

Population range estimation with animal tracking data

Christen H. Fleming

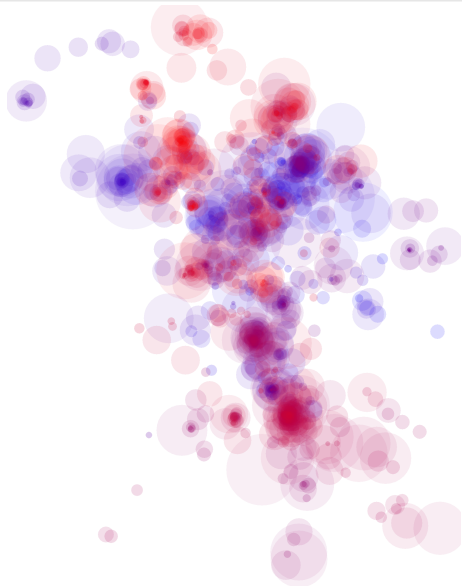
`christen.fleming@ucf.edu`



2023-11-08

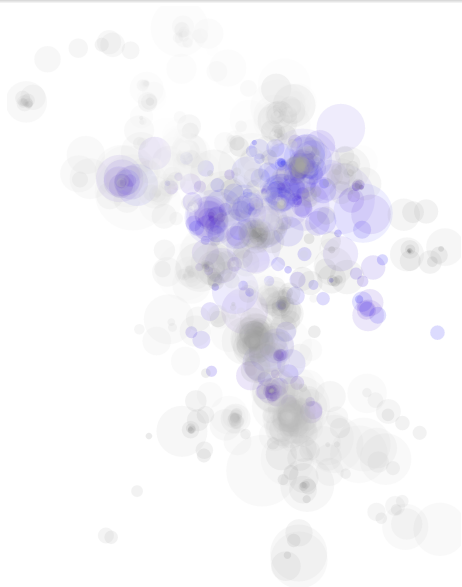
Overview

- Introduction
- Kernel density estimation
- Validation
- Discussion



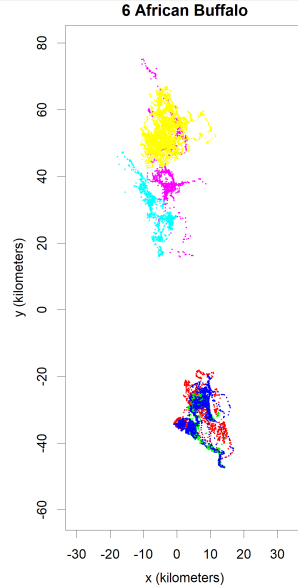
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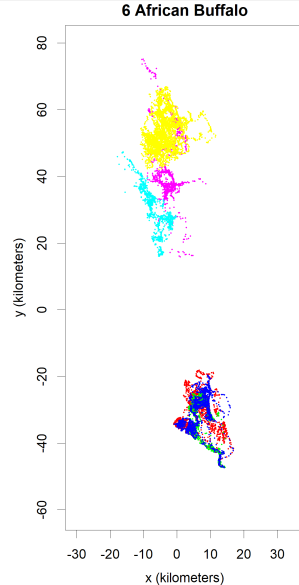
Introduction

- Tracks sampled from populations:
What's the population distribution?



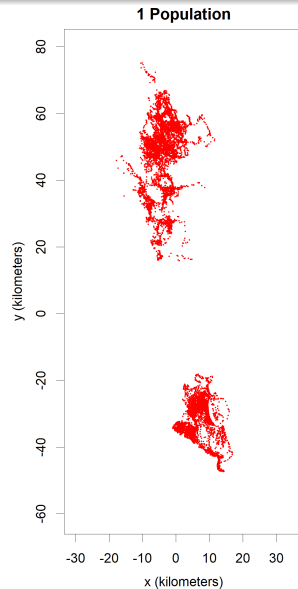
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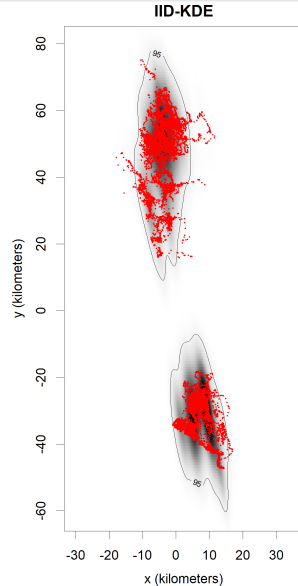
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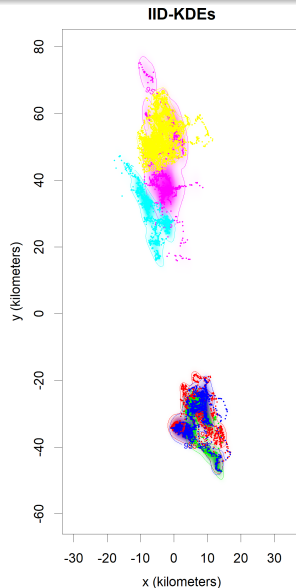
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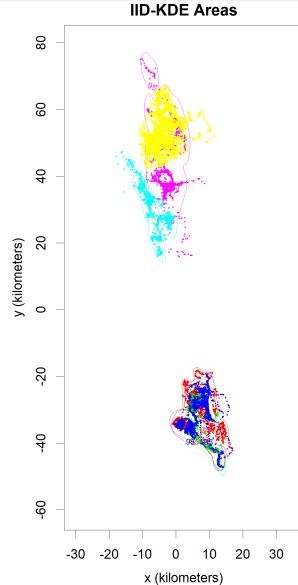
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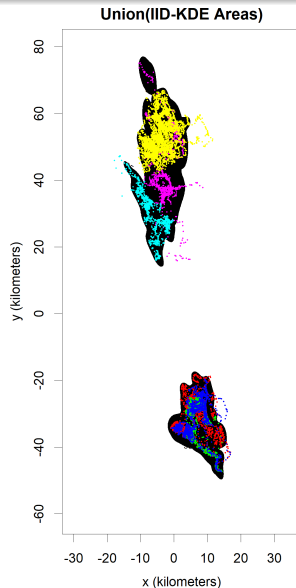
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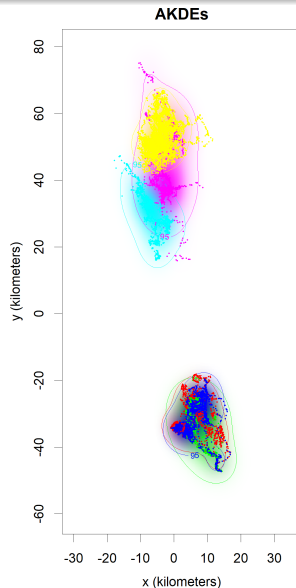
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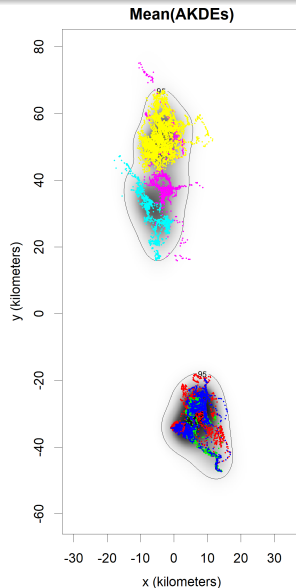
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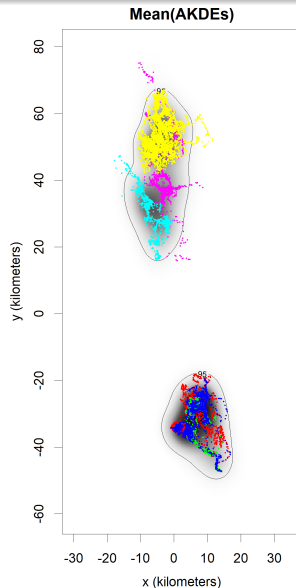
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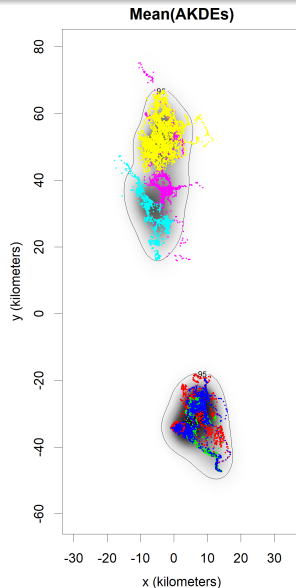
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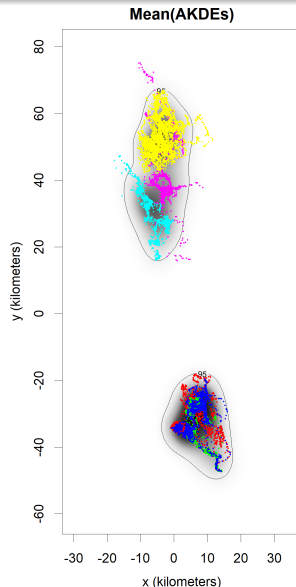
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- Sample size vs. saturation

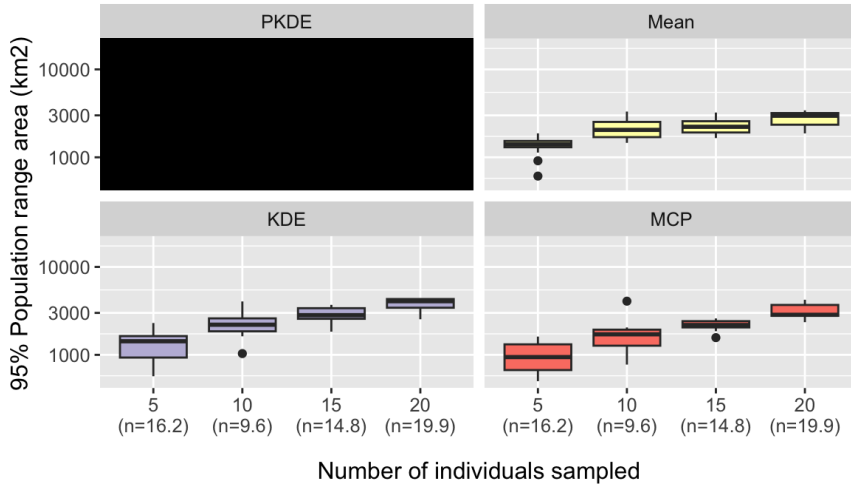


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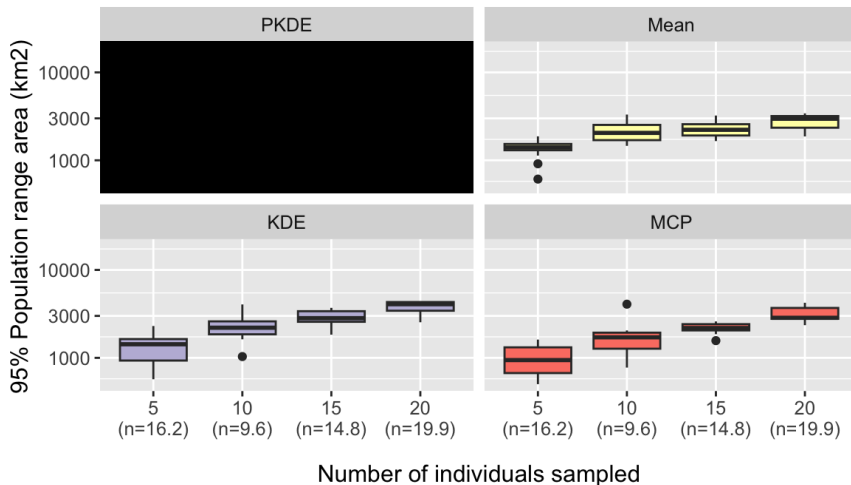
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- Sample size vs. saturation
 - Extrapolation of asymptote



U. a. horribilis Saturation Curve

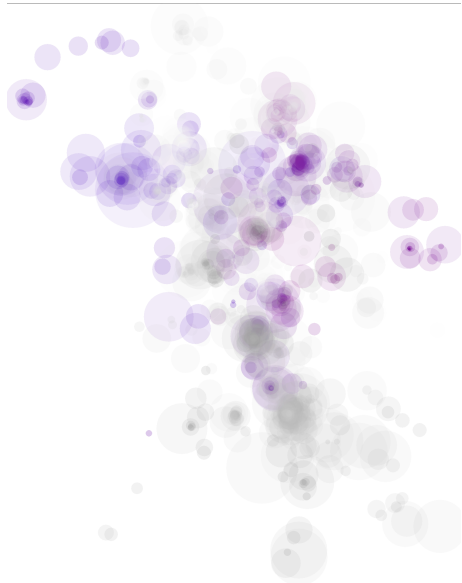


U. a. horribilis Saturation Curve



This is bias from not modeling population variance

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Kernel Kernel Density Estimation

A KDE is a $\hat{p}(\mathbf{x}|\mathbf{H})$:

$$\hat{p}(\mathbf{x}|\mathbf{H}) =$$

$\hat{p}(\cdot)$: probability density estimate

\mathbf{x} : location of interest

Kernel Kernel Density Estimation

A KDE is a weighted average of :

$$\hat{p}(\mathbf{x}|\mathbf{H}) = \sum_t^n w(t)$$

$\hat{p}(\cdot)$: probability density estimate

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$w(t)$: weight at time t

Kernel Kernel Density Estimation

A KDE is a weighted average of kernels:

$$\hat{p}(\mathbf{x}|\mathbf{H}) = \sum_t^n w(t) \kappa(\mathbf{x} - \mathbf{x}(t)|\mathbf{H})$$

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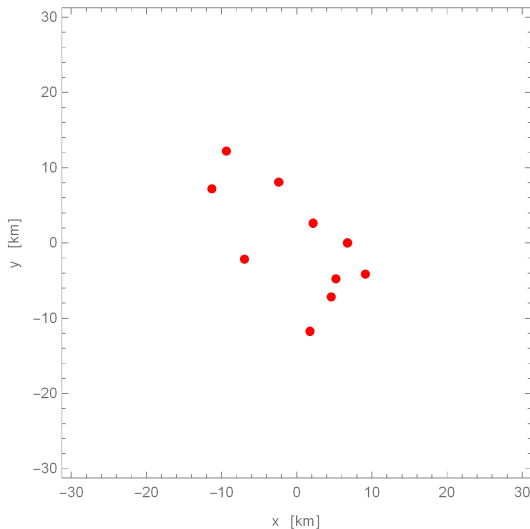
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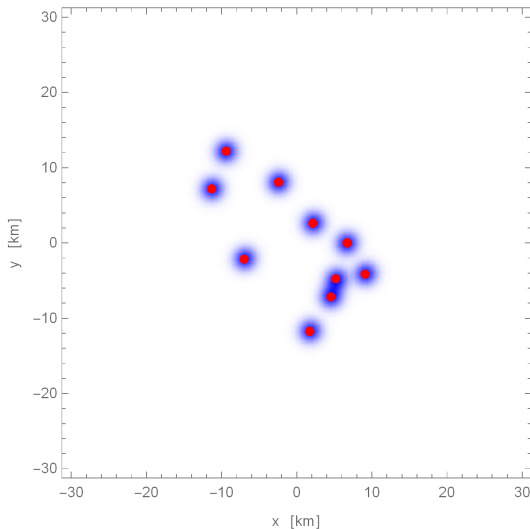
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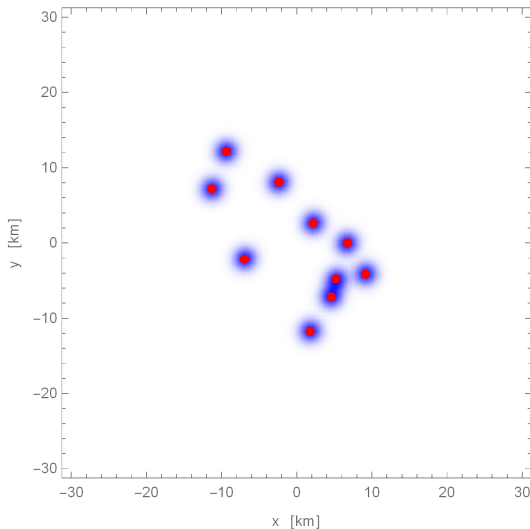
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Where the optimal \mathbf{H} minimizes the MISE:



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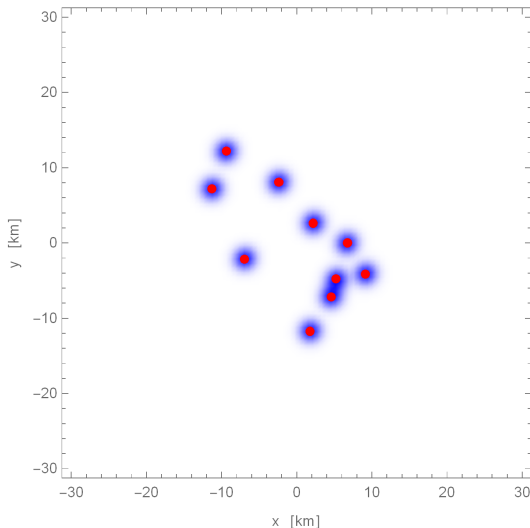
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$$\mathbf{M} \left[\begin{array}{c} \cdot \\ \cdot \\ \cdot \end{array} \right] = \mathbf{E} \left[\begin{array}{c} \cdot \\ \cdot \\ \cdot \end{array} \right]$$



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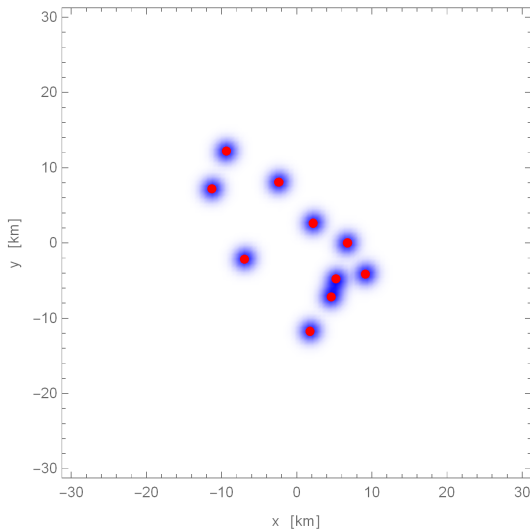
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$$\text{MISE}[\mathbf{H}] = \mathbb{E} \left[\iint \right] d\mathbf{x}$$



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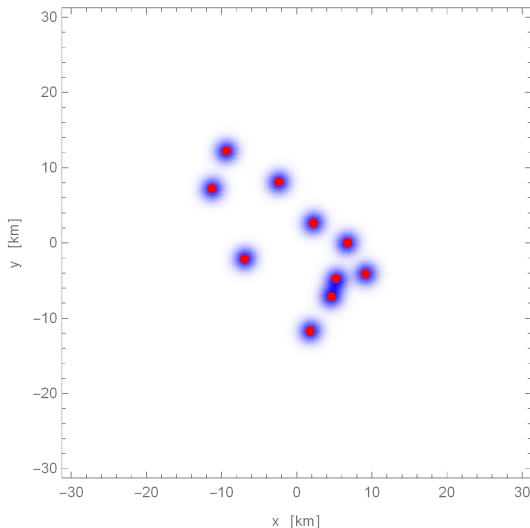
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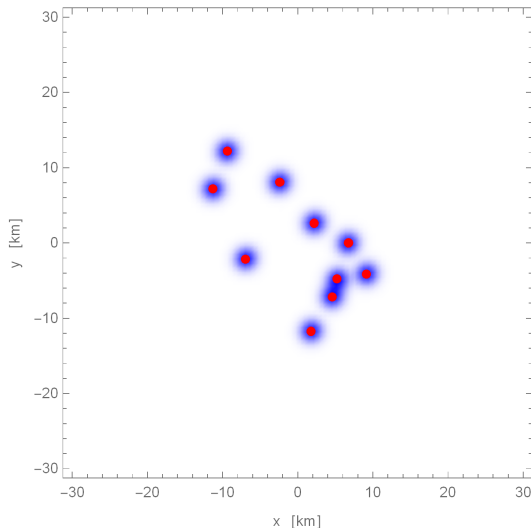
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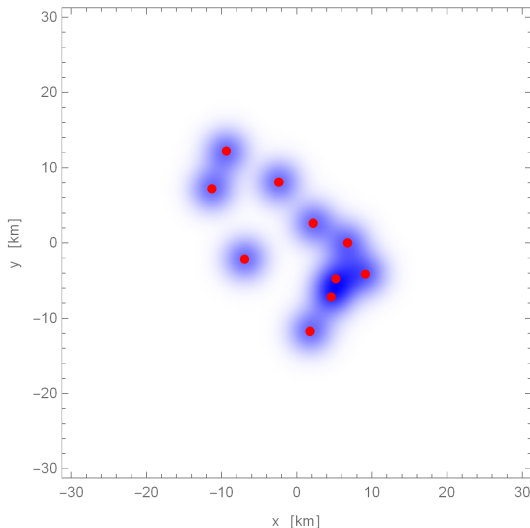
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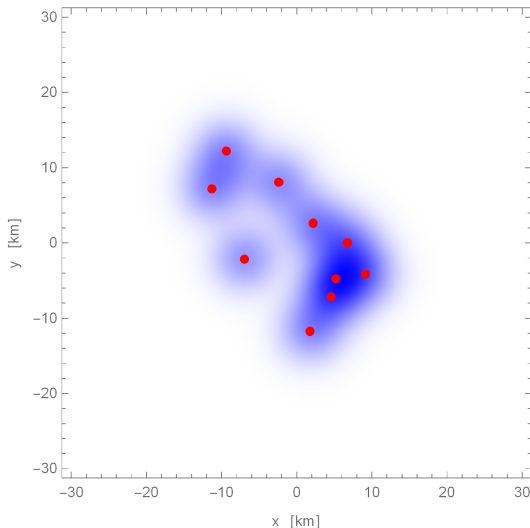
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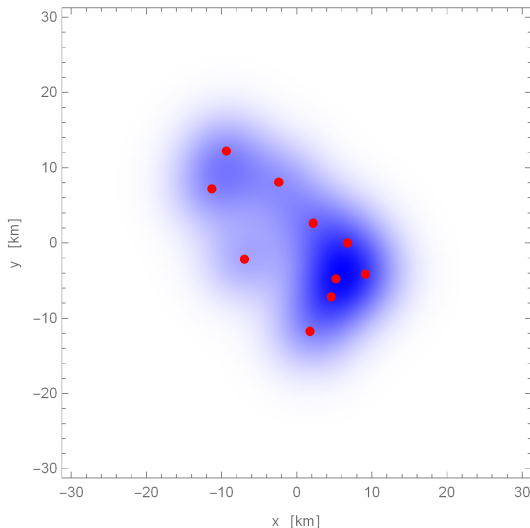
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$p(\mathbf{x})$ = approximation (e.g., Gaussian reference function)

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- bandwidth optimization (Turlach 1993)
- AKDE (Fleming, Fagan, et al. 2015)

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- bandwidth optimization (Turlach 1993)
- AKDE (Fleming, Fagan, et al. 2015)
- bias-corrected AKDEc (Fleming and Calabrese 2017)
- (MISE) optimally weighted wAKDE (Fleming, Sheldon, et al. 2018)

$$\text{MISE}[\mathbf{H}] = \mathbb{E} \left[\iint (\hat{p}_{\text{pop.}}(\mathbf{x}|\mathbf{H}) - p_{\text{pop.}}(\mathbf{x}))^2 d\mathbf{x} \right]$$

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 - Proportional to individual: $\mathbf{H}_{\text{ind.}} = h^2 \text{COV}[\mathbf{x}_{\text{ind.}}]$

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- Bandwidth assumptions:
 - Proportional to individual: $\mathbf{H}_{\text{ind.}} = h^2 \text{COV}[\mathbf{x}_{\text{ind.}}]$ (fast)

Population kernel density estimation

$$\text{MISE}[\mathbf{H}] = \mathbb{E} \left[\iint (\hat{p}_{\text{pop.}}(\mathbf{x}|\mathbf{H}) - p_{\text{pop.}}(\mathbf{x}))^2 d\mathbf{x} \right]$$
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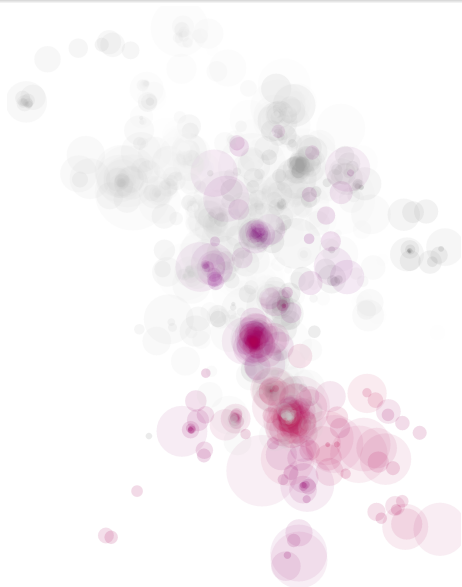
- $p_{\text{pop.}}(\mathbf{x})$ requires a *hierarchical* approximation
- Individual weight assumptions:
 - Uniform: $\sum_t w_{\text{ind.}}(t) = \frac{1}{\sum_{\text{ind.}}}$
 - MISE-optimized $\sum_t w_{\text{ind.}}(t)$ (tends to zero-weight overlapped individuals)
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 - Proportional to individual: $\mathbf{H}_{\text{ind.}} = h^2 \text{COV}[\mathbf{x}_{\text{ind.}}]$ (fast)
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$$\text{MISE}[\mathbf{H}] = \mathbb{E} \left[\iint (\hat{p}_{\text{pop.}}(\mathbf{x}|\mathbf{H}) - p_{\text{pop.}}(\mathbf{x}))^2 d\mathbf{x} \right]$$
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Transition

- Introduction
- Kernel density estimation
- Validation
- Discussion



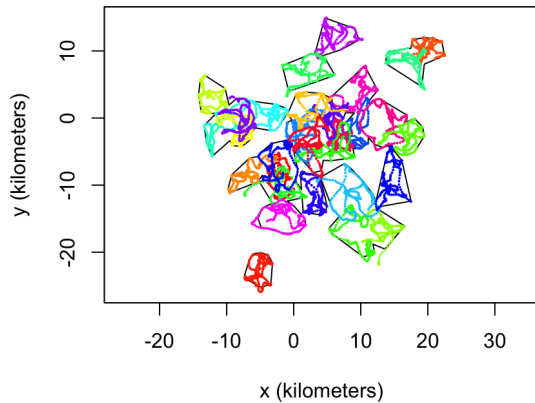
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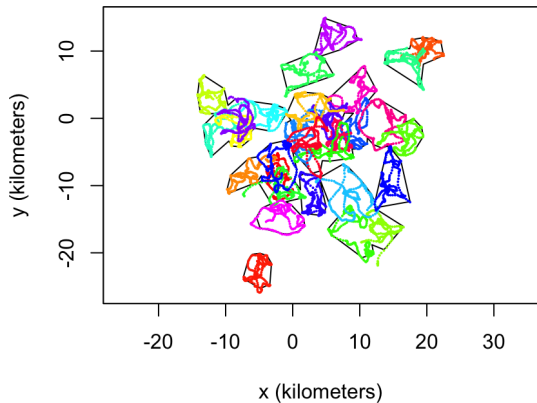
Gayatri Anand (UMD Ph.D.)

Merged MCP estimate

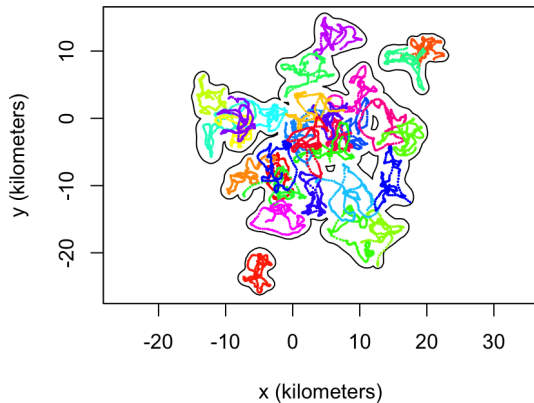


Validation: Simulations

Merged MCP estimate

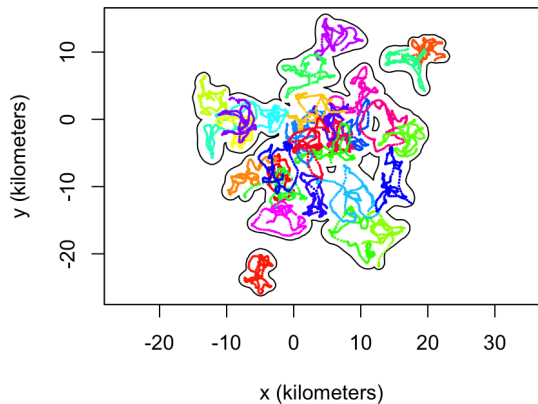


Merged KDE estimate

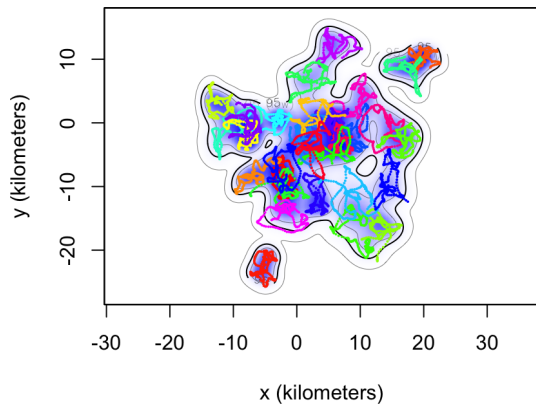


Validation: Simulations

Merged KDE estimate

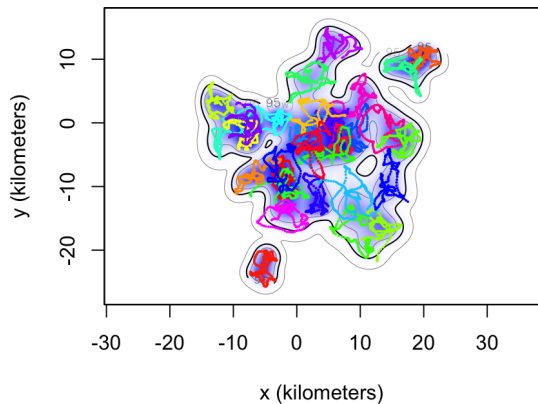


Mean AKDE estimate

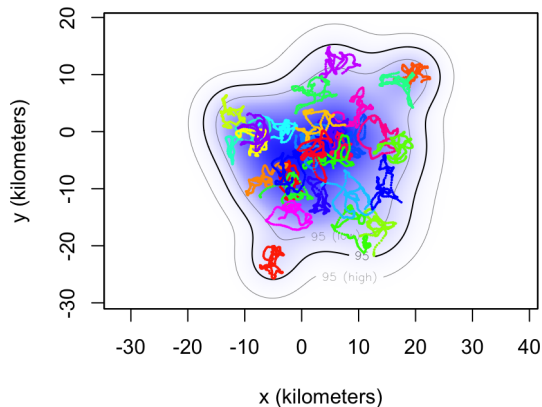


Validation: Simulations

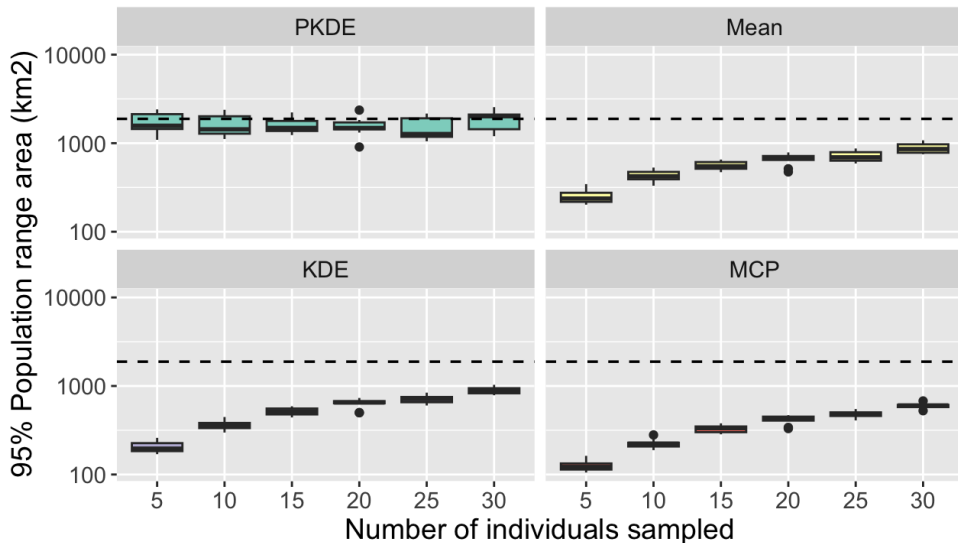
Mean AKDE estimate



PKDE estimate

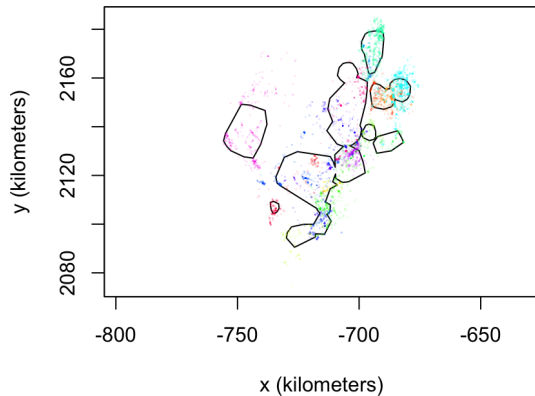


Saturation Curve for Simulated Data



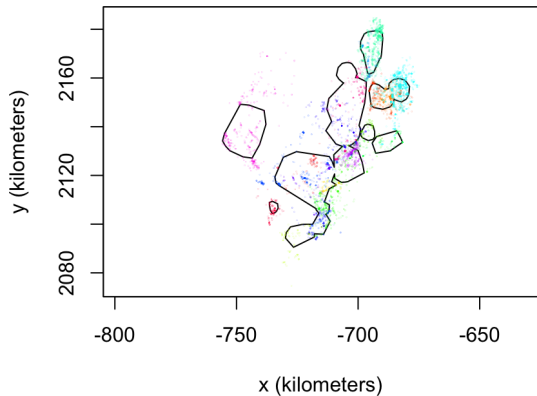
Validation: Empirical (*Ursus arctos horribilis*)

Merged MCP estimate

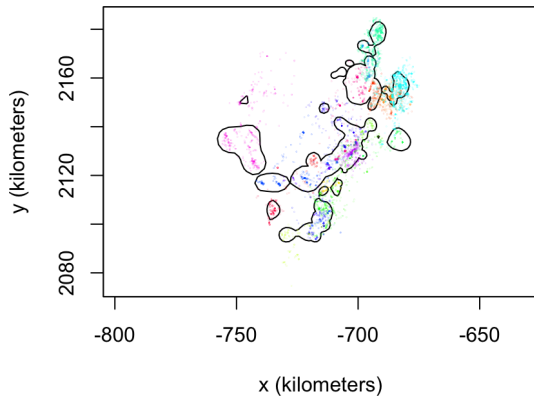


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Merged MCP estimate

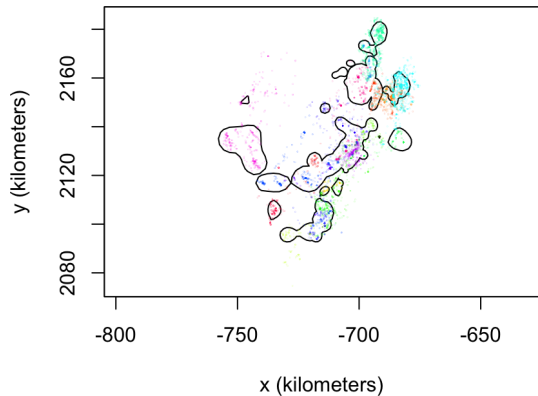


Merged KDE estimate

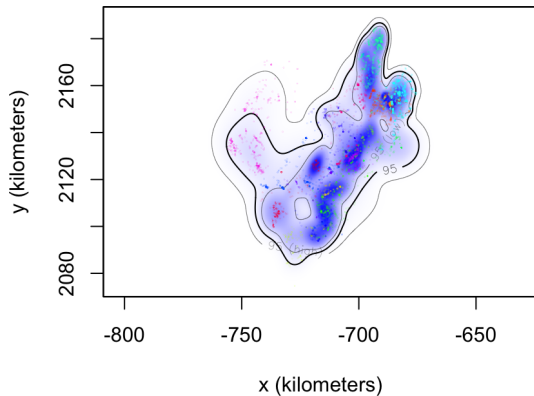


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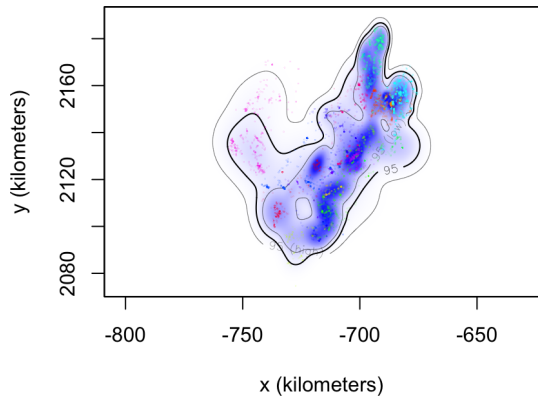


Mean AKDE estimate

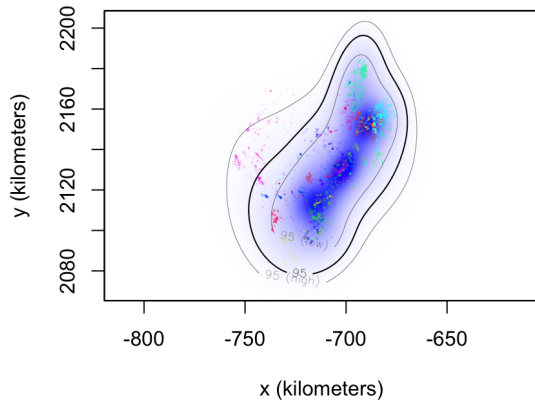


Validation: Empirical (*Ursus arctos horribilis*)

Mean AKDE estimate

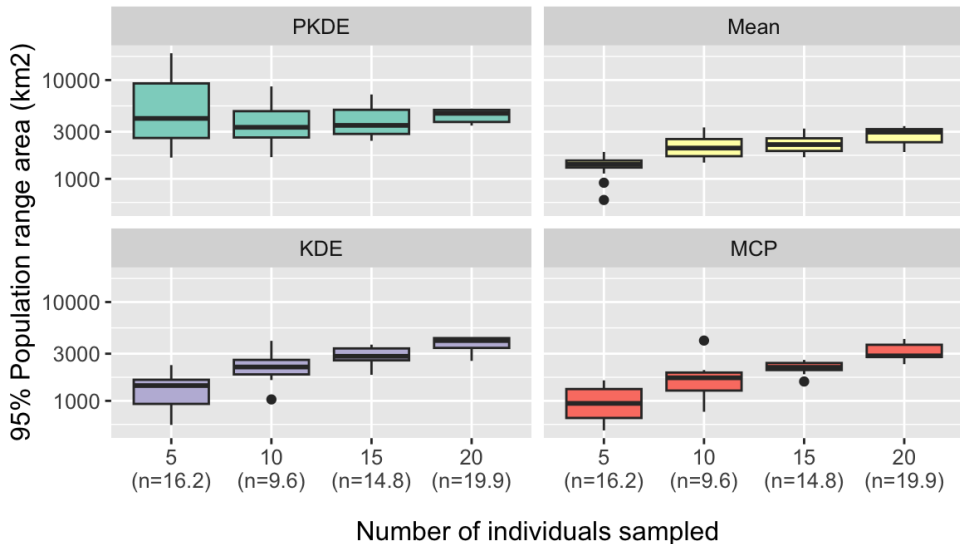


PKDE estimate

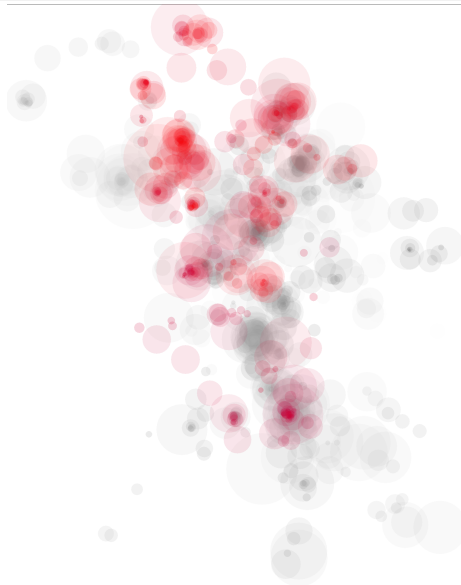


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U. a. horribilis Saturation Curve



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 - Parallels the problem of down-weighting low-quality (large HDOP) locations in home-range estimation

Thank you

- In collaboration with: Gayatri Anand, William F. Fagan, Justin M. Calabrese, etc.
- Funded by and supported by:



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