

# *CryoEM with Spider Kernel Graphs*

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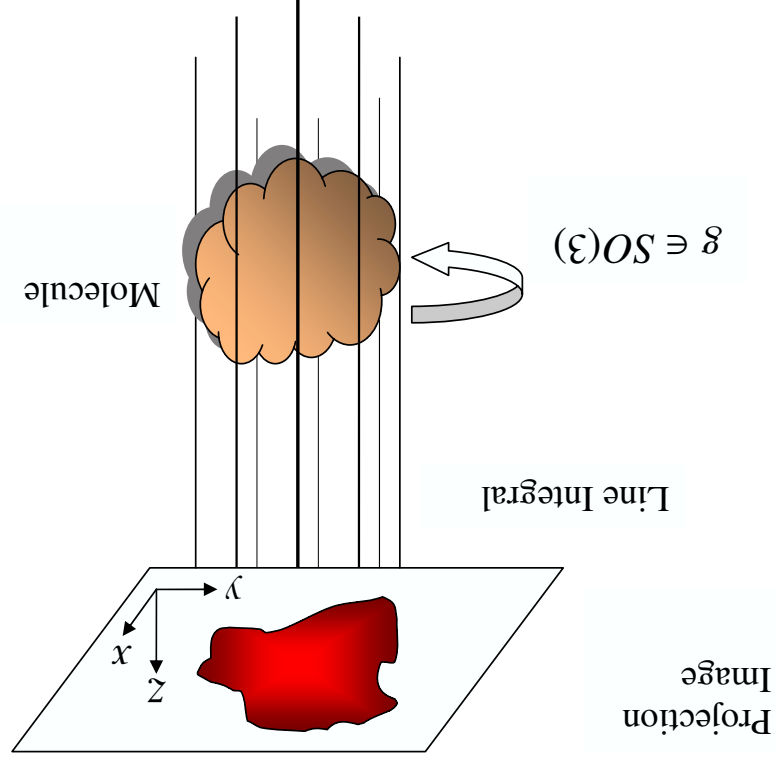
## *Structuring of Protein Channels*

- Rod MacKinnon was co-awarded the 2003 Nobel Prize in Chemistry for structuring the Potassium channel in 1998.
- Proteins were crystallized (all share the same space orientation).
- Classical X-ray Computational Tomography (CT).
- A few other proteins had been structured since.
- However, most channels cannot be crystallized.
- Can a protein be structured without being crystallized?

## *Cryo Electron Microscopy*

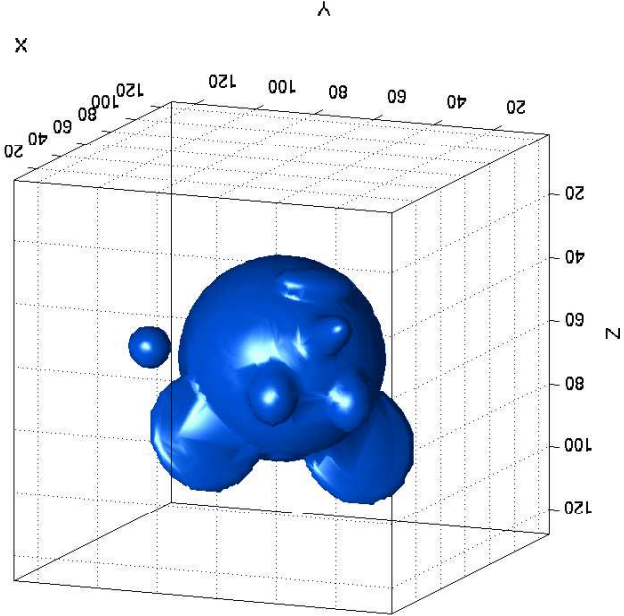
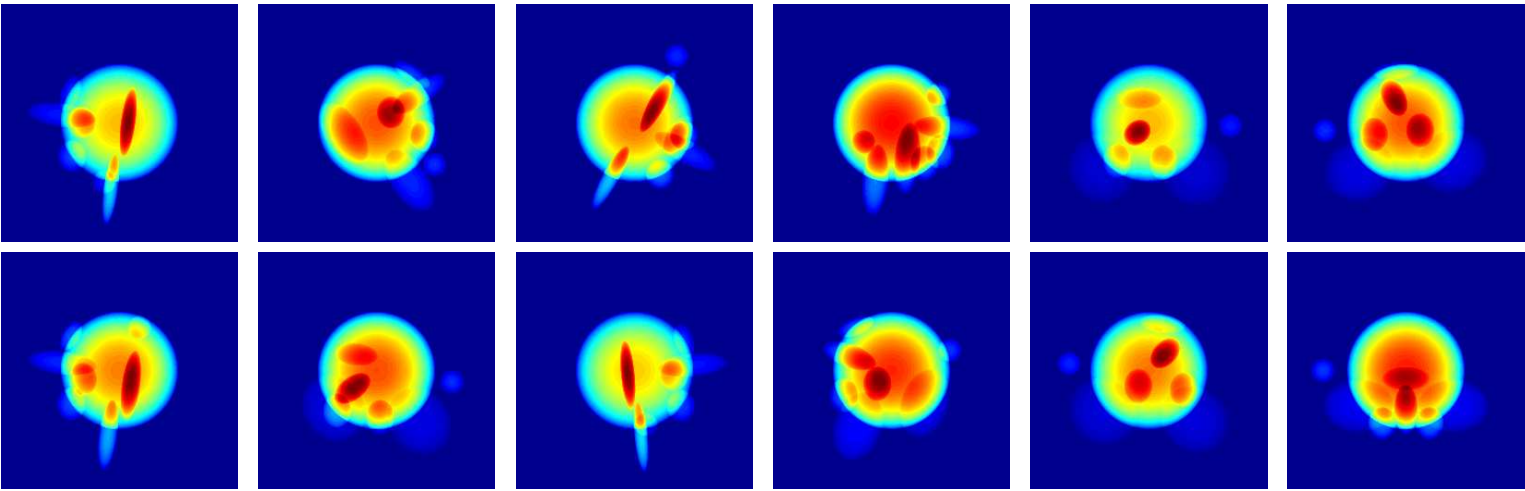
- CryoEM: Electron Microscope imaging of proteins “frozen” in liquid nitrogen.
- Thousands of images: every image corresponds to a different protein frozen in a different space orientation.
- Orientations are random and unknown.
- Highly intense electron beam destroys protein while being imaged: a single protein can be imaged only once.
- Images are very noisy (low SNR)
- Images are  $100 \times 100$  pixels.

## Projection Images



- The projection image is  $P_g(x, y) = \int_{-\infty}^{\infty} \phi^g(x, y, z) dz$ .

- $\phi(r)$  is the electric potential of the molecule,  $\phi^g(r) = \phi(g^{-1}r)$ .



*Projection Images: Toy Example*

## *The Fourier projection-slice theorem*

- $\theta \in S^2$  beaming direction,  $\theta_\perp$  orthogonal plane.

- The 2D FT of the projection image is the double integral

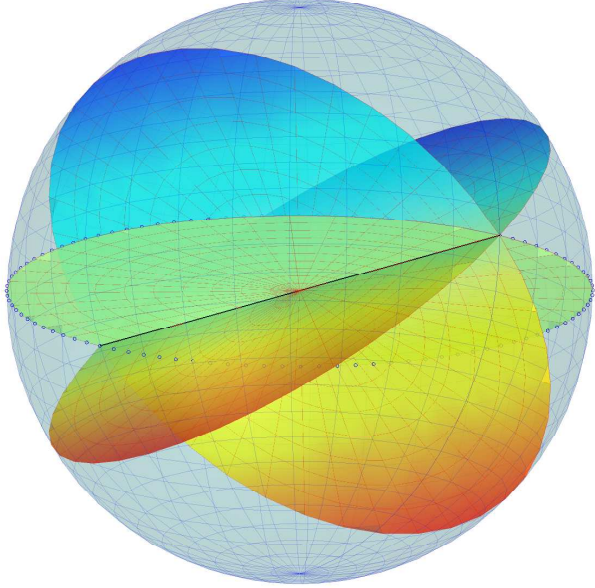
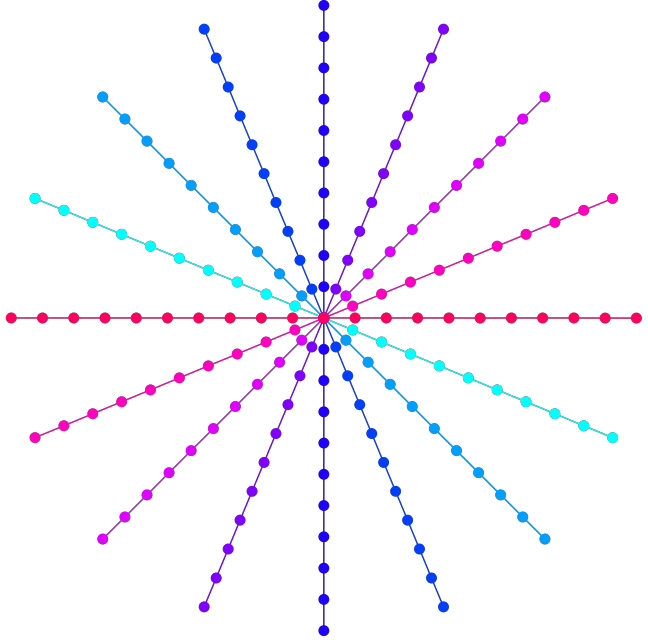
$$\hat{P}_\theta(\xi) = \int_{\theta_\perp} e^{-ir \cdot \xi} P_\theta(r) dr.$$

- The 3D FT of the molecule is the triple integral

$$\hat{\phi}(\xi) = \int_{\mathbb{R}^3} e^{-ir \cdot \xi} \phi(r) dr.$$

- Slice Theorem:  $\hat{P}_\theta(\eta) = \hat{\phi}(\eta)$ ,  $\eta \in \theta_\perp$ .

- Every image is a great circle over  $S^2$ .
- Any pair of images have a common line, or
- Any pair of great circles meet at two antipodal points.



## *The Geometry of the slice theorem*



- The radial lines are the puzzle pieces.
- Every image is a circular chain of pieces.
- Common line: meeting point



*Three Dimensional Puzzle*

## *The Spider Kernel: It's the Network*

- $K$  projection images
- $L$  radial lines

- We build a weighted directed graph  $G = (V, E, W)$ .

- The vertices are the radial lines ( $|V| = KL$ )

$$V = \{(k, l) : 1 \leq k \leq K, 0 \leq l \leq L - 1\}.$$

- The heart of the algorithm is the definition of arrows and weights

$$E = \{(k_1, l_1), (k_2, l_2) : (k_1, l_1) \text{ points to } (k_2, l_2)\}.$$

- $W$  is a sparse weight matrix of size  $KL \times KL$

$$((k_1, l_1), (k_2, l_2)) \notin E \iff W_{(k_1, l_1), (k_2, l_2)} = 0.$$

## *Weights*

- All weights are taken from a single (sparse) symmetric circular weight vector of length  $L$

$$w = (w_0, w_1, \dots, w_{L-1}),$$

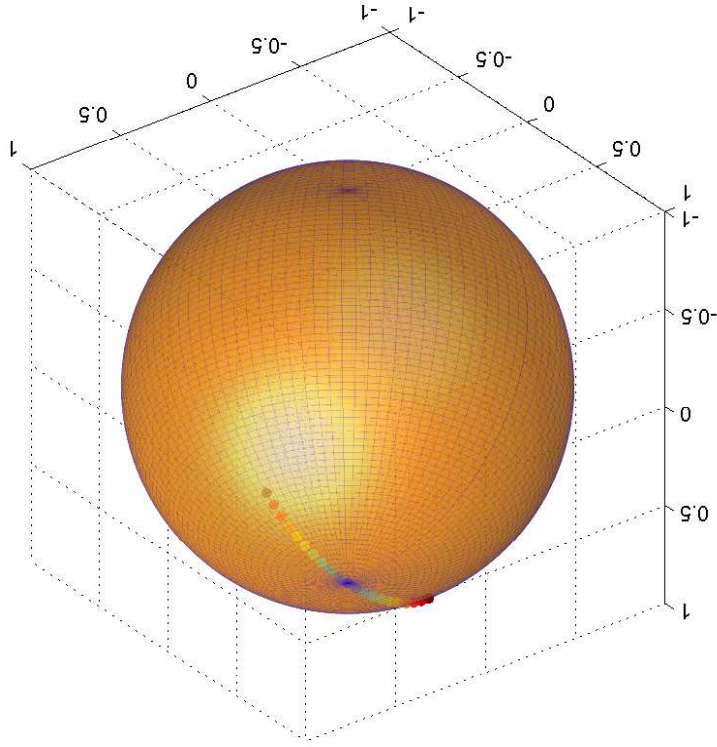
$$w_l = w_{-l}.$$

- Example:

$$w = (1, 1, \dots, 1) = \mathbf{1}$$

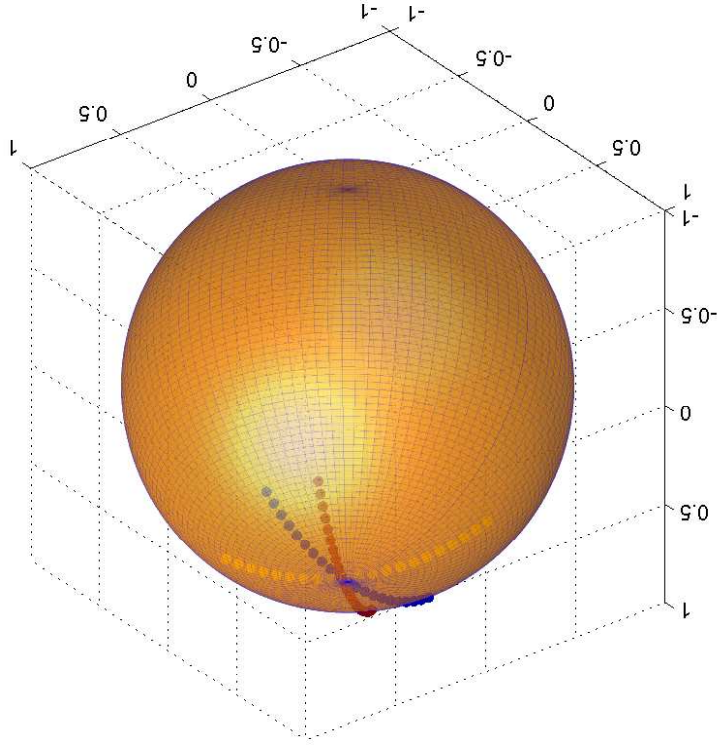
renders  $W$  the adjacency matrix of the graph.

- Blue vertex  $(k_1, l_1)$  is the head of the spider
- Linked vertices:  $(k_1, l_1 + l)$ ,  $-d \leq l \leq d$  (same image radial lines)
- Weights:  $W_{(k_1, l_1), (k_1, l_1 + l)} = w_l \cdot$



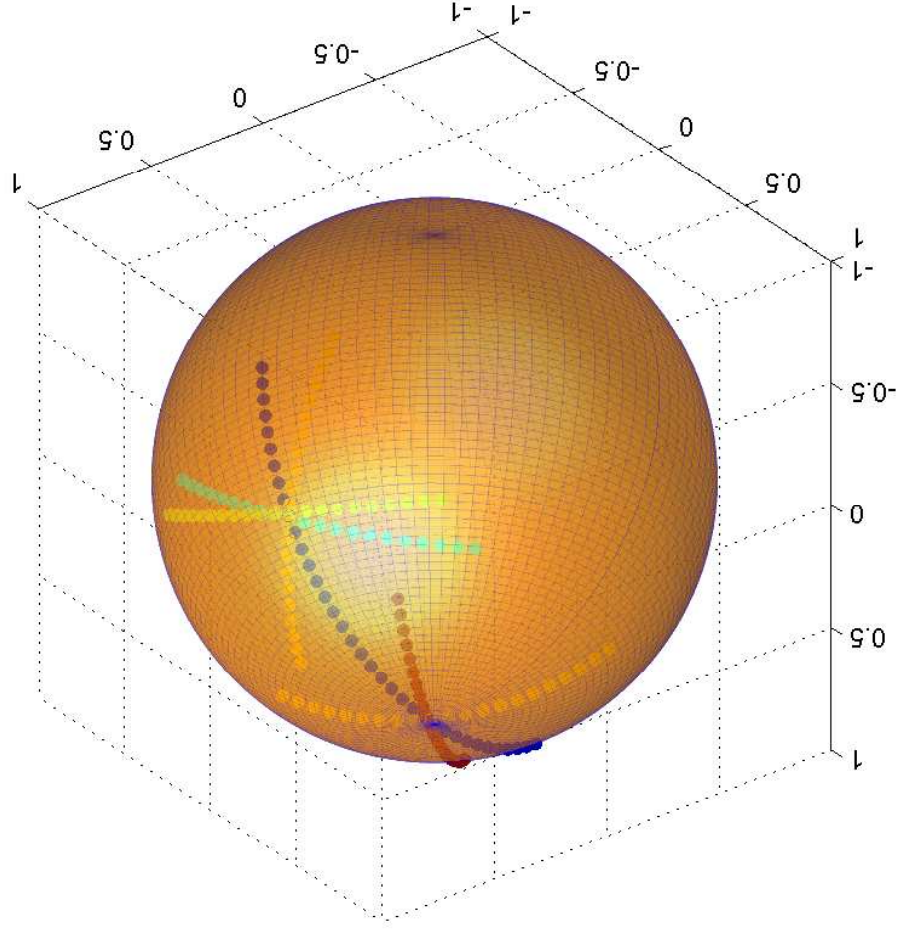
*Spider first pair of legs*

- Weights:  $W^{(k_1, l_1), (k_2, l_2 + l)} = w_l$ .
- Links:  $((k_1, l_1), (k_2, l_2 + l)) \in E$  for  $-d \leq l \leq d$ .
- $(k_1, l_1)$  and  $(k_2, l_2)$  are common radial lines of different images.



*Spider: remaining legs*

# *Communicating Spiders*



## *Sparse weight matrix*

- $W$  is sparse: its number of nonzero entries is only  $|E| = (2d + 1)[KL + 2K(K - 1)]$ .
- There are  $KL$  spiders with first pair legs of size  $2d + 1$ .
- There are  $2\binom{K}{2} = K(K - 1)$  intersection points (with antipodals).  
Every meeting point belongs to two different circles so it appears in two different spiders.
- In every spider it contributes two legs of total length  $2d + 1$ .
- Algorithm is linear in number of lines and intersection points for  $d = O(1)$  (small spiders).

$$\sum_{k',l' \in V} A_{(k,l),(k',l')} = \sum_{k,l \in V} \frac{1}{1+2d} w_{k,l}$$

- The row sums of  $A$  are identical and equal
- $A = D^{-1}W$ , with  $D$  diagonal  $D_{(k,l),(k,l)} = d_{k,l}$   
by dividing each row by its outdegree:
- We normalize the weight matrix  $W$  to have constant row sums

$$d_{k,l} = |\{(k',l') : ((k,l),(k',l')) \in E\}| = M_{k,l}(2d+1).$$

- The outdegree  $d_{k,l}$  of the  $(k,l)$ 'th vertex is

$$\sum_{k',l' \in V} W_{(k,l),(k',l')} = \sum_{k,l} w_{k,l} M_{k,l}$$

- Row sums of  $W$  depend on the number of legs  $M_{k,l}$

### *Averaging operator*



## *Averaging operator*

- $A$  is a spider weighted averaging operator

$$(Af)(k_1, l_1) = \sum_{((k_1, l_1), (k_2, l_2)) \in E} A_{((k_1, l_1), (k_2, l_2))} f(k_2, l_2).$$

- Example:  $w = (1, 1, \dots, 1) = \mathbf{1}$

$A$  is row stochastic, weighted average = non-weighted average

$$(Af)(k_1, l_1) = \frac{1}{d_{k,l}} \sum_{((k_1, l_1), (k_2, l_2)) \in E} f(k_2, l_2).$$

- We call  $A$  the spider kernel.

## *The spectrum of the spider kernel*

- $A$  and  $W$  are not symmetric, their spectrum may be complex.
- $A$  has constant row sums:  $\psi_0 = 1$  is a trivial eigenvector

$$(A\psi_0)(k, l) = \left( \frac{1}{2d+1} \sum_{l=-d}^d w_l \right) \psi_0(k, l), \quad A(k, l) \in V.$$

- Example:  $w = (1, 1, \dots, 1) = \mathbf{1}$   
 $A$  is row stochastic,  $\lambda_0 = 1$ , remaining spectrum  $|\lambda| < 1$ .
- Much more can be said on the spectrum!

## Spherical Harmonics

- The spherical harmonics  $Y_m^l$  are the eigenfunctions of the Laplacian on the sphere

$$\Delta_{S^2} Y_m^l = -l(l+1) Y_m^l, \quad l = 0, 1, 2, \dots, \quad m = -l, \dots, l.$$

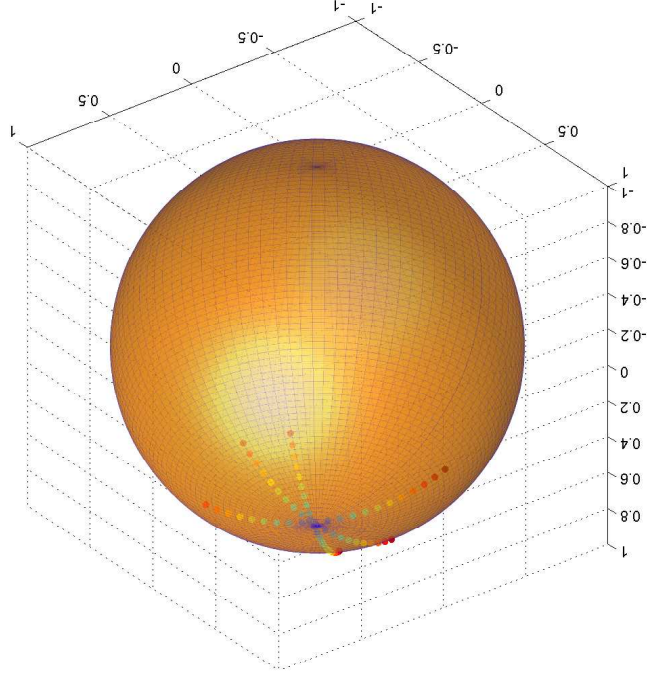
- Funk-Hecke: The spherical harmonics are the eigenfunctions of any integral operator that commutes with rotations:

$$\begin{aligned} (\mathcal{K}f)(\beta) &= \int_{S^2} k(\langle \beta, \beta' \rangle) f(\beta') dS_{\beta'}, \\ \mathcal{K}Y_m^l &= \lambda_l Y_m^l. \end{aligned}$$

- The spider kernel commutes with rotations only on average, so spherical harmonics are not guaranteed.
- The three linear spherical harmonics are exact eigenfunctions of the spider kernel.

## *Linear Eigenfunctions*

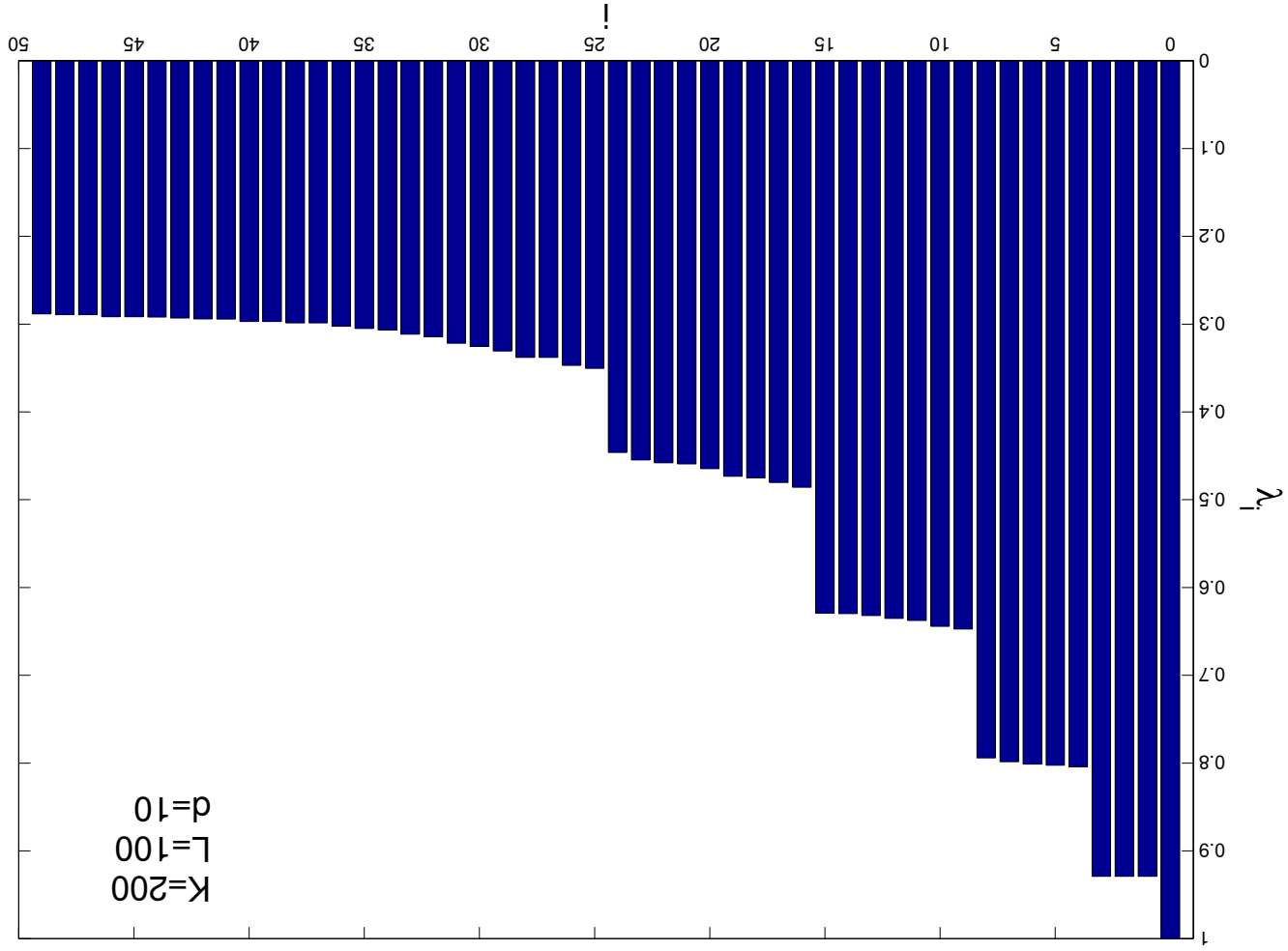
- Linear functions  $f(x, y, z) = a_1x + a_2y + a_3z$  are eigenfunctions
- The center of mass of every spider is beneath the spider's head: any pair of opposite legs balance each other –  $w$  is symmetric.



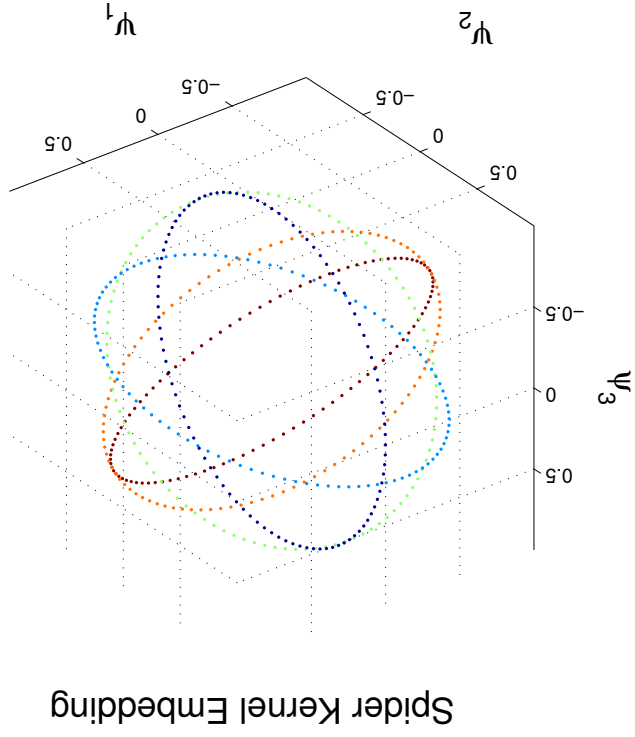
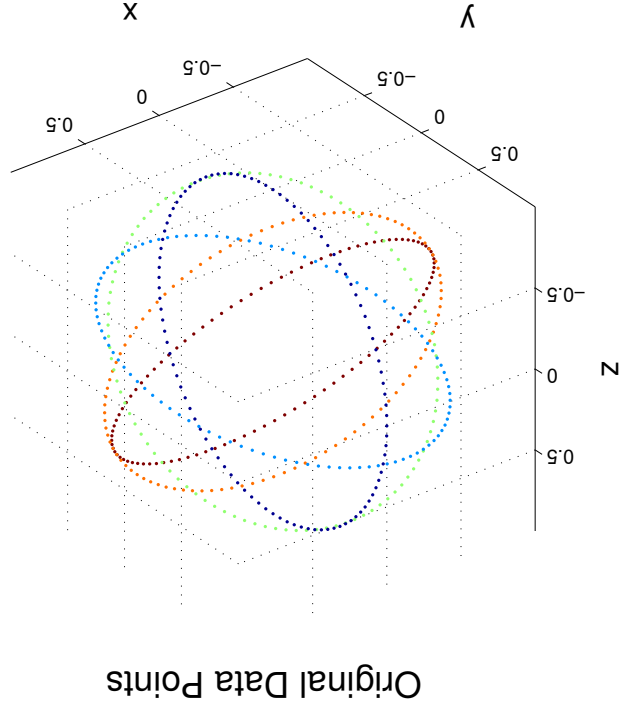
## *Spider kernel embedding and algorithm*

- Find the common lines for all pairs of images.
  - Construct the spider kernel matrix  $A$ .
  - Compute eigenvectors  $A\psi_i = \lambda_i\psi_i$ .
  - Embed the data into the three linear eigenvectors  $(\psi_1, \psi_2, \psi_3)$
- $$(k, l) \mapsto (\psi_1(k, l), \psi_2(k, l), \psi_3(k, l)).$$
- Reveals molecule orientations up to rotation and reflection.
  - Final cosmetics:  
PCA same image radial lines and equally space them.

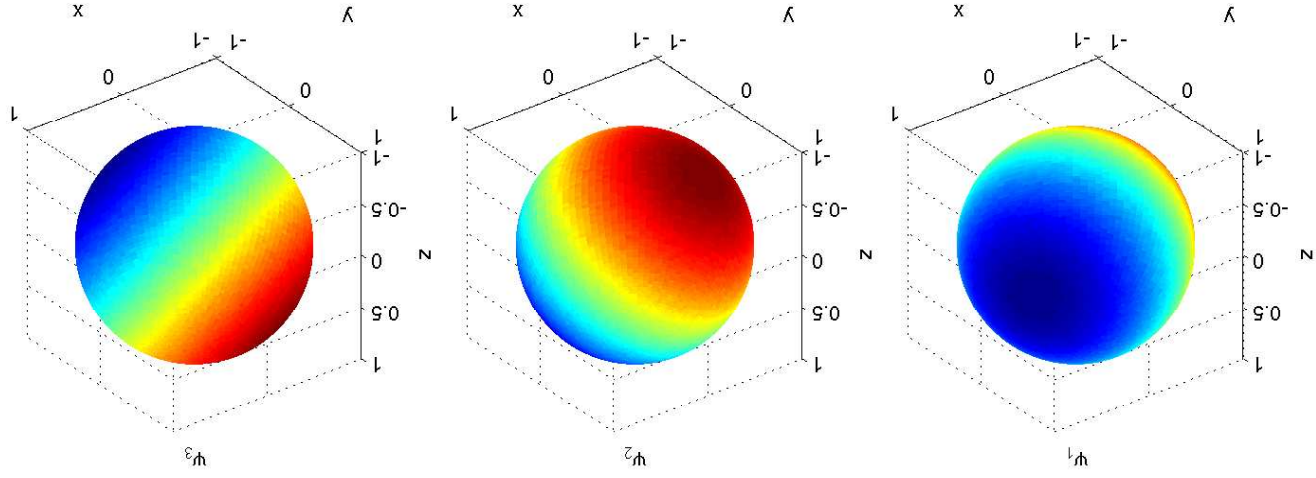
# Numerical Spectrum



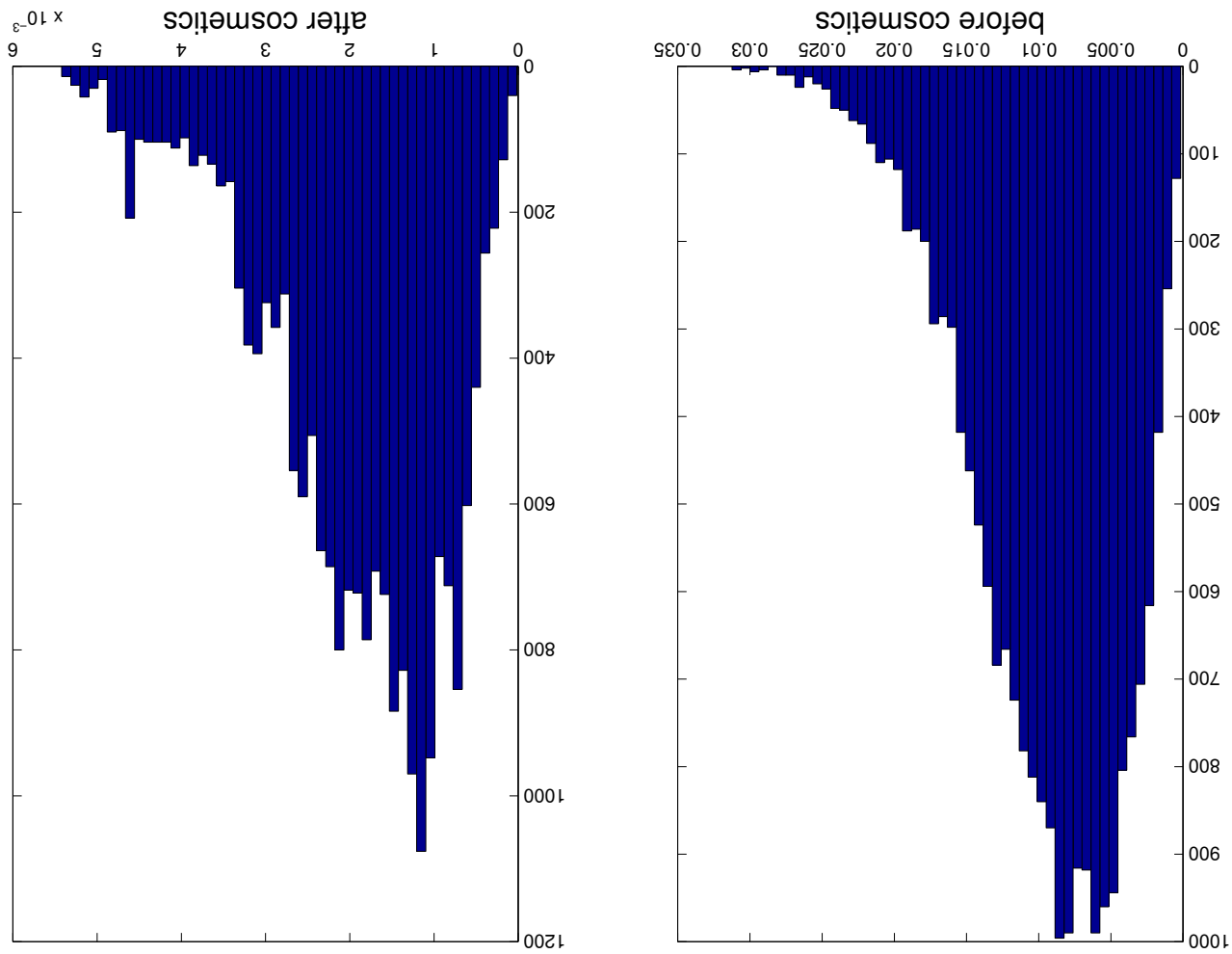
*Data vs. Embedding (only 5 circles are shown)*



*Linear Eigenfunctions*





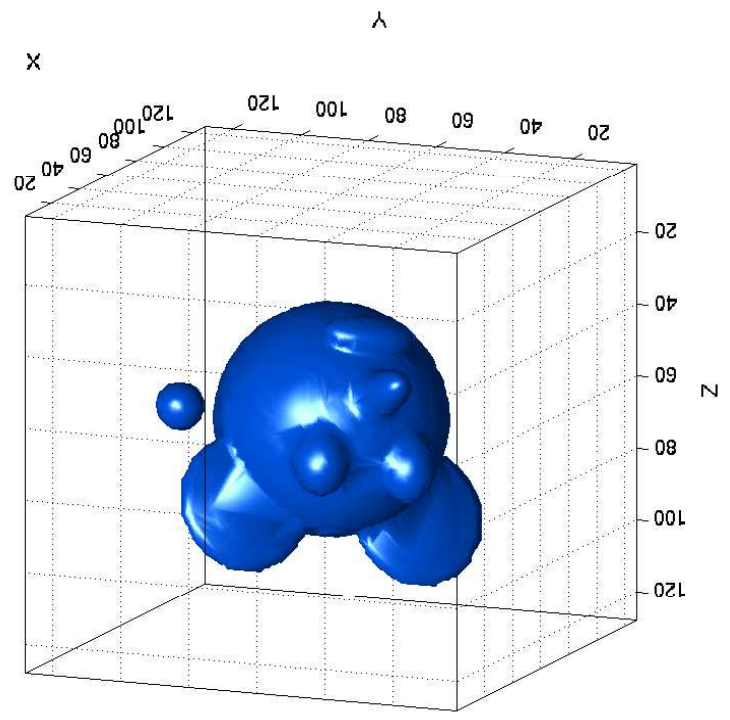


*Angle difference histogram*

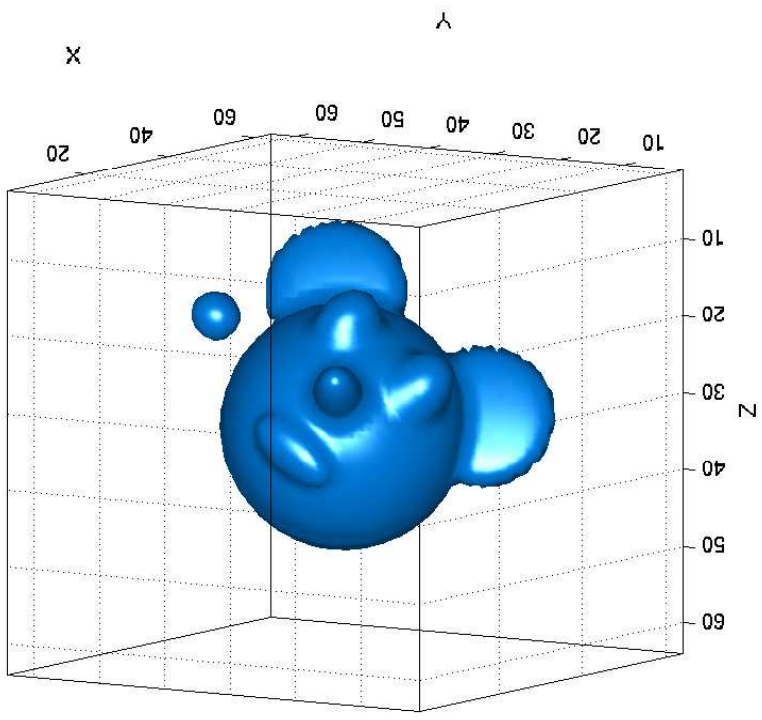
### *Spider kernel advantages*

- Global: all radial lines are linked together.
- Fast: linear in data size  $KL$  and intersection points  $\binom{K}{2}$ .
- Averaging: all geometric information is averaged.
- Robust: errors due to false detections of common lines are smoothed out (can be viewed as matrix perturbation).
- Embedding error decreases like  $1/\sqrt{K}$ .
- Optional: omit uncertain common lines (fewer legs).

(a) original



(b) reconstructed



*Toy Example*