

Fit simulations for top-performing model

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2025-05-02

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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(glmTMB)  
#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.   :0.00000      Min.   :0.000      Min.   :0.000000      Min.   :0.000000  
## 1st Qu.:0.00000      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. :0.00000	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. :0.00000	Min. :0.000000	Min. :0.000000
##	1st Qu.:0.000000	1st Qu.:0.00000	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. :0.0000000	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_shei +
      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 = nbinom2,
              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 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")

```

```

## 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()`).

```

```

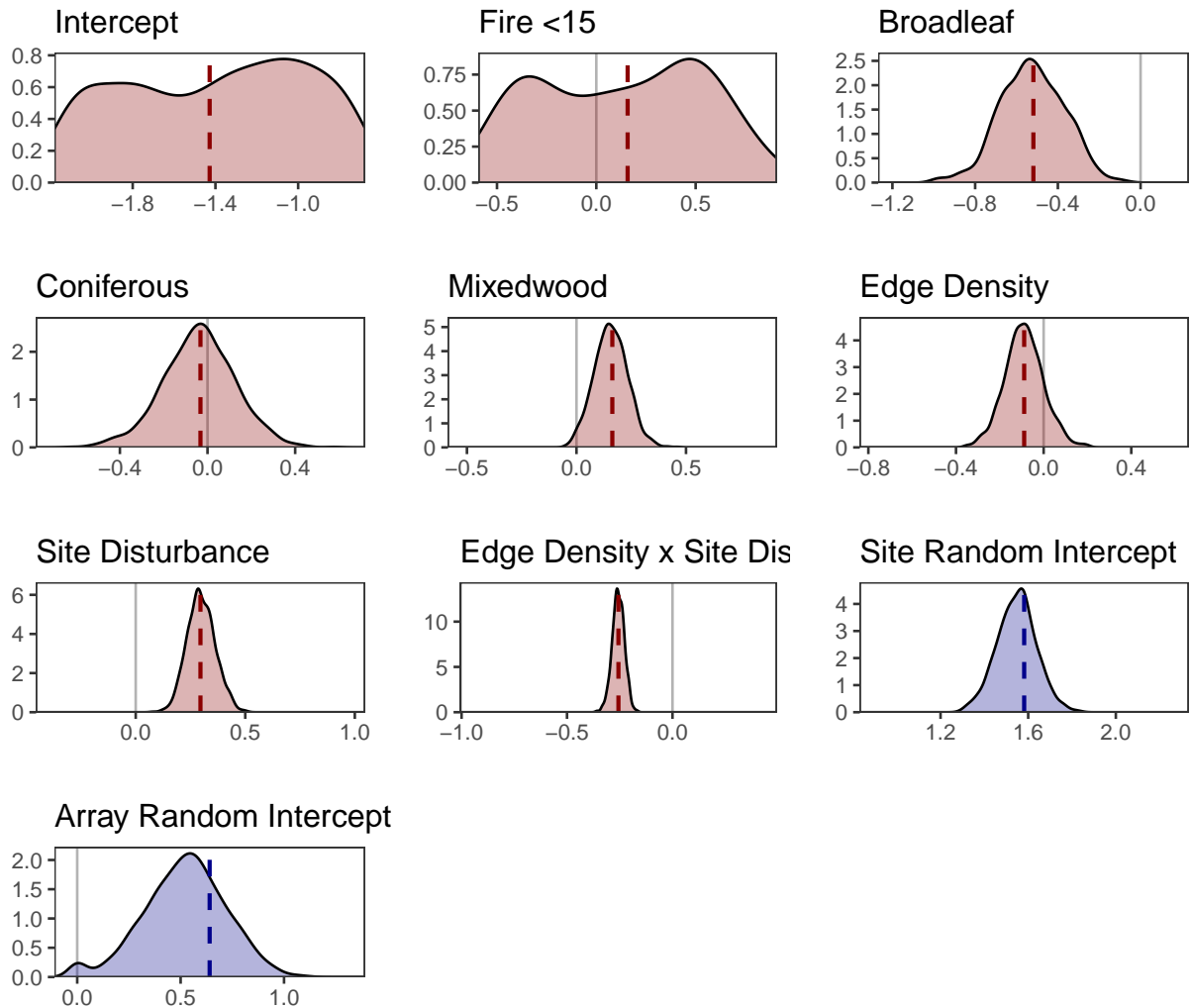
## Warning: Removed 854 rows containing non-finite outside the scale range
## (`stat_density()`).

```

```

## 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
```

```

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