Two Applications of Topological Methods for Neuronal Morphology Analysis

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Joint work with

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Introduction

- Neurons essential to the functioning of life
- Neuronal morphology important in neuron functions
- Understanding 3D morphology of individual neurons
 - Reconstruction from 2D/3D images
 - Characterizing and comparing neuron structures



Based topological methods

Image from https://en.wikipedia.org/wiki/Neuron



Topological methods for:

- Part I:
 - Neuron structures comparison
- Part II:
 - Neuronal Morphology Reconstruction

Neuronal structure 101



Can be considered as a tree structure with augmented information.

Neuron Structures Comparison

- Large number of neuroanatomical data publically available
 e.g, FlyCircuit.org, NeuroMorpho.org
- Efficient algorithms to compare neuron structures
 - E.g. to organize / classify large collection of neurons, to understand variability within a cell type, or to identify features

Related Work

Feature vectorization-based approaches more efficient

better discriminitive power

Alignment and tree distance

L-measure tool

- [Scorcioni et al, 2008]
- Sholl-like analysis
 - [Sholl 1953]
- Arbor density representation
 - [Sümbül et al 2013]
- NBLAST
 - [Costa et al 2016]

Our goal:

- Simple representation to facilitate efficient comparison,
- yet at the same time discriminative, capturing global tree structure

Develop a persistence-based featurevectorization and comparison framework.

Persistence-based feature vectorization framework



A similar persistence-based vectorization method was proposed independently in [Kanari, Dlotko, Scolamiero, Levi, Shillcock, Hess, Markram, arXiv 2016]

Persistence-based feature vectorization framework



Tree representation of neurons

- A set of tree nodes and arcs, where each arc is modeled by a polygonal curve.
- Often assume rooted tree with root r located at soma
- Tree nodes / arc may be associated with other information

Persistence-based feature vectorization framework



- Descriptor function(s) on $T: f: |T| \rightarrow R$
 - Euclidean distance
 - For any $x \in |T|$, f(x) = ||x r||
 - Geodesic distance
 - L-measure based and other morphological descriptors
 - Electrophysiological measures

Persistence-based feature vectorization framework



- Given descriptor function $f: |T| \rightarrow R$
 - Compute the persistence diagram induced by the sub-level set and super-level set filtrations of f as its summary

Persistent Homology 101

- [Edelsbrunner, Letscher, Zomordian 2000], [Zomorodian and Carlsson 2005], Earlier developments: [Frosini 1990], [Robins 1999]
- Given a filtration of a space X
 - $\bullet \ X_1 \subset X_2 \subset \cdots X_i \subset \cdots \subset X_j \subset \cdots X_n = X$
 - Consider this as a lens through which we inspect X
- Capture creation and death of ``features'' by homology
 - $H_*(X_1) \to \cdots \to H_*(X_i) \to \cdots \to H_*(X_j) \to \cdots \to H_*(X_n) = H_*(X)$
 - Summarize the birth/death of homological features in the persistence diagram

Distance Field Filtration Example

• A filtration induced by distance field.



In Neuron Setting

- Assume f is plotted as height function
- Filtration induced by the sub-level set filtration
 - $f^{-1}(-\infty, a_0) \subseteq f^{-1}(-\infty, a_1) \subseteq \cdots \subseteq f^{-1}(-\infty, a_n) = T$



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Remarks

- Depending on the descriptor function $f: |T| \rightarrow R$, a tree may have both down-forks and up-forks.
 - Also consider super-level sets filtration, and its induced persistence diagram Dg_{-f}
- Given a descriptor function f,
 - Obtain persistence diagram summary $Dgf = Dg_f \cup Dg_{-f}$
 - Dg f serves as a summary of T from the perspective of descriptor function f
- Persistence-summary intuitively more discriminative than simply statistics of morphological measures (eg. avg branching angles)

Connection to Sholl-like Analysis

- Sholl function $N: R^+ \to R^+$
 - $N(\lambda) \coloneqq$ number of intersection of T with a circle (sphere) centered at the root r with radius λ



Connection to Sholl-like Analysis

- Sholl function $N: R^+ \to R^+$
 - $N(\lambda) \coloneqq$ number of intersection of T with a circle (sphere) centered at the root r with radius λ
- One can recover full Sholl function from persistence diagrams induced by Euclidean distance function



Persistence-based feature vectorization framework



- To facilitate efficient distance computation
 - Convert persistence diagram Dg f to a featue vector $V_{T,f}$
 - [Bubenik 2012], [Reininghaus et al 2015], [Adams et al 2015],...

Feature Vectorization



Convert diagram D to a 1D density field

$$\rho_D(x) := \sum_{i=1 \in k} m_i \cdot K_t(x, x_i), \text{ for any } x \in \mathbb{R},$$

Discretize it to a *m*-vector

$$\boldsymbol{\nu}_D := [\rho_D(\mathbf{a} + \frac{I}{m}), \ \rho_D(\mathbf{a} + \frac{2I}{m}), \ \cdots, \rho_D(\mathbf{a} + \frac{mI}{m}) = \rho_D(\mathbf{b})].$$

Persistence-based feature vectorization framework



- If there are multiple descriptor functions
 - Concatenate their respective feature vectors
 - Perform dimensionality reduction to reduce dimension

Remarks

Versatile framework

- Can combine multiple type of information of neurons, morphological or electrophysiological measures
- Easy to add new measurements

Discreminative features

- E.g, persistence features from Euclidean function contains more information than Sholl function
- E.g, persistence features from geodesic function encodes global morphological information

Have certain stability guarantees



Three Test Datasets

- Dataset I:
 - 379 neurons taken from neuromorpho.org category Drosophila-Chklovskii, manually categorized into 89 types
 - Fakemura et al, 2013]
- Dataset 2:
 - I 27 neurons from four families: Purkinje, olivocerebellar neurons, Spinal motoneurons and hippocampal interneurons, downloaded also from neuronmorpho.org
- Dataset 3:
 - I268 neurons from Human Brian Project, downloaded from neuromorpho.org. Two primary cell classes: interneurons and principal cells, known for II30 cells
 - [Markram et al 2015]

Leave-one-out classification tests based on k-nearest neighbors

	# neurons for classified correctly out of 346 for all non-singleton classes Dataset 1		
# nearest neighbors	Persist-distance d_P	Persist-vec d_V	Sholl-distance d_S
1	190	164	104
2	221	199	142
3	235	226	160
4	250	235	170
5	262	239	180
	# neurons for classified correctly out of 127 neurons in Dataset 2		
# nearest neighbors	Persist-distance d_P	Persist-vec d_V	Sholl-distance d_S
1	117	111	92
2	121	118	108
3	122	120	113
4	122	120	117
5	123	121	124
	# neurons for classified correctly out of 1130 in Dataset 3		
# nearest neighbors	Persist-distance d_P	Persist-vec d_V	Sholl-distance d_S
1	832	812	763
2	990	985	942
3	1065	1054	1019
4	1093	1083	1058
5	1107	1100	1081

Clustering for Dataset 2





Clustering for Dataset I

Five largest families other than "Tangential"



An interactive visualization tool



This Talk

Part I:

Neuron structures comparison

Part II:

Neuronal Morphology Reconstruction

Neuronal Morphology Reconstruction

 Various imaging techniques produce large number of 2D/3D images



Related Work

DIADEM challenge (2009—2010)

- Diginal Reconstruction of Axonal and Dendritic Morphology
- http://diademchallenge.org/
- BigNeuron (launched in 2015)
 - Large-scale 3D single neuron reconstruction
 - Sponsored by 14 neuroscience-related research organizations and international research groups
 - https://www.alleninstitute.org/bigneuron/about/
- Many algorithms already integrated into platform Vaa3D
 - [Peng et al., 2010] <u>www.vaa3D.org</u>.

The Problem

• On the high level:

 Given a 2D / 3D image data, the goal is to extract one (or multiple) tree-like structure(s) from it.

Some challenges:

- Various types of background ``noise"
- Non-homogeneous distribution of signal in raw data
- Mixture of multiple neurons



• On the high level:

Given a 2D / 3D image data, the goal is to extract one (or multiple) tree-like structure(s) from it.

Previous methods:

- Often rely on local information for decision making, sensitive to noise
- Some thresholding involved, challenging in handling nonuniform signal distribution
- Junction nodes identification challenging
 - E.g, ``growing'' individual branches and ``gluing'' them to obtain tree topology

Morse-based Reconstruction Framework



Morse-based approach

- uses global structure behind data
- junction nodes identification reliable w/o special processing
- robust against noise, small gaps, and non-uniformity in data
- conceptually clean, helps reducing pre-processing of data

Assume input is a scalar field

- $f: I \rightarrow R$, where high value of f indicates high signal value
- Consider graph f as a terrain (mountain range) on I×R
 I can be [0,1]² ⊂ R² or [0,1]³ ⊂ R³



1200

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• Let $f: \mathbb{R}^d \to \mathbb{R}$ be a Morse function

• Gradient of
$$f$$
 at $x: \nabla f(x) = \left[\frac{\partial f}{x_1}, \frac{\partial f}{x_2}, \dots, \frac{\partial f}{x_d}\right]^T$



• Let $f: \mathbb{R}^d \to \mathbb{R}$ be a Morse function

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- Critical points of *f*:
 { x ∈ R^d | ∇f(x) = 0 }



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- Critical points of $f: \{x \in \mathbb{R}^d \mid \nabla f(x) = 0\}$
- An integral line $L: (0, 1) \rightarrow \mathbb{R}^d$:
 - a maximal path in R^d whose tangent vectors agree with gradient of f at every point of the path



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- An integral line $L: (0, 1) \rightarrow R^d$:
 - a maximal path in R^d whose tangent vectors agree with gradient of f at every point of the path
 - has origin and destination at critical points
 - $Dest(L) = \lim_{p \to 1} L(p)$
 - $Ori(L) = \lim_{p \to 0} L(p)$



- Given a critical point x of f
 - Stable manifold $S(x) = \{ y \in \mathbb{R}^d \mid dest(y) = x \}$
 - Unstable manifold $U(x) = \{ y \in \mathbb{R}^d \mid ori(y) = x \}$
- Morse complex, Morse-Smale complex



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1-unstable Manifold



1-unstable manifold (of index d - 1 saddle points) => mountain ridges

1-unstable Manifold



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Simplification





How to decide which pair of critical points to simplify?

Use persistence homology
 [Edelsbrunner, Letscher,
 Zomorodian 2002], [Zomorodian,
 Carlsson 2005], ...

Simplification



Discrete Case

- Input: a piecewise-linear (PL) function defined on a simplicial complex domain
 - Given volumetric data (2D / 3D images), we can first triangulate it and convert it to a simplicial complex domain
- Leverage discrete Morse theory
 - Forman 1998, 2002]
 - [Gyulassy, 2008], [Sousbie 2011] (DisPerSE)

Neuron Reconstruction Overview

- ▶ Input: 2D/3D image $f: I \to R$ with f given at grid points in I
 - (1) Triangulate I to K, and potentially remove background cells to obtain PL function $f: K \rightarrow R$
 - (2) Negate f to $\hat{f} = -f$
 - (3) Compute 1-stable manifold for index-1 saddles
 - (4) Simplify to remove noise
 - ▶ (5) Output Neuron-graph G
 - $\bullet \quad \textbf{(6) Obtain a tree structure } T \text{ from } G$
 - Assign weights to arcs in G as integral of density f along the arc
 - Compute maximum spanning tree T

Neuron Reconstruction Overview



Some DIADEM datasets





OP 1

OP 9

Some DIADEM datasets



OP 9

Diadem Dataset OP2



2011]) as APP2 (from Vaa3D)

Diadem Dataset OP2



Mouse brain LM images from an AAV viral tracer-injection

from Mitra laboratory at CSHL



Remarks

Other advantages of Morse-based framework

- Can be used to merge/integrate multiple reconstructions
- Can be used to provide correction ability



Summary and Remarks

- Two examples of topological methods for neuronal structure analysis
 - Topological methods
 - general and robust
 - capture / leverage global structures
 - tend to be less ad-hoc
- Further develop these applications
- Provide more theoretical justification and understanding