -0.89 1.00
## Intercept[2] 1.75 0.38 1.00 2.55 1.00
## Intercept[3] 4.82 0.49 3.90 5.81 1.00
## Intercept[4] 3.18 0.82 1.62 4.88 1.00
## Survey year2017 1.56 0.35 0.87 2.25 1.00
## Survey year2018 1.43 0.36 0.72 2.16 1.00
## Survey year2019 1.33 0.35 0.63 2.01 1.00
## TreatmentModerate 0.12 0.39 -0.64 0.89 1.00
## TreatmentHeavy 0.64 0.40 -0.16 1.43 1.00
## Site qualitySQ2 -0.46 0.36 -1.17 0.26 1.00
## Survey year2017:TreatmentModerate -0.29 0.48 -1.24 0.63 1.00
## Survey year2018:TreatmentModerate 0.32 0.49 -0.67 1.25 1.00
## Survey year2019:TreatmentModerate -0.47 0.49 -1.45 0.50 1.00
## Survey year2017:TreatmentHeavy -0.40 0.48 -1.34 0.56 1.00
## Survey year2018:TreatmentHeavy -0.62 0.50 -1.59 0.38 1.00
## Survey year2019:TreatmentHeavy -0.99 0.49 -1.95 -0.04 1.00
## Bulk_ESS Tail_ESS
## Intercept[1] 1792 2282
## Intercept[2] 1702 2575
## Intercept[3] 2166 2581
## Intercept[4] 4193 3330
## Survey year2017 1919 2554
## Survey year2018 2002 2828
## Survey year2019 1870 2306
## TreatmentModerate 1749 2235
## TreatmentHeavy 1922 2174
## Site qualitySQ2 2357 2158
## Survey year2017:TreatmentModerate 2106 3004
## Survey year2018:TreatmentModerate 2148 2716
## Survey year2019:TreatmentModerate 2060 2402
## Survey year2017:TreatmentHeavy 2212 2970
## Survey year2018:TreatmentHeavy 2164 2639
## Survey year2019:TreatmentHeavy 2052 2463
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```

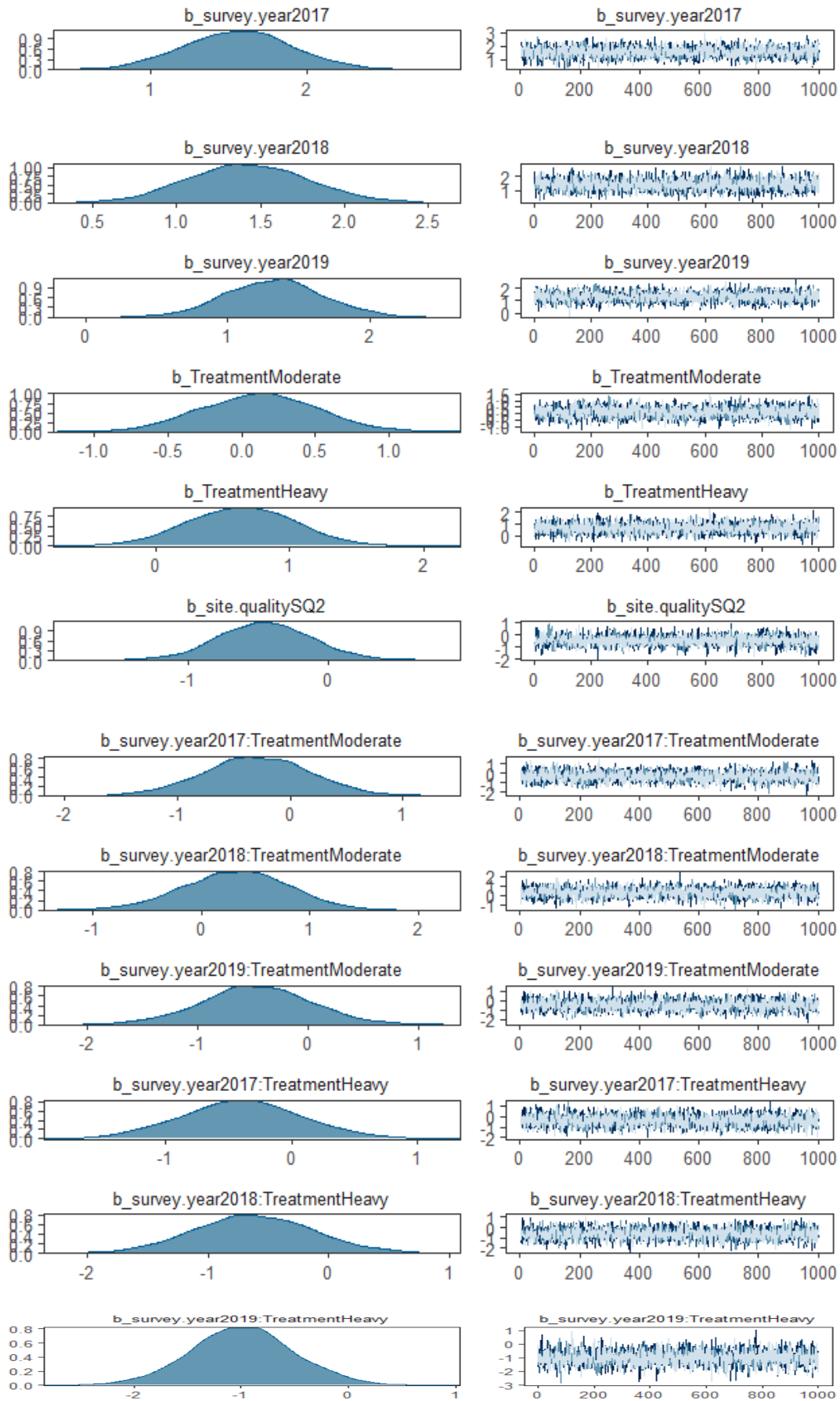


Figure 31 Posterior distributions (left) and chain mixing (right): near surface fuel hazard

## E.4 Combined surface and near surface fuel hazard

**Table 33 Model fitting summary: combined surface and near surface fuel hazard**

<b>Response</b>	Probability of being in each combined surface and near surface fuel hazard category
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	brm from the brms package
<b>Distribution used</b>	Continuation ratio with flexible category thresholds – see explanation for overall fuel hazard above
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	f <sub>site</sub> (a factor over sites), and f <sub>siteplot</sub> (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Not applicable
<b>Confidence comments</b>	Moderate

## Model results summary 19: combined surface and near surface fuel hazard

```
## Family: cratio
## Links: mu = logit; disc = identity
## Formula: risk.surfacecomb.or ~ Survey year * Treatment + Site quality + (1 | fs
iteplot) + (1 | fsite)
## Data: DAT.fire (Number of observations: 789)
## Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup samples = 4000
##
## Group-Level Effects:
## ~fsite (Number of levels: 22)
## Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 0.36 0.14 0.06 0.66 1.00 583 496
##
## ~fsiteplot (Number of levels: 66)
## Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 0.42 0.13 0.15 0.65 1.00 689 788
##
## Population-Level Effects:
## Estimate Est.Error l-95% CI u-95% CI Rhat
## Intercept[1] -2.81 0.30 -3.40 -2.23 1.00
## Intercept[2] -0.91 0.27 -1.43 -0.40 1.00
## Intercept[3] -1.03 0.27 -1.57 -0.51 1.00
## Intercept[4] 2.78 0.30 2.19 3.38 1.00
## Survey year2017 0.41 0.29 -0.14 0.99 1.00
## Survey year2018 0.86 0.29 0.30 1.42 1.00
## Survey year2019 0.84 0.29 0.27 1.43 1.00
## TreatmentModerate -0.15 0.30 -0.75 0.45 1.00
## TreatmentHeavy 0.21 0.31 -0.39 0.82 1.00
## Site qualitySQ2 -0.07 0.24 -0.53 0.39 1.00
## Survey year2017:TreatmentModerate -0.27 0.40 -1.04 0.51 1.00
## Survey year2018:TreatmentModerate 0.21 0.41 -0.57 1.02 1.00
## Survey year2019:TreatmentModerate 0.11 0.41 -0.70 0.90 1.00
## Survey year2017:TreatmentHeavy -0.39 0.41 -1.18 0.39 1.00
## Survey year2018:TreatmentHeavy -0.16 0.40 -0.94 0.62 1.00
## Survey year2019:TreatmentHeavy -0.30 0.41 -1.12 0.49 1.00
## Bulk_ESS Tail_ESS
## Intercept[1] 2099 2609
## Intercept[2] 1770 2544
## Intercept[3] 1745 2362
## Intercept[4] 1855 2240
## Survey year2017 1821 2710
## Survey year2018 1703 2447
## Survey year2019 1874 2478
## TreatmentModerate 1567 2492
## TreatmentHeavy 1456 1829
## Site qualitySQ2 2362 2357
## Survey year2017:TreatmentModerate 1886 2727
## Survey year2018:TreatmentModerate 2126 2744
## Survey year2019:TreatmentModerate 1964 2145
## Survey year2017:TreatmentHeavy 1851 2847
## Survey year2018:TreatmentHeavy 1887 2603
## Survey year2019:TreatmentHeavy 1988 2870
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```

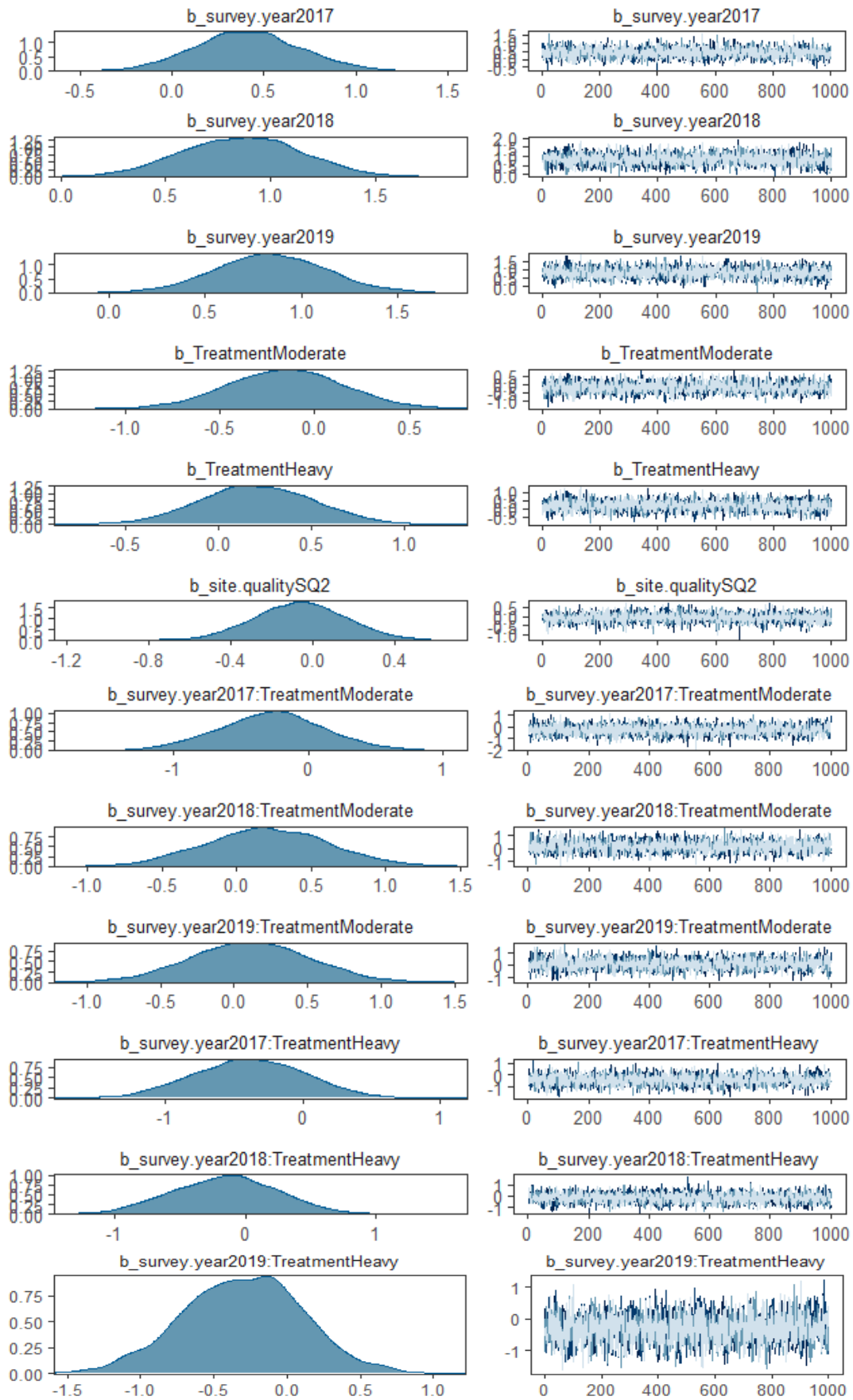


Figure 32 Posterior distributions (left) and chain mixing (right): combined surface and near surface fuel hazard



## E.5 Elevated fuel hazard

### E.5.1 Live elevated vegetation cover

**Table 34 Model fitting summary: live elevated vegetation cover**

<b>Response</b>	Percent live elevated cover visually estimated for three 0.04 ha subplots in each 9 ha plot Percentage: positive continuous bounded variable
<b>Response transformation used</b>	$X + 0.001$
<b>Other transformations compared</b>	$\text{Log}_e(x)$ (with gaussian distribution)
<b>R package and function</b>	glmmTMB function from the glmmTMB package
<b>Distribution used</b>	Beta
<b>Other distributions compared</b>	Gaussian (with $\text{log}_e(x)$ transformation)
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year Also trialled a model with three-way interaction between thinning treatment, survey year and site quality, which was less parsimonious
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	No function is available for calculating R <sup>2</sup> for beta models
<b>Confidence comments</b>	Moderate to high. Passed all Dunn Smyth tests

## Model results summary 20: live elevated vegetation cover

```
summary(model_el_beta)

## Family: beta ( logit )
## Formula:
## (elevated.live + 1e-04) ~ Survey year * Treatment + Site quality +
## (1 | fsite) + (1 | fsiteplot)
## Data: DAT.vegcover
##      AIC      BIC    logLik deviance df.resid
## -4330.7 -4255.9   2181.4  -4362.7     779
##
## Random effects:
##
## Conditional model:
## Groups      Name      Variance Std.Dev.
## fsite      (Intercept) 0.05378  0.2319
## fsiteplot  (Intercept) 0.08282  0.2878
## Number of obs: 795, groups: fsite, 22; fsiteplot, 66
## Overdispersion parameter for beta family (): 38.9
## Conditional model:
##
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   -3.01745    0.13427 -22.473 < 2e-16 ***
## Survey year2017 -0.10834    0.12478  -0.868 0.385228
## Survey year2018  0.09478    0.11989   0.791 0.429195
## Survey year2019  0.72874    0.11156   6.532 6.48e-11 ***
## TreatmentModerate -0.08192    0.15086  -0.543 0.587125
## TreatmentHeavy   -0.29190    0.15447  -1.890 0.058805 .
## Site qualitySQ2  -0.45073    0.13231  -3.407 0.000658 ***
## Survey year2017:TreatmentModerate -0.97090    0.19052  -5.096 3.47e-07 ***
## Survey year2018:TreatmentModerate -0.86713    0.18326  -4.732 2.23e-06 ***
## Survey year2019:TreatmentModerate  0.07530    0.15806   0.476 0.633786
## Survey year2017:TreatmentHeavy  -0.94008    0.19405  -4.844 1.27e-06 ***
## Survey year2018:TreatmentHeavy  -0.98179    0.19013  -5.164 2.42e-07 ***
## Survey year2019:TreatmentHeavy  -0.03564    0.16445  -0.217 0.828421
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

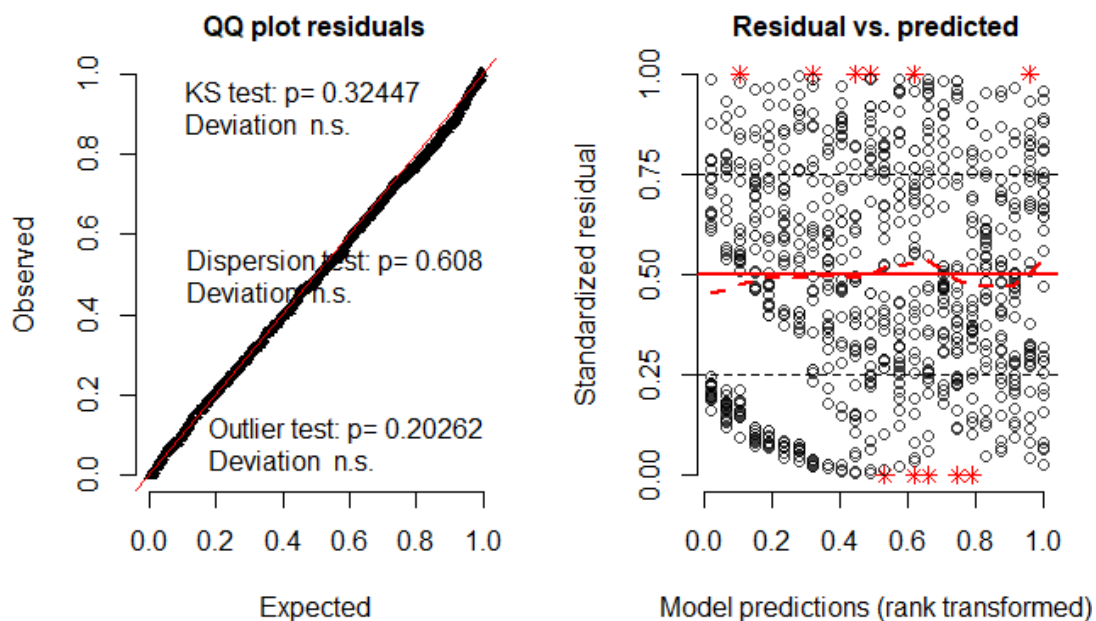


Figure 33 Dunn Smyth simulated residuals: live elevated vegetation

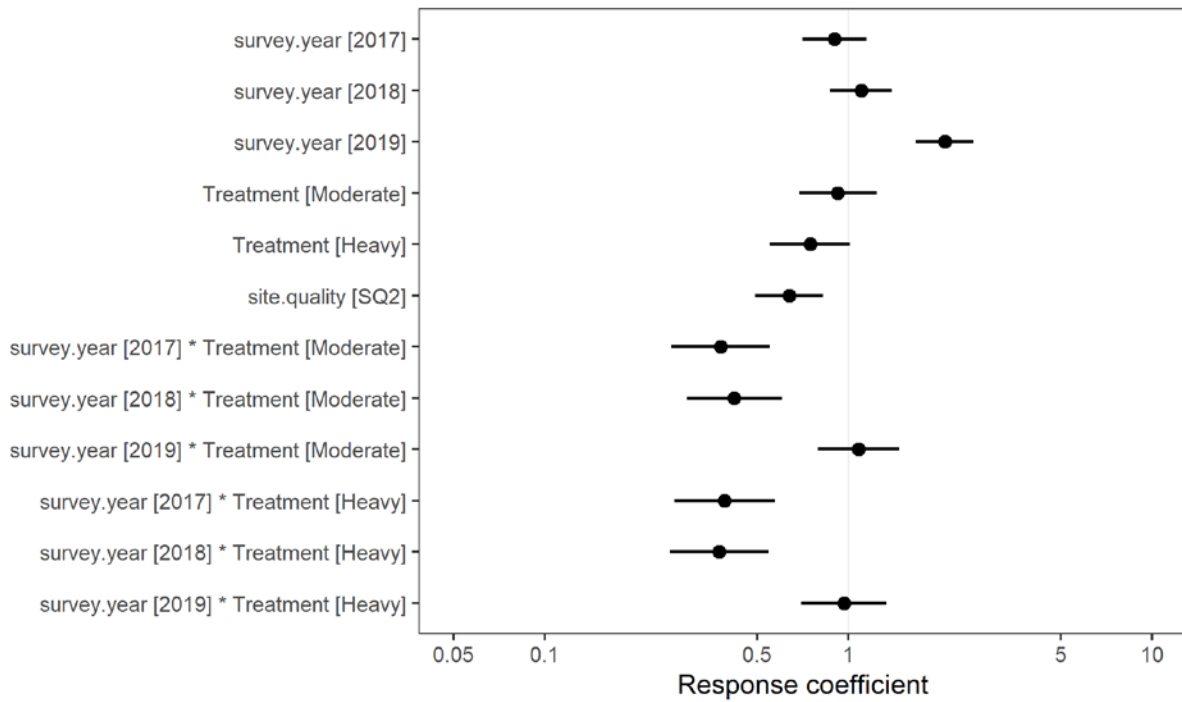


Figure 34 Wald confidence intervals for fixed effects: live elevated vegetation cover

Table 35 Model fitted values for average conditions: live elevated vegetation cover

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	4.6830	4.3299	3.5371
2017	SQ1	4.2215	1.5084	1.2625
2018	SQ1	5.1256	2.0427	1.4821
2019	SQ1	9.2501	9.1946	6.8409
2015	SQ2	3.0320	2.7995	2.2795
2017	SQ2	2.7282	0.9628	0.8046
2018	SQ2	3.3244	1.3077	0.9459
2019	SQ2	6.0953	6.0575	4.4664

## E.5.2 Dead elevated vegetation cover

**Table 36 Model fitting summary: dead elevated vegetation cover**

<b>Response</b>	Percent dead elevated cover visually estimated for three 0.04 ha subplots in each 9 ha plot Percentage: positive continuous bounded variable
<b>Response transformation used</b>	X + 0.001
<b>Other transformations compared</b>	Log <sub>e</sub> (x) (with gaussian distribution)
<b>R package and function</b>	glmmTMB function from the glmmTMB package
<b>Distribution used</b>	Beta
<b>Other distributions compared</b>	Gaussian (with log <sub>e</sub> (x) transformation)
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year Also trialled a model with three-way interaction between thinning treatment, survey year and site quality, which was less parsimonious
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	No function is available for calculating R <sup>2</sup> for beta models
<b>Confidence comments</b>	Low to moderate. Failed Dunn Smyth uniformity and dispersion tests

### Model results summary 21: dead elevated vegetation cover

```
## Family: beta ( logit )
## Formula:
## (elevated.dead + 1e-04) ~ Survey year * Treatment + Site quality +
## (1 | fsite) + (1 | fsiteplot)
## Data: DAT.vegcover2
##
##      AIC      BIC   logLik deviance df.resid
## -7078.2 -7003.4  3555.1  -7110.2      778
##
## Random effects:
##
## Conditional model:
## Groups   Name      Variance Std.Dev.
## fsite    (Intercept) 0.03908  0.1977
## fsiteplot (Intercept) 0.03843  0.1960
## Number of obs: 794, groups: fsite, 22; fsiteplot, 66
##
## Overdispersion parameter for beta family (): 278
```

```
##
## Conditional model:
##
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) -5.78347 0.13199 -43.82 < 2e-16 ***
## Survey year2017 0.71276 0.13025 5.47 4.44e-08 ***
## Survey year2018 0.98834 0.12647 7.82 5.50e-15 ***
## Survey year2019 1.47464 0.12013 12.28 < 2e-16 ***
## TreatmentModerate -0.21372 0.15801 -1.35 0.176189
## TreatmentHeavy -0.27235 0.15806 -1.72 0.084875 .
## Site qualitySQ2 -0.08722 0.10859 -0.80 0.421837
## Survey year2017:TreatmentModerate -0.66048 0.19922 -3.32 0.000915 ***
## Survey year2018:TreatmentModerate -0.41732 0.18937 -2.20 0.027541 *
## Survey year2019:TreatmentModerate 0.12377 0.17123 0.72 0.469795
## Survey year2017:TreatmentHeavy -0.50332 0.19609 -2.57 0.010264 *
## Survey year2018:TreatmentHeavy -0.60633 0.19231 -3.15 0.001617 **
## Survey year2019:TreatmentHeavy 0.07904 0.17233 0.46 0.646461
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

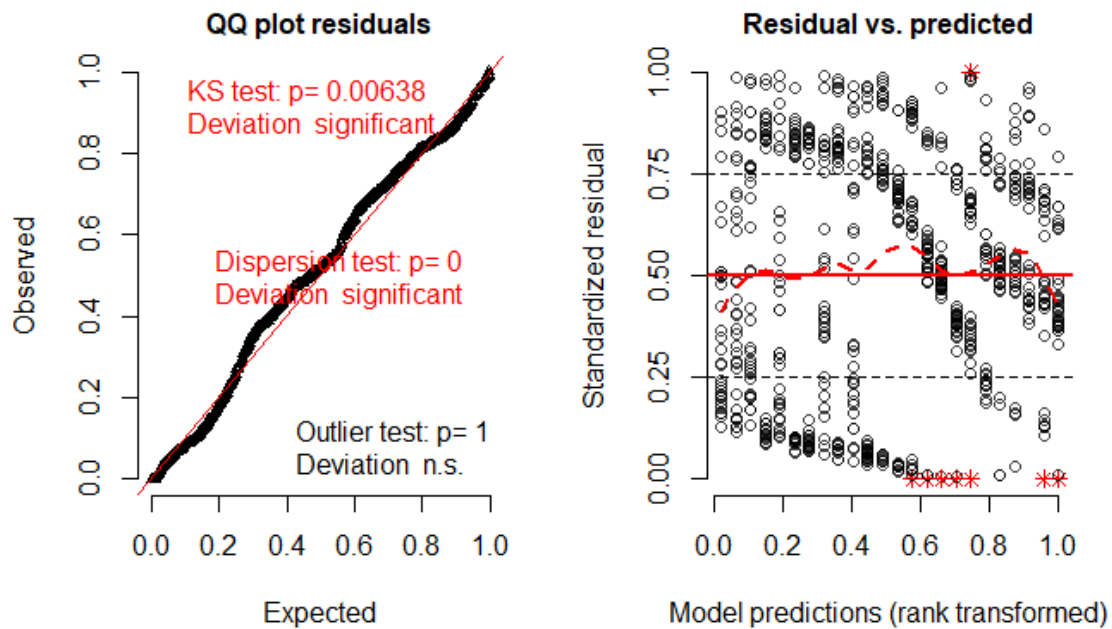
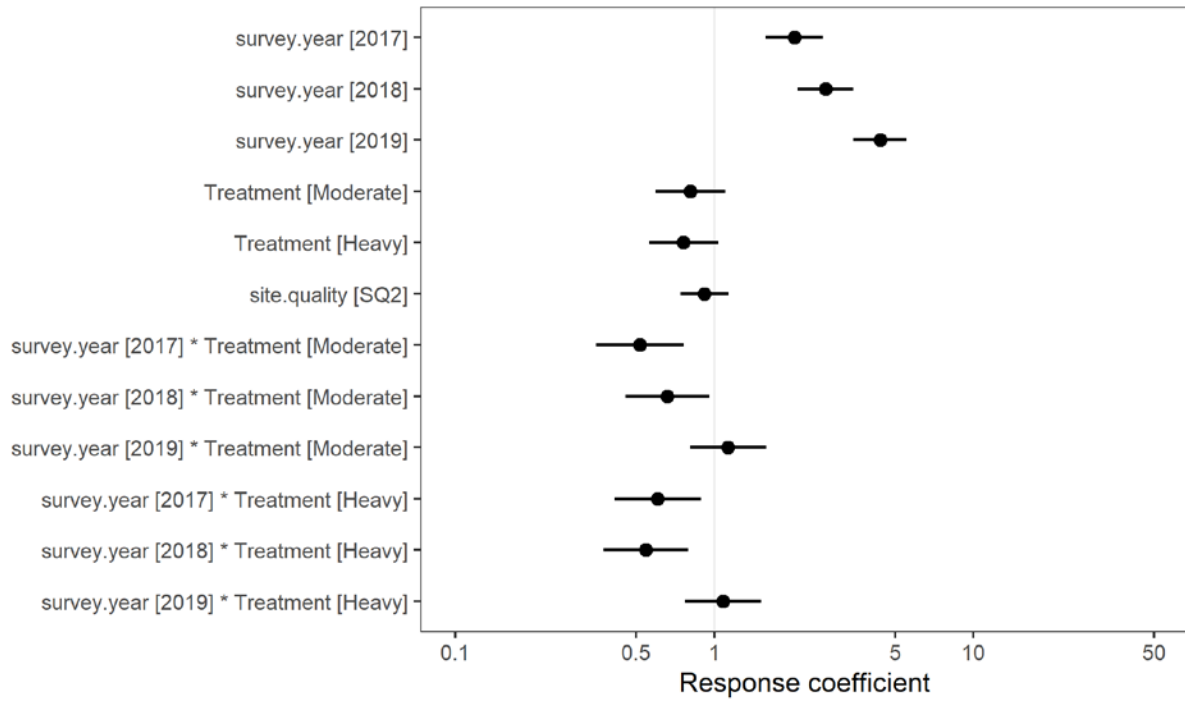


Figure 35 Dunn Smyth simulated residuals: dead elevated vegetation





**Figure 36** Wald confidence intervals for fixed effects: dead elevated vegetation cover

**Table 37** Model fitted values for average conditions: dead elevated vegetation cover

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	0.2973	0.2362	0.2221
2017	SQ1	0.6145	0.2493	0.2765
2018	SQ1	0.8122	0.4276	0.3313
2019	SQ1	1.2841	1.2100	1.0901
2015	SQ2	0.2691	0.2137	0.2009
2017	SQ2	0.5575	0.2255	0.2502
2018	SQ2	0.7373	0.3875	0.3000
2019	SQ2	1.1667	1.0992	0.9900

### E.5.3 Elevated fuel hazard assessment

**Table 38 Model fitting summary: elevated fuel hazard**

<b>Response</b>	Probability of being in each elevated fuel hazard category
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	brm from the brms package
<b>Distribution used</b>	Continuation ratio with flexible category thresholds – see explanation for overall fuel hazard above
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Not applicable
<b>Confidence comments</b>	Moderate

## Model results summary 22: elevated fuel hazard

```
## Family: cratio
## Links: mu = logit; disc = identity
## Formula: risk.elev.or ~ Survey year * Treatment + Site quality + (1 | fsiteplot
) + (1 | fsite)
## Data: DAT.fire (Number of observations: 795)
## Samples: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup samples = 4000
##
## Group-Level Effects:
## ~fsite (Number of levels: 22)
## Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 0.34 0.20 0.02 0.76 1.00 633 1244
##
## ~fsiteplot (Number of levels: 66)
## Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 0.62 0.17 0.25 0.96 1.00 1079 1481
##
## Population-Level Effects:
## Estimate Est.Error l-95% CI u-95% CI Rhat
## Intercept[1] 5.34 1.37 3.25 8.65 1.01
## Intercept[2] 10.63 1.58 8.03 14.27 1.01
## Survey year2017 4.37 1.39 2.28 7.65 1.01
## Survey year2018 4.67 1.38 2.53 7.92 1.01
## Survey year2019 4.66 1.39 2.56 8.01 1.01
## TreatmentModerate 2.29 1.46 0.01 5.68 1.01
## TreatmentHeavy 1.15 1.62 -1.67 4.69 1.01
## Site qualitySQ2 0.48 0.28 -0.07 1.04 1.00
## Survey year2017:TreatmentModerate -2.14 1.51 -5.61 0.30 1.01
## Survey year2018:TreatmentModerate -1.82 1.49 -5.22 0.57 1.01
## Survey year2019:TreatmentModerate -2.72 1.50 -6.19 -0.24 1.01
## Survey year2017:TreatmentHeavy -0.41 1.65 -3.88 2.51 1.01
## Survey year2018:TreatmentHeavy -0.50 1.64 -4.10 2.46 1.01
## Survey year2019:TreatmentHeavy -1.21 1.65 -4.67 1.74 1.01
## Bulk_ESS Tail_ESS
## Intercept[1] 497 651
## Intercept[2] 617 705
## Survey year2017 492 675
## Survey year2018 490 660
## Survey year2019 502 599
## TreatmentModerate 517 587
## TreatmentHeavy 521 706
## Site qualitySQ2 3917 2545
## Survey year2017:TreatmentModerate 520 634
## Survey year2018:TreatmentModerate 513 675
## Survey year2019:TreatmentModerate 537 650
## Survey year2017:TreatmentHeavy 537 665
## Survey year2018:TreatmentHeavy 526 668
## Survey year2019:TreatmentHeavy 526 690
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```

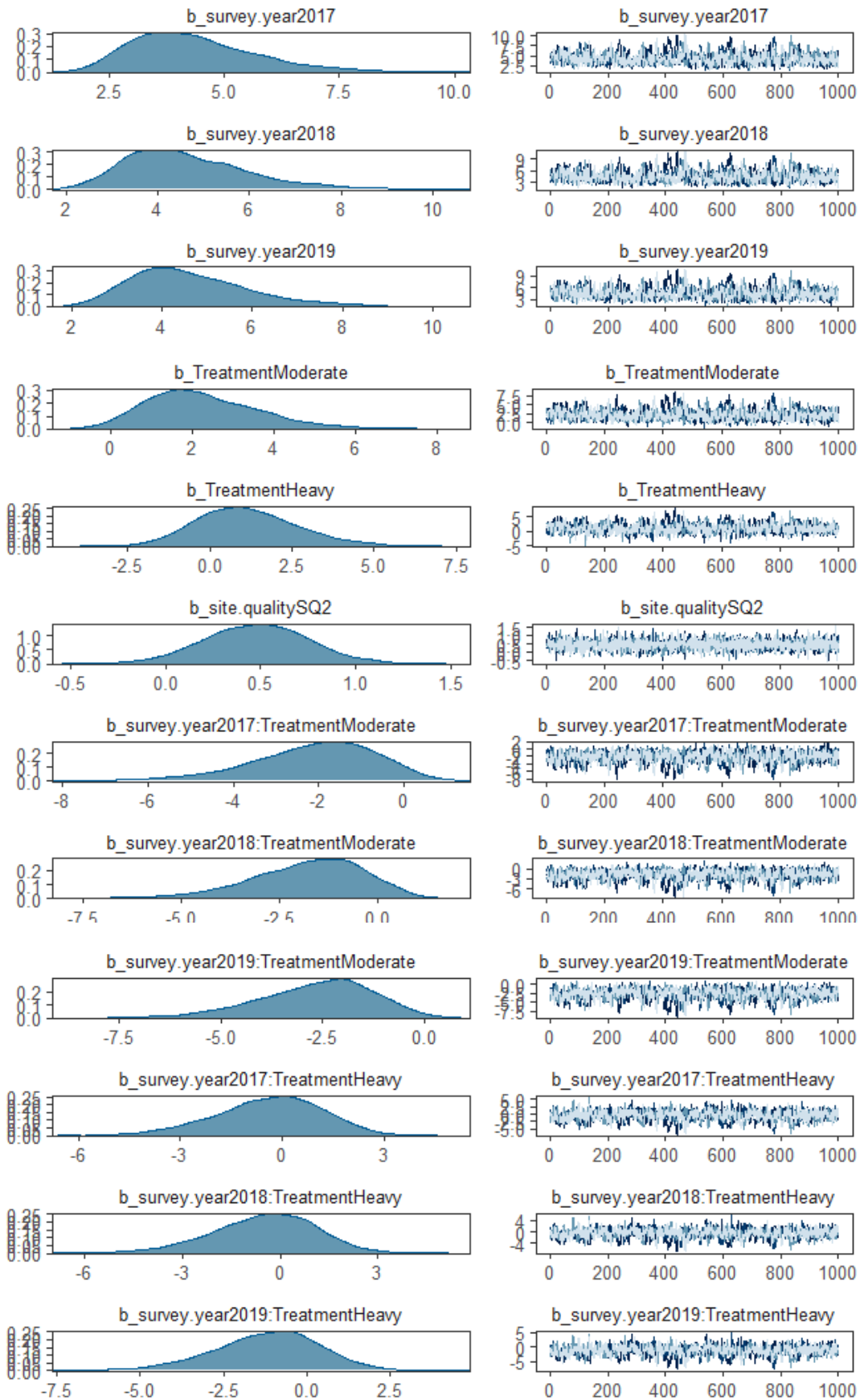


Figure 37 Posterior distributions (left) and chain mixing (right): elevated fuel hazard

# Appendix F: Model summaries – Floristic composition

## F.1 Plant species richness

### F.1.1 Native plant species richness

**Table 39 Model fitting summary: native plant species richness**

<b>Response</b>	Number of native plant species recorded in each 0.04 each subplot
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	Negative binomial Poisson
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year Also trialled a model with three-way interaction between thinning treatment, survey year and site quality, which was less parsimonious
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Marginal = 5.9% Conditional = 42.0%
<b>Confidence comments</b>	High. Passed all Dunn Smyth tests

### Model results summary 23: native plant species richness

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: nat.richness ~ year * Treatment + Site quality + (1 | fsiteplot) +
## (1 | fsite)
## Data: DAT.native.subplot
##
## REML criterion at convergence: 4512.6
##
```

```
## Scaled residuals:
##   Min      1Q  Median      3Q      Max
## -3.2431 -0.6820 -0.0253  0.6847  3.0620
##
## Random effects:
##   Groups      Name      Variance Std.Dev.
## fsiteplot (Intercept)  2.559    1.600
## fsite      (Intercept)  6.956    2.637
## Residual                15.277    3.909
## Number of obs: 795, groups: fsiteplot, 66; fsite, 22
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    16.0324    1.0189  37.9885   15.735 <2e-16 ***
## year2017         1.6970    0.6804  720.4346    2.494  0.0129 *
## year2018        -0.7273    0.6804  720.4346   -1.069  0.2855
## year2019        -0.7576    0.6804  720.4346   -1.113  0.2659
## TreatmentModerate  0.2879    0.8340  159.9892    0.345  0.7304
## TreatmentHeavy   -0.6293    0.8321  158.6163   -0.756  0.4506
## Site qualitySQ2  -0.3678    1.2234   19.9945   -0.301  0.7668
## year2017:TreatmentModerate  0.9091    0.9622  720.4346    0.945  0.3451
## year2018:TreatmentModerate  0.4848    0.9622  720.4346    0.504  0.6145
## year2019:TreatmentModerate  1.4697    0.9622  720.4346    1.527  0.1271
## year2017:TreatmentHeavy  1.8254    0.9586  720.4346    1.904  0.0573 .
## year2018:TreatmentHeavy  1.9362    0.9586  720.4346    2.020  0.0438 *
## year2019:TreatmentHeavy  1.3415    0.9605  720.5062    1.397  0.1630
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##
## Correlation matrix not shown by default, as p = 13 > 12.
## Use print(x, correlation=TRUE) or
##   vcov(x)           if you need it
```

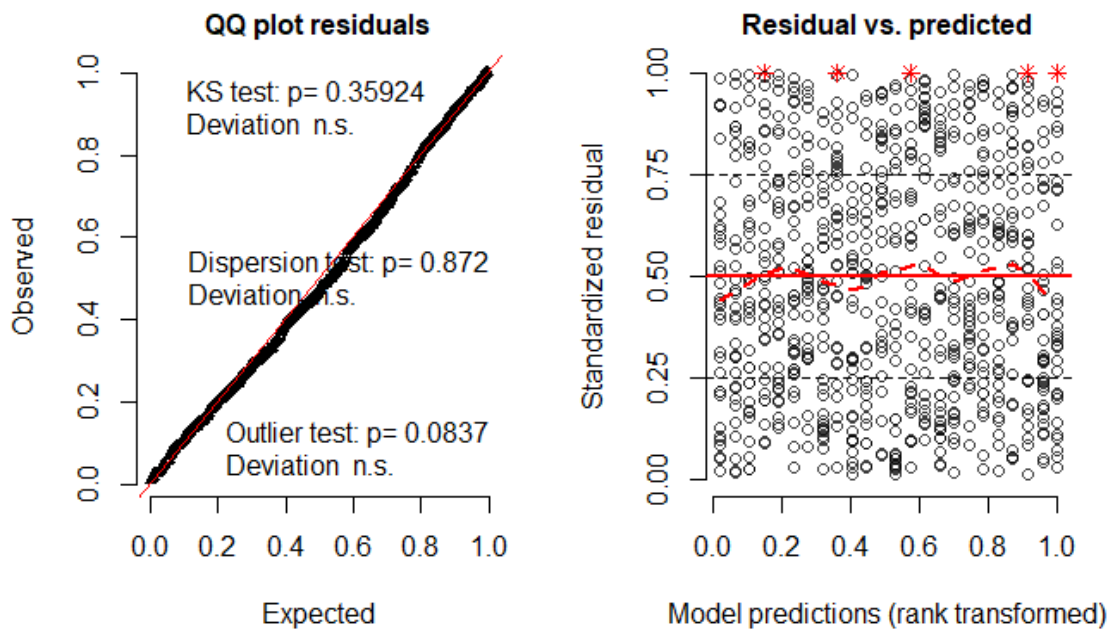
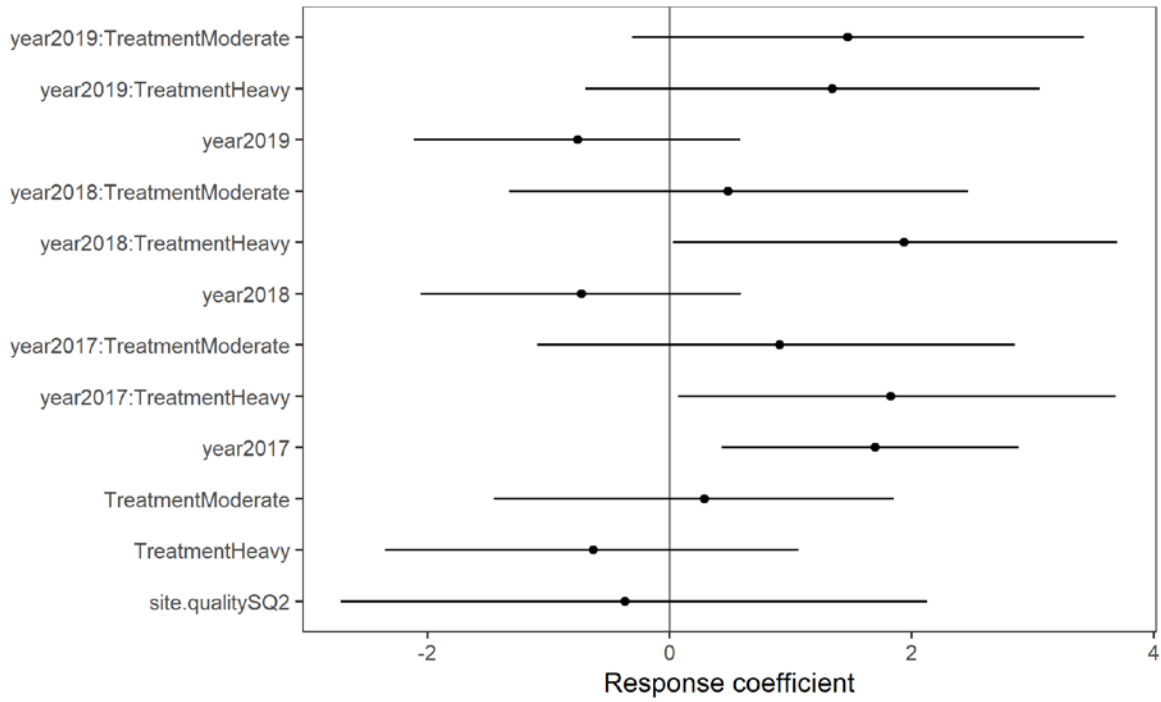


Figure 38 Dunn Smyth simulated residuals: native plant species richness





**Figure 39** Bootstrapped confidence intervals for fixed effects: native plant species richness

**Table 40** Bootstrapped model fitted values for average conditions: native plant species richness

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	16.1329	16.3468	15.5287
2017	SQ1	17.5436	18.8440	18.8474
2018	SQ1	15.1662	15.8765	16.7255
2019	SQ1	14.9979	16.7932	16.2540
2015	SQ2	15.4443	16.0166	15.0460
2017	SQ2	17.2464	18.8272	18.4186
2018	SQ2	15.3681	15.6262	16.4142
2019	SQ2	14.9887	16.8863	15.4330

## F.1.2 Exotic plant species richness

**Table 41 Model fitting summary: exotic plant species richness**

<b>Response</b>	Number of exotic plant species recorded in each 0.04 each subplot
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	Five values with exotic richness above 18 species per subplot
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year Also trialled a model with three-way interaction between thinning treatment, survey year and site quality, which was less parsimonious
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Marginal = 5.5% Conditional = 45.5%
<b>Confidence comments</b>	High. Passed all Dunn Smyth tests

### Model results summary 24: exotic plant species richness

```
DAT.exot.subplot1 <- DAT.exot.subplot %>%
  filter(exot.richness <18) # excludes 5 values

model.exot.rich <- lmer(exot.richness ~ year * Treatment + Site quality + (1|fsite
plot) + (1|fsite), data = DAT.exot.subplot1)

summary(model.exot.rich)

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: exot.richness ~ year * Treatment + Site quality + (1 | fsiteplot) +
## (1 | fsite)
## Data: DAT.exot.subplot1
##
## REML criterion at convergence: 3720.8
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.8099 -0.6751 -0.0513  0.5897  3.3235
##
```

```
## Random effects:
## Groups      Name          Variance Std.Dev.
## fsiteplot (Intercept) 1.434    1.197
## fsite      (Intercept) 3.770    1.942
## Residual                7.094    2.663
## Number of obs: 754, groups: fsiteplot, 66; fsite, 22
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)      5.8585   0.7487  37.6084   7.825 2.08e-09 ***
## year2017          0.7769   0.4840 682.9341   1.605 0.10894
## year2018         -0.7572   0.4962 687.5435  -1.526 0.12743
## year2019          0.2896   0.4879 687.2757   0.594 0.55304
## TreatmentModerate -0.5604   0.6082 152.3110  -0.921 0.35828
## TreatmentHeavy    -0.2922   0.5987 144.9401  -0.488 0.62627
## Site qualitySQ2    0.3251   0.9004  19.9581   0.361 0.72186
## year2017:TreatmentModerate 1.1547   0.6804 682.4435   1.697 0.09013 .
## year2018:TreatmentModerate 1.4788   0.6902 685.0586   2.143 0.03250 *
## year2019:TreatmentModerate 1.5102   0.6843 684.9251   2.207 0.02764 *
## year2017:TreatmentHeavy  1.7733   0.6733 681.5880   2.634 0.00864 **
## year2018:TreatmentHeavy  1.7537   0.6792 683.7054   2.582 0.01003 *
## year2019:TreatmentHeavy  0.7688   0.6746 683.3419   1.140 0.25480
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

##
## Correlation matrix not shown by default, as p = 13 > 12.
## Use print(x, correlation=TRUE) or
## vcov(x) if you need it
```

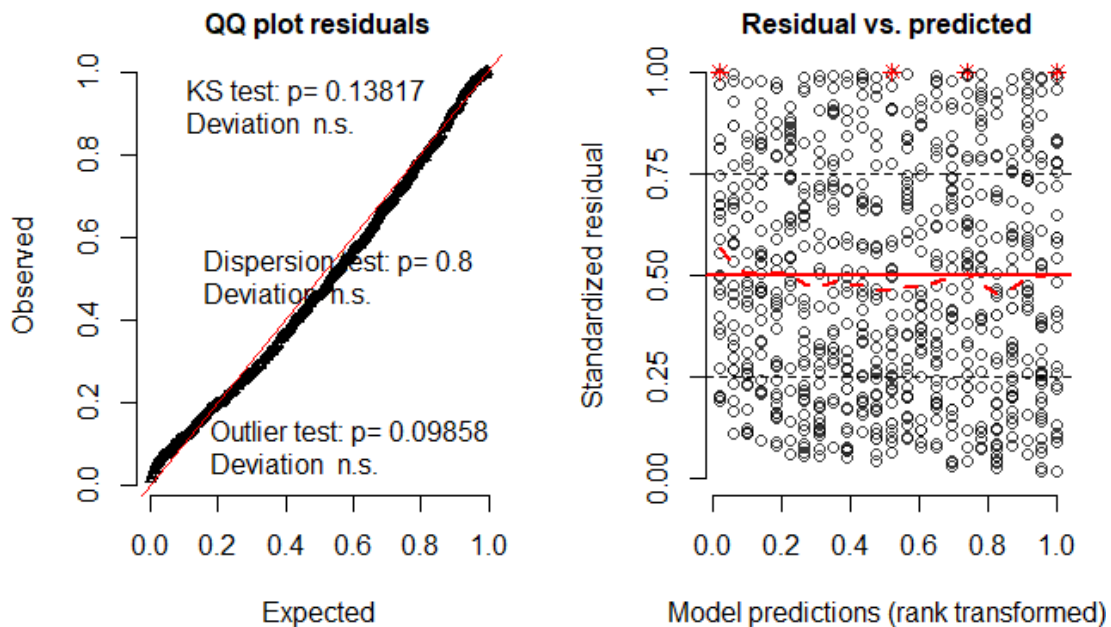
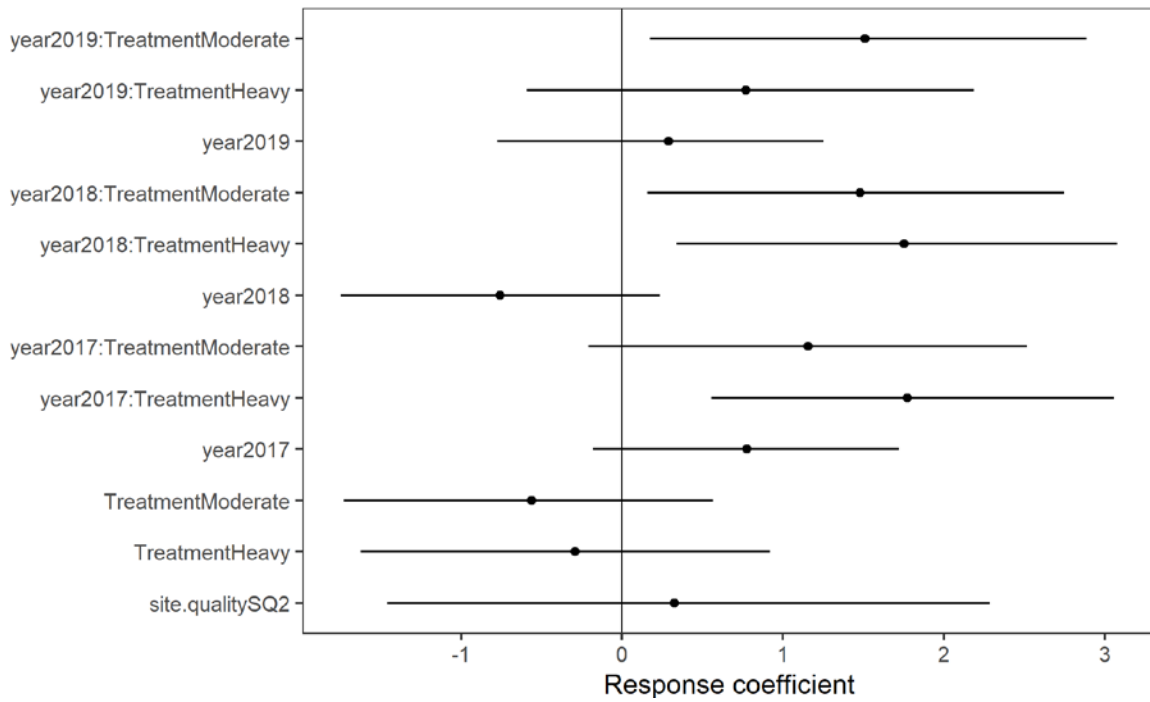


Figure 40 Dunn Smyth simulated residuals: exotic plant species richness



**Figure 41** Bootstrapped confidence intervals for fixed effects: exotic plant species richness

**Table 42** Bootstrapped model fitted values for average conditions: exotic plant species richness

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	5.8783	5.1355	5.6806
2017	SQ1	6.3297	7.2717	8.0660
2018	SQ1	5.0661	5.8675	6.4717
2019	SQ1	6.2661	7.0983	6.2892
2015	SQ2	6.3199	5.7048	5.7941
2017	SQ2	6.7575	7.3935	8.3589
2018	SQ2	5.5951	6.2714	6.9540
2019	SQ2	6.4315	7.4781	6.9973

## F.2 Plant cover

### F.2.1 Native plant cover

**Table 43 Model fitting summary: native plant cover**

<b>Response</b>	Native plant vegetation cover, visually estimated in each 0.04 ha subplot in 2019 only
<b>Response transformation used</b>	Rounding % cover to nearest whole value
<b>Response transformations compared</b>	$\text{Log}_e(x+1)$ transformation attempted on gaussian model ( $X+0.1$ )/100 transformation attempted on beta model
<b>R package and function</b>	glmer function from the lme4 package
<b>Distribution used</b>	Binomial with percentage values rounded up to the nearest 1%
<b>Other distributions compared</b>	Gaussian Beta
<b>Outliers removed</b>	Five values >15% were removed
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and site quality
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Marginal = 2.9% Conditional = 9.5%
<b>Confidence comments</b>	Low. Failed the uniformity test, some under-dispersion in the data, meaning more data than expected fall in the tails of the data

**Model results summary 25: native plant cover**

```

za <- cbind(success = ceiling(DAT.cover.b$nat.live), failure = 100 - ceiling(DAT.cover.b$nat.live))

model.native.binom.b <- glmer(za ~ Site quality * Treatment + (1|fsite) + (1|fsiteplot),
                             data = DAT.cover.b, family = "binomial", glmerControl(
optimizer = "bobyqa"))

summary(model.native.binom.b)

## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula: za ~ Site quality * Treatment + (1 | fsite) + (1 | fsiteplot)
## Data: DAT.cover.b
## Control: glmerControl(optimizer = "bobyqa")
##
##      AIC      BIC   logLik deviance df.resid
##  687.1    713.1  -335.5   671.1     184
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.6624 -0.4601 -0.2319  0.0470  3.9940
##
## Random effects:
## Groups      Name          Variance Std.Dev.
## fsiteplot (Intercept) 0.20007  0.4473
## fsite      (Intercept) 0.03882  0.1970
## Number of obs: 192, groups: fsiteplot, 65; fsite, 22
##
## Fixed effects:
##
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -3.5129    0.1825 -19.251  <2e-16 ***
## Site qualitySQ2 -0.6324    0.2715  -2.329  0.0198 *
## TreatmentModerate -0.0693    0.2514  -0.276  0.7829
## TreatmentHeavy  -0.5369    0.2545  -2.109  0.0349 *
## Site qualitySQ2:TreatmentModerate -0.1315    0.3761  -0.350  0.7266
## Site qualitySQ2:TreatmentHeavy  0.3866    0.3766   1.026  0.3047
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```



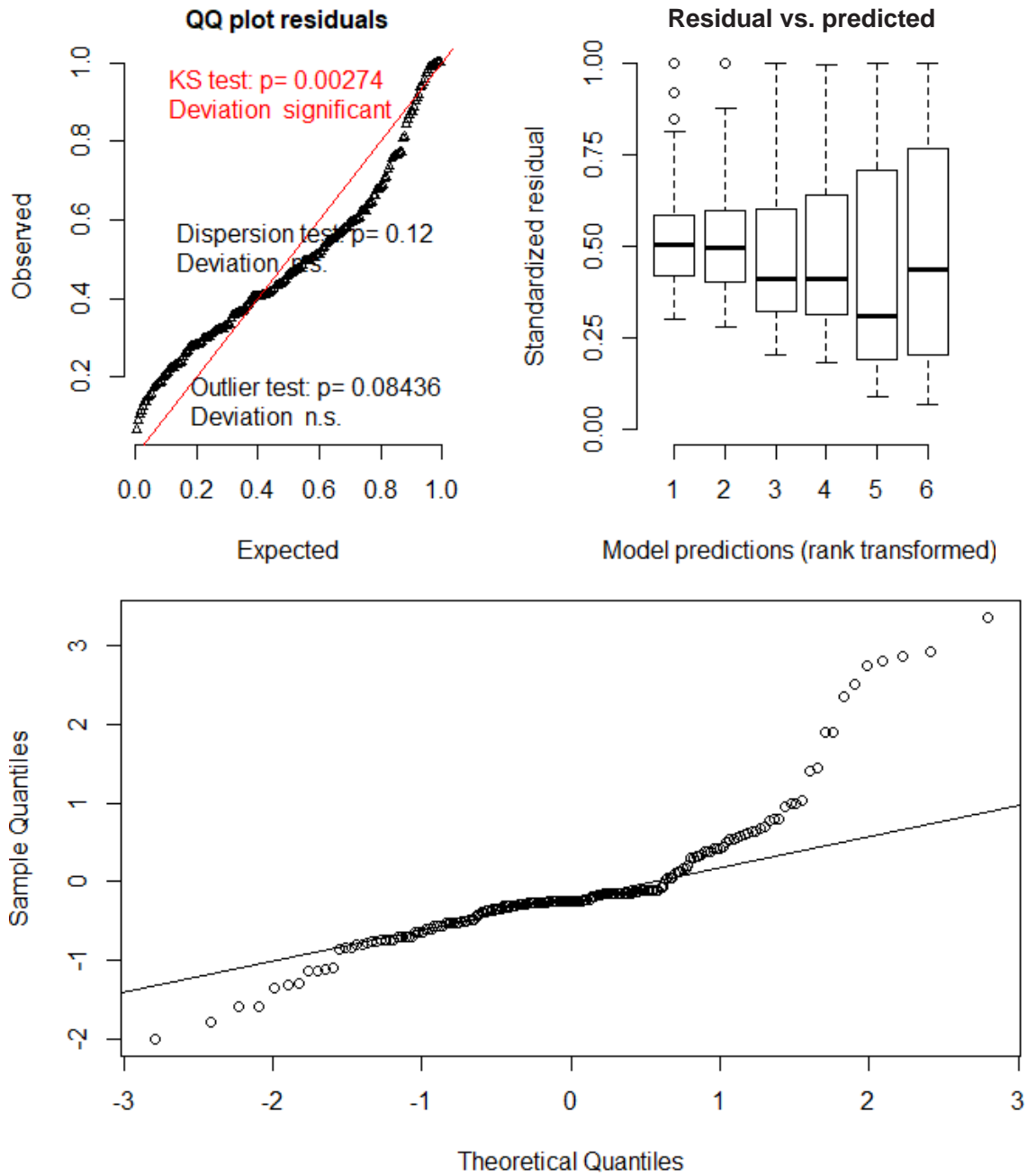
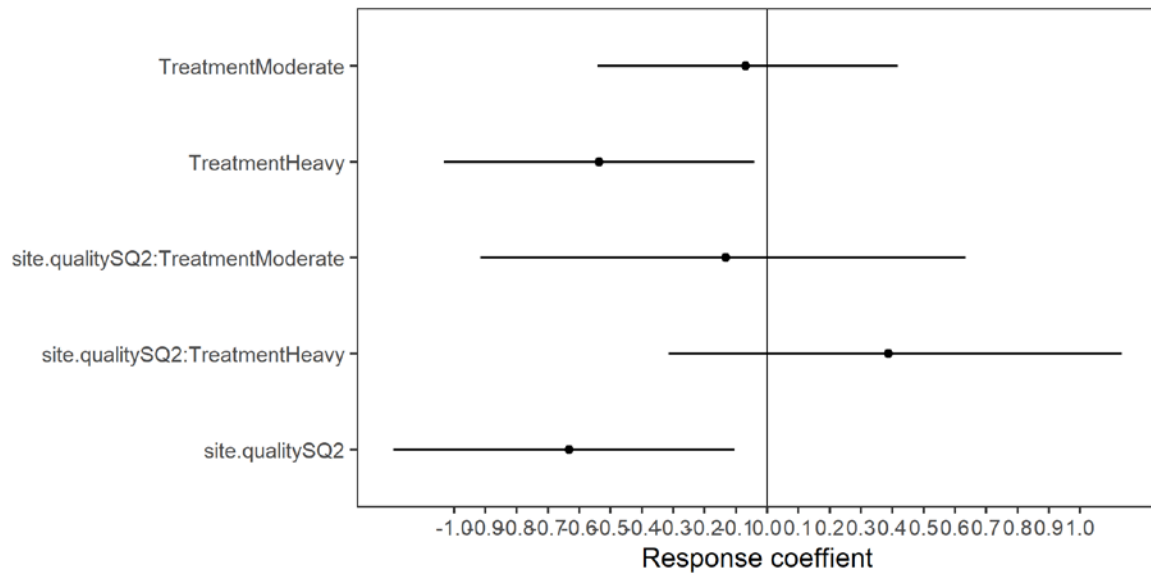


Figure 42 Dunn Smyth simulated residuals (top panels) and normal QQ plot (bottom panel): native plant cover



**Figure 43** Bootstrapped confidence intervals for fixed effects: native plant cover

**Table 44** Bootstrapped model fitted values for average conditions: native plant cover

Survey year	Site quality	Control	Moderate	Heavy
2019	SQ1	2.8519	3.0232	1.7330
2019	SQ2	1.5776	1.2686	1.3522

## F.2.2 Exotic plant cover

**Table 45 Model fitting summary: exotic plant cover**

<b>Response</b>	Exotic plant cover, visually estimated in each 0.04 ha subplot
<b>Response transformation used</b>	Rounding % cover up to nearest whole value
<b>Other transformations compared</b>	$\text{Log}_e(x+1)$ transformation attempted on gaussian model ( $X+0.1$ )/100 transformation attempted on beta model
<b>R package and function</b>	glmer function from the lme4 package
<b>Distribution used</b>	Binomial with data rounded up to nearest whole %
<b>Other distributions compared</b>	Gaussian Beta
<b>Outliers removed</b>	One value >13% was removed
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and site quality
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Marginal = 1.1% Conditional = 7.2%
<b>Confidence comments</b>	Very low. Failed Dunn Smyth uniformity and dispersion tests

### Model results summary 26: exotic plant cover

```

y <- cbind(success = ceiling(DAT.cover.a$exot.live), failure = 100- ceiling(DAT.cover.a$exot.live))

model.exotic.binom <- glmer(y ~ Site quality * Treatment + (1|fsite) + (1|fsiteplot),
                             data = DAT.cover.a, family = "binomial")

summary(model.exotic.binom)

## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula: y ~ Site quality * Treatment + (1 | fsite) + (1 | fsiteplot)
## Data: DAT.cover.a
##
##      AIC      BIC   logLik deviance df.resid
##  582.6   608.8  -283.3   566.6     188
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.2832 -0.3282 -0.0628  0.0581  4.2551

```

```
##
## Random effects:
## Groups      Name          Variance Std.Dev.
## fsiteplot (Intercept) 0.03727  0.1930
## fsite      (Intercept) 0.18197  0.4266
## Number of obs: 196, groups: fsiteplot, 66; fsite, 22
##
## Fixed effects:
##                                     Estimate Std. Error z value Pr(>|z|)
## (Intercept)                        -4.44026   0.21159  -20.985  <2e-16 ***
## Site qualitySQ2                      0.30941   0.28840   1.073   0.2833
## TreatmentModerate                   -0.05626   0.23856  -0.236   0.8136
## TreatmentHeavy                       0.44308   0.21693   2.043   0.0411 *
## Site qualitySQ2:TreatmentModerate    -0.31753   0.32679  -0.972   0.3312
## Site qualitySQ2:TreatmentHeavy      -0.72038   0.31131  -2.314   0.0207 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

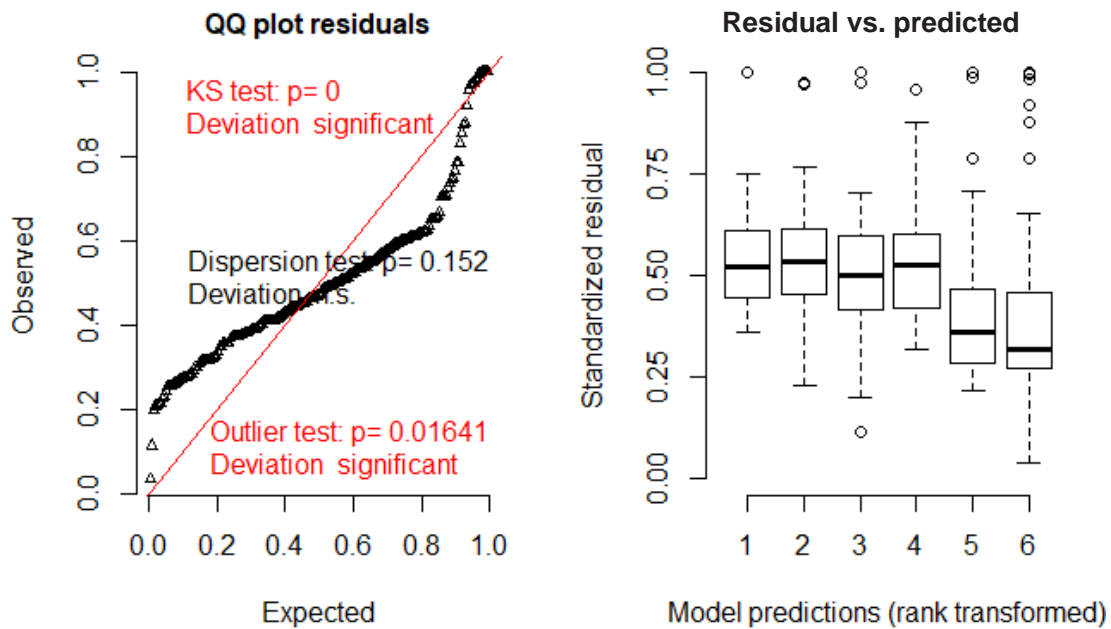


Figure 44 Dunn Smyth simulated residuals: exotic plant cover

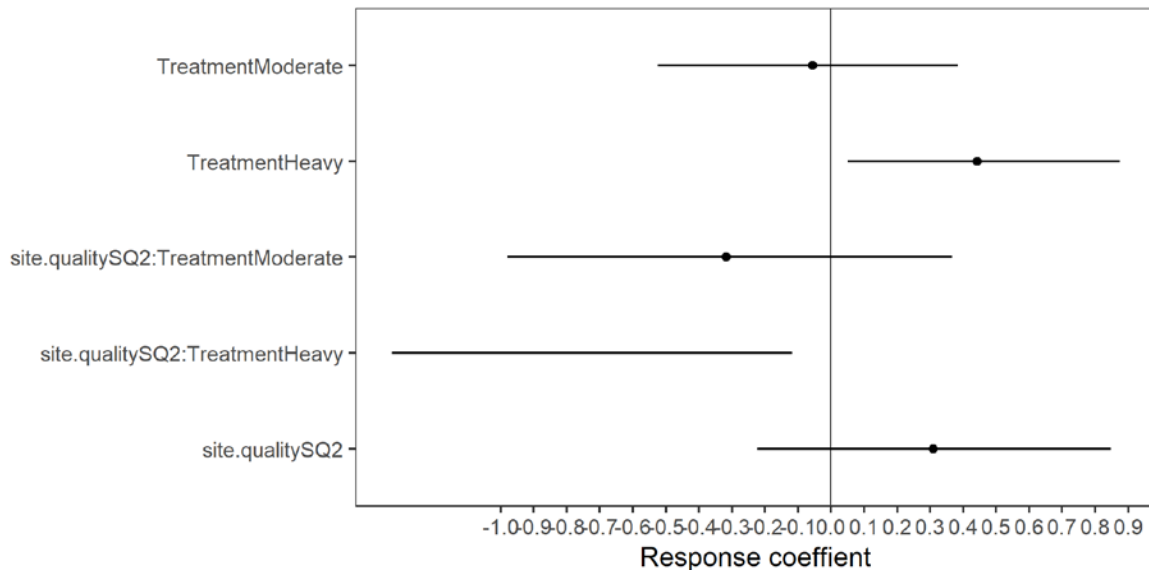


Figure 45 Bootstrapped confidence intervals for fixed effects: exotic plant cover

Table 46 Bootstrapped model fitted values for average conditions: exotic plant cover

Survey year	Site quality	Control	Moderate	Heavy
2019	SQ1	1.2082	1.1428	1.8419
2019	SQ2	1.6823	1.0634	1.2668

### F.3 Floristic community composition 2019–20

This is done by using vegdist and hclust from the vegan package.

### F.4 Threatened plant species

No models were run for threatened plant species.

## Appendix G: Model summaries – Birds

### G.1 Bird species richness

**Table 47 Model fitting summary: bird species richness**

<b>Response</b>	Number of bird species recorded in a 2 ha subplot in each 9 ha plot A positive continuous variable
<b>Response transformation used</b>	None
<b>Other transformations compared</b>	None
<b>R package and function</b>	lmer function from lme4 package
<b>Distribution used</b>	Gaussian
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Ecological thinning treatment (control, moderate, heavy) Survey year (2015, 2017, 2018, 2019) Site quality (SQ1, SQ2)
<b>Fixed factor interactions</b>	Interaction between thinning treatment and survey year Also trialled a model with three-way interaction between thinning treatment, survey year and site quality, which was less parsimonious
<b>Random factors</b>	fsite (a factor over sites), and fsiteplot (a factor over 9 ha treatment plots nested within site)
<b>R<sup>2</sup></b>	Marginal = 5.8% Conditional = 54.5%
<b>Confidence comments</b>	High. Passed all Dunn Smyth tests

### Model results summary 27: bird species richness

```
model.rich3 <- lmer(richness ~ year * Treatment * Site quality + (1|siteplot) + (1|site), data = DAT.richness)
```

```
summary(model.rich)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: richness ~ year * Treatment + Site quality + (1 | siteplot) +
## (1 | site)
## Data: DAT.richness
##
## REML criterion at convergence: 1386.2
##
```



```
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.02929 -0.64965  0.01828  0.64300  2.37681
##
## Random effects:
## Groups   Name            Variance Std.Dev.
## siteplot (Intercept) 2.571     1.603
## site     (Intercept) 7.073     2.660
## Residual                9.006     3.001
## Number of obs: 264, groups: siteplot, 66; site, 22
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    20.8826   1.1146  48.0100  18.736 < 2e-16 ***
## year2017       -2.6364   0.9048 189.0000  -2.914 0.004003 **
## year2018       -3.8636   0.9048 189.0000  -4.270 3.09e-05 ***
## year2019       -3.5909   0.9048 189.0000  -3.969 0.000103 ***
## TreatmentModerate -1.8636   1.0259 168.5657  -1.817 0.071052 .
## TreatmentHeavy   -1.7727   1.0259 168.5657  -1.728 0.085819 .
## Site qualitySQ2  -0.6742   1.2563  20.0000  -0.537 0.597408
## year2017:TreatmentModerate 2.6818   1.2796 189.0000   2.096 0.037433 *
## year2018:TreatmentModerate 2.3182   1.2796 189.0000   1.812 0.071634 .
## year2019:TreatmentModerate 3.7273   1.2796 189.0000   2.913 0.004014 **
## year2017:TreatmentHeavy 2.5909   1.2796 189.0000   2.025 0.044301 *
## year2018:TreatmentHeavy 3.4545   1.2796 189.0000   2.700 0.007571 **
## year2019:TreatmentHeavy 2.5455   1.2796 189.0000   1.989 0.048120 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

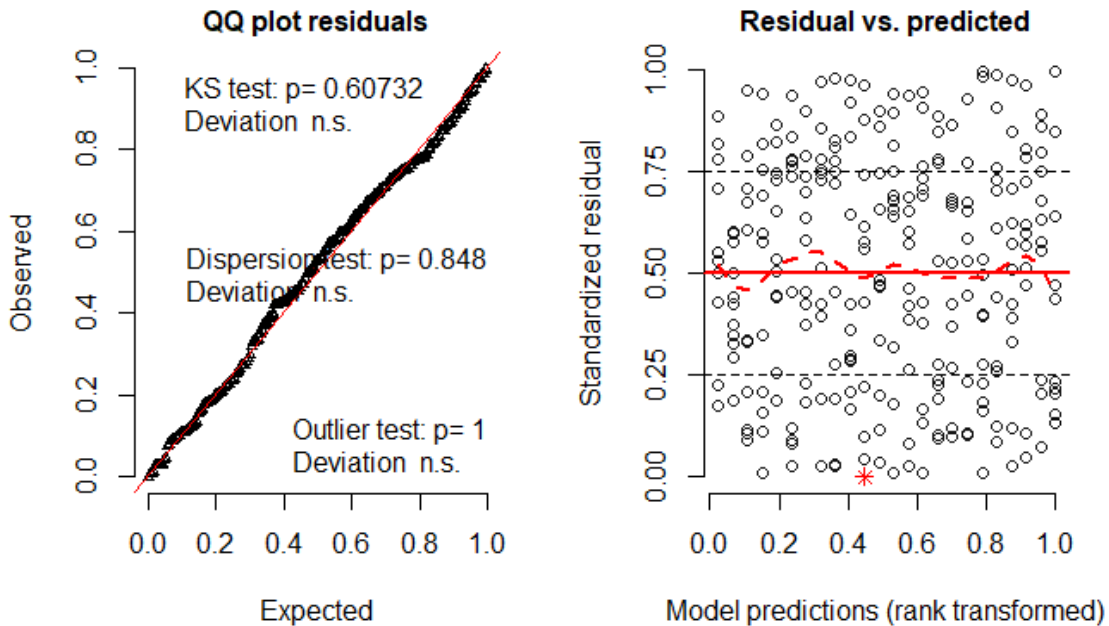


Figure 46 Dunn Smyth simulated residuals: bird species richness

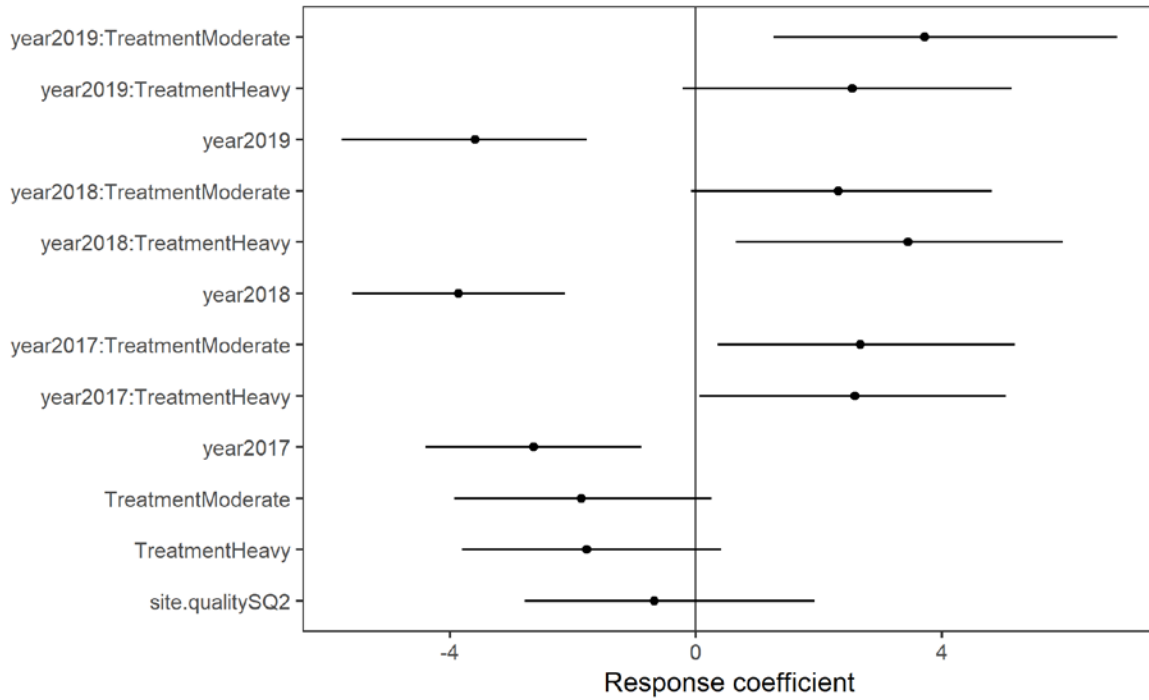


Figure 47 Bootstrapped confidence intervals for fixed effects: bird species richness

Table 48 Bootstrapped model fitted values for average conditions: bird species richness

Survey year	Site quality	Control	Moderate	Heavy
2015	SQ1	20.95207	18.81166	19.09551
2017	SQ1	18.21164	19.12331	19.18590
2018	SQ1	17.20485	17.26662	18.56651
2019	SQ1	17.17184	19.03193	18.19858
2015	SQ2	20.16336	18.40424	18.39405
2017	SQ2	17.49041	18.43281	18.28559
2018	SQ2	16.28156	16.87681	17.91971
2019	SQ2	16.78246	18.44013	17.50177

## G.2 Bird community composition

### G.2.1 Bird community composition 2019–20

See plant community composition above.

### G.3 Threatened bird species

No models were run.

## Appendix H: Model summaries – Bats

### H.1 Overall bat activity

**Table 49 Model fitting summary: overall bat activity**

<b>Response</b>	Total activity of all bat species
<b>Response transformation</b>	None
<b>R package and function</b>	glmer function from lme4 package
<b>Distribution used</b>	Poisson
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Year (2015, 2017, 2018, 2019 and 2020) Thinning treatment (control, moderate and heavy) Site quality (SQ1 and SQ2) Temperature (log of monthly maximum and minimum) and Foliage projective cover (logit)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	plot (a factor over 9 ha treatment plots)
<b>R<sup>2</sup></b>	Marginal = 24.5% Conditional = 47.2%
<b>Confidence comments</b>	Moderate – based on R2 only

### Model results summary 28: overall bat activity

```

Generalized linear mixed model fit by maximum likelihood (Laplace Approximation)
['glmerMod']
Family: poisson ( log )
Formula: Activity ~ Year * Treatment + log(MaxTemp) + log(MinTemp) + qlogis(FPC) +
SQ + (1 | Plot)
Data: Batact

      AIC      BIC   logLik deviance df.resid
63271.6  63368.6 -31615.8  63231.6     924

Scaled residuals:
   Min       1Q   Median       3Q      Max
-15.369  -5.696  -1.310    4.116   44.721

Random effects:
 Groups Name      Variance Std.Dev.
 Plot (Intercept) 0.2024   0.4498
Number of obs: 944, groups: Plot, 66

```

```

Fixed effects:
              Estimate Std. Error z value Pr(>|z|)
(Intercept)    0.13628    0.11634   1.171   0.241
YearY2017     -0.29866    0.01482 -20.146 < 2e-16 ***
YearY2018     -0.12897    0.01193 -10.807 < 2e-16 ***
YearY2019     -0.43427    0.01208 -35.964 < 2e-16 ***
YearY2020     -0.24882    0.01217 -20.453 < 2e-16 ***
TreatmentModerate -0.11424    0.13615  -0.839   0.401
TreatmentHeavy -0.01185    0.13612  -0.087   0.931
log(MaxTemp)   1.17864    0.02017  58.436 < 2e-16 ***
log(MinTemp)   0.53618    0.01013  52.943 < 2e-16 ***
qlogis(FPC)    0.19255    0.00879  21.907 < 2e-16 ***
SQSQ2         -0.23136    0.05767  -4.012 6.03e-05 ***
YearY2017:TreatmentModerate 0.19468    0.01907  10.210 < 2e-16 ***
YearY2018:TreatmentModerate 0.21171    0.01630  12.992 < 2e-16 ***
YearY2019:TreatmentModerate 0.30104    0.01711  17.594 < 2e-16 ***
YearY2020:TreatmentModerate 0.38878    0.01695  22.935 < 2e-16 ***
YearY2017:TreatmentHeavy 0.12462    0.01791   6.960 3.40e-12 ***
YearY2018:TreatmentHeavy 0.39766    0.01570  25.329 < 2e-16 ***
YearY2019:TreatmentHeavy 0.38984    0.01651  23.606 < 2e-16 ***
YearY2020:TreatmentHeavy 0.48772    0.01640  29.739 < 2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
    
```

## H.2 Bat guild activity

### H.2.1 Clutter specialists

**Table 50 Model fitting summary: clutter specialist bat activity**

<b>Response</b>	Total activity of clutter specialist bat species ( <i>Nyctophilus</i> spp.)
<b>Response transformation</b>	None
<b>R package and function</b>	glmer function from lme4 package
<b>Distribution used</b>	Poisson
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Year (2015, 2017, 2018, 2019 and 2020) Thinning treatment (control, moderate and heavy) Site quality (SQ1 and SQ2) Temperature (log of monthly maximum and minimum) and Foliage projective cover (logit)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	plot (a factor over 9 ha treatment plots)
<b>R<sup>2</sup></b>	Marginal = 51.8% Conditional = 56.4%
<b>Confidence comments</b>	Moderate – based on R <sup>2</sup> only

### Model results summary 29: clutter specialist activity

```

Generalized linear mixed model fit by maximum likelihood (Laplace Approximation) [
'glmerMod']
Family: poisson ( log )
Formula: Nyct ~ Year * Treatment + log(MaxTemp) + log(MinTemp) + qlogis(FPC) +
SQ + (1 | Plot)
Data: Batact

      AIC      BIC   logLik deviance df.resid
 9411.9   9508.9  -4685.9   9371.9     924

Scaled residuals:
   Min       1Q   Median       3Q      Max
-8.6559 -1.2750 -0.4383  1.1391 27.9726

Random effects:
 Groups Name      Variance Std.Dev.
 Plot (Intercept) 0.3      0.5477
Number of obs: 944, groups: Plot, 66

```

Fixed effects:					
	Estimate	Std. Error	z value	Pr(> z )	
(Intercept)	1.0803394	0.3691207	2.927	0.00342	**
YearY2017	-0.5224918	0.0609765	-8.569	< 2e-16	***
YearY2018	-2.5263292	0.1006523	-25.100	< 2e-16	***
YearY2019	-0.3136451	0.0441283	-7.108	1.18e-12	***
YearY2020	-2.2014423	0.0833958	-26.398	< 2e-16	***
TreatmentModerate	0.4174611	0.1708359	2.444	0.01454	*
TreatmentHeavy	0.0002503	0.1716795	0.001	0.99884	
log(MaxTemp)	0.2261857	0.1244154	1.818	0.06907	.
log(MinTemp)	0.4765382	0.0531537	8.965	< 2e-16	***
qlogis(FPC)	0.2383790	0.0473708	5.032	4.85e-07	***
SQSQ2	-0.6469071	0.1393483	-4.642	3.44e-06	***
YearY2017:TreatmentModerate	-1.4923440	0.0841234	-17.740	< 2e-16	***
YearY2018:TreatmentModerate	-0.8863092	0.1570616	-5.643	1.67e-08	***
YearY2019:TreatmentModerate	-1.4488981	0.0674051	-21.495	< 2e-16	***
YearY2020:TreatmentModerate	-0.0794297	0.1128784	-0.704	0.48164	
YearY2017:TreatmentHeavy	-0.6573912	0.0838766	-7.838	4.59e-15	***
YearY2018:TreatmentHeavy	0.8166915	0.1291029	6.326	2.52e-10	***
YearY2019:TreatmentHeavy	-0.4339134	0.0705192	-6.153	7.60e-10	***
YearY2020:TreatmentHeavy	0.6143415	0.1168095	5.259	1.45e-07	***
---					
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1					

## H.2.2 Clutter avoiders

**Table 51 Model fitting summary: clutter avoider bat activity**

<b>Response</b>	Total activity of clutter avoider species (all except <i>Nyctophilus</i> spp.)
<b>Response transformation</b>	None
<b>R package and function</b>	glmer function from lme4 package
<b>Distribution used</b>	Poisson
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Year (2015, 2017, 2018, 2019 and 2020) Thinning treatment (control, moderate and heavy) Site quality (SQ1 and SQ2) Temperature (log of monthly maximum and minimum) and Foliage projective cover (logit)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	plot (a factor over 9 ha treatment plots)
<b>R<sup>2</sup></b>	Marginal = 24.6% Conditional = 47.1%
<b>Confidence comments</b>	Moderate – based on R <sup>2</sup> only



**Model results summary 30: clutter avoider activity**

Generalized linear mixed model fit by maximum likelihood (Laplace Approximation) [  
'glmerMod']

Family: poisson ( log )

Formula: NotNyct ~ Year \* Treatment + log(MaxTemp) + log(MinTemp) + qlogis(FPC) +  
SQ + (1 | Plot)

Data: Batact

AIC	BIC	logLik	deviance	df.resid
62902	62999	-31431	62862	924

Scaled residuals:

Min	1Q	Median	3Q	Max
-15.485	-5.486	-1.369	4.249	43.582

Random effects:

Groups Name	Variance	Std.Dev.
Plot (Intercept)	0.221	0.4702

Number of obs: 944, groups: Plot, 66

Fixed effects:

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	-0.214478	0.121008	-1.772	0.07632 .
YearY2017	-0.260091	0.015516	-16.763	< 2e-16 ***
YearY2018	-0.035786	0.012327	-2.903	0.00369 **
YearY2019	-0.458655	0.012798	-35.838	< 2e-16 ***
YearY2020	-0.162403	0.012517	-12.975	< 2e-16 ***
TreatmentModerate	-0.190176	0.142323	-1.336	0.18147
TreatmentHeavy	0.008272	0.142258	0.058	0.95363
log(MaxTemp)	1.223343	0.020690	59.127	< 2e-16 ***
log(MinTemp)	0.561030	0.010431	53.785	< 2e-16 ***
qlogis(FPC)	0.167886	0.009107	18.435	< 2e-16 ***
SQSQ2	-0.240431	0.058873	-4.084	4.43e-05 ***
YearY2017:TreatmentModerate	0.331633	0.019975	16.603	< 2e-16 ***
YearY2018:TreatmentModerate	0.284788	0.016976	16.776	< 2e-16 ***
YearY2019:TreatmentModerate	0.464586	0.018111	25.652	< 2e-16 ***
YearY2020:TreatmentModerate	0.446945	0.017627	25.355	< 2e-16 ***
YearY2017:TreatmentHeavy	0.157800	0.018589	8.489	< 2e-16 ***
YearY2018:TreatmentHeavy	0.366336	0.016178	22.644	< 2e-16 ***
YearY2019:TreatmentHeavy	0.449038	0.017282	25.983	< 2e-16 ***
YearY2020:TreatmentHeavy	0.457768	0.016843	27.179	< 2e-16 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

## H.3 Individual bat species

### H.3.1 White-striped mastiff bat

**Table 52 Model fitting summary: white-striped mastiff bat activity**

<b>Response</b>	Total activity of white-striped mastiff bat
<b>Response transformation</b>	None
<b>R package and function</b>	glmer function from lme4 package
<b>Distribution used</b>	Poisson
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Year (2015, 2017, 2018, 2019 and 2020) Thinning treatment (control, moderate and heavy) Site quality (SQ1 and SQ2) Temperature (log of monthly maximum and minimum) and Foliage projective cover (logit)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	plot (a factor over 9 ha treatment plots)
<b>R<sup>2</sup></b>	Marginal = 24.9% Conditional = 38.9%
<b>Confidence comments</b>	Moderate – based on R <sup>2</sup> only

### Model results summary 31: white-striped mastiff bat activity

```

Generalized linear mixed model fit by maximum likelihood (Laplace Approximation) [
'glmerMod']
Family: poisson ( log )
Formula: Taus ~ Year * Treatment + log(MaxTemp) + log(MinTemp) + qlogis(FPC) +
(1 | Plot)
Data: Batact
Control: glmerControl(optimizer = "bobyqa", optCtrl = list(maxfun = 1e+05))

      AIC      BIC   logLik deviance df.resid
15235.2 15327.4 -7598.6 15197.2     925

Scaled residuals:
   Min     1Q  Median     3Q    Max
-7.084 -2.296 -1.066  1.311 21.954

Random effects:
 Groups Name      Variance Std.Dev.
 Plot   (Intercept) 0.3343   0.5782
Number of obs: 944, groups: Plot, 66

```

Fixed effects:				
	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	-4.76805	0.29096	-16.387	< 2e-16 ***
YearY2017	-1.14396	0.07720	-14.818	< 2e-16 ***
YearY2018	-0.28921	0.04689	-6.168	6.90e-10 ***
YearY2019	-1.22559	0.05416	-22.628	< 2e-16 ***
YearY2020	-0.34234	0.04335	-7.896	2.87e-15 ***
TreatmentModerate	-0.47148	0.18053	-2.612	0.00901 **
TreatmentHeavy	-0.32741	0.17995	-1.819	0.06884 .
log(MaxTemp)	1.76636	0.08908	19.829	< 2e-16 ***
log(MinTemp)	0.43401	0.04434	9.788	< 2e-16 ***
qlogis(FPC)	-0.26652	0.03765	-7.080	1.44e-12 ***
YearY2017:TreatmentModerate	0.88613	0.09616	9.215	< 2e-16 ***
YearY2018:TreatmentModerate	0.16990	0.06688	2.540	0.01108 *
YearY2019:TreatmentModerate	1.25932	0.07242	17.388	< 2e-16 ***
YearY2020:TreatmentModerate	0.53995	0.06250	8.639	< 2e-16 ***
YearY2017:TreatmentHeavy	0.77790	0.09199	8.456	< 2e-16 ***
YearY2018:TreatmentHeavy	0.57415	0.06145	9.343	< 2e-16 ***
YearY2019:TreatmentHeavy	0.99716	0.07155	13.936	< 2e-16 ***
YearY2020:TreatmentHeavy	0.53654	0.06007	8.932	< 2e-16 ***
---				
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1				

### H.3.2 Large forest bat

**Table 53 Model fitting summary: large forest bat activity**

<b>Response</b>	Total activity of large forest bat
<b>Response transformation</b>	None
<b>R package and function</b>	glmer function from lme4 package
<b>Distribution used</b>	Poisson
<b>Other distributions compared</b>	None
<b>Outliers removed</b>	None
<b>Fixed factors</b>	Year (2015, 2017, 2018, 2019 and 2020) Thinning treatment (control, moderate and heavy) Site quality (SQ1 and SQ2) Temperature (log of monthly maximum and minimum) and Foliage projective cover (logit)
<b>Fixed factor interactions</b>	Two-way interaction between thinning treatment and survey year
<b>Random factors</b>	plot (a factor over 9 ha treatment plots)
<b>R<sup>2</sup></b>	Marginal = 16.7% Conditional = 44.6%
<b>Confidence comments</b>	Moderate – based on R <sup>2</sup> only

### Model results summary 32: large forest bat activity

```

Generalized linear mixed model fit by maximum likelihood (Laplace Approximation)
['glmerMod']
Family: poisson ( log )
Formula: Vdar ~ Year * Treatment + log(MaxTemp) + log(MinTemp) + qlogis(FPC) +
(1 | Plot)
Data: Batact
Control: glmerControl(optimizer = "bobyqa", optCtrl = list(maxfun = 1e+05))

      AIC      BIC   logLik deviance df.resid
10569.2 10661.3 -5265.6 10531.2     925

Scaled residuals:
      Min       1Q   Median       3Q      Max
-6.6025 -1.7146 -0.6637  1.0241 18.5381

Random effects:
Groups Name          Variance Std.Dev.
Plot (Intercept) 0.7509  0.8666
Number of obs: 944, groups: Plot, 66

Fixed effects:
              Estimate Std. Error z value Pr(>|z|)
(Intercept)   -4.33555    0.36735 -11.802 < 2e-16 ***
YearY2017     -0.43363    0.07869  -5.510 3.58e-08 ***
YearY2018     -0.14312    0.07143  -2.004 0.045123 *
YearY2019     -0.29834    0.07512  -3.972 7.14e-05 ***
YearY2020     -0.07508    0.08034  -0.935 0.350031
TreatmentModerate  0.44910    0.27163   1.653 0.098264 .
TreatmentHeavy  0.92501    0.27035   3.422 0.000623 ***
log(MaxTemp)   1.43257    0.10841  13.214 < 2e-16 ***
log(MinTemp)   0.49907    0.05385   9.268 < 2e-16 ***
qlogis(FPC)    0.85368    0.04052  21.067 < 2e-16 ***
YearY2017:TreatmentModerate  0.46114    0.09466   4.871 1.11e-06 ***
YearY2018:TreatmentModerate -0.35187    0.09805  -3.589 0.000332 ***
YearY2019:TreatmentModerate  0.74905    0.09427   7.946 1.92e-15 ***
YearY2020:TreatmentModerate  0.17892    0.10265   1.743 0.081334 .
YearY2017:TreatmentHeavy  0.08896    0.08820   1.009 0.313139
YearY2018:TreatmentHeavy -0.37241    0.09104  -4.090 4.31e-05 ***
YearY2019:TreatmentHeavy  0.51986    0.08930   5.822 5.82e-09 ***
YearY2020:TreatmentHeavy  0.73356    0.09252   7.929 2.21e-15 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

# Appendix I: Plant species list

## I.1 Native plant species list

Species name	2015	2017	2018	2019
<i>Acacia dealbata</i>	x	x	x	x
<i>Acaena novae-zelandiae</i>	x	x	x	x
<i>Alternanthera denticulata</i>	x	x	x	x
<i>Amphibromus fluitans</i>	x	x		
<i>Amphibromus nervosus</i>	x	x	x	x
<i>Amyema miquelii</i>	x	x		
<i>Amyema miraculosum</i>	x			
<i>Amyema pendula</i>	x	x		
<i>Arthropodium minus</i>	x	x	x	x
<i>Atriplex semibaccata</i>	x	x	x	x
<i>Austrostipa mollis</i>				x
<i>Austrostipa scabra</i>	x		x	x
<i>Azolla filiculoides</i>	x	x	x	x
<i>Azolla pinnata</i>	x		x	x
<i>Boerhavia dominii</i>		x	x	
<i>Brachyscome basaltica</i>	x	x	x	x
<i>Calotis scapigera</i>	x	x	x	x
<i>Cardamine moirensis</i>	x	x		x
<i>Cardamine paucijuga</i>	x	x	x	x
<i>Carex bichenoviana</i>	x			
<i>Carex inversa</i>	x	x	x	x
<i>Carex tereticaulis</i>	x	x	x	x
<i>Centella asiatica</i>	x		x	x
<i>Centella cordifolia</i>	x	x	x	x
<i>Centipeda cunninghamii</i>	x	x	x	x
<i>Centipeda minima</i>	x	x	x	
<i>Chamaesyce drummondii</i>	x		x	
<i>Chloris truncata</i>		x	x	
<i>Chrysocephalum apiculatum</i>	x	x		
<i>Cotula australis</i>	x	x	x	x
<i>Craspedia paludicola</i>		x	x	x
<i>Crassula colorata</i>		x		
<i>Crassula helmsii</i>	x			x

Species name	2015	2017	2018	2019
<i>Cycnogeton</i> spp.	x	x	x	x
<i>Cynodon dactylon</i>	x	x	x	x
<i>Cyperus exaltatus</i>	x	x	x	
<i>Cyperus gunnii</i>		x		
<i>Damasonium minus</i>	x	x	x	x
<i>Daucus glochidiatus</i>	x	x	x	x
<i>Deyeuxia quadriseta</i>	x	x	x	x
<i>Dianella longifolia</i>	x	x	x	x
<i>Dichelachne crinita</i>		x		x
<i>Dichondra repens</i>	x	x	x	x
<i>Dysphania pumilio</i>	x	x	x	x
<i>Echinochloa colona</i>	x			
<i>Eclipta platyglossa</i>	x	x	x	x
<i>Einadia nutans</i>	x	x	x	x
<i>Elatine gratioloides</i>	x	x		
<i>Eleocharis</i> spp.	x	x	x	x
<i>Elymus scaber</i>	x		x	x
<i>Enchylaena tomentosa</i>	x		x	x
<i>Enteropogon acicularis</i>	x			
<i>Epilobium billardioreanum</i>	x	x	x	x
<i>Epilobium hirtigerum</i>	x	x	x	x
<i>Eragrostis elongata</i>	x			
<i>Eryngium ovinum</i>	x	x	x	x
<i>Eucalyptus camaldulensis</i>	x	x	x	x
<i>Eucalyptus melliodora</i>	x		x	x
<i>Euchiton involucratus</i>	x	x	x	x
<i>Euchiton sphaericus</i>	x	x	x	x
<i>Eulalia aurea</i>	x	x		
<i>Euphorbia drummondii</i>	x	x	x	x
<i>Exocarpos strictus</i>	x	x	x	x
<i>Fimbristylis aestivalis</i>		x		x
<i>Glinus lotoides</i>		x		
<i>Glossostigma elatinoides</i>		x	x	x
<i>Goodenia geniculata</i>	x			
<i>Goodenia gracilis</i>	x	x	x	x
<i>Gratiola peruviana</i>	x	x	x	x
<i>Gratiola pumilo</i>		x	x	



Species name	2015	2017	2018	2019
<i>Haloragis heterophylla</i>	x	x		x
<i>Helichrysum bracteatum</i>	x	x	x	
<i>Hydrocotyle sibthorpioides</i>	x	x	x	x
<i>Hypericum gramineum</i>	x	x	x	x
<i>Isolepis hookeriana</i>	x			
<i>Juncus amabilis</i>	x	x	x	x
<i>Juncus aridicola</i>	x	x	x	
<i>Juncus australis</i>	x			
<i>Juncus flavidus</i>	x	x	x	x
<i>Juncus holoschoenus</i>	x	x	x	x
<i>Juncus ingens</i>	x	x	x	x
<i>Juncus procerus</i>		x		
<i>Juncus radula</i>		x		x
<i>Juncus remotiflorus</i>	x			x
<i>Juncus semisolidus</i>			x	
<i>Juncus</i> sp. D1	x	x		
<i>Juncus</i> sp. H		x		
<i>Juncus</i> sp. I		x	x	
<i>Juncus</i> sp. M		x		
<i>Juncus</i> sp. Q	x	x	x	
<i>Juncus</i> sp. U		x		
<i>Juncus subglaucus</i>	x	x		
<i>Juncus subsecundus</i>	x	x	x	x
<i>Lachnagrostis filiformis</i>	x	x	x	x
<i>Lemna disperma</i>			x	
<i>Lepidium pseudohyssopifolium</i>		x	x	
<i>Linum marginale</i>	x	x	x	x
<i>Ludwigia peploides</i>	x	x	x	x
<i>Lythrum hyssopifolia</i>	x	x	x	x
<i>Maireana enchylaenoides</i>		x		x
<i>Marsilea angustifolia</i>		x		
<i>Marsilea costulifera</i>	x	x	x	x
<i>Marsilea drummondii</i>	x	x	x	x
<i>Mentha australis</i>		x		x
<i>Mentha diemenica</i>	x		x	x
<i>Mentha laxiflora</i>	x			
<i>Mentha satureioides</i>	x		x	x

Species name	2015	2017	2018	2019
<i>Myriophyllum</i> spp.	x	x	x	x
<i>Nymphoides crenata</i>	x	x	x	x
<i>Ottelia ovalifolia</i>	x			
<i>Oxalis perennans</i>	x	x	x	x
<i>Paspalidium jubiflorum</i>	x	x	x	x
<i>Paspalum distichum</i>	x	x		
<i>Persicaria decipiens</i>	x	x	x	x
<i>Persicaria hydropiper</i>	x	x	x	x
<i>Persicaria prostrata</i>	x	x	x	x
<i>Phragmites australis</i>	x	x	x	x
<i>Plantago cunninghamii</i>	x			
<i>Plantago gaudichaudii</i>		x	x	
<i>Poa labillardierei</i>	x	x	x	x
<i>Pogonolepis muelleriana</i>				x
<i>Potamogeton cheesemanii</i>	x	x	x	x
<i>Pratia concolor</i>	x	x	x	x
<i>Pratia pedunculata</i>	x			
<i>Pseudognaphalium luteoalbum</i>	x	x	x	x
<i>Pseudoraphis spinescens</i>	x	x	x	x
<i>Ranunculus inundatus</i>	x	x	x	x
<i>Ranunculus lappaceus</i>		x	x	x
<i>Ranunculus pumilio</i>		x	x	x
<i>Rhodanthe corymbiflora</i>		x	x	x
<i>Rorippa laciniata</i>		x		
<i>Rumex brownii</i>	x	x	x	x
<i>Rumex tenax</i>		x	x	x
<i>Rytidosperma bipartitum</i>		x		
<i>Rytidosperma caespitosum</i>	x		x	x
<i>Rytidosperma duttonianum</i>	x	x		
<i>Rytidosperma erianthum</i>			x	x
<i>Rytidosperma fulvum</i>	x	x	x	
<i>Rytidosperma setaceum</i>	x	x	x	x
<i>Sclerolaena muricata</i>	x			
<i>Senecio bathurstianus</i>				x
<i>Senecio campylocarpus</i>	x	x	x	x
<i>Senecio quadridentatus</i>	x	x	x	x
<i>Senecio runcinifolius</i>	x	x	x	x

Species name	2015	2017	2018	2019
<i>Sigesbeckia orientalis</i>	x	x	x	x
<i>Solenogyne dominii</i>	x	x	x	x
<i>Spergularia brevifolia</i>		x	x	x
<i>Stellaria angustifolia</i>	x	x	x	x
<i>Stellaria caespitosa</i>		x		
<i>Stellaria pungens</i>	x	x		
<i>Stellaria spaff-angustifolia</i>		x		
<i>Typha australis</i>			x	
<i>Typha domingensis</i>	x	x	x	
<i>Typha orientalis</i>	x			
<i>Verbena gaudichaudii</i>	x	x	x	
<i>Vicia disperma</i>				x
<i>Viola betonicifolia</i>	x			
<i>Vittadinia cuneata</i>	x	x	x	x
<i>Vittadinia gracilis</i>	x	x	x	x
<i>Vulpia bromoides</i>	x	x		x
<i>Wahlenbergia</i> spp.	x	x	x	x
<i>Walwhalleya proluta</i>			x	
<i>Xerochrysum bracteatum</i>	x	x	x	x
<i>Xerochrysum viscosum</i>	x			x

## I.2 Exotic plant species list

Species name	2015	2017	2018	2019
<i>Aira cupaniana</i>	x		x	
<i>Aira elegantissima</i>		x		x
<i>Anagallis arvensis</i>	x	x	x	x
<i>Anthoxanthum odoratum</i>	x	x		
<i>Arctotheca calendula</i>	x	x		
<i>Aster subulatus</i>	x	x	x	x
<i>Avena barbata</i>	x	x	x	x
<i>Avena fatua</i>	x	x		x
<i>Avena sativa</i>				x
<i>Briza minor</i>	x	x		
<i>Bromus diandrus</i>	x	x	x	
<i>Bromus hordeaceus</i>	x	x	x	x
<i>Bromus molliformis</i>	x			x

Species name	2015	2017	2018	2019
<i>Bromus rubens</i>		x		x
<i>Callitriche stagnalis</i>		x		
<i>Capsella bursa-pastoris</i>		x	x	x
<i>Carduus tenuiflorus</i>	x	x		x
<i>Centaurea melitensis</i>	x	x	x	x
<i>Centaurium tenuiflorum</i>	x	x	x	x
<i>Chenopodium album</i>	x			x
<i>Chondrilla juncea</i>			x	x
<i>Cirsium vulgare</i>	x	x	x	x
<i>Coryza</i> spp.	x	x	x	x
<i>Cucumis myriocarpus</i>	x			x
<i>Cyperus eragrostis</i>	x	x		x
<i>Dittrichia graveolens</i>		x	x	
<i>Echinochloa crus-galli</i>	x			
<i>Echium plantagineum</i>	x	x	x	x
<i>Echium vulgare</i>	x			x
<i>Ehrharta erecta</i>	x	x	x	x
<i>Ehrharta longiflora</i>	x	x	x	
<i>Euphorbia peplus</i>		x	x	x
<i>Fumaria muralis</i>	x	x	x	x
<i>Fumaria officinalis</i>	x			
<i>Galium aparine</i>	x	x	x	x
<i>Gnaphalium polycaulon</i>		x		
<i>Hedypnois rhagadioloides</i>	x	x		
<i>Heliotropium europaeum</i>	x	x		
<i>Helminthotheca echioides</i>	x	x	x	x
<i>Hordeum glaucum</i>		x	x	x
<i>Hypericum perforatum</i>	x		x	x
<i>Hypochaeris glabra</i>	x	x	x	x
<i>Hypochaeris radicata</i>	x	x	x	x
<i>Kickxia elatine</i>				x
<i>Lactuca saligna</i>	x	x	x	x
<i>Lactuca serriola</i>	x	x	x	x
<i>Leontodon taraxacoides</i>	x	x	x	
<i>Lepidium africanum</i>				x
<i>Lolium loliaceum</i>	x		x	x
<i>Lolium perenne</i>	x	x		x

Species name	2015	2017	2018	2019
<i>Lolium rigidum</i>	x	x	x	x
<i>Lotus subbiflorus</i>	x			
<i>Ludwigia palustris</i>	x	x	x	x
<i>Marrubium vulgare</i>	x	x	x	x
<i>Medicago polymorpha</i>		x	x	x
<i>Mentha pulegium</i>	x	x	x	x
<i>Modiola caroliniana</i>		x		x
<i>Panicum capillare</i>			x	
<i>Petrorhagia dubia</i>	x	x	x	x
<i>Phalaris paradoxa</i>	x	x	x	x
<i>Phyla canescens</i>		x	x	x
<i>Phyla nodiflora</i>			x	
<i>Plantago coronopus</i>	x	x		x
<i>Polycarpon tetraphyllum</i>	x			
<i>Polygonum arenastrum</i>	x			
<i>Polygonum aviculare</i>	x	x	x	x
<i>Rorippa nasturtium-aquaticum</i>	x			
<i>Rorippa palustris</i>		x	x	
<i>Rostraria cristata</i>	x			
<i>Rubus fruticosus</i>		x	x	x
<i>Sagittaria platyphylla</i>	x	x	x	x
<i>Scorzonera laciniata</i>		x		
<i>Sisymbrium irio</i>		x		x
<i>Solanum nigrum</i>	x	x	x	x
<i>Sonchus asper</i>	x	x	x	x
<i>Sonchus oleraceus</i>	x	x	x	x
<i>Spergularia diandra</i>	x			
<i>Stellaria media</i>	x	x	x	x
<i>Stellaria pallida</i>				x
<i>Taraxacum officinale</i>	x	x	x	
<i>Trifolium angustifolium</i>		x	x	
<i>Trifolium arvense</i>	x	x	x	x
<i>Trifolium campestre</i>	x	x	x	x
<i>Trifolium cernuum</i>	x	x	x	
<i>Trifolium dubium</i>		x		
<i>Trifolium globosum</i>			x	
<i>Trifolium glomeratum</i>	x	x	x	x

Species name	2015	2017	2018	2019
<i>Trifolium hirtum</i>	x		x	x
<i>Trifolium subterraneum</i>		x		
<i>Trifolium tomentosum</i>		x		
<i>Urtica urens</i>		x		x
<i>Verbascum virgatum</i>	x	x	x	x
<i>Verbena bonariensis</i>	x	x	x	x
<i>Verbena officinalis</i>		x	x	x
<i>Veronica peregrina</i>		x		
<i>Vicia hirsuta</i>	x		x	
<i>Vicia sativa</i>		x		x
<i>Vulpia muralis</i>	x	x	x	
<i>Vulpia myuros</i>	x	x	x	
<i>Xanthium spinosum</i>	x	x	x	x



## Appendix J: Bird species list

Species name	2015	2017	2018	2019
Australian hobby	x			x
Australian magpie	x	x	x	x
Australian raven	x	x	x	x
Australian reed warbler	x	x		x
Australian shelduck	x			
Australian white ibis	x	x		
Australian wood duck		x		
Azure kingfisher		x		
Black-chinned honeyeater	x			x
Black-faced cuckoo-shrike	x	x	x	x
Blue-faced honeyeater	x	x		x
Brahminy kite			x	
Brown-headed honeyeater	x	x	x	x
Brown falcon		x		
Brown goshawk	x	x	x	x
Brown songlark				x
Brown thornbill	x	x	x	x
Brown treecreeper	x	x	x	x
Buff-rumped thornbill	x	x	x	x
Collared sparrowhawk		x		
Common bronzewing	x	x	x	x
Crested shrike-tit	x	x	x	x
Crimson rosella			x	
Dollarbird	x	x	x	x
Dusky woodswallow	x	x	x	x
Eastern great egret	x			
Eastern rosella	x	x	x	x
Emu	x	x	x	
Fan-tailed cuckoo	x	x		x
Fork-tailed swift			x	
Galah	x	x	x	x
Golden whistler	x	x		x
Great egret	x			
Grey-crowned babbler		x		
Grey butcherbird			x	

Species name	2015	2017	2018	2019
Grey fantail	x	x	x	x
Grey shrike-thrush	x	x	x	x
Grey teal	x			x
Hooded robin	x	x		x
Horsfields bronze-cuckoo	x	x	x	x
Intermediate egret	x			
Jacky winter	x	x	x	x
Laughing kookaburra	x	x	x	x
Leaden flycatcher	x	x		x
Little eagle	x	x		
Little friarbird	x	x	x	x
Little grassbird	x	x		
Little raven	x	x	x	x
Long-billed corella	x	x	x	x
Magpie-lark	x	x		
Mistletoebird	x	x	x	x
Nankeen night heron	x		x	
Noisy friarbird	x	x	x	x
Olive-backed oriole	x			x
Pacific black duck	x	x	x	x
Painted button-quail	x	x		
Painted honeyeater			x	x
Pallid cuckoo		x	x	
Peaceful dove	x	x	x	x
Peregrine falcon	x	x	x	x
Pied butcherbird	x	x		x
Rainbow bee-eater	x	x	x	x
Red-capped robin	x	x	x	x
Red-rumped parrot	x	x	x	x
Red wattlebird	x			
Restless flycatcher	x	x		
Royal spoonbill	x			
Rufous fantail		x		
Rufous songlark	x	x		x
Rufous whistler	x	x	x	x
Sacred kingfisher	x	x	x	x
Satin flycatcher		x		

Species name	2015	2017	2018	2019
Scarlet robin	x	x	x	x
Shining bronze-cuckoo	x	x	x	
Silvereeye	x	x	x	x
Southern boobook	x			
Spotted pardalote	x	x	x	x
Striated pardalote	x	x	x	x
Striated thornbill	x	x	x	x
Sulphur-crested cockatoo	x	x	x	x
Superb fairy-wren	x	x	x	x
Superb parrot	x	x	x	x
Swamp harrier				x
Tree martin	x	x	x	
Varied sittella	x	x	x	x
Wedge-tailed eagle	x	x	x	x
Weebill	x	x	x	x
Welcome swallow	x	x	x	
Western gerygone	x	x	x	x
Whistling kite	x	x	x	x
White-bellied sea-eagle	x		x	x
White-breasted woodswallow	x	x	x	x
White-browed babbler	x	x	x	x
White-browed scrubwren	x	x	x	x
White-browed woodswallow			x	x
White-faced heron	x	x	x	x
White-necked heron	x	x		x
White-plumed honeyeater	x	x	x	x
White-throated treecreeper	x	x	x	x
White-winged chough	x	x	x	x
White-winged triller	x	x	x	x
Willie wagtail	x	x	x	x
Yellow-rumped thornbill	x	x	x	x
Yellow rosella	x	x	x	x
Yellow thornbill	x	x	x	x