

Model selection for Richness as response variable. An AIC comparison was used to test which of the listed generalized linear mixed models best predicted the data. Models with a dAICc score of less than 2 are bold. Season and nested effect of block, plot and trap, were represented as random effect. The most parsimonious model is marked with an asterisk (*).

| Response | Model | AICc | dAICc | df | weight | R ² |
|----------|--|---------------|------------|-----------|---------------|----------------|
| Richness | ~ 1 | 1930.9 | 371.1 | 1 | <0.001 | - |
| | ~ OM + R + SHR + GWF + UWF + TRE + GRA + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1625.2 | 65.5 | 29 | <0.001 | - |
| | ~ OM + R + GWF + BGT + VWC + I(VWC^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1561.5 | 1.7 | 16 | 0.2346 | 0.55 |
| | ~ OM + R + MUL + BGT + VWC + I(VWC ^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1563.9 | 4.1 | 15 | 0.0710 | - |
| | ~ OM + R + GWF + MUL + BGT + VWC + I(VWC ^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1569.7 | 9.9 | 21 | 0.0039 | - |
| | ~ OM + R + BD + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1613.3 | 53.5 | 8 | <0.001 | - |
| | ~ OM + R + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1613.1 | 53.3 | 9 | <0.001 | - |
| | ~ OM + R + GWF + MUL + BGT + VWC + I(VWC ^2) + BD + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1570.3 | 10.6 | 24 | 0.0028 | - |
| | ~ OM + R + PRCP + TMIN + TMAX + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1606.7 | 47.0 | 10 | <0.001 | - |
| | ~ OM + R + SHR + GWF + UWF + TRE + GRA + MUL + BGT + VWC + I(VWC ^2) + PRCP + TAVG + BD + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1587.7 | 28.0 | 42 | <0.001 | - |
| | ~ BD + R + SHR + GWF + UWF + TRE + GRA + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1624.2 | 64.4 | 29 | <0.001 | - |
| | * ~ BD + R + GWF + BGT + VWC + I(VWC^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1559.8 | 0.0 | 16 | 0.5512 | 0.55 |
| | ~ BD + R + MUL + BGT + VWC + I(VWC ^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1562.7 | 3.0 | 15 | 0.1244 | - |
| | ~ BD + R + GWF + MUL + BGT + VWC + I(VWC ^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1568.4 | 8.6 | 21 | 0.0073 | - |
| | ~ BD + R + TCTN + pH + OM + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1613.7 | 53.9 | 10 | <0.001 | - |
| | ~ BD + R + GWF + MUL + BGT + VWC + I(VWC^2) + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1569.3 | 9.5 | 23 | 0.0047 | - |
| | ~ BD + R + PRCP + TMIN + TMAX + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1603.9 | 44.2 | 10 | <0.001 | - |
| | ~ BD + R + SHR + GWF + UWF + TRE + GRA + MUL + BGT + VWC + I(VWC^2) + PRCP + TAVG + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 1586.8 | 27.0 | 41 | <0.001 | - |

Model outputs for the fixed factors of the Richness response variable. Factors that had a significant impact on the response variable are in bold. The numerical covariates are scaled to represent a fair comparison to the effect size of the fixed factors.

| Response | Factor | Estimate | Std. Error | z value | Pr(> z) |
|----------|--|----------|------------|---------|-----------------|
| Richness | Intercept | -1.35840 | 0.21209 | 6.405 | 1.15e-10 |
| | BD: Bulk Density | -0.08304 | 0.04493 | -1.848 | 0.0646 |
| | R1: Weekly irrigation | 0.04782 | 0.09017 | 0.530 | 0.5958 |
| | GWF2: Ground cover wildflowers 1-5% | 0.05497 | 0.21556 | 0.255 | 0.7987 |
| | GWF3: Ground cover wildflowers 6-25% | 0.18151 | 0.18468 | 0.983 | 0.3257 |
| | GWF4: Ground cover wildflowers 26-50% | 0.35748 | 0.18716 | 1.910 | 0.0561 |
| | GWF5: Ground cover wildflowers 51-75% | 0.41348 | 0.18040 | 2.292 | 0.0219 |
| | GWF6: Ground cover wildflowers 76-95% | 0.43953 | 0.18207 | 2.259 | 0.0183 |
| | GWF7: Ground cover wildflowers 96-100% | 0.52883 | 0.20016 | 2.642 | 0.0082 |
| | BGT: Below ground temperature | 0.28846 | 0.06827 | 4.226 | 2.38e-05 |
| | VWC: Soil moisture | 0.07653 | 0.04933 | 1.552 | 0.1207 |
| | VWC^2: Soil Moisture^2 | -0.10882 | 0.02140 | -5.085 | 3.68e-07 |

Model outputs for the random factors of the Richness response variable. The standard deviation for each factor is listed and is scaled to represent a fair comparison to the effect size of the fixed factors.

| Response | Factor | Std. Dev. |
|----------|-----------------|-----------|
| Richness | Season | 0.13255 |
| | Trap:Plot:Block | 0.18615 |
| | Plot:Block | 0.06494 |
| | Block | 0.15737 |

Model selection for the Abundance response variable. An AIC comparison was used to test which of the listed generalized linear mixed models best predicted the data. Models with a dAICc score of less than 2 are bold. Season and nested effect of block, plot and trap, were represented as random effect. The most parsimonious model is marked with an asterisk (*).

| Response | Model | AICc | dAICc | df | weight | R ² |
|-----------|---|--------------|------------|-----------|---------------|----------------|
| Abundance | ~ 1 | 1089.3 | 280.6 | 2 | <0.001 | - |
| | ~ OM + R + SHR + GWF + UWF + TRE + GRA + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 857.2 | 48.5 | 30 | <0.001 | - |
| | ~ OM + R + GWF + BGT + VWC + I(VWC^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 812.1 | 3.4 | 17 | 0.0857 | - |
| | ~ OM + R + MUL + BGT + VWC + I(VWC^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 810.0 | 1.3 | 16 | 0.2414 | 0.60 |
| | ~ OM + R + GWF + MUL + BGT + VWC + I(VWC^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 820.0 | 11.3 | 22 | 0.0016 | - |
| | ~ OM + R + BD + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 829.4 | 20.7 | 9 | <0.001 | - |
| | ~ OM + R + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 832.3 | 23.7 | 10 | <0.001 | - |
| | ~ OM + R + GWF + MUL + BGT + VWC + I(VWC^2) + BD + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 823.0 | 14.4 | 25 | <0.001 | - |
| | ~ OM + R + PRCP + TMIN + TMAX + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 819.3 | 10.6 | 11 | 0.0023 | - |
| | ~ OM + R + SHR + GWF + UWF + TRE + GRA + MUL + BGT + VWC + I(VWC^2) + PRCP + TAVG + BD + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 829.8 | 11.1 | 43 | 0.0018 | - |
| | ~ BD + R + SHR + GWF + UWF + TRE + GRA + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 855.7 | 47.0 | 30 | <0.001 | - |
| | ~ BD + R + GWF + BGT + VWC + I(VWC^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 810.5 | 1.8 | 17 | 0.1868 | 0.62 |
| | * ~ BD + R + MUL + BGT + VWC + I(VWC^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 808.7 | 0.0 | 16 | 0.4644 | 0.61 |
| | ~ BD + R + GWF + MUL + BGT + VWC + I(VWC^2) + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 818.7 | 10.0 | 22 | 0.0032 | - |
| | ~ BD + R + TCTN + pH + OM + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 831.6 | 23.0 | 11 | <0.001 | - |
| | ~ BD + R + GWF + MUL + BGT + VWC + I(VWC^2) + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 821.9 | 13.3 | 24 | 0.0041 | - |
| | ~ BD + R + PRCP + TMIN + TMAX + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 816.7 | 8.0 | 11 | 0.0083 | - |
| | ~ BD + R + SHR + GWF + UWF + TRE + GRA + MUL + BGT + VWC + I(VWC^2) + PRCP + TAVG + TCTN + pH + (1 SEASON) + (1 BLOCK/PLOT/TRAP) | 818.4 | 9.8 | 43 | 0.0035 | - |

Model outputs for the fixed factors of the Abundance response variable. Factors that had a significant impact on the response variable are in bold. The numerical covariates are scaled to represent a fair comparison to the effect size of the fixed factors.

| Response | Factor | Estimate | Std. Error | z value | Pr(> z) |
|-----------|-------------------------------|----------|------------|---------|-----------------|
| Abundance | Intercept | 2.85564 | 0.36541 | 7.815 | 5.50e-15 |
| | BD: Bulk density | -0.12088 | 0.09756 | -1.239 | 0.2153 |
| | R1: Weekly irrigation | 0.19012 | 0.19029 | 0.999 | 0.3177 |
| | MUL2: Mulch 1-5% | -0.11746 | 0.14489 | -0.811 | 0.4175 |
| | MUL3: Mulch 6-25% | -0.32538 | 0.16580 | 0.1962 | 0.0497 |
| | MUL4: Mulch 26-50% | -0.19008 | 0.16634 | -1.143 | 0.2531 |
| | MUL5: Mulch 51-75% | -0.36415 | 0.22734 | -1.602 | 0.1091 |
| | MUL6: Mulch 76-95% | -0.53753 | 0.28647 | -1.876 | 0.0606 |
| | BGT: Below ground temperature | 0.27478 | 0.12202 | 2.252 | 0.0243 |
| | VWC: Soil moisture | 0.34315 | 0.08951 | 3.834 | 0.0001 |
| | VWC^2: Soil Moisture^2 | -0.15303 | 0.03214 | -4.761 | 1.92e-06 |

Model outputs for the random factors of the Abundance response variable. The standard deviation for each factor is listed and is scaled to represent a fair comparison to the effect size of the fixed factors.

| Response | Factor | Std. Dev. |
|-----------|-----------------|-----------|
| Abundance | Season | 0.5464 |
| | Trap:Plot:Block | 0.4302 |
| | Plot:Block | 0.1318 |
| | Block | 0.1997 |

Family and num.lv selection for Gllvm

```
Call:
gllvm(y = y, family = poisson())
family:
[1] "poisson"
method:
[1] "VA"

log-likelihood: -9301.279
Residual degrees of freedom: 32743
AIC: 19242.56
AICc: 19248.83
BIC: 21932.53
> fit_ord <- gllvm(y, family = "negative.binomial")
> fit_ord
```

```
Call:
gllvm(y = y, family = "negative.binomial")
family:
[1] "negative.binomial"
method:
[1] "VA"
```

```
log-likelihood: -7846.125
Residual degrees of freedom: 32636
AIC: 16546.25
AICc: 16557.45
BIC: 20135.68
```

```
fitx1 <- gllvm(y = sp, X = env, family = "negative.binomial", num.lv = 1)
fitx2 <- gllvm(y = sp, X = env, family = "negative.binomial", num.lv = 2)
fitx3 <- gllvm(y = sp, X = env, family = "negative.binomial", num.lv = 3)
```

```
> AIC(fitx1)
[1] 16150.89
> AIC(fitx2)
[1] 16280.9
> AIC(fitx3)
[1] 16602.67
```

Gllvm model selection. Assessment of the contribution of species traits by comparing model fits for trait-environment interaction models against models with only environmental predictors. To evaluate model performance, we conducted likelihood ratio tests (LRTs) using the `anova` function to compare the goodness of fit between nested generalized linear latent variable models (GLLVMs).

```
> anova(gllvm1a, gllvm2a)
Model 1 : y ~ BGT
Model 2 : y ~ (BGT):(Diet + Wings + Body_Size + Larval_Substrate)
  Resid.Df      D Df.diff P.value
1    32737    0.0000      0
2    32719 143.5206     18      0
> anova(gllvm1b, gllvm2b)
Model 1 : y ~ VWC
Model 2 : y ~ (VWC):(Diet + Wings + Body_Size + Larval_Substrate)
  Resid.Df      D Df.diff  P.value
1    32737    0.00000      0
2    32719 69.77604     18 4.93117e-08
> anova(gllvm1c, gllvm2c)
Model 1 : y ~ GWF
Model 2 : y ~ (GWF):(Diet + Wings + Body_Size + Larval_Substrate)
```

```

      Resid.Df      D Df.diff      P.value
1      32737  0.00000      0
2      32719 60.31941     18 1.8166e-06
> anova(gllvm1d, gllvm2d)
Model 1 : y ~ MUL
Model 2 : y ~ (MUL):(Diet + Wings + Body_Size + Larval_Substrate)
      Resid.Df      D Df.diff      P.value
1      32737  0.00000      0
2      32719 77.97844     18 1.93028e-09

```

Gllvm model selection. Alternative GLLVMs were developed using insights from GLMM outcomes, varying in their inclusion of environmental predictors (GWF, MUL, BGT, VWC) and their interactions with functional traits (diet, wing morphology, body size, and larval substrate). Seasonality was included as a random effect to account for pseudo replication. An AIC comparison was used to test which of the listed generalized linear mixed models best predicted the data.

```

gllvm null: y ~ 1
gllvm01: y ~ (GWF + MUL):(Diet + wings + Body Size + Larval Substrate)
gllvm02: y ~ (BGT + VWC):(Diet + wings + Body Size + Larval Substrate)
gllvm03: y ~ (GWF + BGT):(Diet + wings + Body Size + Larval Substrate)
gllvm04: y ~ (GWF + VWC):(Diet + wings + Body Size + Larval Substrate)
gllvm05: y ~ (MUL + BGT):(Diet + wings + Body Size + Larval Substrate)
gllvm06: y ~ (MUL + VWC):(Diet + wings + Body Size + Larval Substrate)
gllvm07: y ~ (GWF + BGT + VWC):(Diet + wings + Body Size + Larval Substrate)
gllvm08: y ~ (MUL + BGT + VWC):(Diet + wings + Body Size + Larval Substrate)
gllvm09: y ~ (GWF + MUL + BGT + VWC):(Diet + wings + Body Size + Larval Substrate)

```

```

      df      AIC
gllvm09  401 15966.16
gllvm08  382 15983.01
gllvm04  363 15988.57
gllvm07  382 15989.61
gllvm05  363 16029.24
gllvm03  363 16029.97
gllvm02  363 16032.51
gllvm06  363 16130.33
gllvm01  363 16157.27
gllvm_null 322 16559.57

```

```

> pseudo_R2 <- 1 - (gllvm09$logL / gllvm_null$logL)
> pseudo_R2
[1] 0.04721206

```

```

> summary(gllvm09)

```

```

Call:
gllvm(y = y, x = env, TR = TR, formula = y ~ (GWF + MUL + BGT +
VWC):(Diet + wings + Body_Size + Larval_Substrate), family = "negative.binomial",
num.lv = 1, row.eff = "random", randomX = ~SEASON, seed = 123,
control.start = list(n.init = 3, jitter.var = 0.01))

```

```

Family: negative.binomial

```

```

AIC: 15966.16 AICc: 15976.03 BIC: 19337.03 LL: -7582 df: 401

```

```

Informed LVs: 0
Constrained LVs: 0
Unconstrained LVs: 1

```

```

Formula:
~GWF:DietCOP+GWF:DietCOP+DET+GWF:DietDET+GWF:DietMYC+GWF:DietMYC+DET+GWF:DietPHY+GWF:DietPOL+GWF:

```

DietPRD+GWF:WingsDIM+GWF:WingsMAC+GWF:Body_Size+GWF:Larval_SubstrateDPL+GWF:Larval_SubstrateDPL+LPL+GWF:Larval_SubstrateFRT+GWF:Larval_SubstrateLPL+GWF:Larval_SubstrateSOI+GWF:Larval_SubstrateSOI+DNG+GWF:Larval_SubstrateSOI+DPL+GWF:Larval_SubstrateSOI+LPL+MUL:DietCOP+MUL:DietCOP+DET+MUL:DietDET+MUL:DietMYC+MUL:DietMYC+DET+MUL:DietPHY+MUL:DietPOL+MUL:DietPRD+MUL:WingsDIM+MUL:WingsMAC+MUL:Body_Size+MUL:Larval_SubstrateDPL+MUL:Larval_SubstrateDPL+LPL+MUL:Larval_SubstrateFRT+MUL:Larval_SubstrateLPL+MUL:Larval_SubstrateSOI+MUL:Larval_SubstrateSOI+DNG+MUL:Larval_SubstrateSOI+DPL+MUL:Larval_SubstrateSOI+LPL+BGT:DietCOP+BGT:DietCOP+DET+BGT:DietDET+BGT:DietMYC+BGT:DietMYC+DET+BGT:DietPHY+BGT:DietPOL+BGT:DietPRD+BGT:WingsDIM+BGT:WingsMAC+BGT:Body_Size+BGT:Larval_SubstrateDPL+BGT:Larval_SubstrateDPL+LPL+BGT:Larval_SubstrateFRT+BGT:Larval_SubstrateLPL+BGT:Larval_SubstrateSOI+BGT:Larval_SubstrateSOI+DNG+BGT:Larval_SubstrateSOI+DPL+BGT:Larval_SubstrateSOI+LPL+VWC:DietCOP+VWC:DietCOP+DET+VWC:DietDET+VWC:DietMYC+VWC:DietMYC+DET+VWC:DietPHY+VWC:DietPOL+VWC:DietPRD+VWC:WingsDIM+VWC:WingsMAC+VWC:Body_Size+VWC:Larval_SubstrateDPL+VWC:Larval_SubstrateDPL+LPL+VWC:Larval_SubstrateFRT+VWC:Larval_SubstrateLPL+VWC:Larval_SubstrateSOI+VWC:Larval_SubstrateSOI+DNG+VWC:Larval_SubstrateSOI+DPL+VWC:Larval_SubstrateSOI+LPL

LV formula: ~ 0

Row effect: ~(1 | site)

Random effects:

| Name | Variance | Std.Dev | Corr |
|----------|----------|---------|---------|
| SEASONSP | 0.3731 | 0.6108 | |
| SEASONSU | 0.0704 | 0.2653 | -0.9955 |

Coefficients predictors:

| | Estimate | Std. Error | z value | Pr(> z) |
|-----------------------------|----------|------------|---------|--------------|
| GWF:DietCOP | 0.36162 | 0.93675 | 0.386 | 0.699468 |
| GWF:DietCOP+DET | 0.46446 | 0.48269 | 0.962 | 0.335932 |
| GWF:DietDET | 0.31025 | 0.56377 | 0.550 | 0.582104 |
| GWF:DietMYC | 0.74134 | 0.63845 | 1.161 | 0.245579 |
| GWF:DietMYC+DET | -0.15747 | 0.51865 | -0.304 | 0.761416 |
| GWF:DietPHY | 0.06659 | 0.62709 | 0.106 | 0.915431 |
| GWF:DietPOL | 0.09468 | 0.61331 | 0.154 | 0.877310 |
| GWF:DietPRD | -0.06633 | 0.56982 | -0.116 | 0.907333 |
| GWF:WingsDIM | -0.76556 | 0.47923 | -1.597 | 0.110157 |
| GWF:WingsMAC | -0.36467 | 0.23508 | -1.551 | 0.120835 |
| GWF:Body_Size | 0.01310 | 0.08561 | 0.153 | 0.878341 |
| GWF:Larval_SubstrateDPL | 0.22035 | 0.44202 | 0.499 | 0.618119 |
| GWF:Larval_SubstrateDPL+LPL | -0.39108 | 0.89598 | -0.436 | 0.662482 |
| GWF:Larval_SubstrateFRT | 0.18605 | 0.60748 | 0.306 | 0.759399 |
| GWF:Larval_SubstrateLPL | -0.27952 | 0.55944 | -0.500 | 0.617328 |
| GWF:Larval_SubstrateSOI | 0.43674 | 0.53395 | 0.818 | 0.413389 |
| GWF:Larval_SubstrateSOI+DNG | 3.97834 | 5.75542 | 0.691 | 0.489419 |
| GWF:Larval_SubstrateSOI+DPL | 0.45544 | 0.51396 | 0.886 | 0.375545 |
| GWF:Larval_SubstrateSOI+LPL | 14.82859 | 27.40944 | 0.541 | 0.588505 |
| MUL:DietCOP | -2.03319 | 1.68319 | -1.208 | 0.227071 |
| MUL:DietCOP+DET | -0.22095 | 0.50858 | -0.434 | 0.663972 |
| MUL:DietDET | -0.50396 | 0.58200 | -0.866 | 0.386543 |
| MUL:DietMYC | -0.83004 | 0.68723 | -1.208 | 0.227125 |
| MUL:DietMYC+DET | -0.86097 | 0.53676 | -1.604 | 0.108710 |
| MUL:DietPHY | -0.66907 | 0.66528 | -1.006 | 0.314564 |
| MUL:DietPOL | -0.40323 | 0.64728 | -0.623 | 0.533305 |
| MUL:DietPRD | -1.02980 | 0.58898 | -1.748 | 0.080385 |
| MUL:WingsDIM | 0.02008 | 0.54137 | 0.037 | 0.970411 |
| MUL:WingsMAC | -0.10613 | 0.25730 | -0.412 | 0.680004 |
| MUL:Body_Size | 0.06020 | 0.09100 | 0.662 | 0.508257 |
| MUL:Larval_SubstrateDPL | 0.63392 | 0.44533 | 1.423 | 0.154600 |
| MUL:Larval_SubstrateDPL+LPL | -0.25298 | 1.09712 | -0.231 | 0.817634 |
| MUL:Larval_SubstrateFRT | 0.59162 | 0.63574 | 0.931 | 0.352059 |
| MUL:Larval_SubstrateLPL | 0.23416 | 0.58896 | 0.398 | 0.690941 |
| MUL:Larval_SubstrateSOI | 0.65540 | 0.54696 | 1.198 | 0.230819 |
| MUL:Larval_SubstrateSOI+DNG | 1.03856 | 4.97461 | 0.209 | 0.834627 |
| MUL:Larval_SubstrateSOI+DPL | 0.69266 | 0.52339 | 1.323 | 0.185698 |
| MUL:Larval_SubstrateSOI+LPL | 8.91151 | 15.48969 | 0.575 | 0.565076 |
| BGT:DietCOP | 2.88157 | 1.12290 | 2.566 | 0.010282 * |
| BGT:DietCOP+DET | 2.21027 | 0.63681 | 3.471 | 0.000519 *** |
| BGT:DietDET | 2.28648 | 0.70160 | 3.259 | 0.001118 ** |
| BGT:DietMYC | 0.36539 | 0.74127 | 0.493 | 0.622068 |
| BGT:DietMYC+DET | 0.97228 | 0.67523 | 1.440 | 0.149891 |
| BGT:DietPHY | 1.96137 | 0.75233 | 2.607 | 0.009132 ** |

| | | | | | |
|-----------------------------|----------|----------|--------|----------|-----|
| BGT:DietPOL | 2.22391 | 0.72877 | 3.052 | 0.002276 | ** |
| BGT:DietPRD | 1.60175 | 0.70818 | 2.262 | 0.023711 | * |
| BGT:wingsDIM | 0.95869 | 0.48811 | 1.964 | 0.049521 | * |
| BGT:wingsMAC | -0.13479 | 0.25858 | -0.521 | 0.602172 | |
| BGT:Body_Size | -0.60399 | 0.11814 | -5.113 | 3.18e-07 | *** |
| BGT:Larval_SubstrateDPL | -0.83257 | 0.60986 | -1.365 | 0.172195 | |
| BGT:Larval_SubstrateDPL+LPL | 1.72582 | 2.55919 | 0.674 | 0.500080 | |
| BGT:Larval_SubstrateFRT | -0.27011 | 0.76122 | -0.355 | 0.722711 | |
| BGT:Larval_SubstrateLPL | -1.53681 | 0.69586 | -2.208 | 0.027210 | * |
| BGT:Larval_SubstratesOI | -1.02461 | 0.66254 | -1.546 | 0.121985 | |
| BGT:Larval_SubstratesOI+DNG | 27.48474 | 24.94146 | 1.102 | 0.270475 | |
| BGT:Larval_SubstratesOI+DPL | -1.14035 | 0.67311 | -1.694 | 0.090237 | . |
| BGT:Larval_SubstratesOI+LPL | -2.37777 | 4.53057 | -0.525 | 0.599703 | |
| VWC:DietCOP | -0.75037 | 0.89218 | -0.841 | 0.400323 | |
| VWC:DietCOP+DET | -0.12114 | 0.39288 | -0.308 | 0.757824 | |
| VWC:DietDET | -1.06204 | 0.46712 | -2.274 | 0.022992 | * |
| VWC:DietMYC | -0.20719 | 0.55276 | -0.375 | 0.707794 | |
| VWC:DietMYC+DET | -0.45443 | 0.42776 | -1.062 | 0.288078 | |
| VWC:DietPHY | -0.58371 | 0.54299 | -1.075 | 0.282382 | |
| VWC:DietPOL | -0.91952 | 0.52544 | -1.750 | 0.080117 | . |
| VWC:DietPRD | -0.65925 | 0.47071 | -1.401 | 0.161353 | |
| VWC:wingsDIM | 0.65323 | 0.45327 | 1.441 | 0.149546 | |
| VWC:wingsMAC | 0.17122 | 0.22591 | 0.758 | 0.448492 | |
| VWC:Body_Size | 0.24012 | 0.07658 | 3.135 | 0.001716 | ** |
| VWC:Larval_SubstrateDPL | -0.02012 | 0.33699 | -0.060 | 0.952394 | |
| VWC:Larval_SubstrateDPL+LPL | -0.29134 | 1.08685 | -0.268 | 0.788656 | |
| VWC:Larval_SubstrateFRT | -0.77251 | 0.64278 | -1.202 | 0.229430 | |
| VWC:Larval_SubstrateLPL | 0.18609 | 0.46606 | 0.399 | 0.689676 | |
| VWC:Larval_SubstratesOI | 0.26434 | 0.43277 | 0.611 | 0.541326 | |
| VWC:Larval_SubstratesOI+DNG | 1.68010 | 3.09078 | 0.544 | 0.586728 | |
| VWC:Larval_SubstratesOI+DPL | 0.53391 | 0.40688 | 1.312 | 0.189450 | |
| VWC:Larval_SubstratesOI+LPL | -0.95516 | 4.33049 | -0.221 | 0.825430 | |

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

Model selection for the FRic response variable. The GLMMs applied here follow the same structure as those used for the richness and abundance models. An AIC comparison was used to evaluate which of the listed models best predicted the data. Season was included as a random effect. The numerical covariates are scaled to represent a fair comparison to the effect size of the fixed factors.

| AICc | dAICc | df | weight |
|------------|--------|-------|-----------|
| BD_GC2 | -547.2 | 0.0 | 13 0.435 |
| OM_GC2 | -547.1 | 0.1 | 13 0.417 |
| OM_GC_soil | -544.5 | 2.7 | 22 0.110 |
| BD_GC_soil | -540.5 | 6.8 | 21 0.015 |
| BD_GC3 | -540.0 | 7.2 | 19 0.012 |
| OM_GC3 | -540.0 | 7.3 | 19 0.012 |
| BD_GC1 | -513.5 | 33.7 | 14 <0.001 |
| OM_GC1 | -512.2 | 35.0 | 14 <0.001 |
| OM_full | -504.7 | 42.6 | 40 <0.001 |
| BD_full | -499.6 | 47.7 | 39 <0.001 |
| BD_chem | -484.1 | 63.1 | 8 <0.001 |
| OM_chem | -480.7 | 66.5 | 7 <0.001 |
| OM_veg | -470.6 | 76.7 | 27 <0.001 |
| OM_phys | -469.5 | 77.7 | 6 <0.001 |
| BD_veg | -469.2 | 78.0 | 27 <0.001 |
| BD_weather | -467.8 | 79.4 | 8 <0.001 |
| OM_weather | -464.1 | 83.1 | 8 <0.001 |
| null | -439.5 | 107.7 | 2 <0.00 |

```

> summary(BD_GC2)
Family: beta ( logit )
Formula:
FRic ~ SBD + fR + fMUL + SBT + SVWC + I(SVWC^2) + (1 | fSEASON)
Data: Sunbridge

```



```
AIC      BIC    logLik deviance df.resid
-548.8   -503.3    287.4   -574.8     232
```

Random effects:

Conditional model:

```
Groups Name      Variance Std.Dev.
fSEASON (Intercept) 1.256e-10 1.121e-05
Number of obs: 245, groups: fSEASON, 3
```

Dispersion parameter for beta family (:): 17.1

Conditional model:

```
Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.35495    0.07307 -18.543 < 2e-16 ***
sBD          -0.02012    0.04153  -0.485  0.62797
fR1          -0.15492    0.08115  -1.909  0.05625 .
fMUL2        -0.16845    0.11515  -1.463  0.14350
fMUL3        -0.34164    0.12894  -2.650  0.00806 **
fMUL4        -0.34911    0.12135  -2.877  0.00402 **
fMUL5        -0.53252    0.21171  -2.515  0.01189 *
fMUL6        -1.62044    0.24592  -6.589  4.42e-11 ***
sBGT         0.30596    0.04753   6.438  1.21e-10 ***
svWC         0.05553    0.06611   0.840  0.40096
I(svWC^2)    -0.08864    0.03080  -2.878  0.00400 **
---
```

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

```
> r.squaredGLMM(BD_GC2)
```

```
      R2m      R2c
[1,] 0.513175 0.513175
```

Model selection for the FEve response variable. The GLMMs applied here follow the same structure as those used for the richness and abundance models. An AIC comparison was used to evaluate which of the listed models best predicted the data. Season was included as a random effect. The numerical covariates are scaled to represent a fair comparison to the effect size of the fixed factors.

```
AICc  dAICc  df weight
BD_weather -259.2    0.0 8  0.3413
OM_weather -258.8    0.4 8  0.2760
OM_chem    -257.6    1.5 7  0.1579
BD_chem    -256.1    3.1 8  0.0739
OM_GC2     -255.2    4.0 13 0.0469
BD_GC2     -254.9    4.2 13 0.0410
OM_GC1     -253.2    6.0 14 0.0170
BD_GC1     -252.8    6.3 14 0.0143
OM_GC_soil -252.5    6.7 22 0.0118
BD_GC_soil -251.2    8.0 21 0.0064
OM_GC3     -250.9    8.3 19 0.0054
OM_phys    -250.4    8.8 6  0.0042
BD_GC3     -250.1    9.0 19 0.0037
OM_veg     -232.4   26.8 27 <0.001
BD_veg     -232.0   27.2 27 <0.001
OM_full    -231.2   28.0 40 <0.001
BD_full    -228.8   30.4 39 <0.001
null       -224.0   35.2 2  <0.001
```

```
> summary(BD_weather)
```

```
Family: beta ( logit )
```

```
Formula: FEve ~ sBD + fR + SPRCP + STMIN + STMAX + (1 | fSEASON)
```

```
Data: Sunbridge
```

```
AIC      BIC    logLik deviance df.resid
-259.8   -231.8    137.9   -275.8     237
```

Random effects:

Conditional model:

```
Groups Name      Variance Std.Dev.
fSEASON (Intercept) 4.752e-11 6.894e-06
Number of obs: 245, groups: fSEASON, 3
```

Dispersion parameter for beta family (): 9.76

Conditional model:

| | Estimate | Std. Error | z value | Pr(> z) |
|-------------|-----------|------------|---------|------------|
| (Intercept) | 0.756102 | 0.061776 | 12.239 | <2e-16 *** |
| sBD | 0.026613 | 0.040704 | 0.654 | 0.5132 |
| fR1 | -0.000594 | 0.082594 | -0.007 | 0.9943 |
| sPRCP | -0.154561 | 0.065534 | -2.358 | 0.0184 * |
| stMIN | -0.193747 | 0.095716 | -2.024 | 0.0430 * |
| stMAX | 0.024615 | 0.097364 | 0.253 | 0.8004 |

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

```
> r.squaredGLMM(BD_weather)
```

```
      R2m      R2c  
[1,] 0.6784597 0.6784597
```

Model selection for the FDiv response variable. The GLMMs applied here follow the same structure as those used for the richness and abundance models. An AIC comparison was used to evaluate which of the listed models best predicted the data. Season was included as a random effect. The numerical covariates are scaled to represent a fair comparison to the effect size of the fixed factors.

| | AICc | dAICc | df | weight |
|------------|--------|-------|----|--------|
| OM_weather | -495.4 | 0.0 | 8 | 0.4831 |
| BD_weather | -495.2 | 0.3 | 8 | 0.4245 |
| OM_chem | -491.0 | 4.5 | 7 | 0.0518 |
| BD_chem | -488.8 | 6.6 | 8 | 0.0178 |
| BD_GC2 | -487.6 | 7.8 | 13 | 0.0096 |
| OM_GC2 | -487.2 | 8.2 | 13 | 0.0080 |
| OM_phys | -485.7 | 9.8 | 6 | 0.0036 |
| BD_GC_soil | -482.9 | 12.5 | 21 | <0.001 |
| OM_GC_soil | -481.5 | 13.9 | 22 | <0.001 |
| BD_GC3 | -478.8 | 16.6 | 19 | <0.001 |
| OM_GC3 | -478.5 | 16.9 | 19 | <0.001 |
| BD_GC1 | -477.1 | 18.4 | 14 | <0.001 |
| OM_GC1 | -476.7 | 18.8 | 14 | <0.001 |
| BD_full | -470.5 | 25.0 | 39 | <0.001 |
| OM_full | -467.7 | 27.8 | 40 | <0.001 |
| OM_veg | -464.6 | 30.8 | 27 | <0.001 |
| BD_veg | -463.1 | 32.3 | 27 | <0.001 |
| null | -417.2 | 78.2 | 2 | <0.001 |

```
> summary(OM_weather)
```

Family: beta (logit)

Formula: FDiv ~ SOM + fR + sPRCP + stMIN + stMAX + (1 | fSEASON)

Data: Sunbridge

| | AIC | BIC | logLik | deviance | df.resid |
|--|--------|--------|--------|----------|----------|
| | -496.1 | -468.0 | 256.0 | -512.1 | 237 |

Random effects:

Conditional model:

| Groups | Name | Variance | Std.Dev. |
|---------|-------------|-----------|-----------|
| fSEASON | (Intercept) | 2.701e-10 | 1.644e-05 |

Number of obs: 245, groups: fSEASON, 3

Dispersion parameter for beta family (): 14.7

Conditional model:

| | Estimate | Std. Error | z value | Pr(> z) |
|-------------|----------|------------|---------|--------------|
| (Intercept) | 1.54100 | 0.06144 | 25.081 | < 2e-16 *** |
| SOM | -0.05011 | 0.04055 | -1.236 | 0.216527 |
| fR1 | -0.06434 | 0.08171 | -0.787 | 0.431029 |
| sPRCP | 0.26126 | 0.06612 | 3.951 | 7.77e-05 *** |
| stMIN | -0.17719 | 0.08813 | -2.010 | 0.044379 * |
| stMAX | 0.32014 | 0.09167 | 3.492 | 0.000479 *** |

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

```
> r.squaredGLMM(OM_weather)
```

```
      R2m      R2c  
[1,] 0.922343 0.922343
```