

# Fit simulations for top-performing model

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## Before you begin

This script is number 6 of 6 in a series of scripts used to replicate the analyses presented in the paper: “Life on the edge: Industrial footprint and edge effects variably affect the distribution of a boreal small mammal”

This script was used to fit models to simulated data and get a sense of the predictive accuracy of our red squirrel models.

When running these scripts, please ensure that you have downloaded the complete GitHub repository. This will ensure you have all the files, data, and proper folder structure you will need to run this code and associated analyses.

Also make sure you open RStudio through the R project (OSM\_red\_squirrel\_distribution.Rproj). This will automatically set your working directory to the correct place (wherever you saved the repository) and ensure you don't have to change the file paths for some of the data. This analysis was initially run in R v4.3.0. If you have any questions or concerns, please contact one of the authors (in order):

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## 0. Setup

```
rm(list = ls())
#library(MASS)
library(glmmTMB)
#library(lme4)
library(tidyverse)
library(MuMIn)
#library(PerformanceAnalytics)
library(ggpubr)
```

## 1. Re-construct the final dataset (best spatial scales)

### 1.1. Reimport the data

```
covs <- read_csv("./data/processed/OSM_all_covariates_HFI_SBFI_final.csv")

response <- read_csv("./data/processed/OSM_monthly_detections_2021_2022_2023.csv") %>%

  # Only species we want is red squirrel
  filter(species == "red squirrel") %>%

  # Only want detections column
  select(-species, -presence) %>%

  rename(squirrel = detections)

  # Add the covariates to the response variable
  data <- response %>%

  left_join(covs, by = c("array", "site"))

  # Make sure there are 20 rows per site/month/year
  data %>%

  group_by(site, month, year) %>%

  summarize(n_obs = n()) %>%

  arrange(n_obs)
```

```

## # A tibble: 4,918 x 4
## # Groups:   site, month [4,754]
##   site     month   year n_obs
##   <chr>    <dbl> <dbl> <int>
## 1 LU13_11     1  2023    22
## 2 LU13_11     2  2023    22
## 3 LU13_11     3  2023    22
## 4 LU13_11     4  2023    22
## 5 LU13_11     5  2023    22
## 6 LU13_11     6  2023    22
## 7 LU13_11     7  2023    22
## 8 LU13_11     8  2023    22
## 9 LU13_11     9  2022    22
## 10 LU13_11    9  2023   22
## # i 4,908 more rows

# 20 for everything. Looks good!!

# z-scaling for variables WITHIN each buffer
data_scaled <- data %>%
  group_by(buffer_dist) %>%
  mutate(across(cfi_site:last_col(), ~ as.numeric(scale(.)))) %>%
  ungroup()

# The mean will be 0 even though we grouped first, since the mean for each buffer is still 0.
summary(data_scaled)

```

	array	site	month	year
##	Length:108196	Length:108196	Min. : 1.000	Min. :2021
##	Class :character	Class :character	1st Qu.: 4.000	1st Qu.:2022
##	Mode :character	Mode :character	Median : 7.000	Median :2023
##			Mean : 6.634	Mean :2023
##			3rd Qu.:10.000	3rd Qu.:2024
##			Max. :12.000	Max. :2024
##				
##	squirrel	array_year	lat	long
##	Min. : 0.0000	Min. :2021	Min. :54.56	Min. :-115.0
##	1st Qu.: 0.0000	1st Qu.:2022	1st Qu.:55.55	1st Qu.:-113.0
##	Median : 0.0000	Median :2022	Median :56.56	Median :-111.8
##	Mean : 0.9339	Mean :2022	Mean :56.30	Mean :-112.2
##	3rd Qu.: 1.0000	3rd Qu.:2023	3rd Qu.:57.10	3rd Qu.:-111.2
##	Max. :70.0000	Max. :2023	Max. :57.56	Max. :-110.1
##				
##	easting_12n	northing_12n	buffer_dist	cfi_site
##	Min. :254793	Min. :6046183	Min. : 50	Min. :-1.0421
##	1st Qu.:375199	1st Qu.:6155882	1st Qu.:1000	1st Qu.:-0.6582
##	Median :448084	Median :6268110	Median :2375	Median :-0.3196
##	Mean :427832	Mean :6241258	Mean :2393	Mean : 0.0000
##	3rd Qu.:487051	3rd Qu.:6330343	3rd Qu.:3750	3rd Qu.: 0.2842
##	Max. :559033	Max. :6379755	Max. :5000	Max. : 6.7755
##				
##	cfi_site_with_harvest	cfi_site_with_vegedges	harvest_0_15	

```

## Min.   :-1.1835      Min.   :-1.1245      Min.   :-0.4446
## 1st Qu.:-0.6693      1st Qu.:-0.6442      1st Qu.:-0.3841
## Median :-0.3013      Median :-0.3122      Median :-0.3240
## Mean    : 0.0000      Mean   : 0.0000      Mean   : 0.0000
## 3rd Qu.: 0.3302      3rd Qu.: 0.2777      3rd Qu.:-0.2203
## Max.    : 6.7241      Max.   : 6.7198      Max.   : 8.8550
##
## harvest_gt_15      harvest_total      osm_industrial      pipe_trans
## Min.   :-0.6256      Min.   :-0.67528     Min.   :-0.50992     Min.   :-0.9652
## 1st Qu.:-0.4893      1st Qu.:-0.56852     1st Qu.:-0.41059     1st Qu.:-0.6878
## Median :-0.3716      Median :-0.41880     Median :-0.30491     Median :-0.3105
## Mean    : 0.0000      Mean   : 0.00000     Mean   : 0.00000     Mean   : 0.0000
## 3rd Qu.:-0.1280      3rd Qu.:-0.03056     3rd Qu.:-0.04273     3rd Qu.: 0.2548
## Max.    : 6.9091      Max.   : 5.08535     Max.   :10.57756     Max.   : 7.4013
##
## railways          roads          seismic          seismic_lines
## Min.   :-0.111      Min.   :-1.3575      Min.   :-1.0230      Min.   :-1.5413
## 1st Qu.:-0.091      1st Qu.:-0.5403      1st Qu.:-0.6049      1st Qu.:-0.6608
## Median :-0.052      Median :-0.1591      Median :-0.3247      Median :-0.2403
## Mean    : 0.000      Mean   : 0.0000      Mean   : 0.0000      Mean   : 0.0000
## 3rd Qu.:-0.038      3rd Qu.: 0.4028      3rd Qu.: 0.2055      3rd Qu.: 0.4071
## Max.    :26.485      Max.   :15.0263      Max.   : 5.7997      Max.   : 5.0866
## NA's    :4918
## seismic_lines_3D    trails          veg_edges          wells_active
## Min.   :-0.4103      Min.   :-1.0576      Min.   :-0.8885      Min.   :-0.56774
## 1st Qu.:-0.3846      1st Qu.:-0.6221      1st Qu.:-0.4756      1st Qu.:-0.47773
## Median :-0.3565      Median :-0.2290      Median :-0.1844      Median :-0.38959
## Mean    : 0.0000      Mean   : 0.0000      Mean   : 0.0000      Mean   : 0.00000
## 3rd Qu.:-0.2986      3rd Qu.: 0.2535      3rd Qu.: 0.1645      3rd Qu.:-0.01219
## Max.    : 6.2937      Max.   :15.8529      Max.   :13.8053      Max.   :19.32623
##
## wells_inactive      wells_total      pct_betu_pap      fire_0_15
## Min.   :-0.9289      Min.   :-0.7841      Min.   :-0.09      Min.   :-0.3601
## 1st Qu.:-0.6250      1st Qu.:-0.5801      1st Qu.:-0.07      1st Qu.:-0.3444
## Median :-0.3368      Median :-0.3392      Median :-0.07      Median :-0.3258
## Mean    : 0.0000      Mean   : 0.0000      Mean   : 0.00      Mean   : 0.0000
## 3rd Qu.: 0.1762      3rd Qu.: 0.1184      3rd Qu.:-0.05      3rd Qu.:-0.3000
## Max.    :13.2921      Max.   : 8.9950      Max.   :20.22      Max.   : 4.5335
## NA's    :59016
## fire_gt_15          pct_lari_lar      lc_broadleaf      lc_coniferous
## Min.   :-0.3240      Min.   :-0.6475      Min.   :-1.0151      Min.   :-2.37235
## 1st Qu.:-0.2827      1st Qu.:-0.5490      1st Qu.:-0.7086      1st Qu.:-0.70739
## Median :-0.2498      Median :-0.3749      Median :-0.4357      Median :-0.07207
## Mean    : 0.0000      Mean   : 0.0000      Mean   : 0.0000      Mean   : 0.00000
## 3rd Qu.:-0.2232      3rd Qu.: 0.1673      3rd Qu.: 0.4674      3rd Qu.: 0.61317
## Max.    : 7.4090      Max.   : 9.4549      Max.   : 4.4843      Max.   : 2.74250
##
## lc_herbs          lc_mixedwood      lc_shrubs          lc_water
## Min.   :-0.7852      Min.   :-1.3024      Min.   :-0.3114      Min.   :-0.5413
## 1st Qu.:-0.6572      1st Qu.:-0.6833      1st Qu.:-0.2894      1st Qu.:-0.3722
## Median :-0.3921      Median :-0.3053      Median :-0.2693      Median :-0.2429
## Mean    : 0.0000      Mean   : 0.0000      Mean   : 0.0000      Mean   : 0.0000
## 3rd Qu.: 0.2136      3rd Qu.: 0.4097      3rd Qu.:-0.2334      3rd Qu.:-0.1295
## Max.    : 6.7377      Max.   : 5.8728      Max.   : 8.9180      Max.   :13.1921

```

```

##      lc_wetland      lc_wetland_treed   pct_pice_gla      pct_pice_mar
##  Min. :-1.0140  Min. :-1.7479  Min. :-0.4873  Min. :-2.1399
##  1st Qu.:-0.6385 1st Qu.:-0.7748  1st Qu.:-0.4199 1st Qu.:-0.9071
##  Median :-0.4023  Median :-0.1654  Median :-0.3262  Median : 0.1614
##  Mean   : 0.0000  Mean   : 0.0000  Mean   : 0.0000  Mean   : 0.0000
##  3rd Qu.: 0.2744  3rd Qu.: 0.6188  3rd Qu.:-0.1272 3rd Qu.: 0.8697
##  Max.   : 6.0380  Max.   : 3.4576  Max.   : 9.5250  Max.   : 1.7085
##
##      pct_pinu_ban      pct_popu_tre  nonanthro_cai_mn  nonanthro_ed
##  Min. :-0.3977  Min. :-1.6340  Min. :-1.6760  Min. :-1.1470
##  1st Qu.:-0.3679 1st Qu.:-0.8626 1st Qu.:-0.7096 1st Qu.:-0.5623
##  Median :-0.3372  Median :-0.1685  Median :-0.1475  Median :-0.2731
##  Mean   : 0.0000  Mean   : 0.0000  Mean   : 0.0000  Mean   : 0.0000
##  3rd Qu.:-0.1929 3rd Qu.: 0.8477  3rd Qu.: 0.6428  3rd Qu.: 0.2039
##  Max.   : 5.0205  Max.   : 2.5358  Max.   : 3.5209  Max.   : 5.1082
##
##      nonanthro_tca      landscape_cai_mn  landscape_ed      landscape_tca
##  Min. :-3.1665  Min.   : NA  Min.   : NA  Min.   : NA
##  1st Qu.:-0.7096 1st Qu.: NA  1st Qu.: NA  1st Qu.: NA
##  Median : 0.1140  Median : NA  Median : NA  Median : NA
##  Mean   : 0.0000  Mean   :NaN  Mean   :NaN  Mean   :NaN
##  3rd Qu.: 0.8431  3rd Qu.: NA  3rd Qu.: NA  3rd Qu.: NA
##  Max.   : 1.6937  Max.   : NA  Max.   : NA  Max.   : NA
##  NA's   :108196  NA's   :108196  NA's   :108196
##
##      nonanthro_cohesion      landscape_cohesion  landscape_contag      landscape_mesh
##  Min. :-17.73901  Min. :-3.90056  Min. :-3.07601  Min. :-1.8860
##  1st Qu.:-0.09964 1st Qu.:-0.69566  1st Qu.:-0.72484 1st Qu.:-0.6640
##  Median : 0.26597  Median :-0.03058  Median :-0.08274  Median :-0.2884
##  Mean   : 0.00000  Mean   : 0.00000  Mean   : 0.00000  Mean   : 0.0000
##  3rd Qu.: 0.50409  3rd Qu.: 0.73933  3rd Qu.: 0.64601  3rd Qu.: 0.3927
##  Max.   : 0.88374  Max.   : 2.95922  Max.   : 4.08776  Max.   : 7.5132
##
##      landscape_np      landscape_shei  landscape_siei
##  Min. :-1.1343  Min. :-4.12918  Min. :-4.3075
##  1st Qu.:-0.5236 1st Qu.:-0.64137  1st Qu.:-0.5770
##  Median :-0.2777  Median : 0.07569  Median : 0.2271
##  Mean   : 0.0000  Mean   : 0.00000  Mean   : 0.0000
##  3rd Qu.: 0.1454  3rd Qu.: 0.70654  3rd Qu.: 0.7397
##  Max.   : 7.0051  Max.   : 3.20495  Max.   : 1.9831
##
##      rm(covs, response)

```

## 1.2. Re-identify top scales:

```

# Pull from the model or specify manually. Manual means we don't need to re-run this every time.
nat_buffer <- 100

comp_buffer <- 4250

config_buffer <- 2250

```

### 1.3. Pull out the final data again

We will pull out the predictors from the submodels **at the appropriate spatial scales** and merge this into one big dataset for final models. This selects a couple extra columns that we don't actually want to model but that's fine, this was efficient. This is the *unscaled* data so that we can run the sampler and apply appropriate constraints before scaling the data again.

```
data_final <- bind_cols(  
  
  # response variables  
  data %>%  
    select(1:squirrel) %>%  
    distinct(),  
  
  # natural data scaled by natural buffer  
  data %>%  
    filter(buffer_dist == nat_buffer) %>%  
    select(fire_0_15:lc_wetland_treed) %>%  
    mutate(natural_buffer = nat_buffer), # won't use this column, just keeping track of the scale  
    ↪ somehow  
  
  # composition data scaled by composition buffer  
  data %>%  
    filter(buffer_dist == comp_buffer) %>%  
    select(harvest_0_15:wells_total) %>%  
    mutate(comp_buffer = comp_buffer), # won't use this column, just keeping track of the scale  
    ↪ somehow  
  
  # configuration data  
  data %>%  
    filter(buffer_dist == config_buffer) %>%  
    select(contains("nonanthro") | contains("landscape") | contains("cfi")) %>%  
    mutate(config_buffer = config_buffer) # won't use this column, just keeping track of the scale  
    ↪ somehow  
) %>%  
  
  relocate(contains("buffer"), .after=squirrel))  
  
summary(data_final)
```

```
##      array          site        month       year  
##  Length:4918      Length:4918     Min.   : 1.000  Min.   :2021  
##  Class :character  Class :character  1st Qu.: 4.000  1st Qu.:2022  
##  Mode  :character  Mode  :character  Median : 7.000  Median :2023  
##                                         Mean   : 6.634  Mean   :2023  
##                                         3rd Qu.:10.000 3rd Qu.:2024  
##                                         Max.   :12.000  Max.   :2024  
##      squirrel      natural_buffer  comp_buffer  config_buffer  
##  Min.   : 0.0000  Min.   :100      Min.   :4250   Min.   :2250  
##  1st Qu.: 0.0000  1st Qu.:100     1st Qu.:4250   1st Qu.:2250  
##  Median : 0.0000  Median :100     Median :4250   Median :2250  
##  Mean   : 0.9339  Mean   :100     Mean   :4250   Mean   :2250  
##  3rd Qu.: 1.0000  3rd Qu.:100     3rd Qu.:4250   3rd Qu.:2250  
##  Max.   :70.0000  Max.   :100     Max.   :4250   Max.   :2250  
##      fire_0_15      fire_gt_15    pct_lari_lar  lc_broadleaf  
##  Min.   :0.000000  Min.   :0.000   Min.   :0.000000  Min.   :0.000000  
##  1st Qu.:0.000000  1st Qu.:0.000  1st Qu.:0.000000  1st Qu.:0.001251
```

```

## Median :0.00000  Median :0.000  Median :0.000000  Median :0.031500
## Mean   :0.05903  Mean   :0.025  Mean   :0.017705  Mean   :0.165230
## 3rd Qu.:0.00000  3rd Qu.:0.000  3rd Qu.:0.002441  3rd Qu.:0.207147
## Max.   :0.99998  Max.   :1.000  Max.   :0.410436  Max.   :0.997400
## lc_coniferous    lc_herbs      lc_mixedwood   lc_shrubs
## Min.   :0.0000  Min.   :0.000000  Min.   :0.000000  Min.   :0.0000000
## 1st Qu.:0.1166  1st Qu.:0.000000  1st Qu.:0.001708  1st Qu.:0.0000000
## Median :0.3411  Median :0.004752  Median :0.024024  Median :0.0000000
## Mean   :0.3693  Mean   :0.030944  Mean   :0.077900  Mean   :0.0254530
## 3rd Qu.:0.5895  3rd Qu.:0.033106  3rd Qu.:0.093809  3rd Qu.:0.0006952
## Max.   :0.9961  Max.   :0.431069  Max.   :0.710038  Max.   :0.9556450
## lc_water        lc_wetland     lc_wetland_treed harvest_0_15
## Min.   :0.000000  Min.   :0.000000  Min.   :0.000000  Min.   :0.000000
## 1st Qu.:0.000000  1st Qu.:0.001763  1st Qu.:0.04007  1st Qu.:0.000000
## Median :0.000000  Median :0.013062  Median :0.15314  Median :0.000000
## Mean   :0.002367  Mean   :0.061577  Mean   :0.23877  Mean   :0.015072
## 3rd Qu.:0.000000  3rd Qu.:0.064013  3rd Qu.:0.38478  3rd Qu.:0.009129
## Max.   :0.180988  Max.   :0.635488  Max.   :0.95080  Max.   :0.190679
## harvest_gt_15   harvest_total   osm_industrial   pipe_trans
## Min.   :0.000000  Min.   :0.000000  Min.   :0.000000  Min.   :0.000000
## 1st Qu.:0.000000  1st Qu.:0.000000  1st Qu.:0.001380  1st Qu.:0.001343
## Median :0.006122  Median :0.01429  Median :0.004808  Median :0.009432
## Mean   :0.034421  Mean   :0.04964  Mean   :0.017392  Mean   :0.013699
## 3rd Qu.:0.037029  3rd Qu.:0.05920  3rd Qu.:0.017930  3rd Qu.:0.019834
## Max.   :0.357731  Max.   :0.38829  Max.   :0.316999  Max.   :0.075590
## railways        roads          seismic       seismic_lines
## Min.   :0.000e+00  Min.   :0.000000  Min.   :0.000000  Min.   :0.000000
## 1st Qu.:0.000e+00  1st Qu.:0.001762  1st Qu.:0.004996  1st Qu.:0.004314
## Median :0.000e+00  Median :0.003128  Median :0.008535  Median :0.006728
## Mean   :1.247e-05  Mean   :0.004132  Mean   :0.012189  Mean   :0.007705
## 3rd Qu.:0.000e+00  3rd Qu.:0.005610  3rd Qu.:0.015080  3rd Qu.:0.009866
## Max.   :1.701e-03  Max.   :0.023507  Max.   :0.075208  Max.   :0.025486
## seismic_lines_3D trails        veg_edges      wells_active
## Min.   :0.000000  Min.   :0.00000000  Min.   :0.000000  Min.   :0.000e+00
## 1st Qu.:0.000000  1st Qu.:0.0005512  1st Qu.:0.002505  1st Qu.:5.372e-05
## Median :0.000000  Median :0.0013095  Median :0.005061  Median :1.047e-03
## Mean   :0.004484  Mean   :0.0018157  Mean   :0.007228  Mean   :4.726e-03
## 3rd Qu.:0.001867  3rd Qu.:0.0024169  3rd Qu.:0.009277  3rd Qu.:5.248e-03
## Max.   :0.064481  Max.   :0.0096659  Max.   :0.062127  Max.   :6.110e-02
## wells_inactive  wells_total    nonanthro_cai_mn nonanthro_ed
## Min.   :0.000000  Min.   :0.000000  Min.   : 0.4909  Min.   : 2.463
## 1st Qu.:0.001104  1st Qu.:0.001783  1st Qu.:21.4830  1st Qu.: 48.377
## Median :0.002435  Median :0.005198  Median :32.7976  Median : 70.769
## Mean   :0.004890  Mean   :0.009951  Mean   :37.0582  Mean   : 96.125
## 3rd Qu.:0.007435  3rd Qu.:0.013065  3rd Qu.:50.4579  3rd Qu.:107.282
## Max.   :0.026174  Max.   :0.091417  Max.   :97.6249  Max.   :503.042
## nonanthro_tca   nonanthro_cohesion landscape_cai_mn landscape_ed
## Min.   : 59.2  Min.   : 94.09  Min.   :0  Min.   :0
## 1st Qu.: 784.2 1st Qu.: 99.31  1st Qu.:0  1st Qu.:0
## Median :1066.2  Median : 99.60  Median :0  Median :0
## Mean   :1014.0  Mean   : 99.37  Mean   :0  Mean   :0
## 3rd Qu.:1297.2  3rd Qu.: 99.77  3rd Qu.:0  3rd Qu.:0
## Max.   :1554.2  Max.   :100.00  Max.   :0  Max.   :0
## landscape_tca  landscape_cohesion landscape_contag landscape_mesh

```

```

## Min.    :0      Min.    :99.08      Min.    :44.10      Min.    : 23.75
## 1st Qu.:0      1st Qu.:99.37      1st Qu.:58.15      1st Qu.: 73.26
## Median :0      Median :99.48      Median :62.90      Median :127.52
## Mean   :0      Mean   :99.49      Mean   :63.57      Mean   :169.60
## 3rd Qu.:0      3rd Qu.:99.62      3rd Qu.:68.46      3rd Qu.:214.44
## Max.   :0      Max.   :99.88      Max.   :90.14      Max.   :1246.62
##   landscape_np  landscape_shei  landscape_siei  cfi_site
## Min.    : 25    Min.    :0.1603    Min.    :0.1494    Min.    :0.00000
## 1st Qu.: 207   1st Qu.:0.5517    1st Qu.:0.6497    1st Qu.:0.02116
## Median : 349   Median :0.6453    Median :0.7651    Median :0.04418
## Mean   : 540   Mean   :0.6334    Mean   :0.7280    Mean   :0.06532
## 3rd Qu.: 580   3rd Qu.:0.7238    3rd Qu.:0.8379    3rd Qu.:0.08529
## Max.   :4255   Max.   :0.9884    Max.   :0.9898    Max.   :0.47178
##   cfi_site_with_harvest cfi_site_with_vegedges
## Min.   :0.00000   Min.   :0.00000
## 1st Qu.:0.02849   1st Qu.:0.02669
## Median :0.05899   Median :0.05142
## Mean   :0.08354   Mean   :0.07503
## 3rd Qu.:0.11090   3rd Qu.:0.09767
## Max.   :0.54318   Max.   :0.47364

```

## 1.4. Re-fit the top model:

Have to fit using the scaled data. Hacky way to deal with scaling and code is a bit repetitive but it works.

```

data_final_scaled <- bind_cols(
  # response variables
  data_scaled %>%
    select(1:squirrel) %>%
    distinct(),

  # natural data scaled by natural buffer
  data_scaled %>%
    filter(buffer_dist == nat_buffer) %>%
    select(fire_0_15:lc_wetland_treed) %>%
    mutate(natural_buffer = nat_buffer), # won't use this column, just keeping track of the scale
    ↴ somehow

  # composition data scaled by composition buffer
  data_scaled %>%
    filter(buffer_dist == comp_buffer) %>%
    select(harvest_0_15:wells_total) %>%
    mutate(comp_buffer = comp_buffer), # won't use this column, just keeping track of the scale
    ↴ somehow

  # configuration data
  data_scaled %>%
    filter(buffer_dist == config_buffer) %>%
    select(contains("nonanthro") | contains("landscape") | contains("cfi")) %>%
    mutate(config_buffer = config_buffer) # won't use this column, just keeping track of the scale
    ↴ somehow
) %>%

relocate(contains("buffer"), .after="squirrel"))

```

```

m_edgeXcfi <- glmmTMB(squirrel ~

  # natural covariates
  fire_0_15 +
  lc_broadleaf +
  lc_coniferous +
  lc_mixedwood +

  # configuration variables
  #landscape_sheli +
  nonanthro_ed +
  cfi_site +
  nonanthro_ed*cfi_site +
  #landscape_mesh +
  (1|array/site),

  data = data_final_scaled,
  family = nbinom2,
  na.action = na.fail)

summary(m_edgeXcfi)

##  Family: nbinom2  ( log )
## Formula:
## squirrel ~ fire_0_15 + lc_broadleaf + lc_coniferous + lc_mixedwood +
##           nonanthro_ed + cfi_site + nonanthro_ed * cfi_site + (1 |      array/site)
## Data: data_final_scaled
##
##          AIC      BIC    logLik deviance df.resid
##  9394.4   9465.9  -4686.2    9372.4     4907
##
## Random effects:
##
## Conditional model:
## Groups      Name        Variance Std.Dev.
## site:array (Intercept) 2.502     1.5816
## array      (Intercept) 0.410     0.6403
## Number of obs: 4918, groups: site:array, 430; array, 10
##
## Dispersion parameter for nbinom2 family (): 0.811
##
## Conditional model:
##                               Estimate Std. Error z value Pr(>|z|)
## (Intercept)             -1.42793   0.23031 -6.200 5.65e-10 ***
## fire_0_15                 0.15770   0.11215   1.406   0.1597
## lc_broadleaf              -0.51842   0.12835  -4.039 5.36e-05 ***
## lc_coniferous              -0.03228   0.11093  -0.291   0.7710
## lc_mixedwood                0.16358   0.09223   1.774   0.0761 .
## nonanthro_ed               -0.08884   0.16631  -0.534   0.5932
## cfi_site                   0.29548   0.12080   2.446   0.0144 *
## nonanthro_ed:cfi_site   -0.25618   0.10011  -2.559   0.0105 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

## 2. Run the simulations

### 2.1. Simulation function

Manually set parameters for debugging the function (eval false)

```
model = m_edgeXcfi
original_data = data_final
n_array = 10
n_site_per_array = 43
n_obs_per_site = 10
sampler_type = "normal"

simulate_top_model <- function(model, original_data, n_array, n_site_per_array, n_obs_per_site,
                                sampler_type){

  # Fixed effects in order of model appearance
  B0 <- fixef(model)$cond[1]
  B1 <- fixef(model)$cond[2]
  B2 <- fixef(model)$cond[3]
  B3 <- fixef(model)$cond[4]
  B4 <- fixef(model)$cond[5]
  B5 <- fixef(model)$cond[6]
  B6 <- fixef(model)$cond[7]
  B7 <- fixef(model)$cond[8]

  # Random effects
  SD_site <- sqrt(as.numeric(summary(model)$varcor$cond[["site:array"]]))
  SD_array <- sqrt(as.numeric(summary(model)$varcor$cond[["array"]]))

  if(sampler_type == "normal"){
    # Generate data values based on normal distribution
    fire_0_15 <- rnorm(100000, mean = mean(original_data$fire_0_15),
                        sd = sd(original_data$fire_0_15))
    lc_broadleaf <- rnorm(100000, mean = mean(original_data$lc_broadleaf),
                           sd = sd(original_data$lc_broadleaf))
    lc_coniferous <- rnorm(100000, mean = mean(original_data$lc_coniferous),
                            sd = sd(original_data$lc_coniferous))
    lc_mixedwood <- rnorm(100000, mean = mean(original_data$lc_mixedwood),
                           sd = sd(original_data$lc_mixedwood))
    nonanthro_ed <- rnorm(100000, mean = mean(original_data$nonanthro_ed),
                           sd = sd(original_data$nonanthro_ed))
    cfi_site <- rnorm(100000, mean = mean(original_data$cfi_site),
                      sd = sd(original_data$cfi_site))
  }

  if(sampler_type == "uniform"){
    # Generate data values based on uniform distribution
    fire_0_15 <- runif(100000, min = min(original_data$fire_0_15),
                        max = max(original_data$fire_0_15))
    lc_broadleaf <- runif(100000, min = min(original_data$lc_broadleaf),
                           max = max(original_data$lc_broadleaf))
    lc_coniferous <- runif(100000, min = min(original_data$lc_coniferous),
                            max = max(original_data$lc_coniferous))
    lc_mixedwood <- runif(100000, min = min(original_data$lc_mixedwood),
                           max = max(original_data$lc_mixedwood))
    nonanthro_ed <- runif(100000, min = min(original_data$nonanthro_ed),
```

```

            max = max(original_data$nonanthro_ed))
cfi_site <- runif(100000, min = min(original_data$cfi_site),
                  max = max(original_data$cfi_site))
}

data_sim <- tibble(
  fire_0_15,
  lc_broadleaf,
  lc_coniferous,
  lc_mixedwood,
  nonanthro_ed,
  cfi_site
) %>%

mutate(landcover = lc_broadleaf + lc_coniferous + lc_mixedwood) %>%
filter(if_all(everything(), ~ . >= 0),
      landcover<1) %>%
filter(row_number(.) <= n_array * n_site_per_array) %>%

# Scale the data using the original scaled values
mutate(
  fire_0_15 = (fire_0_15 - mean(original_data$fire_0_15, na.rm = TRUE)) /
    sd(original_data$fire_0_15, na.rm = TRUE),
  lc_broadleaf = (lc_broadleaf - mean(original_data$lc_broadleaf, na.rm = TRUE)) /
    sd(original_data$lc_broadleaf, na.rm = TRUE),
  lc_coniferous = (lc_coniferous - mean(original_data$lc_coniferous, na.rm = TRUE)) /
    sd(original_data$lc_coniferous, na.rm = TRUE),
  lc_mixedwood = (lc_mixedwood - mean(original_data$lc_mixedwood, na.rm = TRUE)) /
    sd(original_data$lc_mixedwood, na.rm = TRUE),
  nonanthro_ed = (nonanthro_ed - mean(original_data$nonanthro_ed, na.rm = TRUE)) /
    sd(original_data$nonanthro_ed, na.rm = TRUE),
  cfi_site = (cfi_site - mean(original_data$cfi_site, na.rm = TRUE)) /
    sd(original_data$cfi_site, na.rm = TRUE)
) %>%

mutate(array = paste0("Array", rep(1:n_array, each = n_site_per_array)),
      site = paste0(array, "_", 1:n_site_per_array)) %>%

crossing(obs = 1:n_obs_per_site, .) %>%
group_by(array) %>%
mutate(ranef_array = rnorm(n = 1, mean = 0, sd = SD_array)) %>%
ungroup() %>%
group_by(array, site) %>%
mutate(ranef_site = rnorm(n = 1, mean = 0, sd = SD_site)) %>%
ungroup() %>%
mutate(eta = B0 +
      B1 * fire_0_15 +
      B2 * lc_broadleaf +
      B3 * lc_coniferous +
      B4 * lc_mixedwood +

```

```

    B5 * nonanthro_ed +
    B6 * cfi_site +
    B7 * nonanthro_ed * cfi_site +
    ranef_array +
    ranef_site,

    mu = exp(eta),

    squirrel = rnbinom(n = n(), mu = mu, size = sigma(model)))

sim_naive_occ <- data_sim %>%
  group_by(site) %>%
  summarize(dets = max(squirrel)) %>%
  mutate(presence = ifelse(dets > 0, 1, 0)) %>%
  summarize(sum(presence)) %>%
  as.numeric()

sim_indet <- sum(data_sim$squirrel)

sim <- glmmTMB(squirrel ~

  # natural covariates
  fire_0_15 +
  lc_broadleaf +
  lc_coniferous +
  lc_mixedwood +

  # configuration variables
  #landscape_shei +
  nonanthro_ed +
  cfi_site +
  nonanthro_ed*cfi_site +
  #landscape_mesh +

  (1|array/site),

  data = data_sim,
  family = nbnom2,
  na.action = na.fail)

sim_summary <- tibble(cov = names(fixef(sim)$cond),
  value = fixef(sim)$cond) %>%

pivot_wider(names_from = cov) %>%

mutate(`site:array` = sqrt(as.numeric(VarCorr(sim)$cond[[1]])),
  array = sqrt(as.numeric(VarCorr(sim)$cond[[2]])),
  convergence = sim$fit$convergence,
  n_array = n_array,
  n_site_per_array = n_site_per_array,
  n_obs_per_site = n_obs_per_site,
  sim_indet = sim_indet,
  sim_naive_occ = sim_naive_occ,
  sampler_type = sampler_type,
  sim_date = as.character(Sys.Date()),
  )

# Only return a model if it converged

```

```

  #if(sim$fit$convergence==0){
  #  return(sim_summary)
  #}

}

```

## 2.2. Run simulations

Erase all previous simulations:

```

sim_master_results <- list()
save(sim_master_results, file = "./data/raw/simulation_checkpoint.RData")

```

Run a fresh set of 1000 simulations (in batches of 10) for each combination of parameters that we want.

```

# 1000 simulations with normal sampler
for(i in 1:100) {

  cat("\r\r Working on `normal` batch", i)

  load("./data/raw/simulation_checkpoint.RData")

  sim_results <- purrr::map_dfr(1:10, ~
    simulate_top_model(model = m_edgeXcfi,
                        original_data = data_final,
                        n_array = 10,
                        n_site_per_array = 43,
                        n_obs_per_site = 10,
                        sampler_type = "normal"
    )
  )

  sim_master_results[[length(sim_master_results)+1]] <- sim_results

  save(sim_master_results, file = "./data/raw/simulation_checkpoint.RData")
}

# 1000 simulations with uniform sampler
for(i in 1:100) {

  cat("\r\r Working on `uniform` batch", i)

  load("./data/raw/simulation_checkpoint.RData")

  sim_results <- purrr::map_dfr(1:10, ~
    simulate_top_model(model = m_edgeXcfi,
                        original_data = data_final,
                        n_array = 10,
                        n_site_per_array = 43,
                        n_obs_per_site = 10,
                        sampler_type = "uniform"
    )
  )

  sim_master_results[[length(sim_master_results)+1]] <- sim_results
}

```

```

save(sim_master_results, file = "./data/raw/simulation_checkpoint.RData")
}

# 1000 simulations with reduced sites per array
for(i in 1:100) {

  cat("\r\r Working on `reduced sites` batch", i)

  load("./data/raw/simulation_checkpoint.RData")

  sim_results <- purrr::map_dfr(1:10, ~
    simulate_top_model(model = m_edgeXcfi,
                        original_data = data_final,
                        n_array = 10,
                        n_site_per_array = 22,
                        n_obs_per_site = 10,
                        sampler_type = "uniform"
    )
  )

  sim_master_results[[length(sim_master_results)+1]] <- sim_results

  save(sim_master_results, file = "./data/raw/simulation_checkpoint.RData")
}

```

### 3. Visualize results of the simulations

Let's take a look at the master results

```

load("./data/raw/simulation_checkpoint.RData")

# Fetch the simulation results
sim_master_results_df <- sim_master_results %>%
  bind_rows() %>%
  distinct() # Make sure there are no duplicate simulations.

```

Let's also import the pretty names

```

pretty_names <- read_csv("./tables/OSM_all_covariates_formatted_names.csv")

## Rows: 44 Columns: 2
## -- Column specification -----
## Delimiter: ","
## chr (2): Covariate, PrettyName
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.

```

Plot using the normal (Gaussian) sampler

```

true_vals_fixef <- enframe(fixef(m_edgeXcfi)$cond,
                            name = "term", value = "value")

```

```

true_vals_ranef <- enframe(VarCorr(m_edgeXcfi)$cond,
                           name = "term", value = "value") %>%
  mutate(value = sqrt(as.numeric(value)))

true_vals <- bind_rows(true_vals_fixef,
                      true_vals_ranef)

# Run simulations
sim_coefficient_plots <- purrr::map(true_vals$term, ~{

  # Pretty label for the plot
  label <- (pretty_names %>% filter(Covariate == .x))$PrettyName

  true_val <- (true_vals %>% filter(term==.x))$value

  # Fetch the simulation results
  sim_master_results_df %>%

    # For a specific subset of simulations
    filter(sampler_type == "normal",
           n_site_per_array == 43,
           convergence == 0) %>%

    # Pivot to long format to make our lives easier in purrr
    pivot_longer(., cols = 1:10) %>%

    # Coefficient values for just the variable we want
    filter(name == .x) %>%

    # Plot it
    ggplot(., aes(x = value)) +
    # Dotted vertical line at x = 0
    geom_vline(xintercept = 0, color = "grey70") +
    geom_density(fill = ifelse(str_detect(label, "Random")==TRUE,
                               "darkblue", "darkred"), alpha = 0.3) +
    geom_vline(xintercept = true_val,
               linetype = "dashed",
               color = ifelse(str_detect(label, "Random")==TRUE,
                             "darkblue", "darkred"),
               linewidth=0.8) + # Dotted vertical line at x = 0

    # Crop the axis since model weights are low
    scale_y_continuous(expand = expansion(mult = c(0, 0.05))) +
    scale_x_continuous(limits = c(true_val-0.75, true_val+0.75),
                       expand = c(0,0)) +
    labs(y = "",
         x = "",
         title = label) +
    theme_bw() +
    theme(panel.grid = element_blank()) # Remove background panel grid
})

```

```

ggpubr::ggarrange(plotlist = sim_coefficient_plots, ncol = 3, nrow = 4) +
  bgcolor("white")

ggsave("./figures/top_model_simulations_coefficient_density_gaussian_sampler.png",
       width = 9, height = 8)

```

Plot using the uniform distribution sampler

```

true_vals_fixef <- enframe(fixef(m_edgeXcfi)$cond,
                            name = "term", value = "value")

true_vals_ranef <- enframe(VarCorr(m_edgeXcfi)$cond,
                            name = "term", value = "value") %>%
  mutate(value = sqrt(as.numeric(value)))

true_vals <- bind_rows(true_vals_fixef,
                       true_vals_ranef)

# Run simulations
sim_coefficient_plots <- purrr::map(true_vals$term, ~{

  # Pretty label for the plot
  label <- (pretty_names %>% filter(Covariate == .x))$PrettyName

  true_val <- (true_vals %>% filter(term==.x))$value

  # Fetch the simulation results
  sim_master_results_df %>%

    # For a specific subset of simulations
    filter(sampler_type == "uniform",
           n_site_per_array == 43,
           convergence == 0) %>%

    # Pivot to long format to make our lives easier in purrrr
    pivot_longer(., cols = 1:10) %>%

    # Coefficient values for just the variable we want
    filter(name == .x) %>%

    # Plot it
    ggplot(., aes(x = value)) +

      geom_vline(xintercept = 0, color = "grey70") +
      geom_density(fill = ifelse(str_detect(label, "Random")==TRUE,
                                 "darkblue", "darkred"), alpha = 0.3) +
      geom_vline(xintercept = true_val,
                 linetype = "dashed",
                 color = ifelse(str_detect(label, "Random")==TRUE,
                               "darkblue", "darkred"),
                 linewidth=0.8) + # Dotted vertical line at x = 0

    # Crop the axis since model weights are low
    scale_y_continuous(expand = expansion(mult = c(0, 0.05))) +

```

```

scale_x_continuous(limits = c(true_val-0.75, true_val+0.75),
                   expand = c(0,0)) +
  labs(y = "",
       x = "",
       title = label) +
  theme_bw() +
  theme(panel.grid = element_blank()) # Remove background panel grid
}

ggpubr::ggarrange(plotlist = sim_coefficient_plots, ncol = 3, nrow = 4) +
  bgcolor("white")

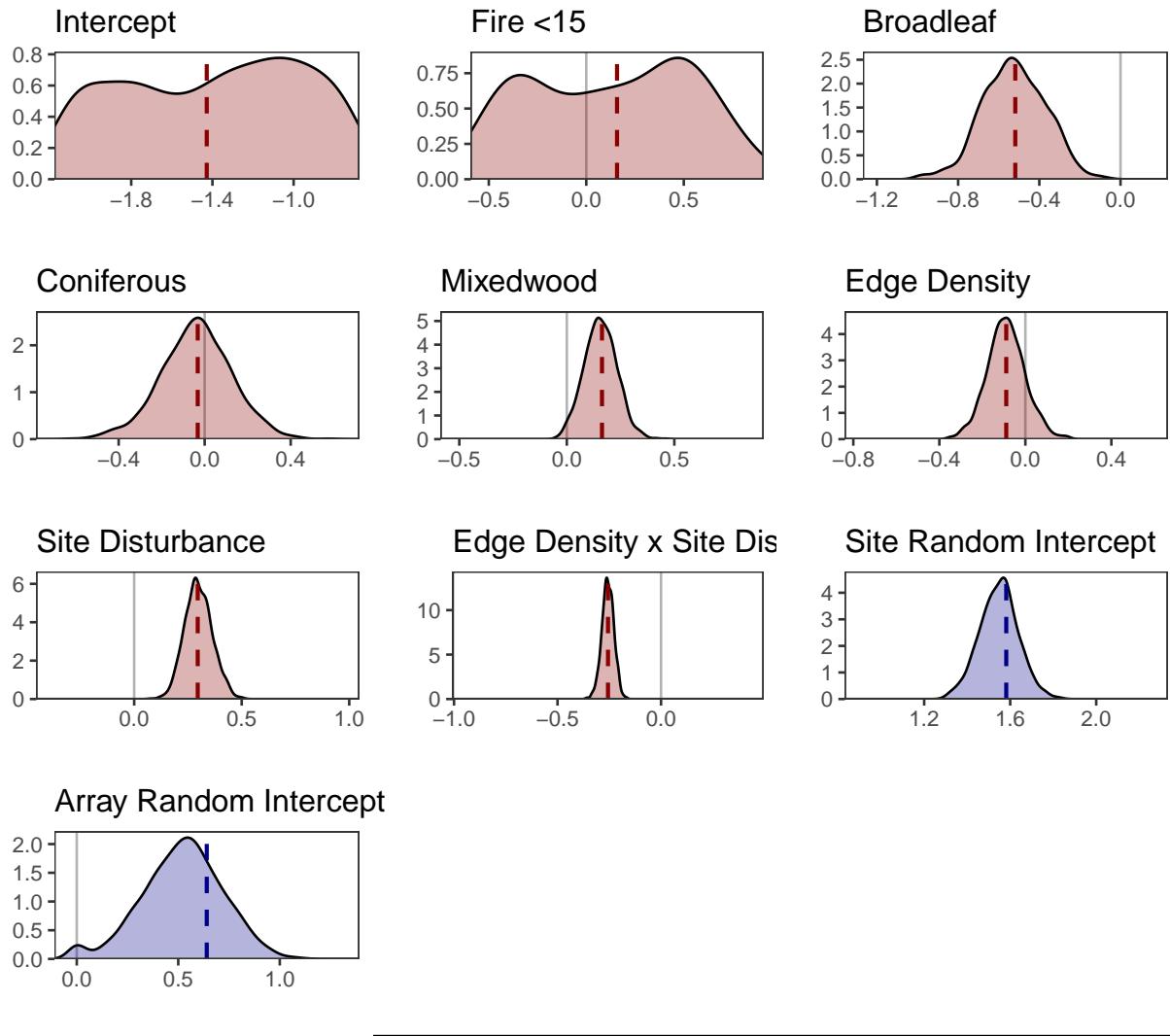
## Warning: Removed 428 rows containing non-finite outside the scale range
## (`stat_density()`).

## Warning: Removed 1 row containing missing values or values outside the scale range
## (`geom_vline()`).

## Warning: Removed 854 rows containing non-finite outside the scale range
## (`stat_density()`).

## Warning: Removed 1 row containing missing values or values outside the scale range
## (`geom_vline()`).

```



```
ggsave("./figures/top_model_simulations_coefficient_density_uniform_sampler.png",
       width = 9, height = 8)
```

Plot using the uniform sampler *and* reduced sample size

```
true_vals_fixef <- enframe(fixef(m_edgeXcfi)$cond,
                           name = "term", value = "value")

true_vals_ranef <- enframe(VarCorr(m_edgeXcfi)$cond,
                           name = "term", value = "value") %>%
  mutate(value = sqrt(as.numeric(value)))

true_vals <- bind_rows(true_vals_fixef,
                      true_vals_ranef)

# Run simulations
sim_coefficient_plots <- purrr::map(true_vals$term, ~{

  # Pretty label for the plot
  # ... (rest of the code is omitted for brevity)
})
```

```

label <- (pretty_names %>% filter(Covariate == .x))$PrettyName
true_val <- (true_vals %>% filter(term==.x))$value

# Fetch the simulation results
sim_master_results_df %>%
  # For a specific subset of simulations
  filter(sampler_type == "uniform",
         n_site_per_array == 22,
         convergence == 0) %>%
  # Pivot to long format to make our lives easier in purrr
  pivot_longer(., cols = 1:10) %>%
  # Coefficient values for just the variable we want
  filter(name == .x) %>%
  # Plot it
  ggplot(., aes(x = value)) +
    geom_vline(xintercept = 0, color = "grey70") +
    geom_density(fill = ifelse(str_detect(label, "Random")==TRUE,
                               "darkblue", "darkred"), alpha = 0.3) +
    geom_vline(xintercept = true_val,
               linetype = "dashed",
               color = ifelse(str_detect(label, "Random")==TRUE,
                             "darkblue", "darkred"),
               linewidth=0.8) + # Dotted vertical line at x = 0
    # Crop the axis since model weights are low
    scale_y_continuous(expand = expansion(mult = c(0, 0.05))) +
    scale_x_continuous(limits = c(true_val-0.75, true_val+0.75),
                       expand = c(0,0)) +
    labs(y = "",
         x = "",
         title = label) +
    theme_bw() +
    theme(panel.grid = element_blank()) # Remove background panel grid
}

ggpubr::ggarrange(plotlist = sim_coefficient_plots, ncol = 3, nrow = 4) +
  bgcolor("white")

ggsave("./figures/top_model_simulations_coefficient_density_reduced_sample_size.png",
       width = 9, height = 8)

```

Let's look at some summary statistics for our empirical samples. Do they match the simulations?

```

true_vals <- enframe(fixef(m_edgeXcfi)$cond,
                      name = "term", value = "mean_model")
true_se <- enframe(summary(m_edgeXcfi)$coefficients$cond[, "Std. Error"],

```

```

        name = "term", value = "se_model")

# Fetch the simulation results
sim_uniform_results <- sim_master_results_df %>%
  # For a specific subset of simulations
  filter(sampler_type == "uniform",
         n_site_per_array == 43,
         convergence == 0) %>%
  select(1:8) %>%
  mutate(sim_id = row_number()) %>%
  pivot_longer(., cols = 1:8, names_to = "term") %>%
  group_by(term) %>%
  summarize(
    mean_estimate = mean(value),
    se_estimate = sd(value)
  ) %>%
  left_join(true_vals, by = "term") %>%
  left_join(true_se, by = "term") %>%
  mutate(mean_bias = mean_estimate - mean_model,
         se_ratio = se_estimate / se_model)%>%
  select(term, contains("mean"), contains("se")) %>%
  arrange(mean_bias)

sim_uniform_results

## # A tibble: 8 x 7
##   term      mean_estimate mean_model mean_bias se_estimate se_model se_ratio
##   <chr>        <dbl>       <dbl>     <dbl>      <dbl>      <dbl>      <dbl>
## 1 lc_mixedwood    0.157      0.164   -0.00647     0.0760     0.0922     0.824
## 2 lc_coniferous   -0.0384     -0.0323  -0.00608     0.164      0.111      1.48 
## 3 lc_broadleaf    -0.524      -0.518   -0.00548     0.155      0.128      1.21 
## 4 nonanthro_ed    -0.257      -0.256   -0.000673    0.0293     0.100      0.293
## 5 nonanthro_ed    -0.0890     -0.0888  -0.000197    0.0889     0.166      0.535
## 6 cfi_site        0.299       0.295    0.00344     0.0641     0.121      0.530
## 7 (Intercept)     -1.40       -1.43     0.0310      0.949      0.230      4.12 
## 8 fire_0_15        0.289       0.158    0.131       3.72       0.112     33.1 

summary(m_edgeXcfi)

## Family: nbinom2  ( log )
## Formula:
## squirrel ~ fire_0_15 + lc_broadleaf + lc_coniferous + lc_mixedwood +
##           nonanthro_ed + cfi_site + nonanthro_ed * cfi_site + (1 | array/site)
## Data: data_final_scaled
##
```

```

##      AIC      BIC  logLik deviance df.resid
##  9394.4   9465.9  -4686.2    9372.4      4907
##
## Random effects:
##
## Conditional model:
## Groups      Name        Variance Std.Dev.
## site:array (Intercept) 2.502     1.5816
## array       (Intercept) 0.410     0.6403
## Number of obs: 4918, groups: site:array, 430; array, 10
##
## Dispersion parameter for nbinom2 family (): 0.811
##
## Conditional model:
##                               Estimate Std. Error z value Pr(>|z|)
## (Intercept)             -1.42793   0.23031 -6.200 5.65e-10 ***
## fire_0_15                0.15770   0.11215  1.406  0.1597
## lc_broadleaf            -0.51842   0.12835 -4.039 5.36e-05 ***
## lc_coniferous            -0.03228   0.11093 -0.291  0.7710
## lc_mixedwood              0.16358   0.09223  1.774  0.0761 .
## nonanthro_ed             -0.08884   0.16631 -0.534  0.5932
## cfi_site                  0.29548   0.12080  2.446  0.0144 *
## nonanthro_ed:cfi_site -0.25618   0.10011 -2.559  0.0105 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

End script