## Computing logarithmic Voronoi cells

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## Logarithmic Voronoi cells

- A probability simplex is defined as

$$
\Delta_{n-1}=\left\{\left(p_{1}, \ldots, p_{n}\right): p_{1}+\cdots+p_{n}=1, p_{i} \geq 0 \text { for } i \in[n]\right\}
$$



- A statistical model is a subset of $\Delta_{n-1}$. An algebraic statistical model is a subset $\mathcal{M}=\mathcal{V} \cap \Delta_{n-1}$ for some variety $\mathcal{V} \subseteq \mathbb{C}^{n}$.
- For an empirical data point $u=\left(u_{1}, \ldots, u_{n}\right) \in \Delta_{n-1}$, the log-likelihood function defined by $u$ assuming distribution $p=\left(p_{1}, \ldots, p_{n}\right) \in \mathcal{M}$ is

$$
\ell_{u}(p)=u_{1} \log p_{1}+u_{2} \log p_{2}+\cdots+u_{n} \log p_{n} .
$$

## Maximum likelihood estimation

Fix an algebraic statistical model $\mathcal{M} \subseteq \Delta_{n-1}$
(1) The maximum likelihood estimation problem (MLE):

Given a sampled empirical distribution $u \in \Delta_{n-1}$, which point $p \in \mathcal{M}$ did it most likely come from? In other words, we wish to maximize $\ell_{u}(p)$ over all points $p \in \mathcal{M}$.

## Maximum likelihood estimation

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(2) Computing logarithmic Voronoi cells:

Given a point $q \in \mathcal{M}$, what is the set of all points $u \in \Delta_{n-1}$ that have $q$ as a global maximum when optimizing the function $\ell_{u}(p)$ over $\mathcal{M}$ ?

The set of all such elements $u \in \Delta_{n-1}$ is the logarithmic Voronoi cell at $q$.

## Proposition (A., Heaton)

Logarithmic Voronoi cells are convex sets.
The log-normal space at $q$ is the space of possible data points $u \in \mathbb{R}^{n}$ for which $q$ is a critical point of $\ell_{u}(p)$. It is a linear space.

Intersecting this space with the simplex $\Delta_{n-1}$, we obtain a polytope, which we call the log-normal polytope at $q$.

The log-normal polytope at $q$ contains the logarithmic Voronoi cell at $q$.

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## Example (The twisted cubic.)

The curve is given by $p \mapsto\left(p^{3}, 3 p^{2}(1-p), 3 p(1-p)^{2},(1-p)^{3}\right)$.


## The Hardy-Weinberg curve

Consider a model parametrized by

$$
p \mapsto\left(p^{2}, 2 p(1-p),(1-p)^{2}\right) .
$$

Performing implicitization, we find that the model $\mathcal{M}=\mathcal{V}(f)$ where

$$
f=\left[\begin{array}{c}
4 x_{1} x_{3}-x_{2}^{2} \\
x_{1}+x_{2}+x_{3}-1
\end{array}\right]
$$

The augmented Jacobian is given by:

$$
A=\left[\begin{array}{ccc}
4 x_{3} & -2 x_{2} & 4 x_{1} \\
1 & 1 & 1 \\
u_{1} / x_{1} & u_{2} / x_{2} & u_{3} / x_{3}
\end{array}\right]
$$

Fix a point $q \in \mathcal{M}$ and substitute $x_{i}$ for $q_{i}$ in $A$. All points $u \in \mathbb{R}^{3}$ at which the determinant vanishes define the log-normal space at $q$.

## The Hardy-Weinberg curve

$\operatorname{det} A=4 u_{1}-4 u_{3}-4 u_{2} \cdot \frac{x_{1}}{x_{2}}+2 u_{1} \cdot \frac{x_{2}}{x_{1}}-2 u_{3} \cdot \frac{x_{2}}{x_{3}}+4 u_{2} \cdot \frac{x_{3}}{x_{2}}$
For example, at $p=0.2$, we get a point $q=(0.04,0.32,0.64) \in \mathcal{M}$. The log-normal space at $q$ is the plane

$$
20 u_{1}+7.5 u_{2}-5 u_{3}=0 .
$$

Sampling more points, we get the following pictures:

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Log-normal polytopes $=$ Log-Voronoi cells

## Polytopal cells

The maximum likelihood degree (ML degree) of $\mathcal{M}$ is the number of complex critical points when optimizing $\ell_{u}(x)$ over $\mathcal{M}$ for generic data $u$.

Theorem (A., Heaton)
If $\mathcal{M}$ is a finite model, a linear model, a toric model, or a model of ML degree 1, the logarithmic Voronoi cell at any point $p \in \mathcal{M}$ is equal to the log-normal polytope at $p$.

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What about logarithmic Voronoi cells that are not polytopes?

## Numerical algebraic geometry

Logarithmic Voronoi cells that are not polytopes are mysterious. Are they semialgebraic sets? How to describe their boundary?

## Conjecture

The boundary of logarithmic Voronoi cells are, in general, not algebraic.
Numerical algebraic geometry provides an alternative approach to logarithmic Voronoi cells for more complicated models.

## Example (Mixture model)

Consider the 3-dimensional model in $\Delta_{5}$ parametrized as:

$$
(a, s, t) \mapsto\left(a s^{5}+(1-a) t^{5}, \ldots, a(1-s)^{5}+(1-a)(1-t)^{5}\right)
$$

## Mixture model, continued

Implicitly, it is described by

$$
\begin{aligned}
& f_{0}(x)=x_{1}+x_{2}+x_{3}+x_{4}+x_{5}+x_{6}-1 \\
& f_{1}(x)=20 x_{1} x_{3} x_{5}-10 x_{1} x_{4}^{2}-8 x_{2}^{2} x_{5}+4 x_{2} x_{3} x_{4}-x_{3}^{3} \\
& f_{2}(x)=100 x_{1} x_{3} x_{6}-20 x_{1} x_{4} x_{5}-40 x_{2}^{2} x_{6}+4 x_{2} x_{3} x_{5}+2 x_{2} x_{4}^{2}-x_{3}^{2} x_{4} \\
& f_{3}(x)=100 x_{1} x_{4} x_{6}-40 x_{1} x_{5}^{2}-20 x_{2} x_{3} x_{6}+4 x_{2} x_{4} x_{5}+2 x_{3}^{2} x_{5}-x_{3} x_{4}^{2} \\
& f_{4}(x)=20 x_{2} x_{4} x_{6}-8 x_{2} x_{5}^{2}-10 x_{3}^{2} x_{6}+4 x_{3} x_{4} x_{5}-x_{4}^{3} .
\end{aligned}
$$

- Pick a point $p_{\text {fav }}=\left(\frac{518}{9375}, \frac{124}{625}, \frac{192}{625}, \frac{168}{625}, \frac{86}{625}, \frac{307}{9375}\right) \in \mathcal{M}$.
- The log-normal space at $p_{\mathrm{fav}}$ is 3 -dimensional.
- Intersecting with $\Delta_{5}$, we get the 2-dimensional log-normal polytope at $p_{\mathrm{fav}}$, which is a hexagon.


## Mixture model, continued



## Mixture model, continued

- Sample 60000 points $u_{i}$ from within the log-normal hexagon.
- We will color each point green if $p_{\mathrm{fav}}$ is its MLE and pink otherwise.
- Let $d f=\left[\nabla f_{0} \nabla f_{1} \ldots \nabla f_{4}\right]^{T}$ be a $5 \times 6$ matrix.
- Multiply by a random $3 \times 5$ matrix $A$ on the left, then $A \cdot d f$ is a $3 \times 6$ matrix with the same rowspace. This is dimension reduction.
- Add the 4 th row $\nabla \ell_{u}=\left(u_{i} / p_{i}\right)$.
- Introducing new variables $\lambda_{1}, \lambda_{2}, \lambda_{3}$ we will require that some linear combination of the first 3 rows is equal to the last row.
- This gives 6 equations:

$$
G:=\left[\begin{array}{lll}
\lambda_{1} & \lambda_{2} & \lambda_{3}-1
\end{array}\right]\left[\begin{array}{c}
A \cdot d f \\
\nabla \ell_{u}
\end{array}\right] \operatorname{diag}\left(p_{i}\right)=0
$$

- In addition, we add the original 5 equations $f_{0}, f_{1}, f_{2}, f_{3}, f_{4}$, so we have 11 equations in 9 variables (rather than 225 minors in 6 variables).


## HomotopyContinuation.jl and monodromy

We could start with a random point $u_{\text {start }}$, solve the system, and use these starting solutions to track solutions for the 60000 points $u_{i}$. But solve(G) reports that we have to track 110592 paths... Use monodromy!

## HomotopyContinuation.jl and monodromy

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- The ML degree of our model is 39 .
- Start with a random $u_{0}$ in the hexagon; we know one solution: $p_{\text {fav }}$ !
- Perturb our system of equations around a loop in parameter space, tracking this solution along the way.
- When we return to the original system, the original solution likely has moved to another one of the 39 solutions. Now we have two!
- Repeating this process, we discover all 39 solutions.
- For the 60000 other parameter points $u_{i}$, perturb the system of equations at $u_{0}$, and track all solutions along the way. This way we can find the 39 solutions for each sample point in the hexagon.
- We check which one of them maximizes log-likelihood. If it is $p_{\mathrm{fav}}$, we color it green. If not, we color it pink.


## In general

Theorem (A., Heaton)
Let $\mathcal{M}$ be an irreducible algebraic model in $\Delta_{n-1}$ described implicitly by $m$ polynomials $f$. Let $u \in \Delta_{n-1}$ be fixed and generic. With probability 1, all points $p \in \mathcal{M}$ such that $u \in \log N_{p} \mathcal{M}$ are among the finitely many isolated solutions to the square system

$$
\underbrace{\left[\begin{array}{l}
{[\lambda-1]}
\end{array}\left[\begin{array}{c}
A \cdot d f \\
\nabla \ell_{u}
\end{array}\right] f\right]}_{1 \times(n+m)}\left[\begin{array}{c}
I_{n+c} \\
B
\end{array}\right]=\underbrace{[0 \cdots 0]}_{1 \times(n+c)} .
$$

## Thanks!

