| group |  | Estimate | Standard Error | df | statistic | p-value |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| Fixed effects | | | | | | | |
|  | (Intercept) | 5.772 | 0.240 | 11 | 24.008 | 0.0000 | \*\*\* |
|  | Colony\_fEI | -0.368 | 0.442 | 5 | -0.832 | 0.4417 |  |
|  | Colony\_fDI | -0.913 | 0.256 | 161 | -3.562 | 0.0005 | \*\*\* |
|  | Colony\_fNI | -0.591 | 0.155 | 112 | -3.820 | 0.0002 | \*\*\* |
|  | MonsoonSE | -0.294 | 0.166 | 69 | -1.778 | 0.0798 | . |
|  | SexM | -0.500 | 0.140 | 184 | -3.563 | 0.0005 | \*\*\* |
|  | Sexunknown | 0.042 | 0.159 | 176 | 0.266 | 0.7902 |  |
|  | Breed\_Stageincubation | 0.204 | 0.156 | 216 | 1.310 | 0.1916 |  |
|  | Breed\_Stagepre-egg | 0.227 | 0.308 | 184 | 0.736 | 0.4629 |  |
|  | Breed\_Stagebreeding | -0.275 | 0.487 | 279 | -0.565 | 0.5723 |  |
| Random effects | | | | | | | |
| BirdID | sd\_\_(Intercept) | 0.296 |  |  |  |  |  |
| year\_f | sd\_\_(Intercept) | 0.316 |  |  |  |  |  |
| Residual | sd\_\_Observation | 0.966 |  |  |  |  |  |
| Signif. codes: 0 <= '\*\*\*' < 0.001 < '\*\*' < 0.01 < '\*' < 0.05 | | | | | | | |
|  | | | | | | | |
| square root of the estimated residual variance: 1.0 | | | | | | | |
| data's log-likelihood under the model: -698.3 | | | | | | | |
| Akaike Information Criterion: 1,422.6 | | | | | | | |
| Bayesian Information Criterion: 1,477.0 | | | | | | | |