

# **BIRDS BIODIVERSITY TEMPORAL TRENDS ANALYSIS**

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Technical Report

This report analyses 12 years of bird monitoring data from Martinique (2014-2025) to quantify temporal trends in biodiversity indicators. The dataset comes from a standardised breeding bird monitoring program that follows a consistent protocol: observers conduct 5-minute point counts at 10 fixed observation points along each transect, with sites visited at most twice during the monitoring period. This structured approach ensures comparability across sites and years, though sampling effort varies both temporally and spatially, a factor we'll need to account for in trend analyses.

Our analysis focuses on three complementary goals: first, to thoroughly understand the dataset's structure and quality, second, to quantify multi-year trends in key biodiversity indicators with proper uncertainty estimates, and third, to examine species-level patterns for a subset of ecologically interesting species.

## **1. DATASET FAMILIARISATION AND DESCRIPTIVE ANALYSIS**

### **1.1 Dataset Structure and Scope**

The monitoring data comes from an Excel workbook with three interconnected sheets. The main observation table (NOM FRANÇAIS) contains 114,497 records spanning 2014 to 2025, with each row representing a single point visit. After cleaning (removing 4 records with zero counts and handling data quality issues), we retained 114,493 observations across 102 unique bird species, 72 transects, and 42 observers.

The species lookup table (ESPECES) provides taxonomy and migration status for 86 species, leaving 16 observed species without metadata. The GPS table (GPS-MILIEU) maps 651 observation points to habitat types and coordinates, though we didn't use precise GPS data for this analysis. Instead, we focused on transect-level spatial patterns, which proved sufficient for understanding coverage variation.

The temporal coverage spans from March 29, 2014 to July 3, 2025. However, the 2014 data only has 5,377 observations. Complete years (2015-2025) show much more consistent effort, ranging from 8,162 to 11,010 observations annually throughout the year. We decided to keep the partial year in our exploratory analyses but flag them explicitly in visualisations, since they still provide useful context even if we can't directly compare them to full years.

One key aspect of the protocol: each transect consists of exactly 10 observation points where observers spend 5 minutes recording all birds detected. This standardisation is helpful because it means we can normalise counts by dividing by the number of points surveyed, giving us a consistent metric even when sampling intensity varies.

### **1.2 Critical Data Quality Issues**

Before diving into any analysis, we had to tackle some serious data quality problems. We found 5 observations with negative wind values, which is obviously impossible. These appear to be data entry errors, so we set those wind values to NaN while keeping the bird observations. Only 5 records out of 114,000+ isn't a huge deal, but it's the kind of thing we think that can break statistical models.

The distance sampling data columns were mostly empty, which makes sense. Most observers just recorded total counts rather than breaking them down by distance bands. So we chose to ignore any analysis requiring distance-specific information and just stick with total counts per point.

There's also a significant observer effect that we need to keep in mind. One person, CONDE Beatriz, conducted 36.5% of all observations and surveyed 46 of the 72 transects (63.9% of spatial coverage). That's a huge concentration of effort, and it means observer-specific biases (differences in detection skill, preferred sites, recording habits) could influence our results. For any trend modeling, we'll probably want to include observer as a random effect or at minimum check whether patterns hold up when excluding this single observer.

### **1.3 Environmental Conditions Summary**

Weather conditions were recorded for nearly all observations, which is good news since weather can affect detection rates. The data shows that surveys generally occurred under favorable conditions:

- Wind: recorded for 99.98% of observations, with most surveys (75th percentile) occurring at wind level 2 or below on a 0-4 scale. The median was 1, suggesting calm to light breeze conditions.
- Rain: 97% of surveys occurred without rain (coded as 1 = no rain). Only 3% had rain present during observations.
- Cloud cover: recorded for 99.99% of observations, with a median of 2 on a 1-3 scale. Most surveys had partial cloud coverage rather than completely overcast or completely clear skies.
- Visibility: recorded for 99.99% of observations, median of 1, indicating generally good visibility conditions.

The consistency here suggests the monitoring program followed a protocol of only conducting surveys under suitable weather conditions, which is standard practice. Since weather conditions don't vary dramatically and are nearly complete, we probably don't need to worry too much about weather as a confounding factor in temporal trends. Though if we were being really thorough, we could include it as a covariate in models.

### **1.4 Missing Values Analysis**

The missing data pattern tells a clear story about what this dataset can and can't support. Distance bands (columns Unnamed: 13-20) are 65-99% empty, confirming what I mentioned earlier, distance sampling analysis isn't something we want to do analysis with. The original distance category column is also 48.6% missing.

But here's the important part in our opinion: all essential columns are complete. The eight core fields we need for biodiversity trend analysis (observer name, transect name, species name, date, year, individual count, point number, and visit number) have zero missing values. That's exactly what we need for our temporal modeling.

The detection method columns (auditory, visual, audio-visual combinations) have minimal missing data (0.0-0.2%). Instead of a single categorical "detection\_type" column, counts are split across multiple columns by detection method and then summed to get total individuals. We didn't focus on this for our analysis, but it's nice to know the information is there if someone wants to explore detection patterns later. We chose to just treat all detections equally for our analysis, looking at total count

### **1.5 Outlier Detection**

Using the standard IQR method ( $Q3 + 1.5 \times IQR$ ), we identified 3,364 observations (2.94%) as statistical outliers, counts exceeding 18 individuals per observation. The distribution of bird counts is heavily right-skewed, with a median of 6 birds per observation but a maximum of 600. That maximum isn't actually a mistake as we initially thought: it comes from surveys of tern colonies (*Sterna de Dougall*, *Sterna fuligineuse*) at marine sites in 2025 and 2018-2022, where large flocks are ecologically normal.

Should we remove these outliers? We argue not. The high counts aren't data errors, they represent real biological phenomena (colonial seabirds, flocking behavior, potentially habitat-specific aggregations). Dropping them would bias our diversity and abundance estimates downward. Instead, we'll keep them but use statistical methods that are robust to skewness, or log-transform counts if we need to meet model assumptions. The key is being aware they exist and interpreting results appropriately.

That said, if a specific analysis is sensitive to extreme values, we can always do sensitivity checks by comparing results with and without outliers above some threshold.

### **1.6 Temporal Patterns and Sampling Effort**

The observation effort varies considerably over time, and this matters for trend analysis. The partial year 2014 had just 5,377 observations, then effort jumped to 8,162 in 2015 and peaked at 11,010 in 2016. From 2017 onwards, effort stabilized between roughly 9,200 and 10,800 observations per year. The coefficient of variation for temporal effort is 15.9%, which isn't trivial.

What this means: we can't just compare raw species counts or richness across years and call it a trend. A year with more observations will naturally detect more species, even if true biodiversity hasn't changed. We'll need to either standardise by effort (e.g., species per observation, counts per point) or use statistical models that explicitly account for varying sampling intensity.

Interestingly, the number of active observers dropped from 24-28 in early years down to 13-15 in 2020-2022, then rebounded to 24-26 in 2024-2025. But total observation effort didn't drop proportionally, probably because the remaining observers (especially that one super-active observer) compensated by increasing their individual workload.

### **1.7 Spatial Coverage Variation**

Spatial coverage, the percentage of all 72 transects surveyed in a given year, ranges from 56.9% in 2014 to 90.3% in 2018. This does make sense with 2014 being a partial year. No year achieved complete coverage. The average across all years is 83.9%, meaning we're typically missing about 16% of transects each year.

The spatial coefficient of variation is 15.9%, matching the temporal variation. This means that not only does effort vary over time, but which specific transects get surveyed changes from year to year. This is actually more problematic than simple temporal variation in effort, because different transects have different habitat types and species compositions.

For example, if we surveyed mostly forest transects in one year and mostly agricultural transects the next, observed changes in species composition might just reflect sampling location rather than real biodiversity shifts. Any trend analysis needs to account for this, either by including transect as a fixed/random effect in models, or by focusing on trends within transects that were consistently surveyed.

The fact that the best year only reached 90.3% coverage suggests there are practical constraints on surveying all sites (accessibility, time limitations, observer availability). This incomplete coverage is something we'll need to explicitly acknowledge as a limitation in our final synthesis.

## **1.8 Species Composition and Dominance**

The bird community is dominated by a small number of common species. The top 20 species (out of 102 total) account for 90.5% of all individual birds recorded, 676,289 out of 747,639 total. The single most abundant species, Quiscale merle (Greater Antillean Grackle), makes up 15.3% of all observations with 114,356 individuals. The top 5 species alone account for more than half of all detections.

This extreme dominance is typical of ecological communities but has implications for analysis. Common species will drive aggregate metrics like total abundance and diversity indices. Rare species contribute to richness but not much to abundance-based measures. If we want to understand the fate of less common species, we'll need to look at them individually rather than relying solely on community-level indices.

The species accumulation curve shows how the monitoring program has built up its species list over time. Starting with 63 species in 2014, the curve rises rapidly to 94 species by 2018, then flattens out dramatically. From 2018 to 2024, only 6 new species were added. The 2025 year detected 2 new bringing the cumulative total to 102.

This pattern suggests the sampling has largely saturated, we've detected most of the species that regularly occur in the monitored habitats. The slowing accumulation rate is good news for the monitoring program because it means we're unlikely to miss major biodiversity shifts just due to detection failures. New species additions at this point probably represent either genuine colonizers, rare vagrants, or occasionally detected species at the edge of their range.

## **1.9 Observer Effort Distribution**

The observer effort distribution is remarkably skewed. CONDE Beatriz alone contributed 41,779 observations (36.5%) and surveyed 46 unique transects (63.9% of all transects). The second-highest observer contributed only 8,913 observations (7.8%). The top 10 observers account for 75.6% of all data.

This concentration raises important methodological questions. Observer identity could be confounded with year, site, or season if particular observers consistently worked at particular times and places. Detection rates might vary between observers due to differences in identification skills, hearing ability, or diligence in recording all detections. If that dominant observer has systematic biases, those biases will heavily influence our dataset-wide patterns.

For robust trend analysis, we should probably do a few things: (1) include observer as a random effect in mixed models to account for individual-level variation; (2) check whether temporal trends hold up when we exclude the top observer or restrict to only observers who surveyed across multiple years; (3) test whether species accumulation patterns differ between high-effort and low-effort observers as a proxy for detection skill.

The fact that one observer covered nearly two-thirds of all transects is both good and bad. Good because it provides consistency in methodology and reduces inter-observer variability at those sites. Bad because it means that observer's specific biases (whatever they might be) have outsized influence on spatial patterns.

### **Synthesis and Recommendations for Data Familiarisation and Descriptive Statistics**

- **Key Insights from Exploratory Analysis:**

The monitoring dataset is fundamentally sound for temporal trend analysis, but comes with important caveats. The core observation data, species identity, counts, locations, dates, is complete and reliable after correcting the datetime encoding issue. The standardised protocol (10 points per transect, 5 minutes per point) provides a consistent framework for comparing observations across space and time.

However, three major sources of variation will constrain how we interpret downstream results:

First, sampling effort varies substantially both temporally (15.9% CV) and spatially (average 16% of transects missed each year). This isn't fatal, we can standardise metrics or model effort explicitly, but it means raw counts or richness values aren't directly comparable across years.

Second, spatial coverage is incomplete and varies, meaning different habitat types and species pools get sampled in different years. Trends could potentially reflect sampling artifact rather than real biological change if we're not careful about controlling for site-level variation.

Third, observer identity is heavily skewed, with one individual contributing over a third of all data. Observer effects are notoriously difficult to disentangle from real patterns in ecological monitoring, so any trend that relies heavily on this single observer's data needs to be interpreted cautiously.

On the plus side, species accumulation has largely saturated, suggesting we're detecting the vast majority of regularly occurring species. The weather data shows surveys occurred under suitable conditions. And the outlier analysis reveals that extreme counts are biologically plausible rather than data errors.

- **Data Quality Limitations:**

The 5 negative wind values and 4 zero-count records were minor issues easily handled by setting values to NaN or removing records.

The missing species metadata (16 species without scientific names or migration status) is annoying but not crippling. We can still analyze those species; we just can't link them to external databases or assign ecological traits. Future data collection should close this gap.

Distance sampling data is too sparse to use (65-99% missing). This limits our ability to account for detection probability varying with distance from the observer, which is a standard concern in point count surveys. We're essentially assuming perfect detection within the survey area, which is unrealistic. But given the data we have, that's the best we can do.

The most serious limitation is the incomplete spatial coverage combined with observer concentration. We can't definitively separate "CONDE Beatriz's sites changed" from "the whole island changed" without additional controls. Subsequent analyses should include sensitivity checks removing this observer or restricting to transects with multi-observer coverage.

- Recommendations for Future Monitoring:

1. Aim for complete spatial coverage (100% of transects annually). If resource constraints prevent this, at least ensure a consistent core set of transects gets surveyed every year to enable within-site trend analysis.
2. Balance observer effort more evenly. Heavy reliance on a single observer creates vulnerability (what if they leave?) and makes observer effects hard to disentangle from real trends. Consider paired surveys where multiple observers sample the same transect to quantify and correct for observer effects.
3. Continue recording weather conditions, but also consider adding effort predictors like start time, survey duration, and completion status (did the observer finish all 10 points?). These could help model detection probability more explicitly.
4. Close the species metadata gap. Get scientific names and trait information for the 16 species currently missing from the lookup table.
5. If feasible, revisit the distance sampling protocol. Either train observers to consistently record distance bands (and enforce it) or drop the requirement and acknowledge we're just doing unlimited-distance point counts. The current approach gives us empty columns that create confusion without adding information.
6. For trend detection, focus on complete years (2015-2025) as the primary analysis window, using 2014 only for context. This avoids biases from partial year coverage.

- Final Thoughts

This dataset is far from perfect, the datetime encoding issue, incomplete coverage, observer concentration, and partial years all create analytical challenges. But it's still a valuable resource for quantifying biodiversity trends in Martinique's breeding birds. The key is being explicit about limitations and using statistical methods that are robust to the messiness of real-world monitoring data.

The exploratory analysis here gives us confidence that the core data are reliable and sufficiently rich to support indicator-based trend analysis and species-level investigation. With appropriate caution about observer effects and effort variation, we should be able to produce meaningful inferences about how this bird community has changed over the past decade.

## **2. Indicator Trends**

### **2.1 Indicators Utilized**

Each indicator captures a unique facet of biodiversity change. To maintain interpretability and scientific rigor, we computed and analyzed each using established ecological formulas and visualization techniques.

#### **2.1.1 Species Richness**

Species richness was defined as the number of unique species observed within a given year across all sites. This metric provides a direct measure of species turnover and potential local extinction or colonization events. Richness was computed annually, and a simple linear or generalized additive model (GAM) was fitted to visualize multi-year trends.

### **2.1.2 Shannon Diversity**

The Shannon Index ( $H'$ ) integrates both the number and relative abundance of species:

$$H' = -\sum_i p_i \ln(p_i)$$

is the proportion of individuals belonging to species  $i$ . This indicator reflects how evenly individuals are distributed among species — a stable or increasing Shannon value suggests balanced community composition.

### **2.1.3 Total Abundance**

Total abundance was measured as the sum of all individual bird detections recorded per year. This provides a coarse measure of total population density across the monitored sites. Since sampling effort varies, trends were interpreted with caution, focusing on relative year-to-year shifts rather than absolute counts.

### **2.1.4 Spatial Coverage**

Spatial coverage quantifies how many distinct sites recorded bird activity each year, offering a proxy for distributional range or occupancy. It is particularly informative for detecting shifts in geographic spread or habitat use.

Each indicator was visualized as a time series, and smoothed trend lines were fitted to identify consistent temporal patterns. Model diagnostics were later used to assess residual variability and uncertainty across years.

## **2.2 Results and Interpretation**

The four biodiversity indicators reveal nuanced but interpretable temporal patterns in Martinique's breeding bird community between 2014 and 2025 (Figure 2.1). While individual indicators vary in direction and statistical significance, together they suggest a modest restructuring of bird communities over the 12-year period.

For consistency across all indicator analyses, statistical significance was evaluated based on the  $p$ -value of the fitted linear trend slope. Trends with  $p < 0.05$  were considered statistically significant, indicating strong evidence for a directional change over time. Trends with  $0.05 \leq p < 0.10$  were treated as marginal, reflecting suggestive but inconclusive evidence of change. Results with  $p \geq 0.10$  were classified as not significant, implying that observed variation is likely attributable to interannual noise rather than a consistent long-term trend.

### 2.2.1 Species Richness

Species richness shows a slight but consistent decline over the study period (slope =  $-0.60$  species  $\cdot$  year $^{-1}$ ,  $R^2 = 0.39$ ,  $p = 0.0535$ ). The downward slope suggests that the number of species detected each year has decreased by roughly 7–8% since 2014. Although the trend is only marginally significant, it indicates potential attrition in local species diversity — possibly due to habitat degradation or reduced detection probabilities in later years.

The 95% confidence interval widens toward 2025, reflecting greater interannual variability in recent surveys.

### 2.2.2 Shannon Diversity

In contrast, Shannon diversity remains effectively stable across the same period (slope =  $+0.0007$   $\cdot$  year $^{-1}$ ,  $R^2 = 0.01$ ,  $p = 0.7826$ ). Despite minor fluctuations, the absence of a clear directional trend implies that community evenness has remained relatively constant, even as richness declined slightly. This stability suggests that dominant species maintained similar relative abundances, and that community composition may be restructuring without major losses in evenness.

### 2.2.3 Total Abundance

Total abundance exhibits a weak increasing trend (slope =  $+1,195$  individuals  $\cdot$  year $^{-1}$ ,  $R^2 = 0.36$ ,  $p = 0.0684$ ). Although not statistically significant at the 0.05 level, this pattern hints at overall increases in the number of individuals recorded per year, possibly driven by a few abundant generalist species. The upward shift in total counts despite declining richness supports the hypothesis of community homogenization, where fewer species dominate a larger share of detections.

### 2.2.4 Spatial Coverage

Spatial coverage displays the strongest and most significant trend among the indicators (slope =  $+0.92$  sites  $\cdot$  year $^{-1}$ ,  $R^2 = 0.56$ ,  $p = 0.0125$ ). This increase in the number of occupied transects suggests a broader spatial distribution of bird detections, potentially reflecting range expansions or improved detectability over time. The positive coverage trend contrasts with the modest decline in richness, implying that birds are being observed across more sites, but with fewer species represented overall.

### 2.2.5 Summary of Indicator Trends

Indicator	Direction	R <sup>2</sup>	p-value	Significance	Ecological Interpretation
Species Richness	↓	0.390	0.0535	Marginal	Fewer species detected annually; possible biodiversity loss
Shannon Diversity	→	0.010	0.7826	Not significant	Stable community evenness



Total Abundance	↑	0.357	0.0684	Marginal	Rising total counts; possible dominance by common species
Spatial Coverage	↑	0.562	0.0125	Significant	More sites occupied; broader distribution

### 2.2.6 Integrated Interpretation

Collectively, these results suggest a decoupling between species composition and spatial occupancy. While bird detections are spreading spatially (increased coverage), richness is trending downward and abundance upward, indicating that communities may be becoming less diverse but more widespread and numerically dominated by a few adaptable species.

This pattern is consistent with regional ecological shifts such as habitat simplification or expansion of generalist species, both of which can maintain high overall abundance while reducing diversity at the community level. The moderate  $R^2$  values across models (0.36–0.56 for three indicators) suggest that these patterns are biologically meaningful but still influenced by residual variation, likely due to sampling effort differences and environmental fluctuations across years.

### 2.2.7 Model Diagnostics and Residual Analysis

To ensure that linear trend assumptions were valid for each indicator, residual distributions were evaluated visually using Q–Q plots and statistically using the Shapiro–Wilk test for normality. In all cases, residuals exhibited approximately normal distributions ( $p \geq 0.05$ ), indicating that linear regression provided an appropriate and unbiased fit for temporal trend estimation.

Indicator	Residual Mean	Residual SD	Shapiro–Wilk W	Shapiro p	Normality Conclusion
Species Richness	2.8422e-15	2.1541	0.913	0.301	Normal
Shannon Diversity	−4.4409e-17	0.0190	0.957	0.755	Normal
Total Abundance	9.4587e-12	4612.6554	0.954	0.712	Normal
Spatial Coverage	2.1316e-15	2.3196	0.940	0.552	Normal

All four models passed the Shapiro–Wilk test ( $p \geq 0.05$ ), suggesting no major deviations from normality in residuals. This supports the robustness of the linear trend estimates reported above. Slightly wider residual variance in Total Abundance and Species Richness models reflects the higher natural variability of those indicators, consistent with their moderate  $R^2$  values.

## 2.3 Interpretation & Synthesis

To assess coherence among biodiversity indicators and potential methodological influences, we examined relationships among annual species richness, Shannon diversity, total abundance, and spatial coverage between 2015 and 2024.

### **2.3.1 Richness vs Shannon Diversity**

The weak negative correlation ( $r = -0.19$ ,  $p = 0.60$ ) indicates that variation in species richness was largely independent of community evenness. This suggests that years with more species did not necessarily exhibit higher evenness, possibly reflecting the dominance of a few abundant species.

### **2.3.2 Abundance and Biodiversity Indicators**

Relationships between total abundance and both richness ( $r = -0.12$ ,  $p = 0.74$ ) and Shannon diversity ( $r = 0.09$ ,  $p = 0.80$ ) were very weak and non-significant. This implies that changes in total individual counts were not strongly coupled with diversity patterns, suggesting population growth in common species rather than broad community expansion.

### **2.3.3 Sampling Effects (Spatial Coverage)**

Spatial coverage was strongly correlated with total abundance ( $r = 0.57$ ,  $p = 0.086$ ) and moderately negatively correlated with Shannon diversity ( $r = -0.48$ ,  $p = 0.16$ ). The abundance–coverage relationship approaches marginal significance, indicating that years with higher sampling coverage tended to yield higher total counts—a likely methodological rather than ecological effect. However, richness did not covary significantly with coverage ( $r = -0.26$ ,  $p = 0.48$ ), suggesting that interannual richness variation reflects genuine ecological signal rather than sampling bias.

### **2.3.4 Temporal Patterns and Indicator Coherence**

When standardized, the indicators show moderate synchrony through time: abundance and spatial coverage rise steadily, while richness exhibits a mild downward trend (slope =  $-0.60$ ,  $p = 0.053$ ). Shannon diversity remains stable. Together, these patterns indicate stable community structure despite modest declines in the number of recorded species.

### **2.3.5 Ecological Synthesis**

Overall, the data suggest a stable but slightly homogenizing community: species richness shows a weak decline, but evenness and total abundance remain stable or increase. The weak correlation among biodiversity indicators further supports that community dynamics are decoupled from sampling effort in most years. The positive abundance–coverage link, however, highlights the need to adjust for survey effort in subsequent species-level trend models.

## **3. Species-Level Population Dynamics**

### **3.1.1 Strategic Species Selection**

The analysis began with a systematic identification of ecologically informative species through multiple complementary criteria:

Abundance-based selection: Identification of the most frequently observed species to ensure statistical reliability

Variability assessment: Focus on species exhibiting high inter-annual variation, potentially indicating sensitivity to environmental changes

Consistency evaluation: Inclusion of species with stable presence across the study period to establish baseline patterns

### **3.1.2 Robust Trend Analysis with Uncertainty Quantification**

#### **Annual Population Estimation:**

- Aggregated individual counts by year, treating missing years as true zeros (accounting for detection absence)
- Maintained consistent temporal coverage (2015-2024) for comparative analysis

#### **Statistical Trend Modeling:**

- Applied ordinary least squares regression to quantify directional changes
- Incorporated significance testing ( $p < 0.05$  threshold) to distinguish meaningful trends from random variation
- Implemented data quality checks (minimum 3 non-zero years for trend estimation)

### **3.1.3 Visual Communication of Complex Patterns**

#### **Dynamic Plot Configuration:**

- Implemented adaptive subplot arrangements based on species count
- Maintained consistent scaling for cross-species comparisons
- Included confidence intervals as shaded regions to communicate estimation uncertainty

#### **Informative Annotation:**

- Displayed key metrics (total counts, presence years) directly on plots
- Highlighted statistically significant trends with dashed trend lines
- Provided p-values for transparent statistical reporting

### **3.1.4 Contextualized Species Assessment**

#### **Metadata Integration from ESPECES Database:**

- Scientific nomenclature: Linked common French names to standardized scientific classifications for taxonomic precision
- Conservation status: Incorporated species-specific designations (Endemic, Resident, Migratory, Introduced) to inform expectation setting
- Regional ecological knowledge: Applied Martinique-specific habitat associations and behavioral patterns

#### **Structured Interpretation Framework:**

- Abundance classification: Categorized species as common, moderate (20-100), or rare based on detection frequency
- Statistical significance hierarchy: Distinguished strong evidence from suggestive patterns for nuanced interpretation
- Caribbean-specific drivers: Considered regional factors including hurricane impacts, invasive species dynamics, and habitat fragmentation

#### **Limitations-Aware Reporting:**

- Data quality flags: Automatically identified species with limited detection history or low abundance
- Statistical power assessment: Highlighted cases where trend detection was constrained by sample size or variability
- Contextual caveats: Differentiated methodological limitations from ecological uncertainties

### **3.1.5 Results Synthesis and Export**

The analytical outputs were systematically compiled to support both scientific reporting and conservation applications:

#### **Comprehensive Metrics Compilation:**

- Temporal coverage: Years of detection relative to study period
- Population metrics: Total abundance, mean annual counts, and inter-annual variability (coefficient of variation)
- Trend statistics: Slope coefficients with associated p-values for directional change assessment

### **3.1.6 Presence-Absence Dynamics Analysis**

Complementing abundance-based trends, we implemented occurrence probability modeling to assess distributional changes:

#### **Methodological Approach:**

- Binary occurrence encoding: Transformed count data to presence-absence matrices for each species-year combination
- Logistic regression framework: Modeled detection probability as function of time to assess range expansion/contraction
- Robust error handling: Implemented exception management for sparse detection histories

#### **Analytical Advantages:**

- Reduced zero-inflation bias: Less sensitive to variation in count magnitude than abundance models
- Occupancy trend detection: Identified changes in geographic or temporal distribution independent of population density
- Complementary perspective: Provided insights into species persistence patterns beyond abundance fluctuations

#### **3.1.7 Community-Level Contextualization**

To situate individual species trends within broader ecological patterns, we conducted comparative analyses between species-specific trajectories and community-wide richness dynamics:

##### **Community Richness Benchmarking:**

- Annual richness calculation: Quantified unique species counts per year across all monitoring sites
- Linear trend modeling: Applied ordinary least squares regression to assess directional changes in community composition
- Comparative framework: Established community trend as reference point for interpreting individual species patterns

##### **Pattern Classification System:**

- Concordant trends: Species showing directional changes aligned with community patterns (both increasing or both decreasing)
- Divergent trends: Species exhibiting trajectories opposite to community directionality
- Neutral patterns: Species with stable trends amidst changing community composition

##### **Ecological Interpretation Framework:**

- Contributor identification: Species with positive trends during community richness increases likely represent successful adapters or beneficiaries of environmental conditions
- Sentinel value: Species declining during community stability may indicate specific vulnerabilities or habitat specialists at risk
- Compensatory dynamics: Divergent trends may reflect ecological replacement or niche partitioning within the avian community

### **3.1.8 Habitat-Specific Population Dynamics**

#### **Habitat Classification Integration:**

- GPS-MILIEU database linkage: Merged observational records with habitat type classifications from spatial metadata
- Habitat-specific trend analysis: Computed annual abundance patterns within distinct habitat categories
- Visualization of niche dynamics: Plotted parallel trajectories to identify habitat preferences and shifts

#### **Analytical Implementation:**

- Data validation: Verified habitat classification completeness and consistency across transects
- Missing data management: Implemented graceful degradation when habitat metadata was incomplete
- Output optimization: Generated publication-ready multi-panel figures for each target species

#### **Conservation Applications:**

- Habitat association strength: Identification of species with strong habitat specialization versus generalist strategies
- Refugia identification: Detection of habitats supporting stable populations amid broader declines
- Management prioritization: Guidance for habitat-specific conservation interventions based on species-habitat relationships

### **3.1.9 Observer Detection Bias Assessment**

#### **Detection Rate Quantification:**

- Observer effort normalization: Calculated detection rates relative to total observation effort per observer
- Quality filtering: Applied minimum observation thresholds ( $\geq 100$  records) to ensure statistical reliability
- Performance ranking: Identified observers with consistently high detection rates for target species

#### **Methodological Quality Control:**

- Benchmark establishment: Provided reference detection rates for data quality assessment
- Expert observer identification: Flagged consistently high-performing observers for potential validation roles

- Bias awareness: Documented inter-observer variability to contextualize abundance estimates

### **3.1.10 Multi-Metric Visualization Suite**

#### **Dual-Metric Plotting Framework:**

- Absolute abundance tracking: Primary y-axis displays raw count data for population trajectory assessment
- Relative abundance context: Secondary y-axis shows proportional representation within annual community totals
- Temporal synchronization: Maintained consistent annual scaling across both metrics for direct comparison

#### **Visual Design Optimization:**

- Publication standards: High-resolution output (300 DPI) with professional formatting
- Accessibility features: Clear differentiation of data series through line styles and colors
- Informative annotation: Integrated statistical summaries directly within plot titles

#### **Analytical Value-Added:**

- Dominance assessment: Species maintaining consistent proportional representation despite community fluctuations indicate ecological stability
- Contextual interpretation: Absolute increases with declining proportions may reflect general community growth rather than specific success
- Conservation prioritization: Species showing both absolute and relative declines represent highest conservation concern

## **3.2 Synthesis and Recommendations**

### **3.2.1 Key Pattern Synthesis**

Our species-level analysis revealed three distinct population trajectories: stable residents maintaining consistent presence, notable decliners showing concerning decreases, and successful adapters demonstrating population growth. These patterns operated somewhat independently, with species showing individualistic responses rather than synchronized community-wide trends. Observer detection rates varied significantly, indicating that some species require specialized identification skills while others are reliably detected across observers.

### **3.2.2 Methodological Insights**

The 10-year monitoring period proved adequate for detecting trends in common species but lacked statistical power for rare species with fewer than 50 observations. Habitat-specific analyses revealed that generalist species showed consistent patterns across environments, while specialists exhibited habitat-dependent trajectories. The integration of spatial metadata significantly enhanced our ability to interpret population changes within ecological context.

### **3.2.3 Data Quality Assessment**

We identified several data quality considerations: observer expertise varied substantially, affecting detection rates for challenging species; certain habitats appeared underrepresented in sampling; and rare species data remained insufficient for robust trend analysis. The treatment of missing years as true absences proved reasonable for most species but may overestimate declines for cryptic or rarely detected taxa.

### **3.2.4 Recommended Monitoring Improvements**

We recommend implementing targeted protocols for rare species monitoring, standardized observer training for consistent detection, and improved habitat stratification in sampling design. Future analyses would benefit from incorporating climate covariates and land-use change data to better understand environmental drivers. Multi-model inference approaches could enhance trend estimation reliability.

## **4. Conclusion**

From a statistical perspective, this analysis demonstrates that the linear modeling framework applied to the Martinique bird monitoring dataset provides consistent, interpretable, and diagnostically sound results. All four biodiversity indicators exhibited residuals that met normality assumptions, confirming the suitability of linear trend estimation over the 2015–2024 period. Although only spatial coverage displayed a statistically significant trend ( $p = 0.0125$ ), both species richness ( $p = 0.0535$ ) and total abundance ( $p = 0.0684$ ) showed marginal evidence for directional change, while Shannon diversity remained stable. Moderate  $R^2$  values (0.36–0.56 for key indicators) indicate that temporal variation is partially but not fully captured by linear models, reflecting underlying biological and sampling variability. Correlation analyses further revealed weak interdependence among indicators, validating their treatment as distinct dimensions of community structure. Overall, the statistical workflow—comprising data cleaning, normalization, regression modeling, residual diagnostics, and cross-indicator correlation analysis—proved robust, transparent, and reproducible, providing a solid quantitative foundation for subsequent ecological interpretation and future modeling extensions.