DENSITY TREE AND DENSITY RANKING IN SINGULAR MEASURES

Yen-Chi Chen

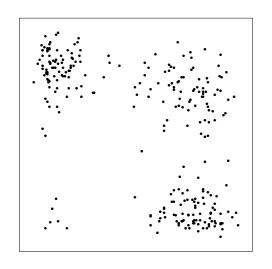
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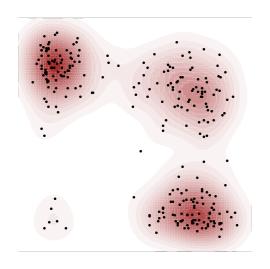


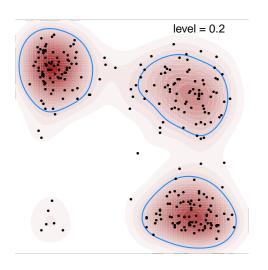
Outline

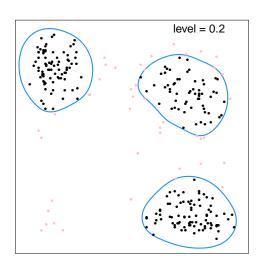
- Density Trees
- Density Ranking
- o Density Ranking: Multiple Datasets
- Summary

DENSITY TREES



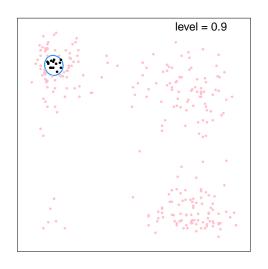


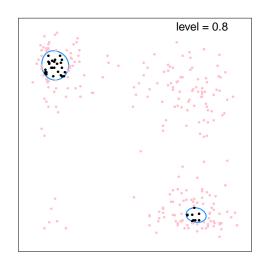


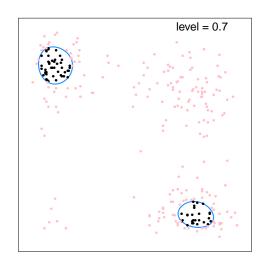


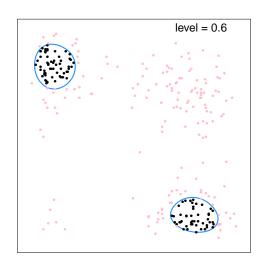
- The idea of using a density level (threshold) λ leads to clusters representing high density regions.
- Thus, the level λ has an effect on the clustering result.

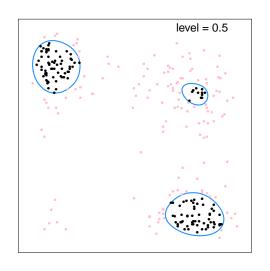
- The idea of using a density level (threshold) λ leads to clusters representing high density regions.
- Thus, the level λ has an effect on the clustering result.
- Varying the level λ may lead to a creation of a new cluster or a merging of existing clusters.

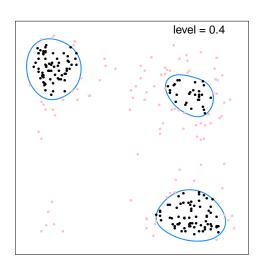


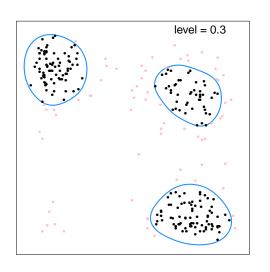


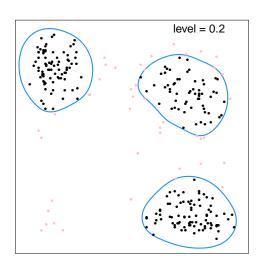


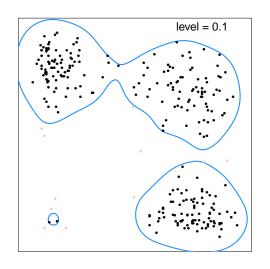


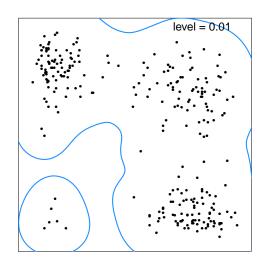








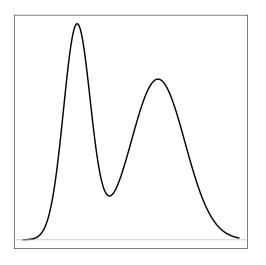


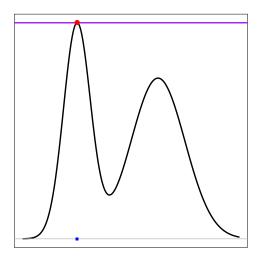


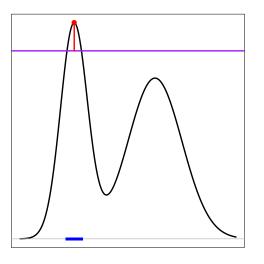
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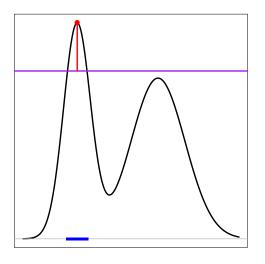
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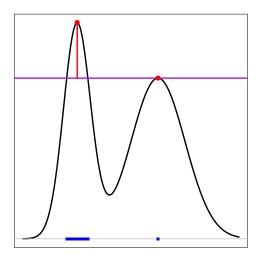
- When the level changes, we see the *evolution* of clusters.
- Cluster tree (Stuetzle 2003) is to summarize such an evolution process by a tree.
- When applied to a density function, a cluster tree is also called a density tree (Klemelä 2004).

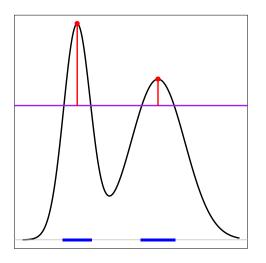


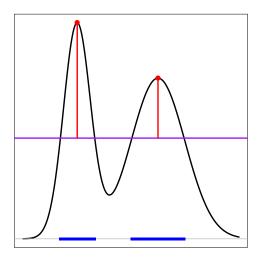


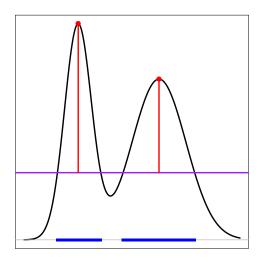


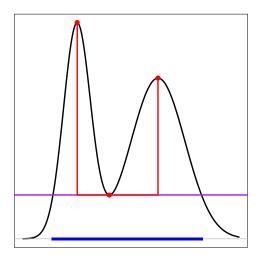


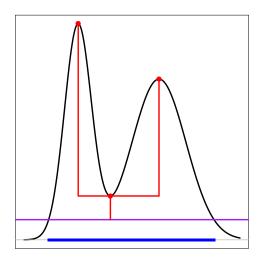


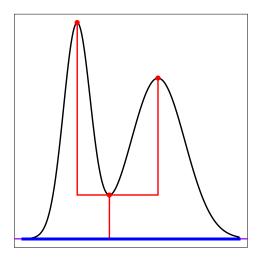




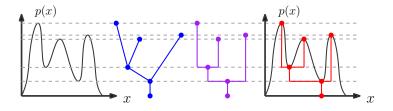






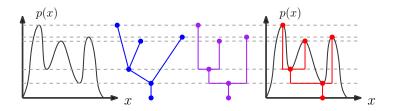


Density Trees - 1



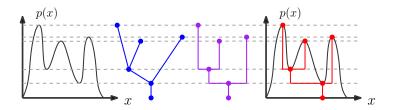
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- The merging of connected components is often associated with local minima or saddle points.

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- We define a collection $T_p = \bigcup_{\lambda \geq 0} \{C_{\lambda,1}, \cdots, C_{\lambda,J(\lambda)}\}$. Namely, T_p is the collection of all connected components from every level.
- \circ Then the elements of T_p admits a tree structure this tree structure is the density tree.

- In statistics, we often do not know the true density function *p*.
- Instead, we observe a random sample $X_1, \dots, X_n \in \mathbb{R}^d$ that are IID from p.
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- Here we use the kernel density estimator (KDE):

$$\widehat{p}_n(x) = \frac{1}{nh^d} \sum_{i=1}^n K\left(\frac{X_i - x}{h}\right),$$

where $K(\cdot)$ is the kernel function that is often a smooth function like a Gaussian, and h > 0 is the smoothing bandwidth that controls the amount of smoothing.

• To measure the estimation error, a simple metric is

$$d_{\infty}(\widehat{T_p}, T_p) = \sup_{x} \|\widehat{p}_n(x) - p(x)\|,$$

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• Under smoothness conditions and $n \to \infty$, $h \to 0$,

$$P_n \geq 1 - e^{-nh^{d+4} \cdot C_p}$$

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for some constant C_p depending on the density function p.

• There are other notions of convergence/consistency of a tree estimator.

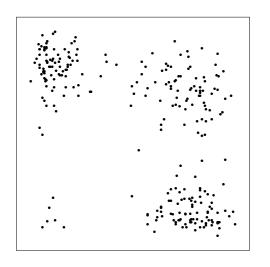
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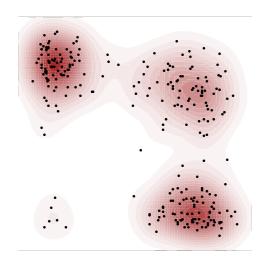
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 Balakrishnan et al. 2013) is another way to measure the consistency of a tree estimator.
- Note: density tree can also be recovered by a kNN approach; see Chaudhuri and Dasgupta (2010) and Chaudhuri et al. (2014) for more details.

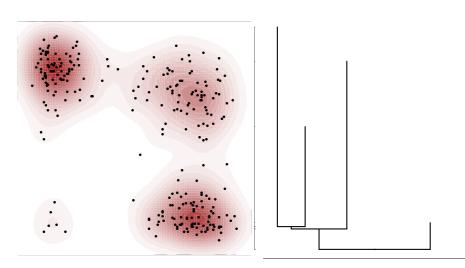
Kernel Density Estimator: an Example



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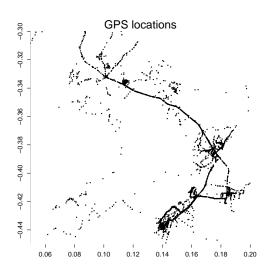
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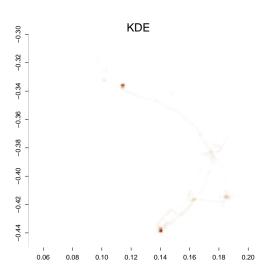
- Density trees provide topological information about the density function and they can be transformed into the persistent diagrams easily.
- When using a density level sets to define clusters, the density tree contains the information about the evolution and stability of clusters.
- Moreover, density trees can always be displayed in 2D plane. So they are good tools for visualizing multivariate functions.

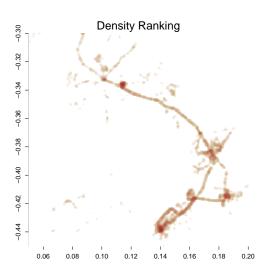
DENSITY RANKING

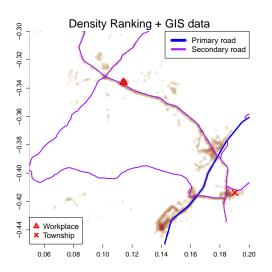
Failure of Density Trees and KDE

- Although density trees and KDE are good approaches, sometimes they may fail.
- In particular, when the PDF does not exist, we cannot use the usual definition for density trees and the KDE to analyze our data.









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- However, density ranking still works!

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- The density ranking at point *x* is

$$\widehat{\alpha}(x) = \frac{1}{n} \sum_{i=1}^{n} I\left(\widehat{p}(x) \ge \widehat{p}(X_i)\right)$$

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= ratio of observations' density below the density of point x.

• Namely, $\widehat{\alpha}(x) = 0.3$ implies that the (estimated) density of point x is above the (estimated) density of 30% of all observations.

Property of Density Ranking

• For an observation X_{max} with $\widehat{\alpha}(X_{\text{max}}) = 1$, then it means

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- If an observation X_{ℓ} satisfies $\widehat{\alpha}(X_{\ell}) = 0.25$, this means that the ranking of density at X_{ℓ} is the 25%.
- Moreover, for any pairs of data points X_i , X_j ,

$$\widehat{p}(X_i) > \widehat{p}(X_j) \Longrightarrow \widehat{\alpha}(X_i) > \widehat{\alpha}(X_j)$$

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- Density ranking $\widehat{\alpha}(x)$ can be viewed as an estimator to certain characteristics of the underlying population distribution.
- When the distribution function has a PDF, the population version of density ranking is defined as:

$$\alpha(x) = P(p(x) \ge p(X_1)).$$

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- Let C_d be the volume of a d dimensional unit ball and $B(x, r) = \{y : ||x y|| \le r\}.$
- For any integer *s*, we define

$$\mathcal{H}_s(x) = \lim_{r \to 0} \frac{P(B(x, r))}{C_s r^s}.$$

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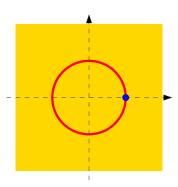
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• For a point x, we then define

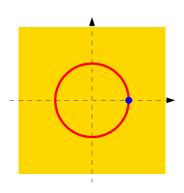
$$\tau(x) = \max\{s \le d : \mathcal{H}_s(x) < \infty\}, \quad \rho(x) = \mathcal{H}_{\tau(x)}(x).$$

Hausdorff Density: Example - 1

• Assume the distribution function P is a mixture of a 2D uniform distribution within $[-1,1]^2$, a 1D uniform distribution over the ring $\{(x,y): x^2 + y^2 = 0.5^2\}$, and a point mass at (0.5,0), then the support can be partitioned as follows:



Geometric Hausdorff: Example - 2



- Orange region: $\tau(x) = 2$.
- Red region: $\tau(x) = 1$.
- Blue region: $\tau(x) = 0$.

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- \circ We can use τ and ρ to compare any pairs of points and construct a ranking.
- For two points x_1, x_2 , we define an ordering such that $x_1 \succ_{\tau, \rho} x_2$ if

$$\tau(x_1) < \tau(x_2), \quad \text{or} \quad \tau(x_1) = \tau(x_2), \quad \rho(x_1) > \rho(x_2).$$

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$$\tau(x_1) < \tau(x_2),$$
 or $\tau(x_1) = \tau(x_2),$ $\rho(x_1) > \rho(x_2).$

 Namely, we first compare the dimension of the two points, the lower dimensional structure wins. If they are on regions of the same dimension, we then compare the density of that dimension.

Constructing Density Ranking using Hausdorff Density

• Using the ordering $\succ_{\tau,\rho}$, we then define the population density ranking as

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• Using the ordering $\succ_{\tau,\rho}$, we then define the population density ranking as

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• When the PDF exists, the ordering $\succ_{\tau,\rho}$ equals to $\succ_{d,p}$ so

$$\alpha(x) = P(x \ge_{d,p} X_1) = P(p(x) \ge p(X_1)),$$

which recovers our original definition.

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- To generalize density trees, we use the cluster tree of density ranking.
- We call this tree the ranking tree.
- Formally, the ranking tree is the set

$$T_{\alpha} = \bigcup_{\lambda} \{A_{\lambda,1}, \cdots, A_{\lambda,J(\lambda)}\}$$

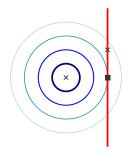
where

$$A_{\lambda,1},\cdots,A_{\lambda,J(\lambda)}$$

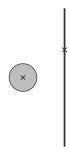
are the connected components of the λ -level set of $\alpha(x)$.

- In singular measure, there is a new type of critical points. We call them the *dimensional critical points*.
- These critical points contribute to the change of topology of level sets as the usual critical points but they cannot be defined by setting gradient to be 0.

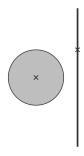
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- Note: this is a mixture of 2D distribution and a 1D distribution on the black line (maximum value occurs at the cross).



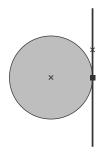
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- However, the speed of diverging depends on $\tau(x)$. The smaller $\tau(x)$, the faster (actually the diverging rate is $O(h^{\tau(x)-d})$).
- So eventually, we can separate different dimensional structures.

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- Despite the pointwise convergence and convergence in $L_2(P)$, there no guarantee for the uniform convergence $\sup_x |\widehat{\alpha}(x) \alpha(x)|$.
- Example of non-convergence of supreme norm: consider a sequence of points on a higher dimensional space but moving toward a lower dimensional space within distance $\frac{h}{2}$.

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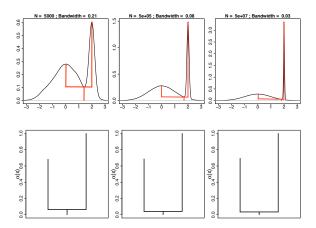
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- Although we do not have uniform convergence, we can still recover the topology of the tree.
- In addition, the height of each branch of the tree will also converge.

Density Ranking and Cluster Tree: Example

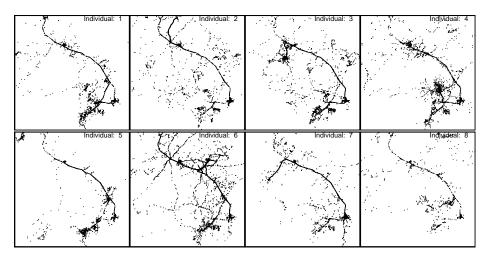
Here the population distribution function is a mixture of a 1*D* standard normal distribution and a point mass at 2. We consider three sample sizes: $n = 5 \times 10^3$, 5×10^5 , 5×10^7 .



DENSITY RANKING: MULTIPLE

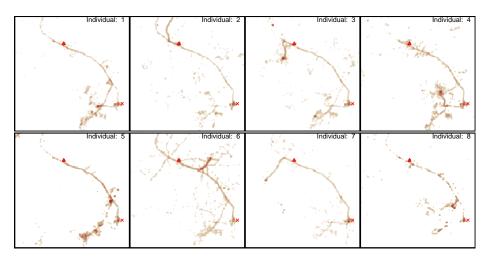
DATASETS

Application of Density Ranking: GPS dataset - 1



Joint work with Adrian Dobra and Zhihang Dong.

Application of Density Ranking: GPS dataset - 2



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- In the above example, we have multiple GPS datasets and each of them yields one density ranking.
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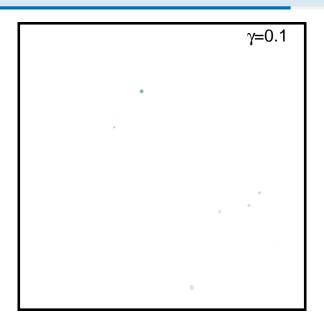
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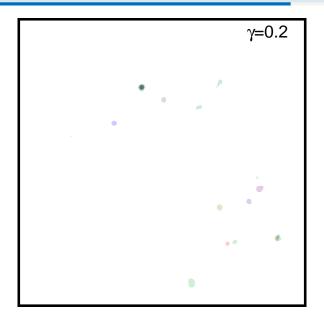
 We can compare the density ranking of each individual by overlapping their level sets at different levels.

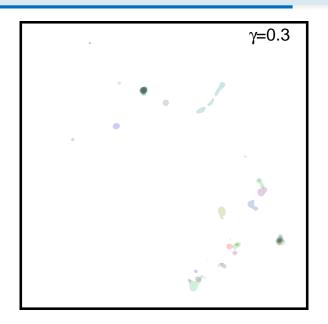
- Note that we use 1γ as the level in the set \widehat{A}_{γ} .
- This is because such a set has a natural interpretation in activity space.
- Activity space: the spatial regions where an individual undertakes his/her daily life.

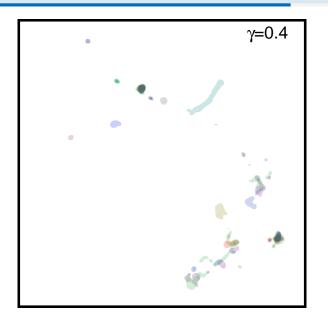
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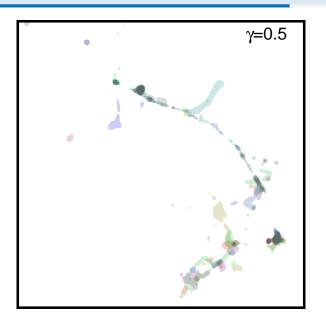
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- We can interpret \widehat{A}_{γ} as the (top) $\gamma \cdot 100\%$ activity space because they are regions containing at least $\gamma \cdot 100\%$ GPS records.
- Namely, $\widehat{A}_{\gamma=0.3}$ is the (top) 30% activity space.

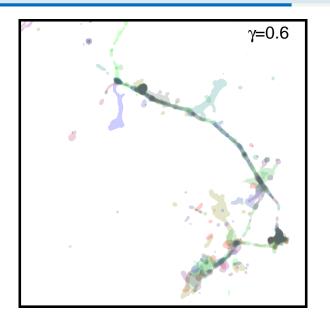


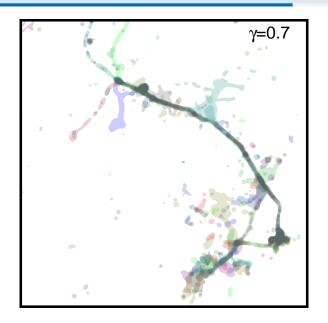


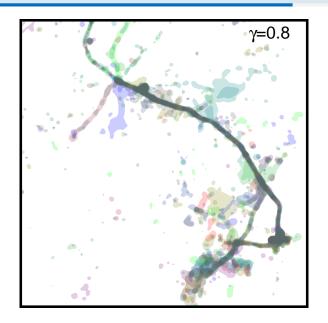


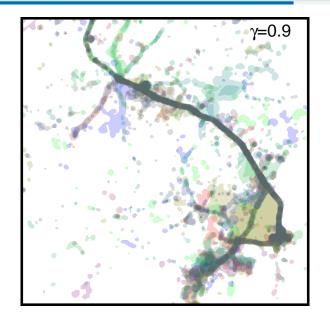


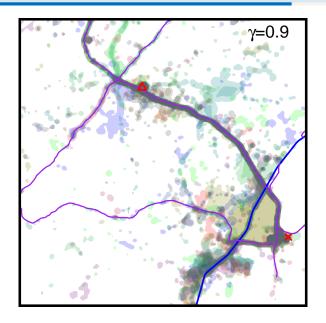












Summary Curves of Density Ranking

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- However, it has two drawbacks:
 - When we have many individuals, this approach might not work (too many contours).
 - We often need to choose a level γ to show the plot but which level to be chosen is unclear.
- Here we introduce a few curves to summarize geometric and topological features of density ranking.

Mass-Volume Curve

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• Namely, we are plotting the size of set \widehat{A}_{γ} at various level.

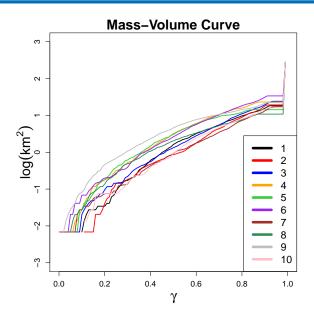
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- Namely, we are plotting the size of set \widehat{A}_{γ} at various level.
- In practice, we often plot γ versus $\log Vol(\widehat{A}_{\gamma})$.

Mass-Volume Curve: Example



Betti Number Curve

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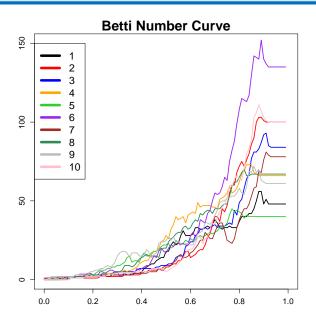
$$\left(\gamma, \mathsf{Betti}_0(\widehat{A}_\gamma)\right) : \gamma \in [0,1],$$

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 Note that the number of connected component is called the oth order Betti number (oth order topological structure); one can generalize this idea to higher order topological structures.

Betti Number Curve: Example

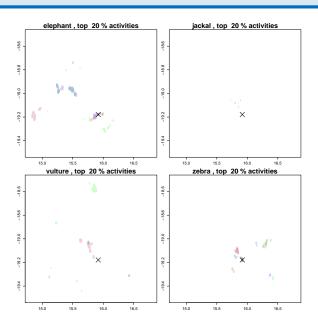


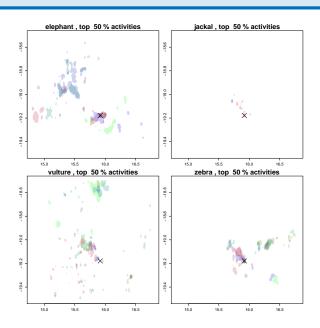
Applying to African Animal Datasets

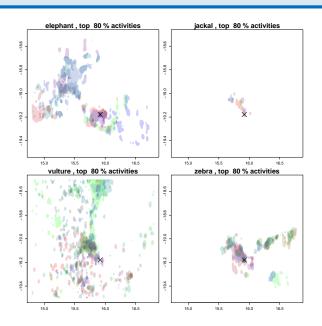
- We apply our methods to a GPS data about African animals.
- This data is from the Movebank Data Repository¹ and was analyzed in Abrahms et al. (2017).
- Here we compare 4 different types of animals: elephants, jackals, vultures, and zebras.
- In this data, we have 8 elephants, 15 jackals, 10 vultures, and 9 zebras.
- Each animal has a set of GPS records.

https://www.datarepository.movebank.org/

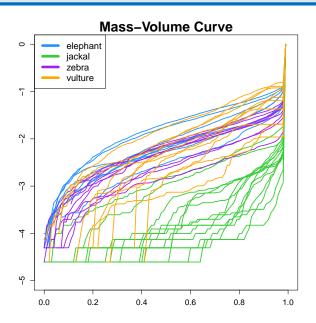




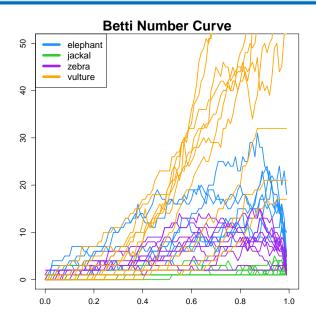




Mass-Volume Curve: Animal Example



Betti Number Curve: Animal Example



SUMMARY

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- But we can use density ranking to analyze data.
- Density ranking defines a ranking tree that act as a density tree.
- When multiple GPS datasets are available, we can summarize them by functional summaries of density ranking.

Thank You!

An R script for density ranking: https://github.com/yenchic/density_ranking

More details can be found in http://faculty.washington.edu/yenchic/

References

- Chen, Yen-Chi, Christopher R. Genovese, and Larry Wasserman. "Density level sets: Asymptotics, inference, and visualization." Journal of the American Statistical Association (2017): 1-13.
- Jisu, K. I. M., Yen-Chi Chen, Sivaraman Balakrishnan, Alessandro Rinaldo, and Larry Wasserman. "Statistical inference for cluster trees." In Advances In Neural Information Processing Systems, pp. 1839-1847. 2016.
- 3. Chen, Yen-Chi. "Generalized Cluster Trees and Singular Measures." arXiv preprint arXiv:1611.02762 (2016).
- Chen, Yen-Chi, and Adrian Dobra. "Measuring Human Activity Spaces With Density Ranking Based on GPS Data." arXiv preprint arXiv:1708.05017 (2017).
- Stuetzle, Werner. "Estimating the cluster tree of a density by analyzing the minimal spanning tree of a sample." Journal of classification 20, no. 1 (2003): 025-047.
- Klemelä, Jussi. "Visualization of multivariate density estimates with level set trees." Journal of Computational and Graphical Statistics 13, no. 3 (2004): 599-620.
- Chaudhuri, Kamalika, and Sanjoy Dasgupta. "Rates of convergence for the cluster tree." In Advances in Neural Information Processing Systems, pp. 343-351. 2010.
- Chaudhuri, Kamalika, Sanjoy Dasgupta, Samory Kpotufe, and Ulrike von Luxburg. "Consistent procedures for cluster tree estimation and pruning." IEEE Transactions on Information Theory 60, no. 12 (2014): 7900-7912.
- Eldridge, Justin, Mikhail Belkin, and Yusu Wang. "Beyond hartigan consistency: Merge distortion metric for hierarchical clustering." In Conference on Learning Theory, pp. 588-606. 2015.
- Balakrishnan, Sivaraman, Srivatsan Narayanan, Alessandro Rinaldo, Aarti Singh, and Larry Wasserman. "Cluster trees on manifolds." In Advances in Neural Information Processing Systems, pp. 2679-2687. 2013.
- 11. Abrahms B, Seidel DP, Dougherty E, Hazen EL, Bograd SJ, Wilson AM, McNutt JW, Costa DP, Blake S, Brashares JS, Getz
 - WM~(2017)~"Suite~of~simple~metrics~reveals~common~movement~syndromes~across~vertebrate~taxa."~Movement~Ecology~compared to the compared to t
 - 5:12. doi:10.1186/s40462-017-0104-2

Assumptions for Regular Distributions

- **(R1)** The density function p has a compact support \mathbb{K} .
- (R2) The density function is a Morse function and is in BC^3 .
- **(K1)** The kernel function K is in \mathbf{BC}^2 and integrable.
- **(K2)** *K* satisfies the VC-type class condition.

Kernel Conditions

(K₂) Let

$$\mathcal{K}_r = \left\{ y \mapsto K^{(\alpha)} \left(\frac{x-y}{h} \right) : x \in \mathbb{R}^d, |\alpha| = r \right\},\,$$

where $K^{(\alpha)}$ is the α -th derivative and let $\mathcal{K}_l^* = \bigcup_{r=0}^l \mathcal{K}_r$. We assume that \mathcal{K}_2^* is a VC-type class. i.e. there exists constants A, v and a constant envelope b_0 such that

$$\sup_{Q} N(\mathcal{K}_{2}^{*}, \mathcal{L}^{2}(Q), b_{0}\epsilon) \leq \left(\frac{A}{\epsilon}\right)^{v}, \tag{1}$$

where $N(T, d_T, \epsilon)$ is the ϵ -covering number for an semi-metric set T with metric d_T and $\mathcal{L}^2(Q)$ is the L_2 norm with respect to the probability measure Q.

Assumptions for Singular Distributions

(S1) The support can be partitioned into

$$K=K_0\bigcup K_1\bigcup\cdots\bigcup K_d,$$

where $K_{\ell} = \{x \in \mathbb{K} : \tau(x) = \ell\}.$

- **(S2)** There exist ρ_{\min} , ρ_{\max} such that $0 < \rho_{\min} \le \rho(x) \le \rho_{\max} < \infty$ for every $x \in \mathbb{K}$.
- **(S₃)** Restricted to each \mathbb{K}_{ℓ} where $\ell > 0$, $\rho(x)$ is a Morse function.
- **(K1')** The kernel function K is in \mathbf{BC}^2 , integrable, and supported in [-1,1].
- **(K2)** K satisfies the VC-type class condition.