Calculate normalized averages from trap data for each county subdivision and week

Integrate data from each trap type into a single estimate of *Aedes aegypti* density

Estimate relative risk of dengue

Stage 1: Calculate Summer Maximum Abundance by Trap Type

Stage 2: Calculate Relative Abundance as Proportion of Summer Maximum per Trap Type (m₁, m₂)

Stage 3: Integrate Weighted Abundances per Trap Type (m)

Stage 4: Estimate Dengue Relative Risk

Calculate Mosquito Abundance

- Include CO₂-baited and Biogents (BG)
 Sentinel trap collections per week during
 Aug & Sep from the last 5 years
- Include only county subdivisions with ≥ 100 collections during Aug-Sep of the 5year period
- Divide total mosquitoes by total trap-nights per trap type

Log-Transform Abundance Data

 Apply logarithmic transformation [log(mosquitoes/trap-night+1)] per trap collection to stabilize data & reduce impact of outliers

Calculate the geometric mean of mosquito abundance for each subcounty and week for "summer" (Aug-Sep)

- Calculate averages of log-transformed *Ae*. *aegypti* per trap-night for each subcounty and week for which trapping data are available during the 5-year period
- Average weekly numbers for all available weeks over the 5-year period to get a universal average for "summer" for each county subdivision
- Back-transform the summer average of logtransformed abundance to revert it to original scale
- Extract maximum value for the backtransformed summer average per trap type across all county subdivisions to serve as an upper limit representing "high" mosquito abundance

Output: Maximum summer statistics per trap type (m_{S1}, m_{S2})

Calculate Abundance use CO₂-baited and BG Sentinel Trap Collections

- Include all weeks with trap data for each county subdivision
- Divide total mosquitoes by total trapnights per trap type

Log-Transform Abundance

 Apply logarithmic transformation [log(mosquitoes/trap-night+1)] per trap collection to stabilize data & reduce impact of outliers

Calculate 4-Week Moving Average

- Calculate temporal moving average of log abundance
- Incorporate current week and previous 3 weeks to smooth out fluctuations and highlight trends

Calculate Geometric Mean with Weighted Approach

- Calculate spatially weighted average with 2/3 weight assigned to each county subdivision and 1/3 weight to neighboring county subdivisions

Normalize Spatially Weighted Average

- Back-transform the weekly average of logtransformed abundance to revert it to original scale
- Divide back-transformed weekly values (m_{W1}, m_{W2}) by maximum summer abundance per trap type (m_{S1}, m_{S2})
 - $m_1 = m_{W1} / m_{S1}$
 - $m_2 = m_{W2}/m_{S2}$
- Limit m_1 and m_2 to [0,1] range

Output: Normalized Abundance per Trap Type (m_1, m_2)

Calculate Average of m₁ and m₂ Weighted by Trapping Effort for Each Trap Type

 $m^* = (m_1 \times \text{proportion of CO}_2\text{-baited trap$ $nights}) + (m_2 \times \text{proportion BG Sentinel trap-nights})$

Scale m* to a realistic limit for *Aedes*aegypti Density

- $m = m^* \times 5$
- Scaled to constrain density within [0,5] interval, which assumes an upper limit of 5 mosquitoes per human as the highest density across all county subdivisions and weeks

Estimate Survival Rate (p)

Function of temperatures: minimum (tmin), maximum (tmax), and mean (tmean) temperatures

Estimate Biting Rate (a)

- Function of mean temperature (tmean)

Estimate Extrinsic Incubation Period (n)

- Function of mean temperature (tmean)

Estimate Relative Risk using the Formula

Relative Risk =
$$\frac{ma^2bcp^2}{-ln(p)r}$$

- m: Mosquito density (mosquitoes per human)
- a: Biting rate (bites per mosquito per day)
- b: Transmission efficiency (mosquito → human)
- c: Transmission efficiency (human → mosquito)
- p: Mosquito survival (proportion surviving per day)
- n: Extrinsic incubation period (# of days for virus to become infectious in mosquitoes)
- r: Human recovery rate from dengue infection (1/infectious period)

Final Output: Estimated Relative Risk of Dengue

Temperature-dependent parameters