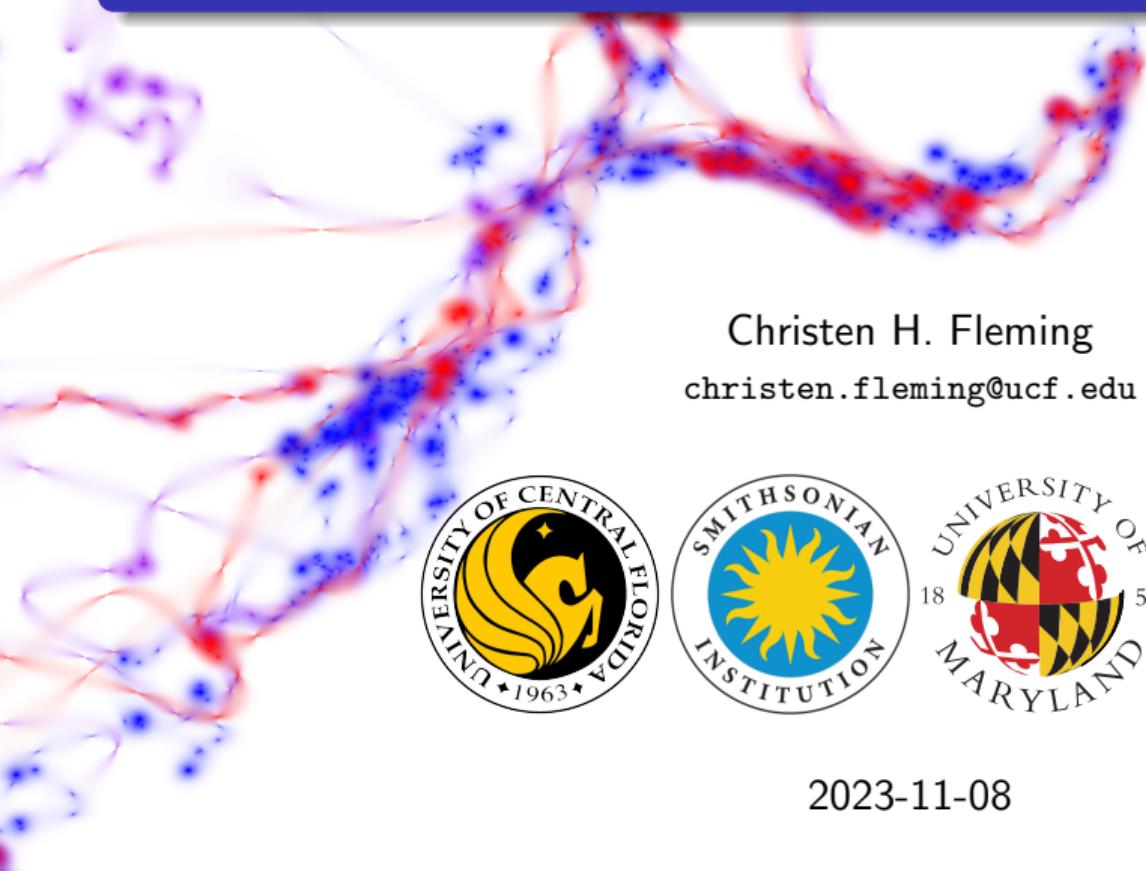


# Population range estimation with animal tracking data



Christen H. Fleming  
[christen.fleming@ucf.edu](mailto:christen.fleming@ucf.edu)



2023-11-08

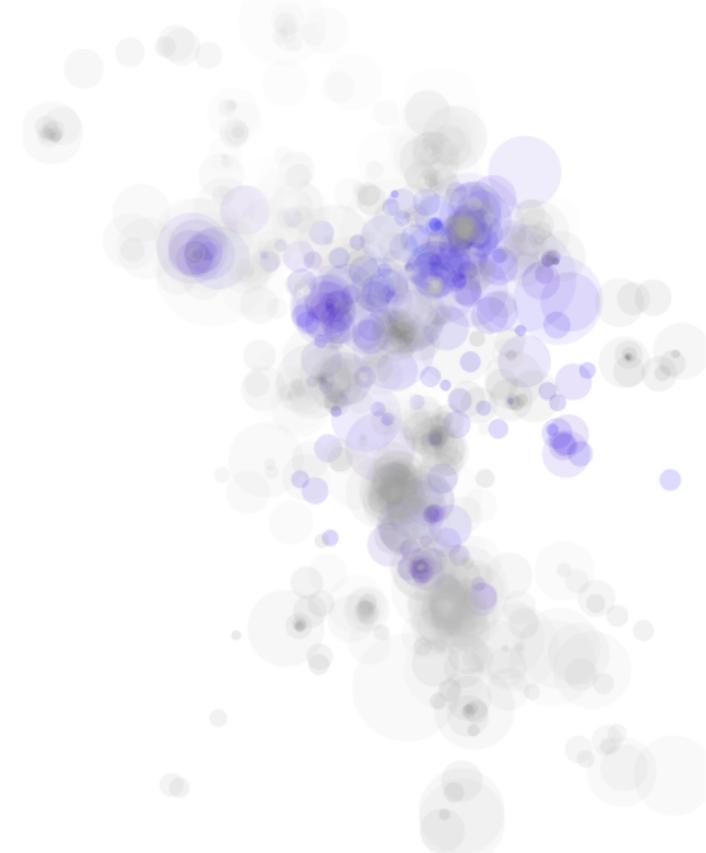
# Overview

- Introduction
- Kernel density estimation
- Validation
- Discussion



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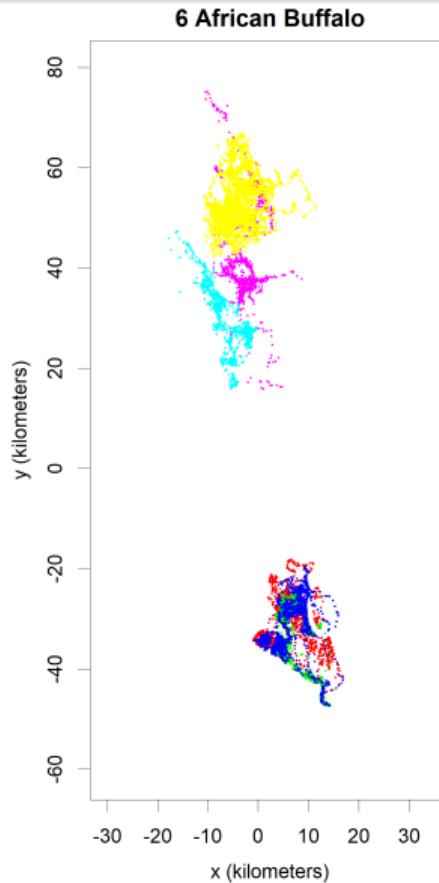
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# Introduction

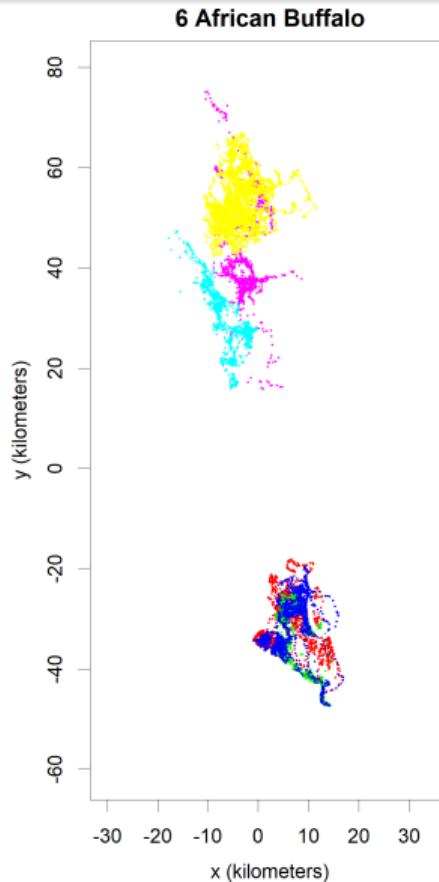
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- Tracks sampled from populations:  
What's the population distribution?



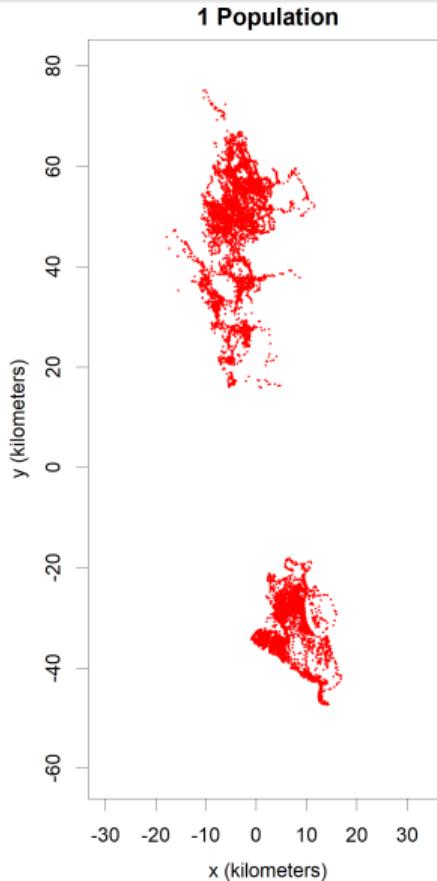
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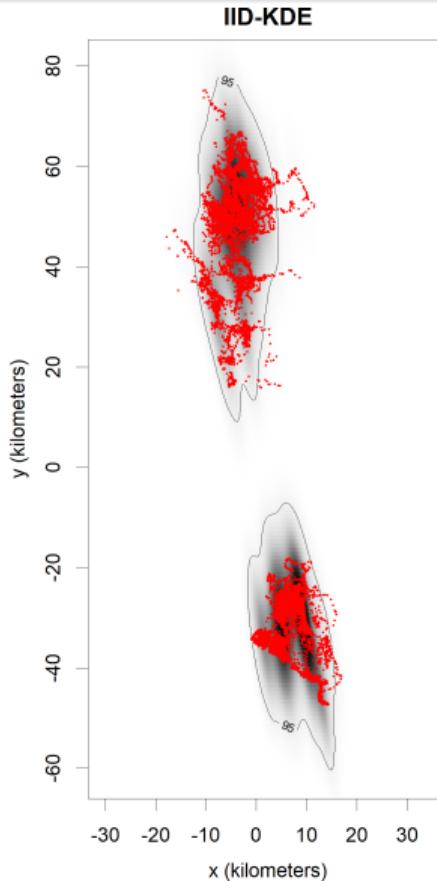
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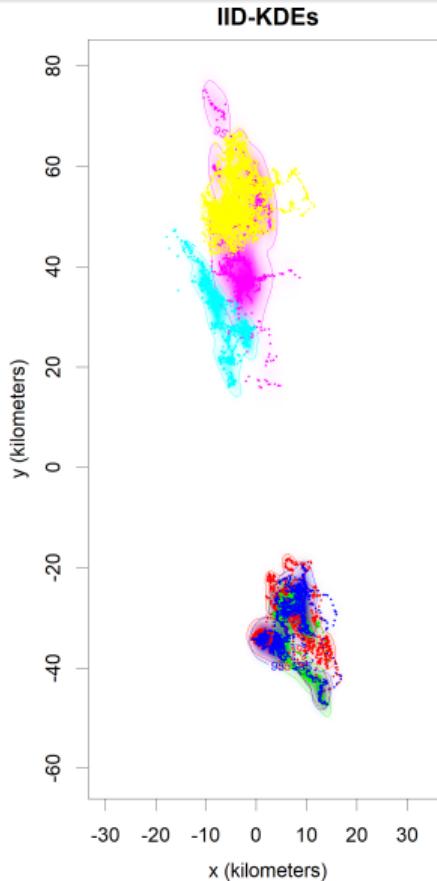
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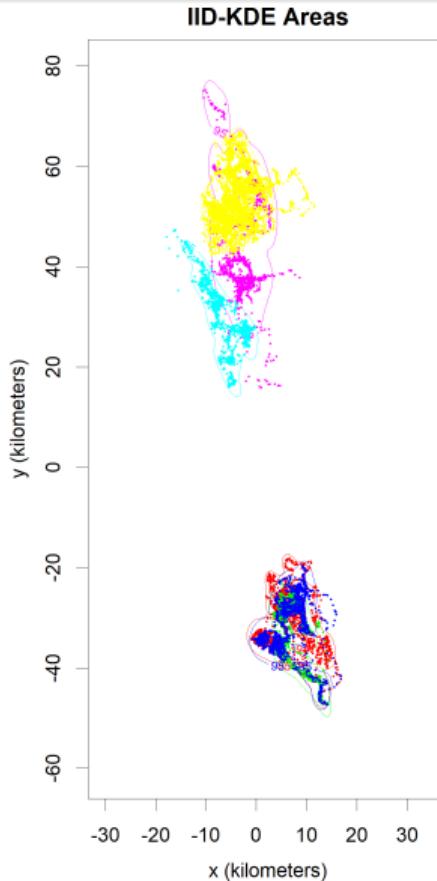
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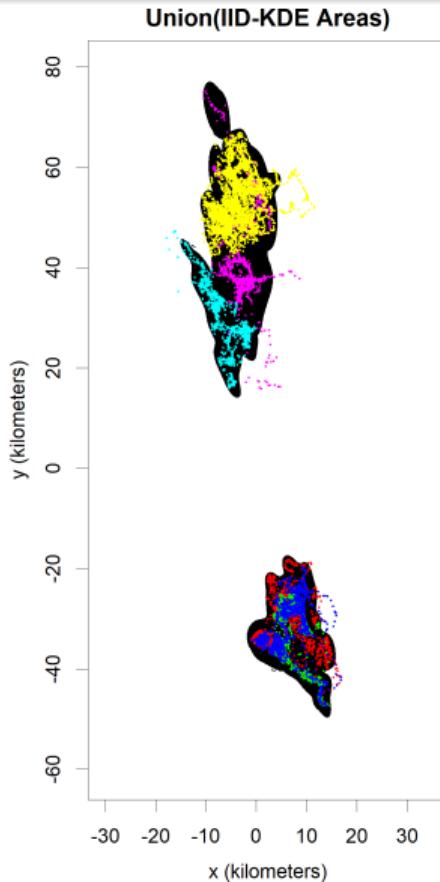
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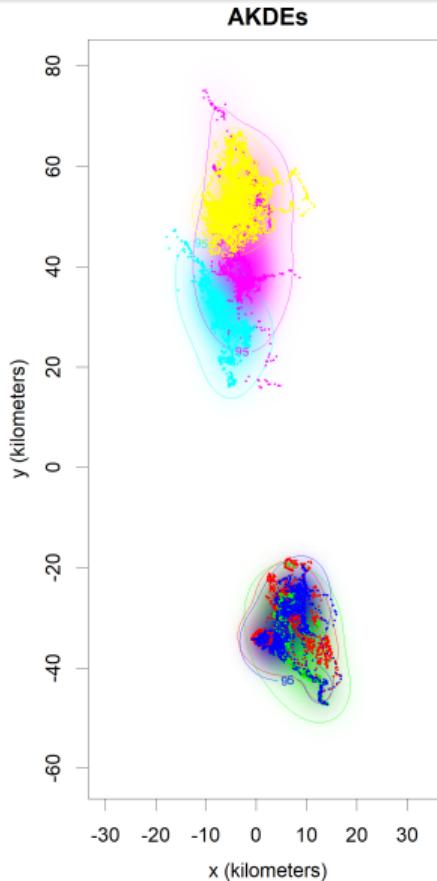
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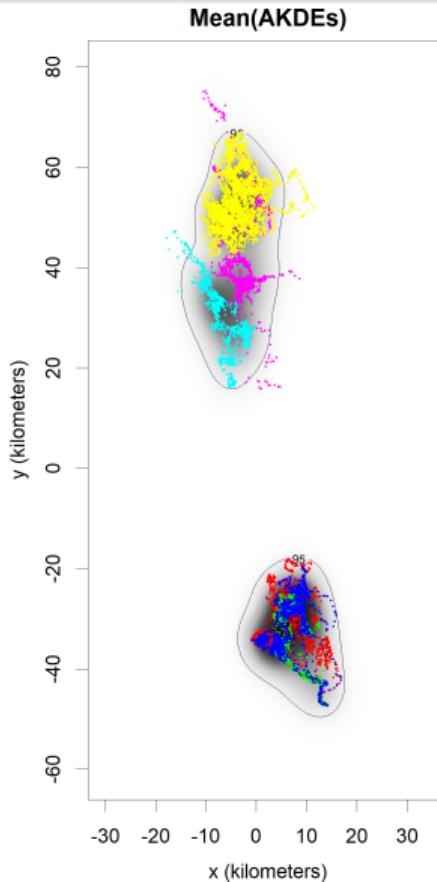
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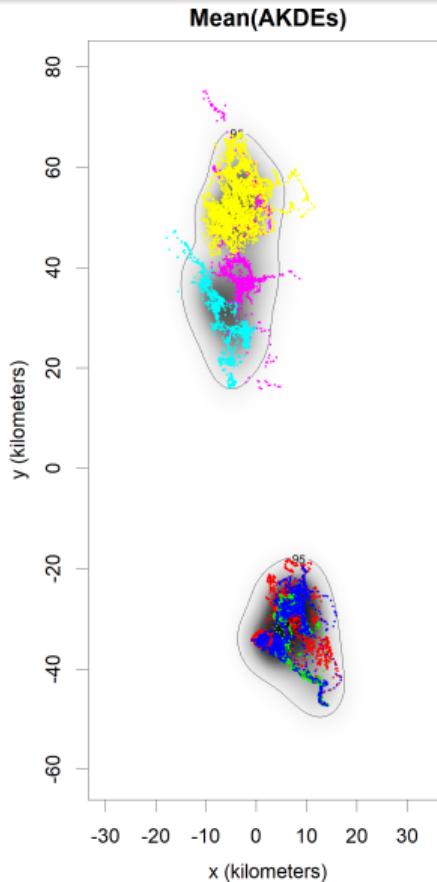
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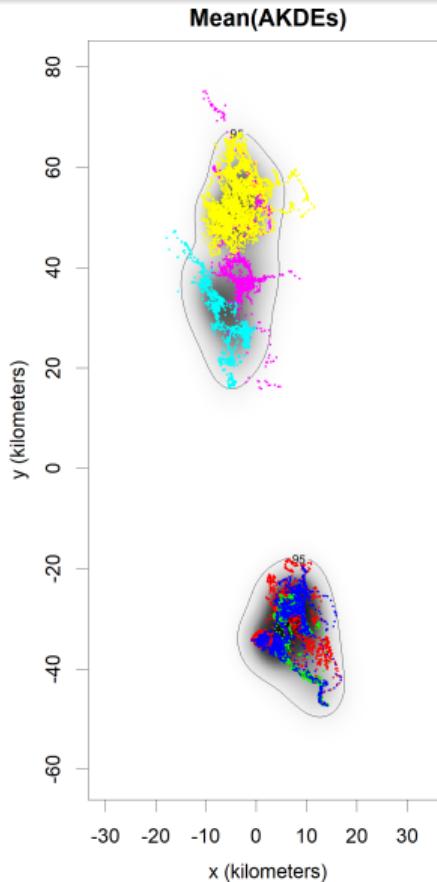
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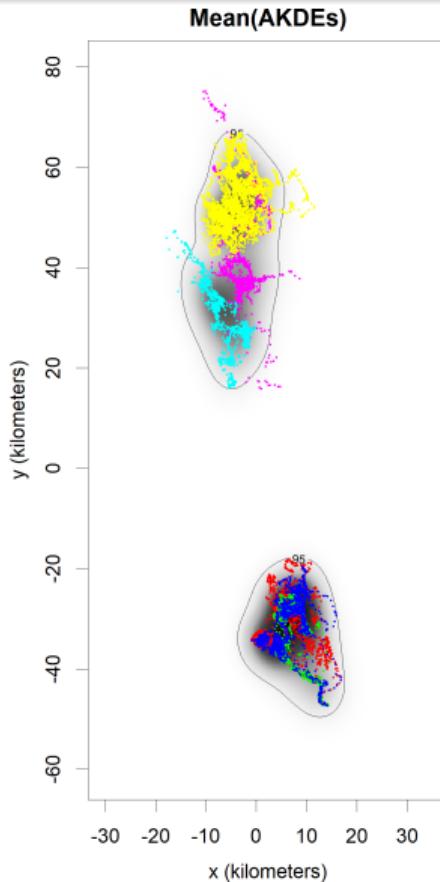
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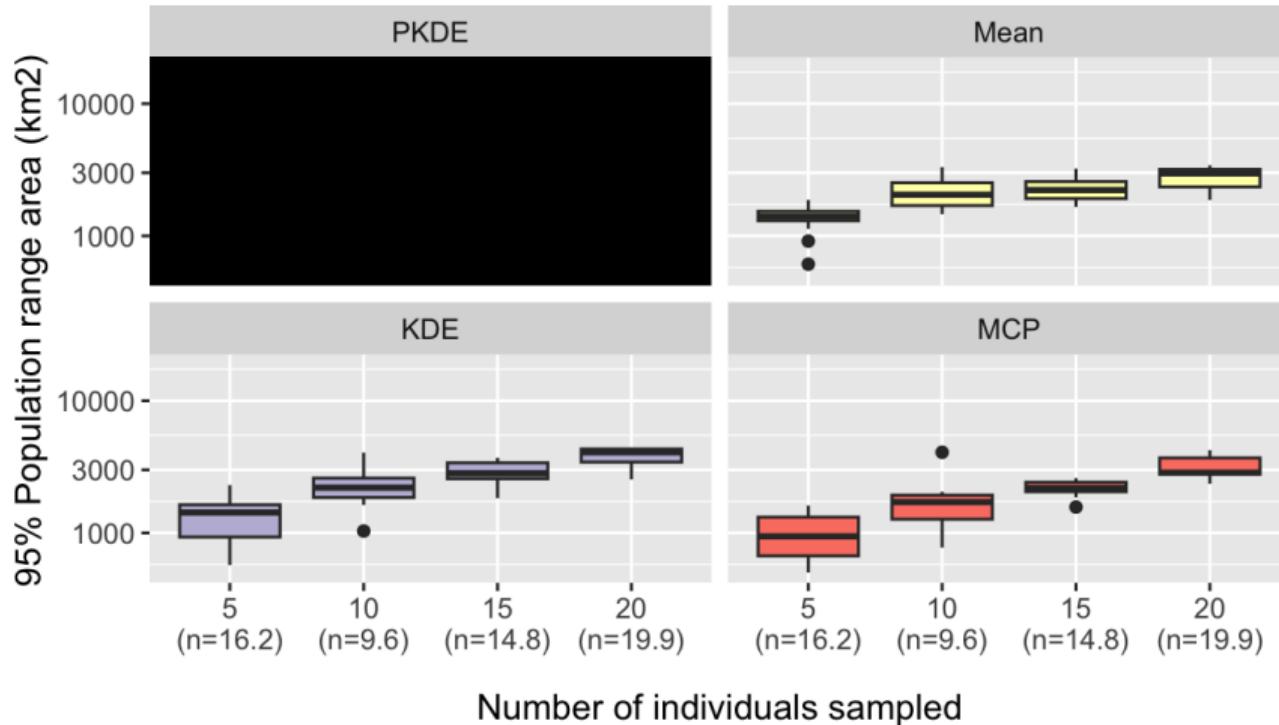
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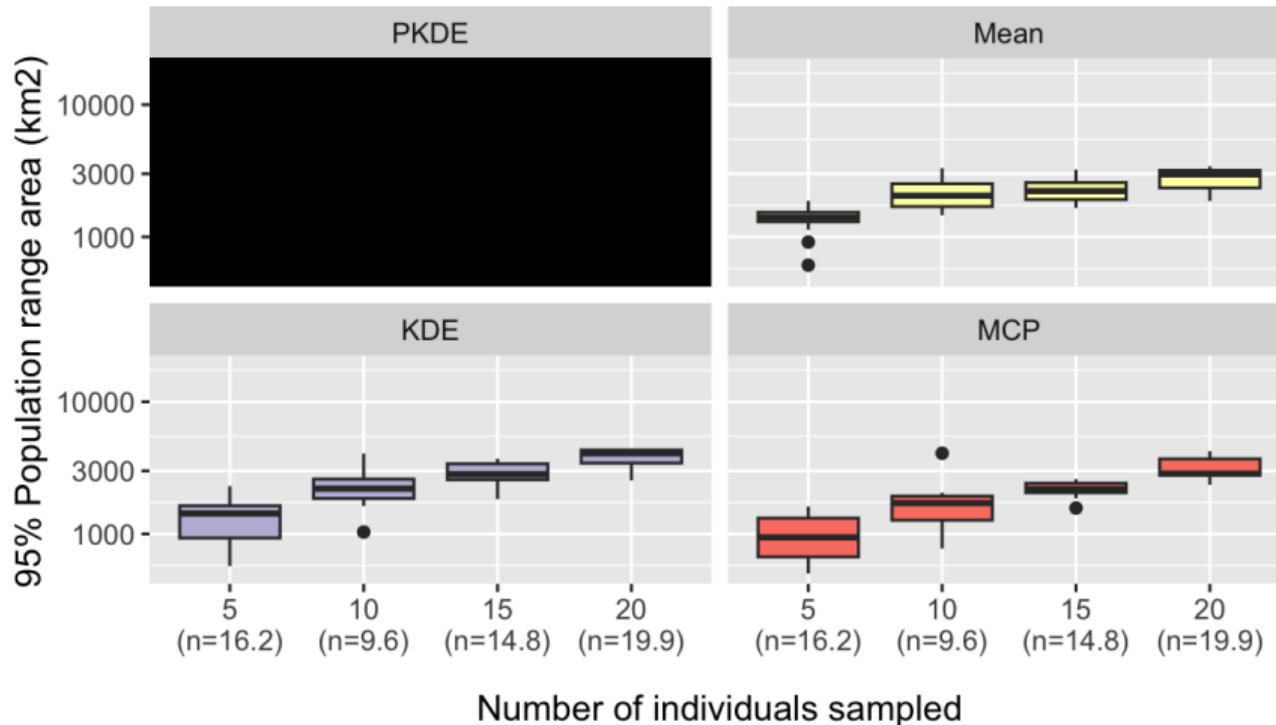
# Introduction

## *U. a. horribilis* Saturation Curve



# Introduction

## *U. a. horribilis* Saturation Curve



**This is bias from not modeling population variance**

- Introduction
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# Kernel Kernel Density Estimation

A KDE is a

:

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

$\hat{p}(\cdot)$ : 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  $t$

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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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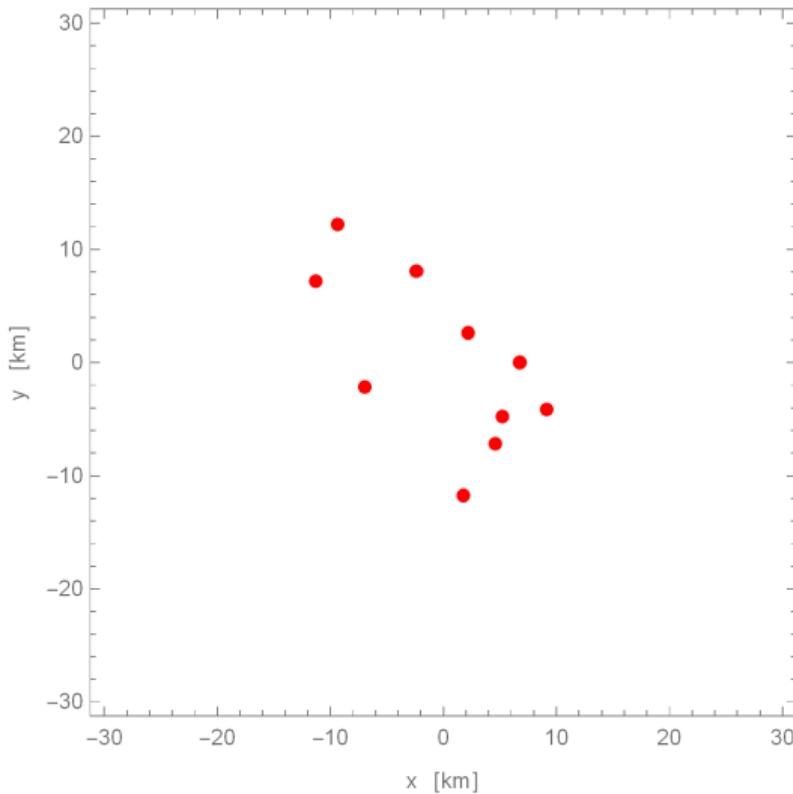
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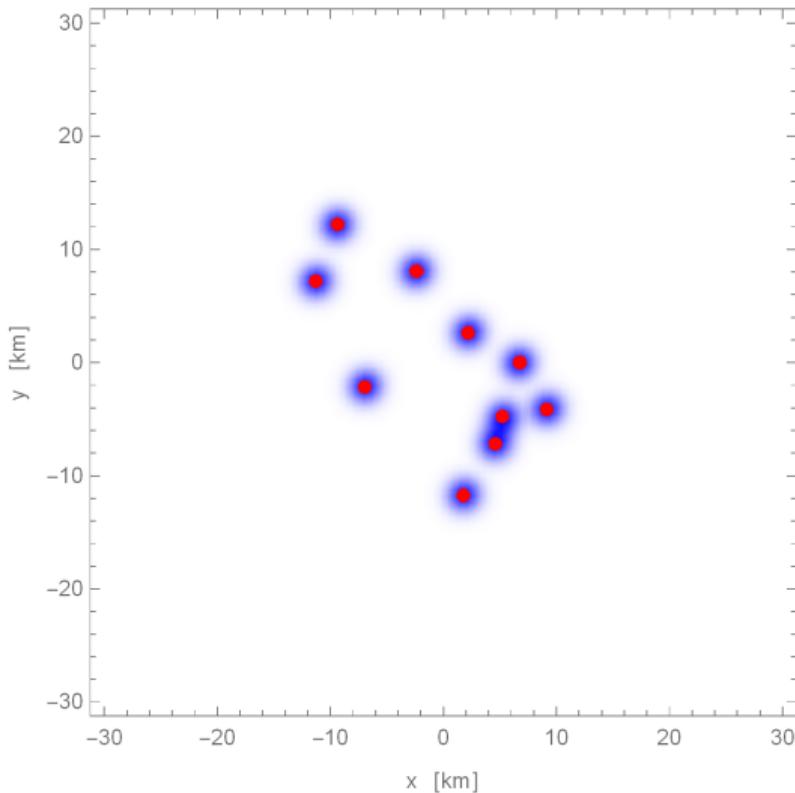
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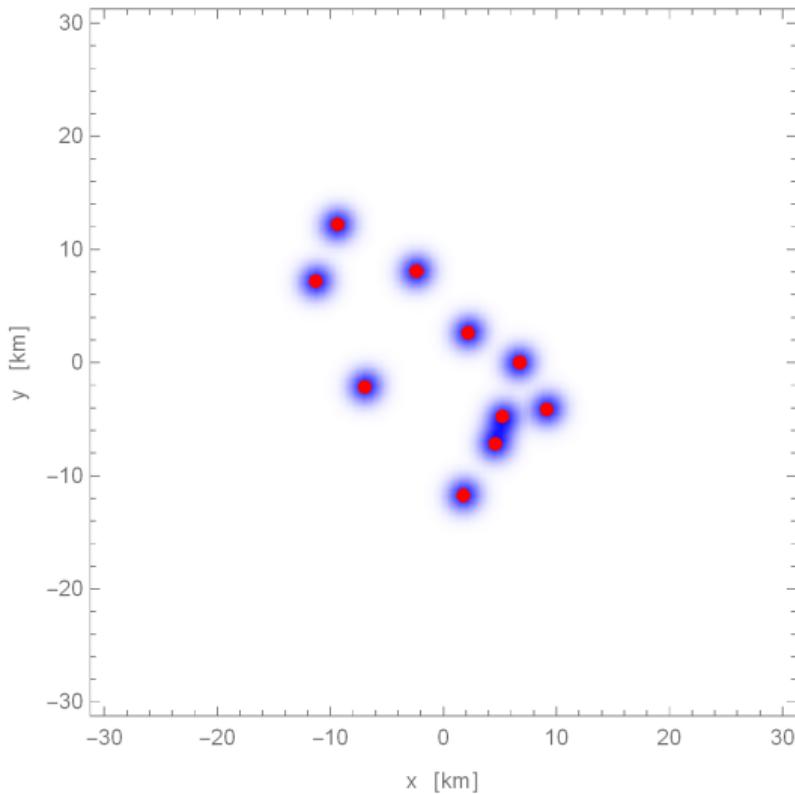
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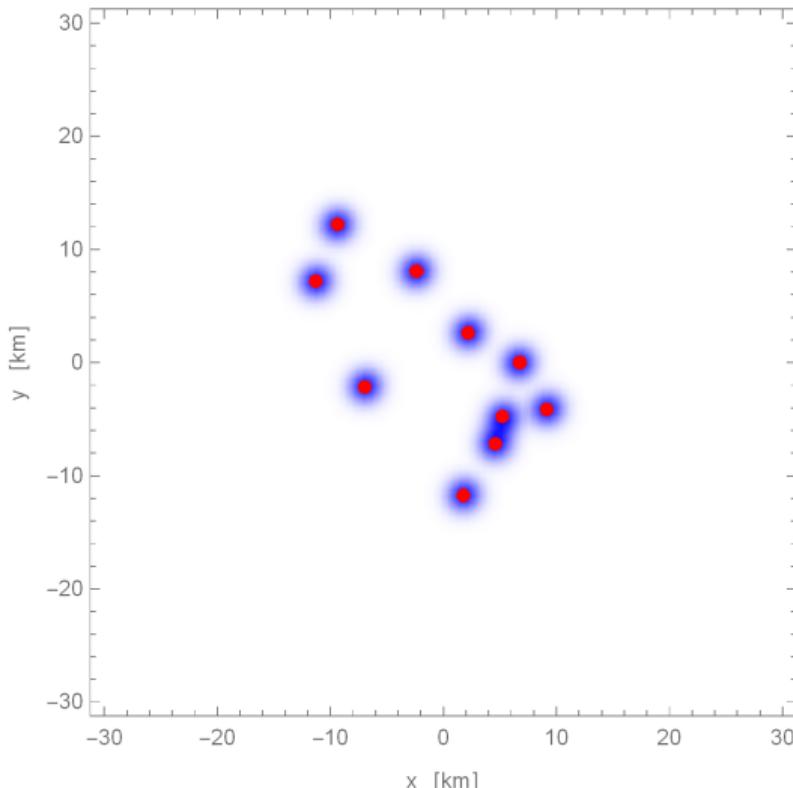
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$$M[\quad] = E[\quad]$$



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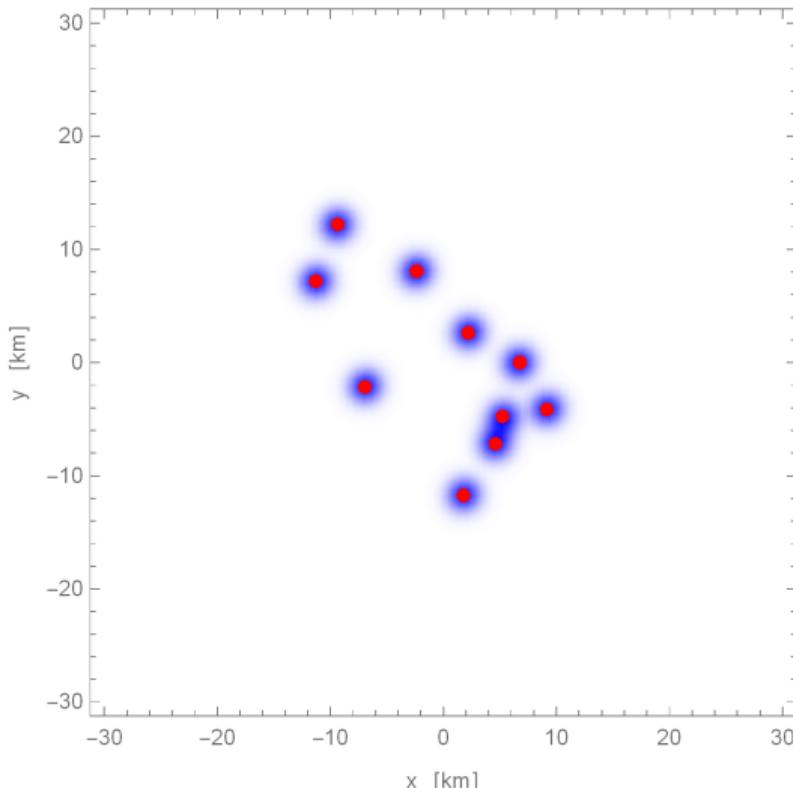
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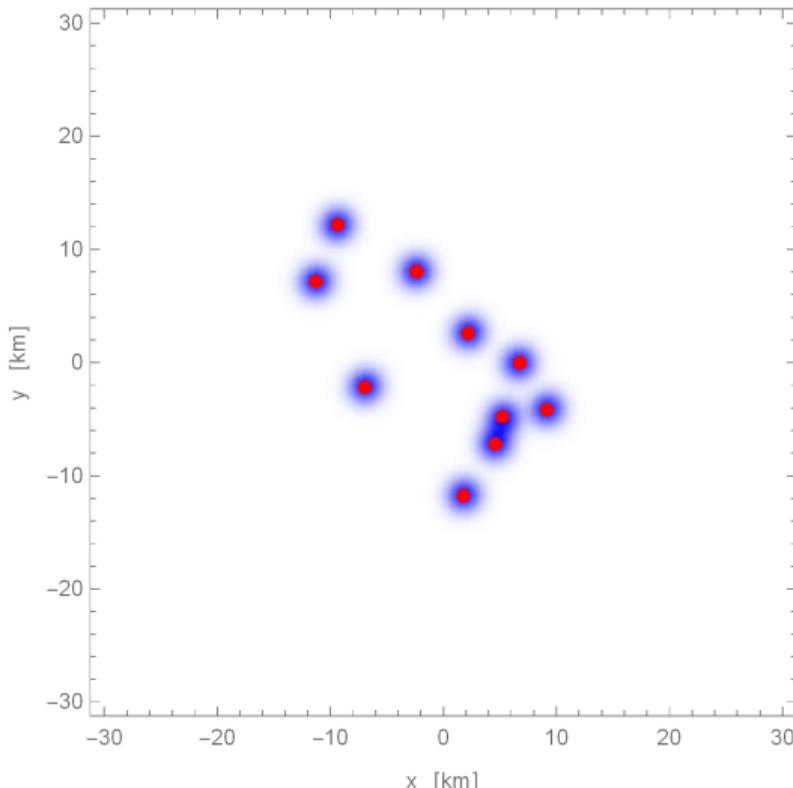
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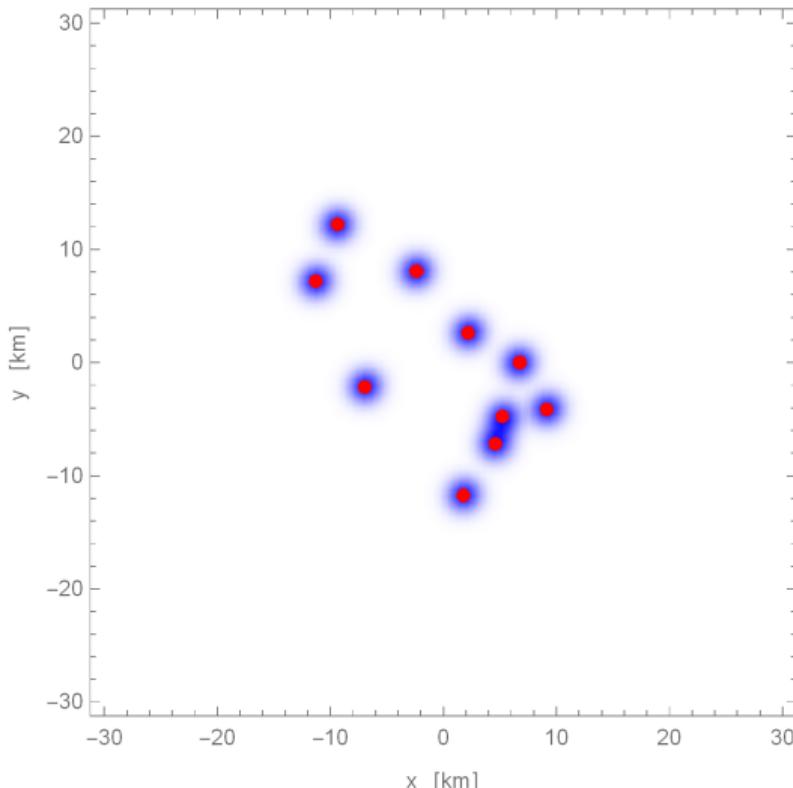
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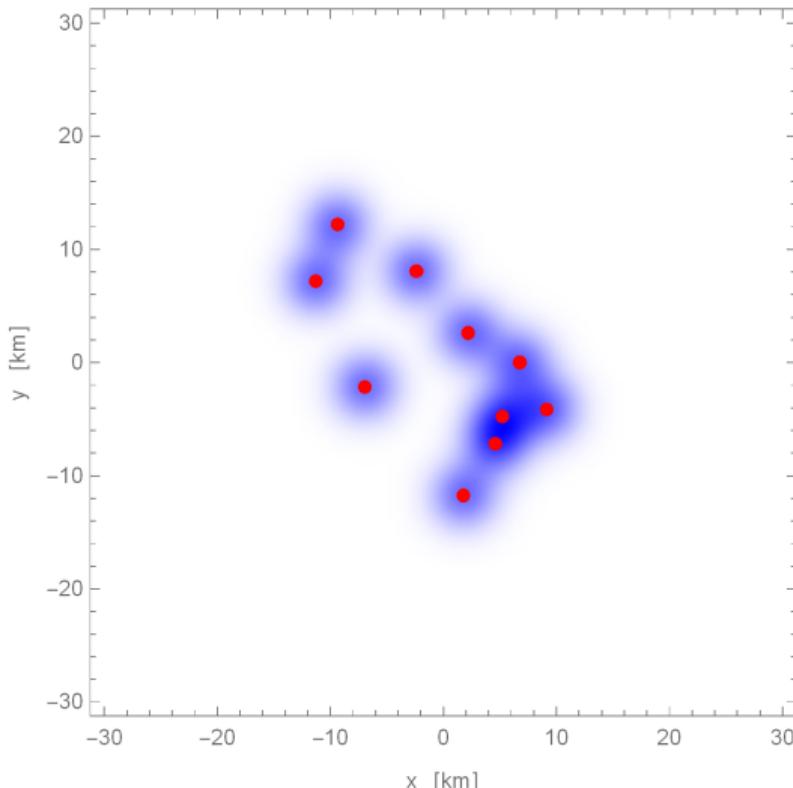
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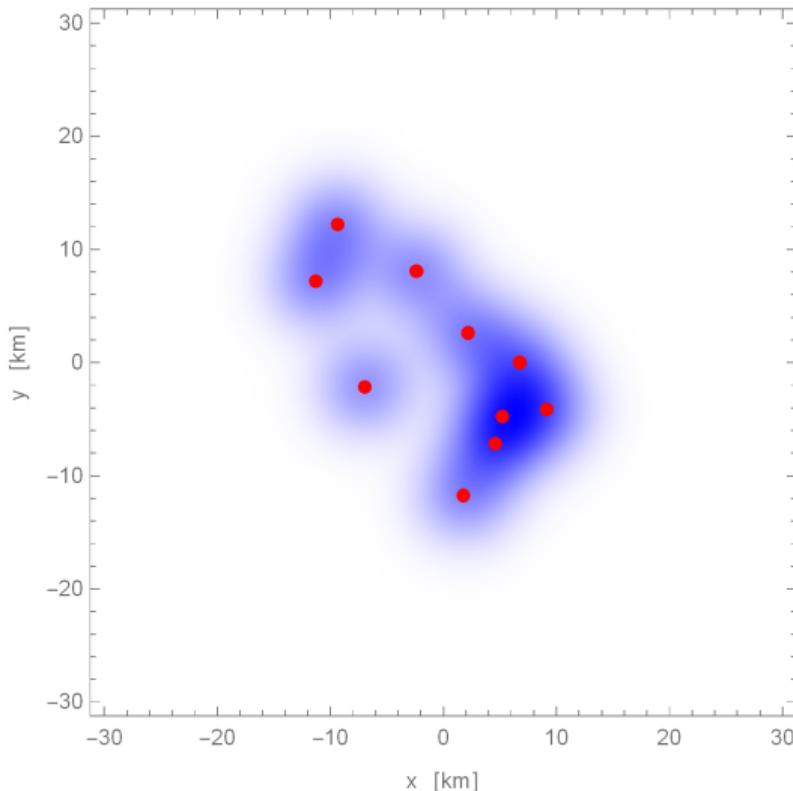
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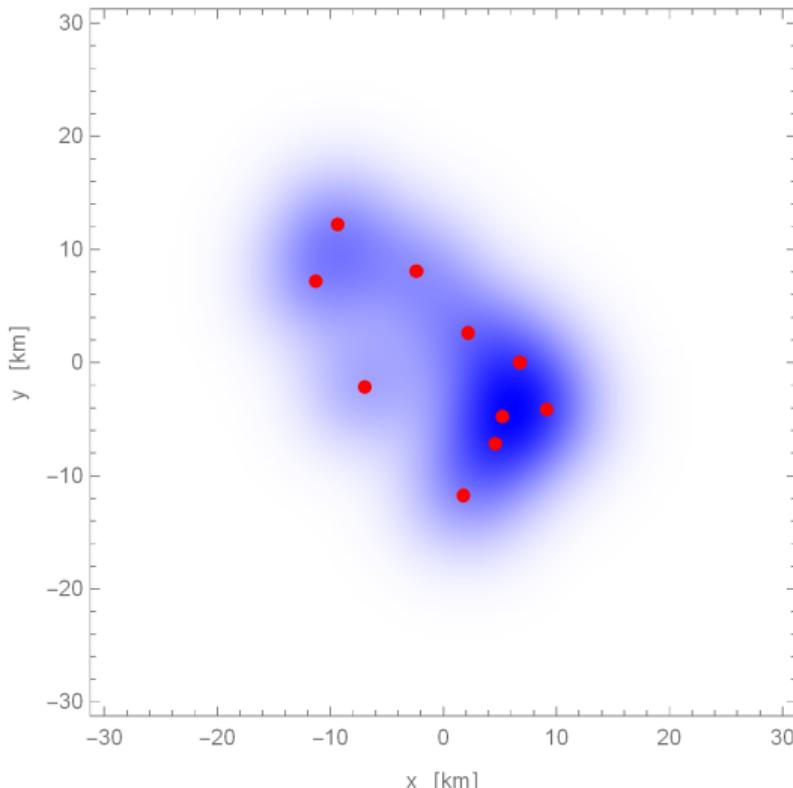
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$$\text{MISE}[\mathbf{H}] = \mathbb{E} \left[ \iint (\hat{p}(\mathbf{x}|\mathbf{H}) - p(\mathbf{x}))^2 d\mathbf{x} \right]$$

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$\mathbf{x}(t) \sim \text{ACF}$  (autocorrelated KDE)

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- (MISE) optimally weighted wAKDE (Fleming, Sheldon, et al. 2018)

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

# Population kernel density estimation

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- Individual weight assumptions:

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

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

$$\hat{p}_{\text{pop.}}(\mathbf{x}|\mathbf{H}) = \sum_{\text{ind.}} \sum_t w_{\text{ind.}}(t) \kappa(\mathbf{x} - \mathbf{x}_{\text{ind.}}(t)|\mathbf{H}_{\text{ind.}})$$

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

# Population kernel density estimation

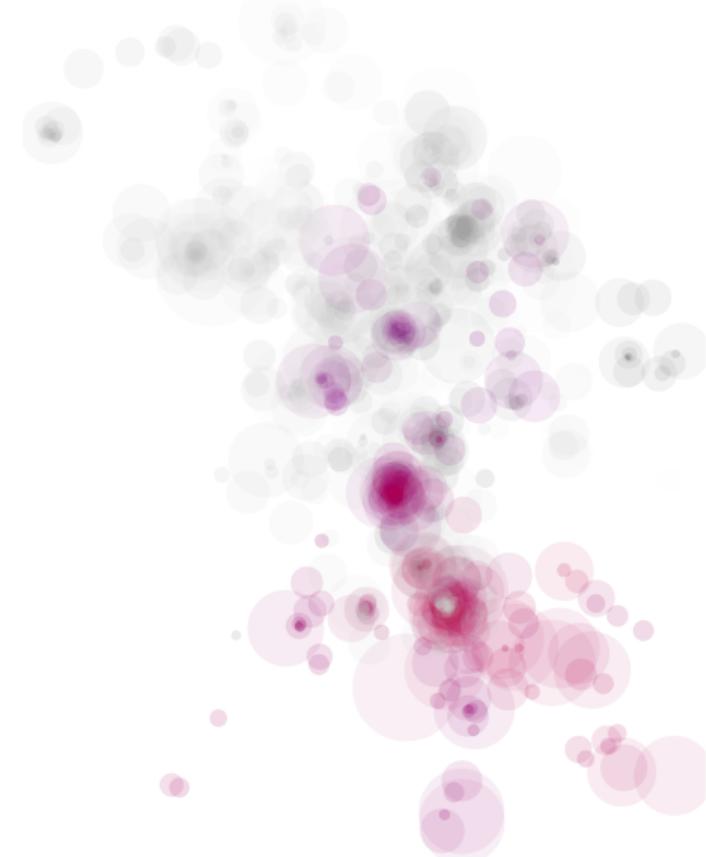
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# Transition

- Introduction
- Kernel density estimation
- Validation
- Discussion



# Transition

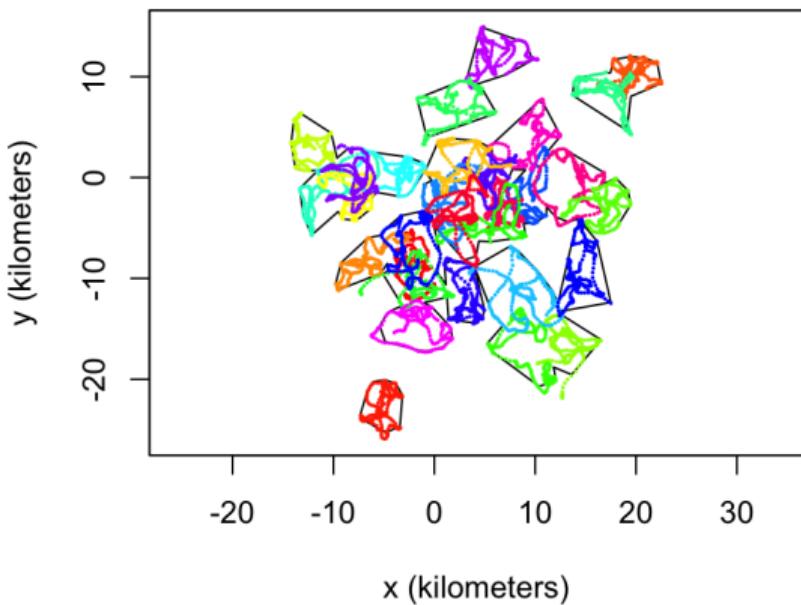
- Introduction
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Gayatri Anand (UMD Ph.D.)

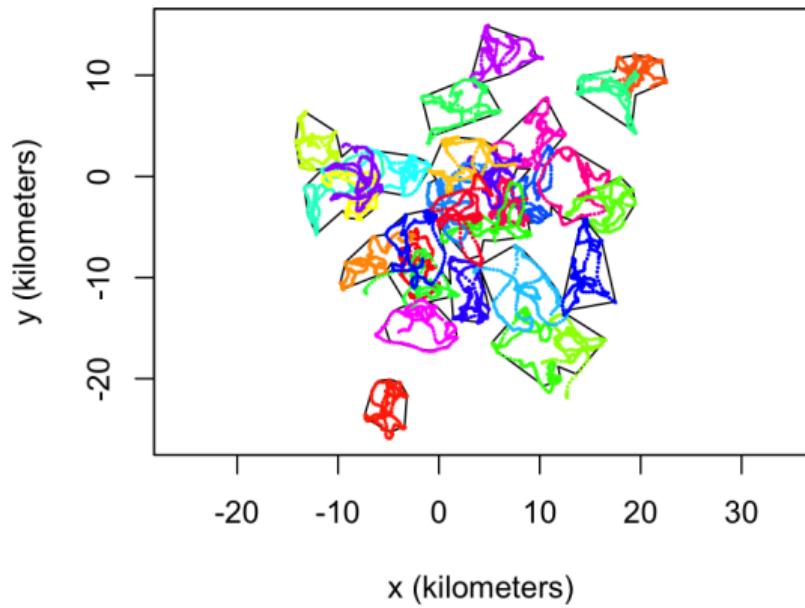
# Validation: Simulations

Merged MCP estimate

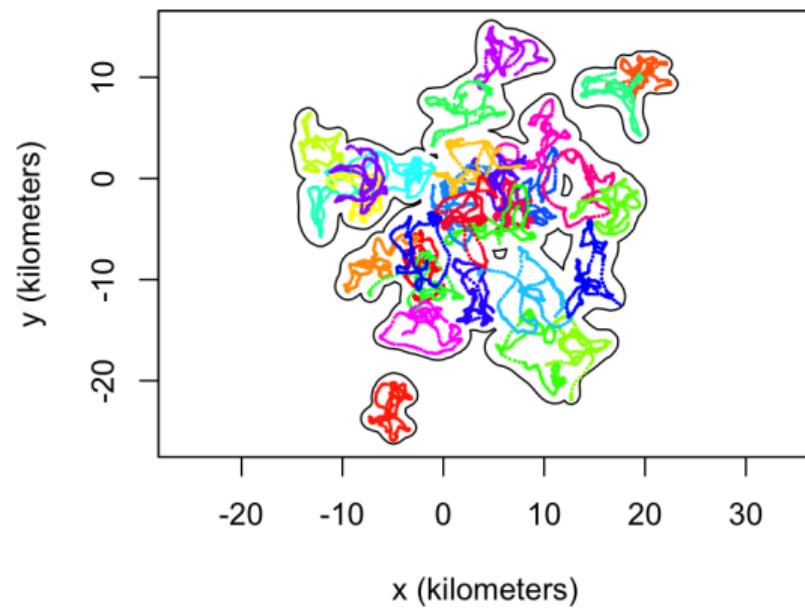


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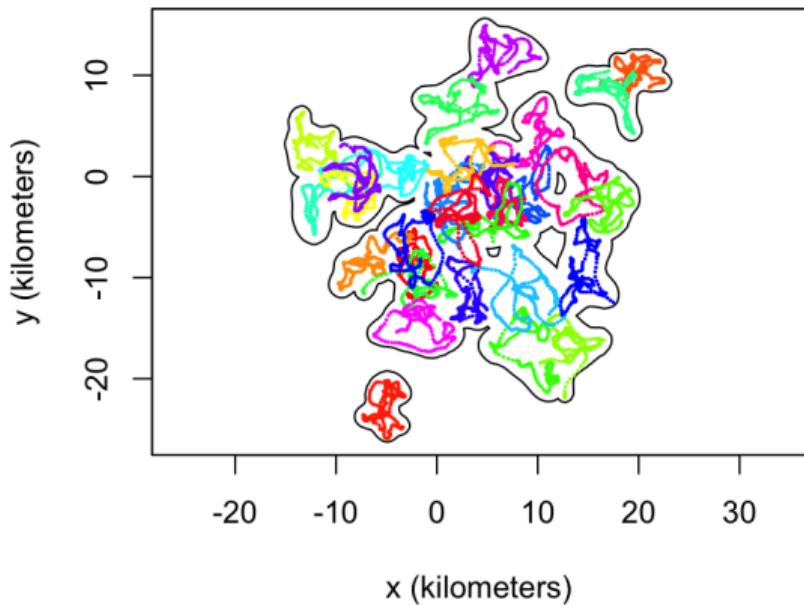


Merged KDE estimate

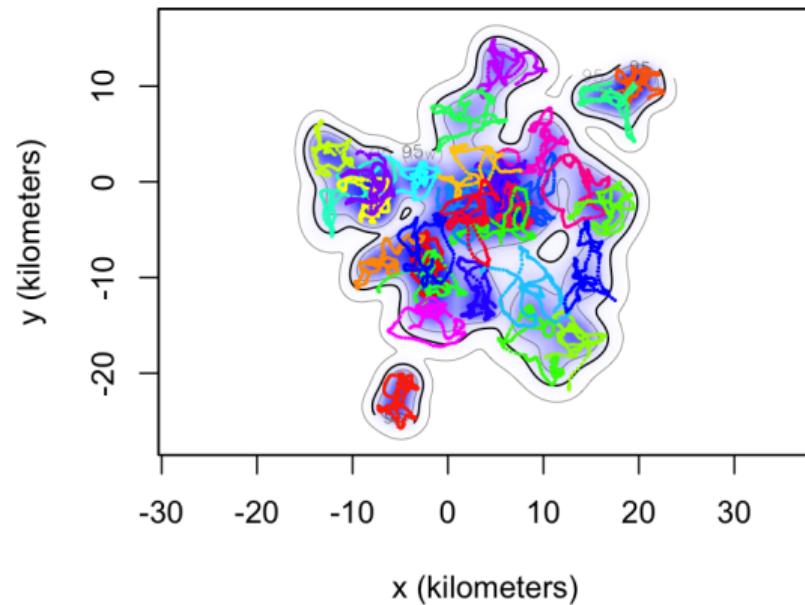


# Validation: Simulations

Merged KDE estimate

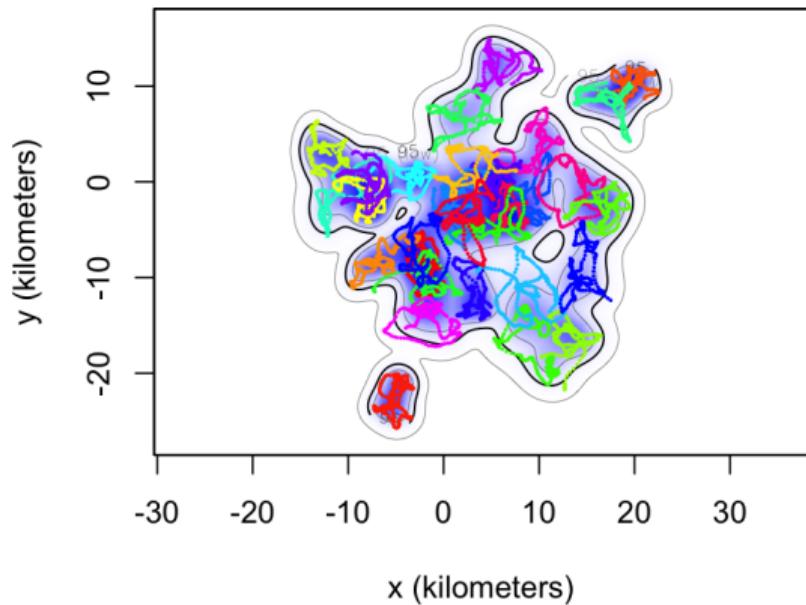


Mean AKDE estimate

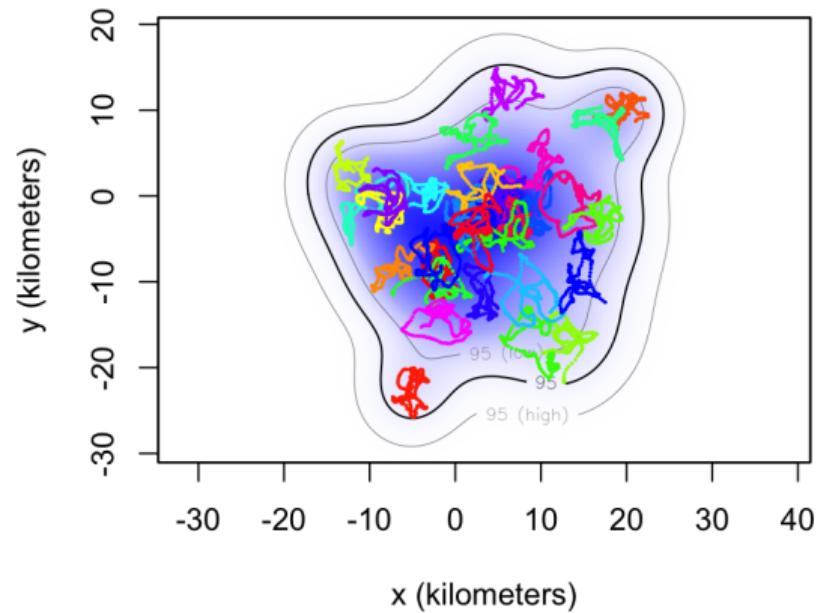


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Mean AKDE estimate

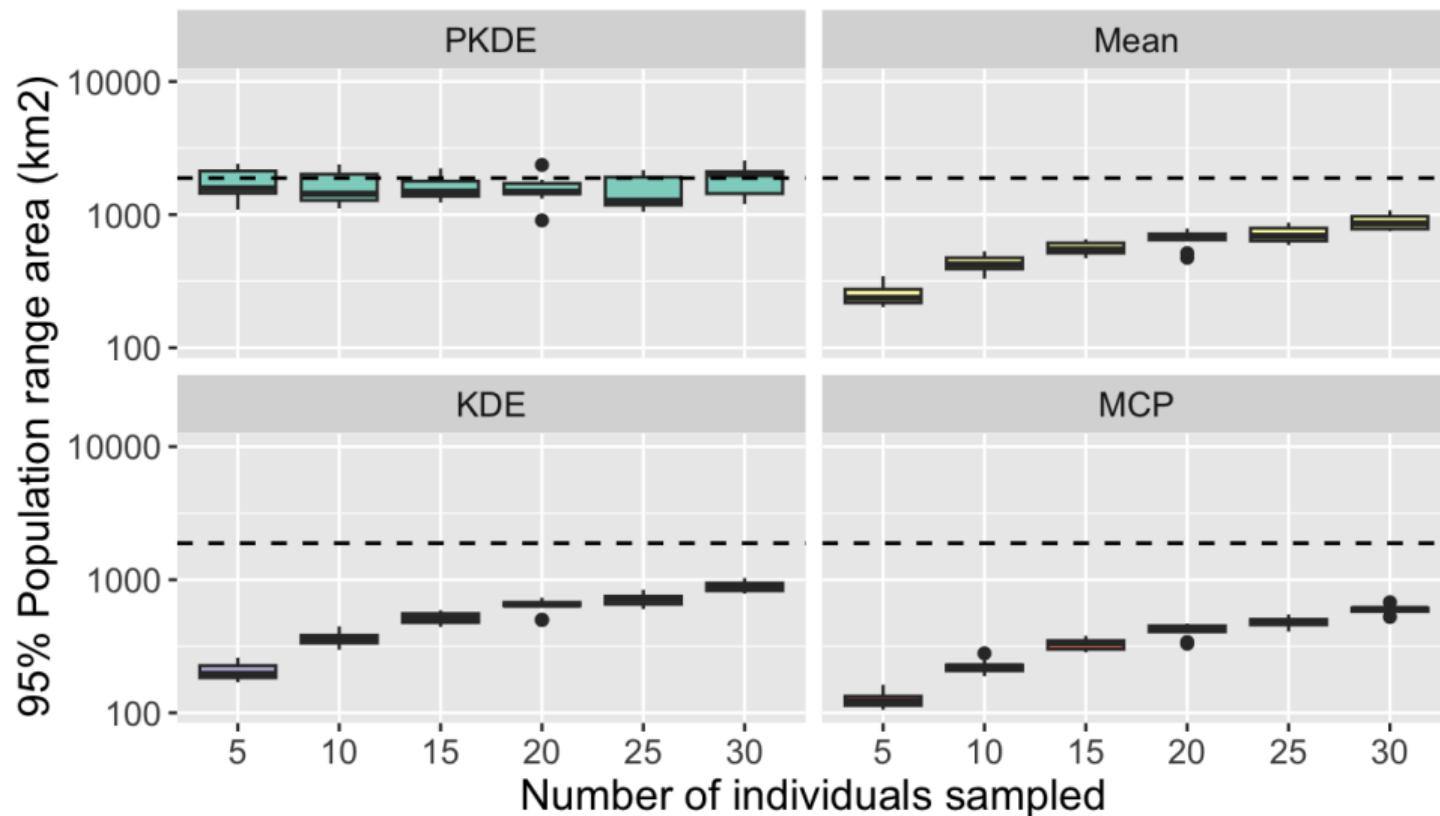


PKDE estimate



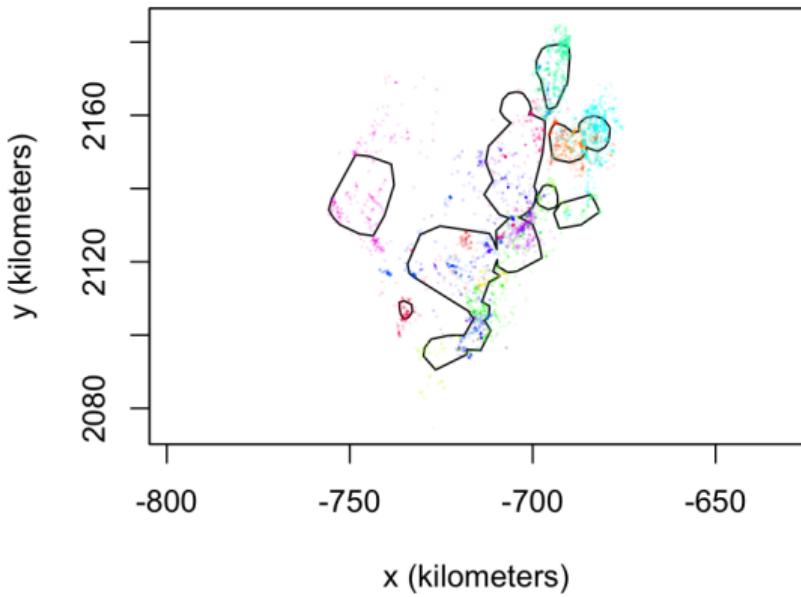
# Validation: Simulations

## Saturation Curve for Simulated Data



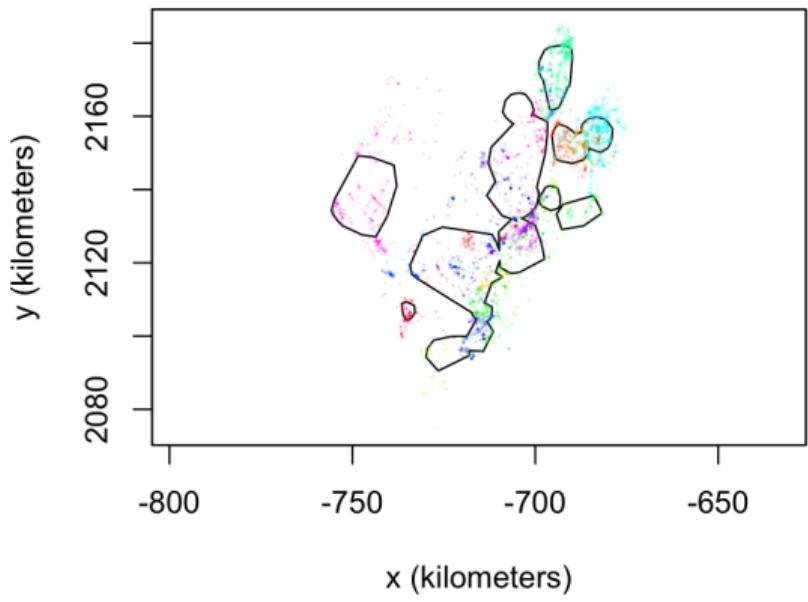
# Validation: Empirical (*Ursus arctos horribilis*)

Merged MCP estimate

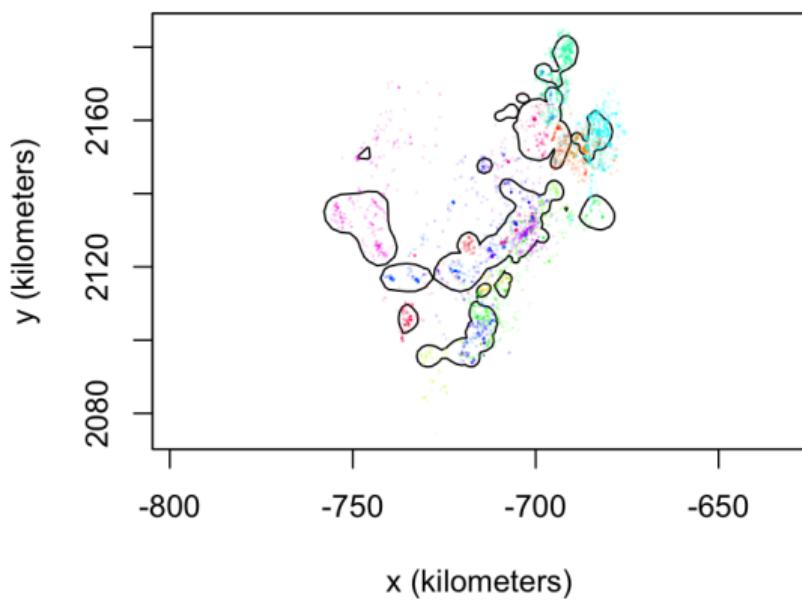


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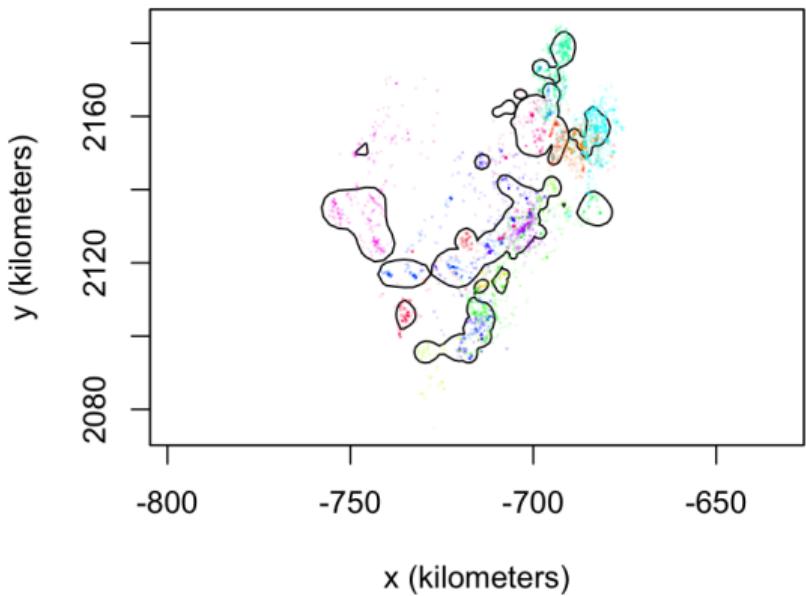


Merged KDE estimate

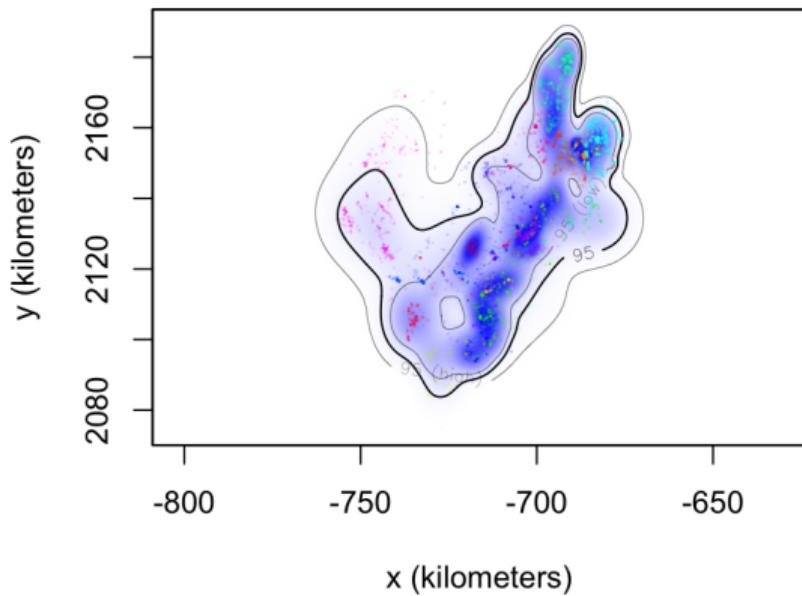


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Merged KDE estimate

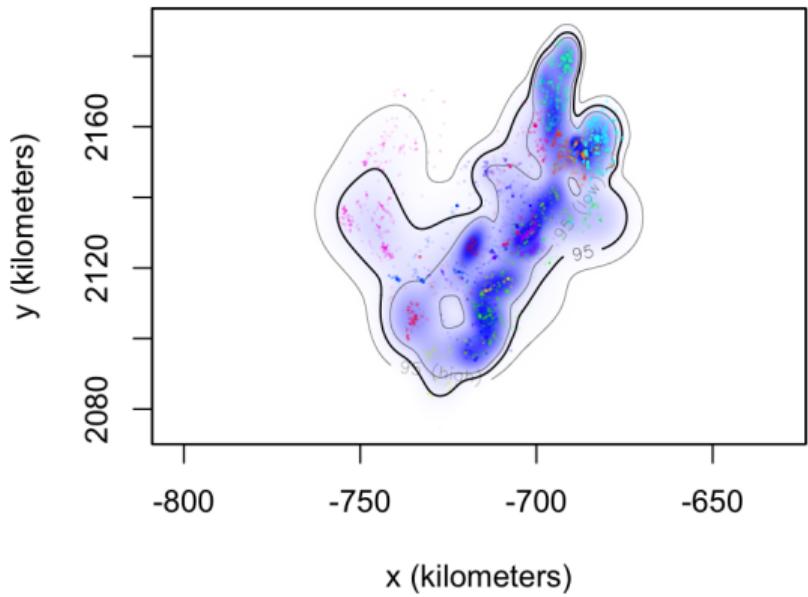


Mean AKDE estimate

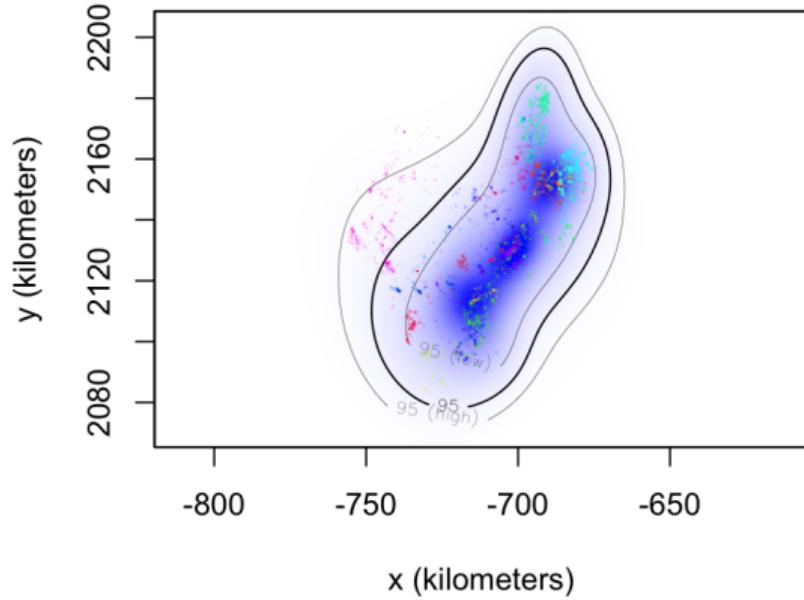


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Mean AKDE estimate

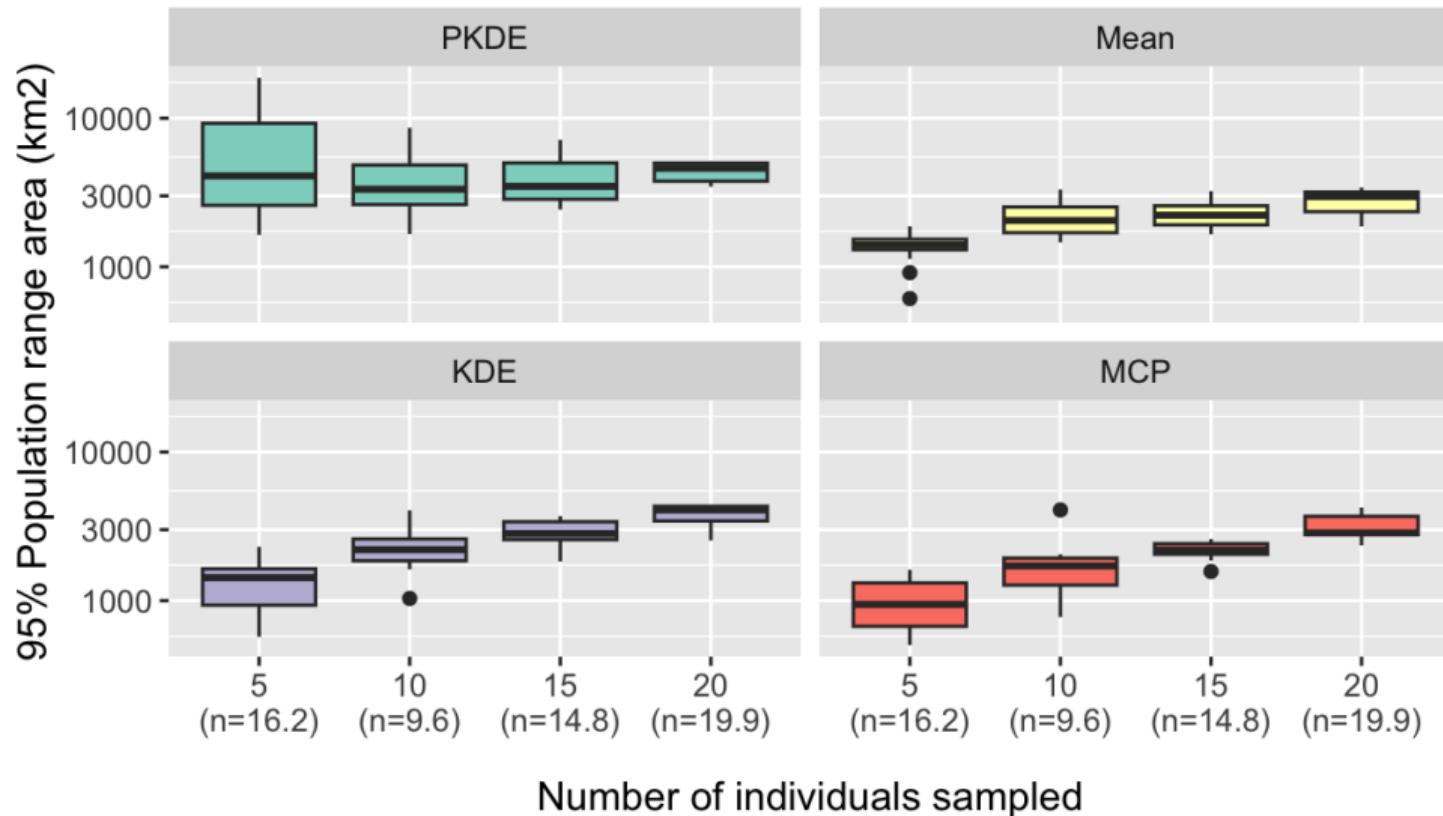


PKDE estimate

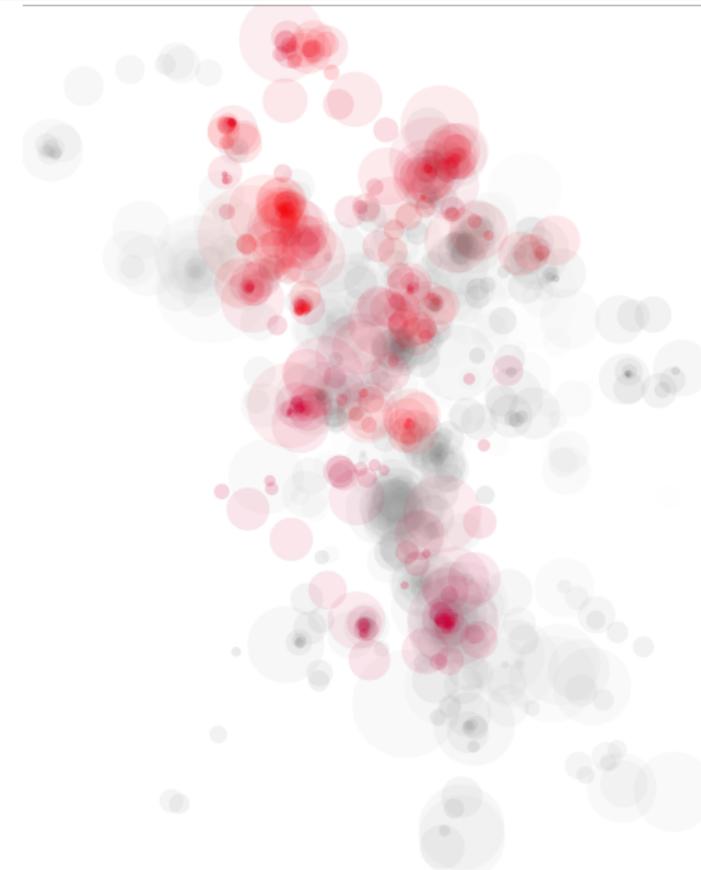


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## U. a. horribilis Saturation Curve



- Introduction
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# Discussion

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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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