

STRUCTURAL AND DYNAMICAL ASPECTS OF NEURAL  
NETWORKS WITH ANISOTROPIC TISSUE GEOMETRY

FELIX HOFFMANN

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under joint supervision of*

Prof. Dr. Peter Pfaffelhuber  
Department of Mathematical Stochastics,  
University of Freiburg,

*and*

Prof. Dr. Stefan Rotter  
Bernstein Center Freiburg & Faculty of Biology,  
University of Freiburg.

*University of Freiburg*

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## A B S T R A C T

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Short summary of the contents in English...



## CONTENTS

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|     |   |    |
|-----|---|----|
| 1   | INTRODUCTION                                  | 1  |
| 2   | BIOLOGY                                       | 3  |
| 3   | GRAPH THEORY                                  | 5  |
| 3.1 | Directed graphs . . . . .                     | 5  |
| 3.2 | Network measures . . . . .                    | 8  |
| 3.3 | Random Graph Theory . . . . .                 | 10 |
| 3.4 | Geometric Graphs . . . . .                    | 12 |
| 4   | NETWORK MODEL                                 | 17 |
| 4.1 | Overview . . . . .                            | 17 |
| 4.2 | Anisotropy in Neural Connectivity . . . . .   | 18 |
| 4.3 | Anisotropic geometric network model . . . . . | 23 |
| 4.4 | Numerical Implementation . . . . .            | 25 |
| 4.5 | Distance dependent connectivity . . . . .     | 27 |
| 4.6 | Rewiring . . . . .                            | 29 |
| 4.7 | Anisotropy Measure . . . . .                  | 34 |
| 4.8 | Summary and Discussion . . . . .              | 35 |
| 5   | STRUCTURAL ASPECTS                            | 37 |
| 5.1 | Introduction . . . . .                        | 37 |
| 5.2 | Degree distribution . . . . .                 | 40 |
| 5.3 | Small World Properties . . . . .              | 43 |
| 5.4 | Two Neuron Connections . . . . .              | 44 |
| 5.5 | Tuning distance-dependency . . . . .          | 47 |
| 5.6 | Motifs . . . . .                              | 50 |
| 5.7 | Common neighbor rule . . . . .                | 60 |
| 5.8 | Discussion . . . . .                          | 63 |
| A   | APPENDIX                                      | 65 |
| A.1 | Mathematica . . . . .                         | 65 |
| A.2 | Supplementary figures . . . . .               | 66 |
|     | REFERENCES                                    | 69 |
|     | INDEX   | 73 |



# 1

## INTRODUCTION

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# 2

BIOLOGY

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# 3

## GRAPH THEORY

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A review of graph theory lays the foundation for the mathematical considerations in this work. Following the discussion of directed graphs as a basis for the networks introduced in this work, common network measures are reviewed. Random graph models are integral to this work and are discussed in Sections 3.3 and 3.4, the latter introducing geometric random graphs and performing geometric, probabilistic computations essential for the analytical discussions in later chapters.

### 3.1 DIRECTED GRAPHS

Here we introduce the various categories of directed graphs. The main reference for this section is [Bang-Jensen and Gutin \(2008\)](#), for the formal definition below however, we follow [nLab \(2014\)](#).

**Definition 3.1** (Directed graphs). A *directed pseudograph*  $G$  consists of two finite sets  $V(G)$ , the *set of vertices* of  $G$ , and  $E(G)$ , the *set of edges* of  $G$ , and two maps  $s, t : E \rightarrow V$ , the *source* and *target functions* of  $G$ . A *directed multigraph* is a directed pseudograph without loops, that is the map  $d = (s, t) : E \rightarrow V^2$  already maps to  $V^2 \setminus \Delta_V$ , where  $V^2 = V \times V$  denotes the cartesian product and  $\Delta_V = \{(x, x) \mid x \in V\} \subseteq V^2$  the diagonal. Similarly, a *directed loop graph* is a directed pseudograph where  $d$  is injective. Finally, a *simple directed graph* can be defined as a directed pseudograph where  $d$  is both injective and already maps to  $V^2 \setminus \Delta_V$ .

short:  
 $V(G)$  vertex set  
 $E(G)$  edge set

Thus, in simple directed graphs, neither parallel edges nor loops (edges between the same vertex) are allowed, whereas directed multigraphs and directed loop graphs admit one of them respectively. We refer to any of the four graph types simply as a directed graph and only specify the type when needed.

A **morphism**  $\phi : G \rightarrow H$ , between two directed graphs  $G = (V_G, E_G, s_G, t_G)$  and  $H = (V_H, E_H, s_H, t_H)$ , consists of a pair of maps  $\phi_V : V_G \rightarrow V_H$  and  $\phi_E : E_G \rightarrow E_H$ , such that

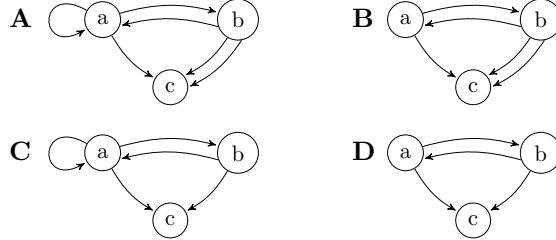
$$s_H \circ \phi_E = \phi_V \circ s_G \quad \text{and} \quad t_H \circ \phi_E = \phi_V \circ t_G,$$

that is such that the following diagram commutes:

$$\begin{array}{ccc} E_G & \xrightarrow{\phi_E} & E_H \\ s_G \downarrow t_G & & \downarrow s_H t_H \\ V_G & \xrightarrow{\phi_V} & V_H \end{array}$$

A morphism  $\varphi : G \rightarrow H$ , between two directed pseudographs  $G$  and  $H$  is an **isomorphism**, if the maps  $\varphi_V : V_G \rightarrow V_H$  and  $\varphi_E : E_G \rightarrow E_H$  are bijections. Two directed pseudographs are called *isomorphic* if there exists an isomorphism inbetween them.

*Remark.* The last definition implies that, if there exists an isomorphism  $\varphi : G \rightarrow H$ , an isomorphism  $\psi : H \rightarrow G$  can be found. This isomorphism is, of course, easily constructed via  $\psi_V : V_H \rightarrow V_G, v \mapsto \varphi_V^{-1}(v)$ ,  $\psi_E : E_H \rightarrow E_G, e \mapsto \varphi_E^{-1}(e)$ .



**Figure 3.1: Examples of the directed graph types** **A)** directed pseudograph **B)** directed multigraph **C)** directed loop graph **D)** simple directed graph.

The subobjects in the directed graph category are subgraphs. Containing only a subset of the vertices of the original graph, it may contain any number of eligible edges from  $G$ .

**Definition 3.2** (Subgraph). Let  $G$  be a directed graph. A *subgraph*  $H$  of  $G$  is a directed graph with vertex set  $V(H) \subseteq V(G)$  and edge set  $E(H) \subseteq E(G)$  with source and target functions from  $G$  restricted to  $E(H)$ , such that for every  $e \in E(H)$  the source and target  $s(e), t(e)$  are in  $V(H)$ . We call the subgraph *full* if

$$E(H) = \{e \in E(G) \mid s(e), t(e) \in V(H)\}.$$

that is if  $H$  contains all edges of  $G$  with source and target in  $V(H)$ .

A **motif**  $H$  in  $G$  is then a full, connected subgraph of  $G$ , that is a subgraph such that every vertex in  $H$  is source or target of at least one edge in  $E(H)$ . An  $n$ -motif then refers to a motif  $H$  with  $|V(H)| = n$ . Usually

motifs with only few vertices are considered and are often understood as the “building blocks” of graphs. In neuronal networks, specific connection patterns are associated with specific dynamical functionality, making motifs an import aspect in the theory of neural network dynamics and a central object of interest in this thesis.

**Definition 3.3** (Weighted directed graphs). An *edge-weighted directed graph* is a directed graph  $G$  along with a mapping  $\omega : E(G) \rightarrow \mathbb{R}$ , called the *weight function*. Similarly, a *vertex-weighted directed graph* is a directed graph with a mapping  $\nu : V(G) \rightarrow \mathbb{R}$ .

*Remark.* A directed graph  $G$  can be equivalently defined as a pair of finite sets  $V$ , the *set of vertices* of  $G$ , and  $E \subseteq V^2$  the *set of edges* of  $G$ . For an edge  $(x, y) \in E$ , we call  $x$  the *source* and  $y$  the target of the edge  $(x, y)$ . Source and target functions are then uniquely determined as the projections on the first and second component,

$$s = \text{pr}_1, t = \text{pr}_2 : E(G) \rightarrow V.$$

Conversely, the edge set  $E(G) \subseteq V^2$  can be determined from the source and target functions as  $E := \{(s(e), t(e)) \mid e \in E\}$ . The trivial identities  $(x, y) = (\text{pr}_1(x, y), \text{pr}_2(x, y))$  and  $\text{pr}_1(s(e), t(e)) = s(e)$  with  $\text{pr}_2(s(e), t(e)) = t(e)$  quickly verify the equivalence of the definitions. Given a directed loop graph  $G$ , we often assume the graph to be given in this form and write edges as  $e = (x, y)$ . Note that this concept is more complicated to introduce for directed pseudographs, since parallel edges  $e$  and  $e'$  should be differentiated in the egde set of  $G$ , establishing the need for  $E(G)$  to be a multi- or indexed set, notions we are trying to avoid in this document.

Equivalent  
definiton for  
directed loop  
graphs

From now on any directed graph is assumed to be a directed loop graph. Although most, if not all, concepts work for directed pseudographs just as well, we want to start to heavily use the canonical edge representation, which when talking about pseudograpgs makes problems as mentioned before.

For a pair of vertex sets  $X, Y \subseteq V(G)$  of a directed graph  $G$  we write

$$(X, Y)_G = \{(x, y) \in E(G) \mid x \in X, y \in Y\}$$

for the set of edges with source in  $X$  and target in  $Y$ . Specifically we write  $T(x) = (x, V(G))_G$  for the set of *targets* for edges originating from the vertex  $x$  and  $S(x) := (V(G), x)_G$  for the set of *sources* for edges projecting to  $x$ .

Notation for  
target and source  
sets

**Definition 3.4** (In- and out-degree). For a directed graph  $G$  the **in-degree**  $d_G^-(x)$  of a vertex  $x$  is defined as the number of edges of  $G$  with target  $x$ , that is

$$d_G^-(x) = |S(x)|.$$

Similarly, the **out-degree**  $d_G^+(x)$  of  $x$  is defined as

$$d_G^+(x) = |T(x)|,$$

the number of edges in  $G$  with source  $x$ .

A basic property of the in- and out-degree in directed graphs is that number of in-degrees of every vertex, as well the sum of every out-degree, equal the total number of edges:

**Proposition 3.5.** *In every directed graph  $G$ , we have*

$$\sum_{x \in V(G)} d^-(x) = \sum_{x \in V(G)} d^+(x) = |E(G)|.$$

*Proof.* Since  $(V(G), x)_G \cap (V(G), y)_G = \emptyset$  for  $x \neq y$ , we can write

$$\sum_{x \in V(G)} d^-(x) = \left| \bigcup_{x \in V(G)} (V(G), x)_G \right| = |(V(G), V(G))_G| = |E(G)|.$$

Analogously for the out-degree.  $\square$

### 3.2 NETWORK MEASURES

Let  $G$  be a (simple) directed graph. A **walk**  $W$  in  $G$  is an alternating sequence  $(x_1, e_1, x_2, e_2, x_3, \dots, x_{n-1}, e_{n-1}, x_n)$  of vertices  $x_i$  and edges  $e_i$  from  $G$ , such that

$$s(e_i) = x_i \quad \text{and} \quad t(e_i) = x_{i+1}, \quad \text{for } i = 1, \dots, n-1,$$

that is, such that the vertices are connected by the edges inbetween them. We denote the set of vertices  $(x_1, \dots, x_n)$  of  $W$  as  $V(W)$  and the sequence of edges  $(e_1, \dots, e_{n-1})$  as  $E(W)$ .

The vertices  $x_1$  and  $x_n$  are called the *end vertices* of  $W$  and we also say that  $W$  is an  $(x, y)$ -walk. The **length** of  $W$  is defined as the length of the sequence of edges; a walk consisting of only one vertex has length zero.

**Definition 3.6** (Distance). The **distance** of two vertices  $x, y$  in a directed graph  $G$ , is defined as the minimum length of an  $(x, y)$ -walk, if any such walk exists, otherwise  $\text{dist}(x, y) = \infty$ . In short,

$$\text{dist}(x, y) = \inf\{|E(W)| \mid W \text{ is } (x, y)\text{-walk}\}.$$

**Proposition 3.7.** *The distance function  $\text{dist} : V(G) \times V(G) \rightarrow \mathbb{N}$  of a directed graph  $G$  satisfies the triangle equality,*

$$\text{dist}(x, z) \leq \text{dist}(x, y) + \text{dist}(y, z), \quad \text{for } x, y, z \in V(G).$$

*Proof.* Let  $x, y, z$  be vertices in  $G$ . If either no  $(x, y)$ -walk or  $(y, z)$ -walk exists, the inequality holds by definition. Otherwise, let  $W$  be an  $(x, y)$ -walk of minimal length and let  $U$  be a  $(y, z)$ -walk of minimal length. Certainly, by concatenating  $W$  and  $U$  we obtain an  $(x, z)$ -walk of length  $|E(W)| + |E(U)| = \text{dist}(x, y) + \text{dist}(y, z)$ , proofing that

$$\text{dist}(x, z) \leq \text{dist}(x, y) + \text{dist}(y, z).$$

□

**Definition 3.8** (Average path length). The *average path length* of a directed graph  $G$  with  $V(G) = n$  is defined as

$$l = \frac{1}{n(n-1)} \sum_{x \neq y \in V(G)} \text{dist}(x, y).$$

In practice, vertex pairs with  $\text{dist}(x, y) = \infty$ , that is pairs that are not connected by a walk, are disregarded in the computation and the average path length is determined in the connected components of the graph, ensuring that  $l$  is finite.

The concept of a small-world property in graphs was introduced by [Watts and Strogatz \(1998\)](#). Networks associated with the property are characterized by a small average path length, while however most nodes are organized in “cliques”, connecting to nodes that are themselves neighbors. A measure capturing this property is the (local) clustering coefficient:

**Definition 3.9** (Clustering coefficient). The *clustering coefficient* of a vertex  $x$  in a directed graph  $G$  is defined as ratio of realized and possible edges between the neighbors of  $x$ . If  $N_x$  is the neighborhood of all vertices reciprocally connected to  $x$ ,

$$N_x = \{v \mid v \in T(x) \wedge v \in S(x)\},$$

then the clustering coefficient is given by

$$\text{clust}(x) = \frac{|(N_x, N_x)_G|}{|N_x|(|N_x| - 1)}.$$

Small-worldness is then described by a low average path length and high clustering coefficient, usually considered as the mean of all vertices.

*focus on labeled,  
simple directed  
graphs*

### 3.3 RANDOM GRAPH THEORY

From this chapter on, as it is common and practical when talking about random graph models, we move away from the abstract notion of graphs and their equivalence classes and consider *labeled graphs*, where the vertex set of a graph with  $n$  vertices takes the form  $V = \{1, \dots, n\}$ . Simple directed graphs constitute the fundamental mathematical object underlying the concepts developed in this work and if not specified otherwise, all graphs are assumed to be labeled and simple directed.

The concept of a random graph was first formally introduced by Erdős and Rényi (1959). In their  $G(n, N)$  model, an undirected graph with  $n$  vertices and  $N$  edges is randomly and with equal probability selected from the set of such possible graphs. In the same year Gilbert (1959) independently introduced his  $G(n, p)$  model, realizing edges between vertex pairs with a fixed probability  $p$ . The two models are closely related (Luczak 1990) and overlap in literature, both at times being referred to as *Erdős-Rényi graphs*. Here we focus on the  $G(n, p)$  model, as it closer in concept to a computational implementation of a random graph. Defining it in detail in 3.10, we will refer to the random graph model as the *Gilbert random graph model*.

In general, a random graph model is a probability space over a set of graphs (Janson et al. 2000). Rather than specifying the sample space and probability measure explicitly, random graph models are often defined by a random process that generates such graphs, leaving probability measure and sample space implicit (Bollobás 2001). The term *random graph*, in the graph theoretical context, refers to the random graph model itself. Especially in the computational context however, a random graph often refers to a single graph generated by a random process. Here we try to avoid this ambiguity and strictly refer to the mathematical object as a random graph model.

Keeping in mind that the term *graph* now refers to labeled, simple directed graphs if not otherwise specified we define  $G^n$  to be the set of simple directed graphs with  $n$  vertices,

$$G^n := \{G \mid G \text{ graph}, |V(G)| = n\}.$$

We first introduce Gilbert's random graph model  $G(n, p)$  by explicitly defining a probability space over  $G^n$  and show later how the model may be realized as a random process.

**Definition 3.10** (Gilbert random graph model). Let  $n \in \mathbb{N}$  and  $0 \leq p \leq 1$ . The *Gilbert random graph model*  $G(n, p)$  is a discrete probability space over  $G^n$  with event space  $\mathcal{P}(G^n)$  and probability measure  $P$ ,

such that every graph  $G$  with  $|E(G)| = k$  edges appears with equal probability

$$P(G) = p^k(1-p)^{n(n-1)-k},$$

for  $0 \leq k \leq n(n-1)$ .

*Remark.* Clearly  $G(n, p)$  is well-defined, as there exist  $\binom{n(n-1)}{k}$  distinct labeled graphs with  $n$  vertices and  $k$  edges and thus

$$\sum_{G \in G^n} P(G) = \sum_{k=0}^{n(n-1)} \binom{n(n-1)}{k} p^k (1-p)^{n(n-1)-k} = 1,$$

after the binomial theorem.

Equivalently, the Gilbert random graph model can be defined as a stochastic process; to an empty graph with  $n$  vertices, for each of the  $n(n-1)$  vertex pairs an edge is added at random and independently with probability  $p$ . The probability to obtain a specific graph  $G$  with  $k$  edges is then obviously  $p^k(1-p)^{n(n-1)-k}$ , already proofing the equivalence, since assuming a process as above with edge probability  $p'$  such that the induced probability measure on  $G^n$  equals  $P$  from 3.10, already yields  $p = p'$  in the choice of  $n = 2$  and  $k = 1$ .

equivalent  
definition as  
random process

**Proposition 3.11.** *In- and out-degree distribution of vertices in the Gilbert random graph model are binomial.*

*Proof.* Let  $X$  be a random variable on the random graph model, mapping to the in-degree (out-degree)  $d_G(v)$  of a vertex  $v$  of a graph  $G \in G^n$ . There are  $n-1$  other vertices that, with probability  $p$ , project to  $v$  (receive input from  $v$ ), thus

$$P(X = k) = \binom{n-1}{k} p^k (1-p)^{n-1-k},$$

showing that  $P^X = \mathcal{B}_{n-1,p}$ . □

The Gilbert random graph model is therefore also often referred to as *binomial random graph*. As typical neuronal networks are large ( $n \geq 10^3$ ) with sparse connectivity ( $p \approx 0.1$ ), in- and out-degree distribution can be approximated by a Poisson distribution,  $P^X(k) \approx \text{Pois}_\lambda(k)$ , with  $\lambda = (n-1)p$ , after the Poisson limit theorem.

Most results in the study of random graph models consider  $n \rightarrow \infty$ . In this study we are mostly interested in patterns of connectivity that arise in local circuits, leaving behind limit considerations and employ the Gilbert random graph model as a reference for the development of more detailed and specific random graph models.

### 3.4 GEOMETRIC GRAPHS

The theory of geometric graphs addresses the embedding of graphs in  $\mathbb{R}^d$ . Planar graphs are graphs that can be drawn on a surface with their edges drawn as straight lines between the vertices, such that no two lines intersect (Diestel 2000). Here we are interested in graphs embedded on surfaces. In the models introduced, connectivity then depends on geometric properties, such as distance between nodes or relative orientation between vertices. The basic graph type to allow for such connectivity rules is a geometric graph.

**Definition 3.12** (Geometric directed graph). A *geometric directed graph*  $G_\Phi$  is a directed graph  $G$  paired with a map

$$\Phi : V(G) \rightarrow [0, 1]^2,$$

representing vertex positions on the unit square.

This definition diverges from the usual notion of a geometric graph, which determines the existence of edges only between nodes within a spatial distance  $x$  in a specified norm (Penrose 2003). Moreover, geometric graphs are usually only discussed in the context of random graph models, a concept first introduced by Gilbert (1961). Denote the set of geometric graphs with  $n$  edges by  $G_\Phi^n$ . The spatial embedding of geometric graphs allows us to define (random) connectivity depending on vertex positions and inter-vertex distances. Central to this study is the distance-dependent random graph model, in which edges are added with probability  $p(x)$  depending on the distance  $x$  between vertex pairs:

**Definition 3.13** (Distance-dependent random graph model). Let  $n \in \mathbb{N}$  and  $C : [0, \sqrt{2}] \rightarrow [0, 1]$  a continuous map. A *distance-dependent geometric random graph model*  $G_\Phi(n, C)$  is a random graph model over  $G_\Phi^n$ , generated by distributing uniformly at random the  $n$  vertices on the unit square and adding an edge from  $v$  to  $w$  for each vertex pair  $(v, w) \in V(G_\Phi)^2 \setminus \Delta_{V(G_\Phi)}$  with a probability  $C(x)$ , depending on the distance  $x = \|\Phi(v) - \Phi(w)\|$  between the vertices.

We call the function  $C$  the graph's *distance-dependent connection probability profile*. Note that the connection profile is *not* a probability density but gives a probability of connection between a vertex pair at distance  $x$ . Clearly, connectivity statistics in the distance-dependent graph model intrinsically depend on the choice of  $C$ . To develop a thorough understanding of connectivity in the model however, mapping the distribution of inter-vertex distances is equally important. Here we calculate the distribution of the distance between two random points in

a square of side-length  $s$ . Being able to identify distributions of transformed random variables is integral to the calculation:

**Lemma 3.14.** *Let  $X, Y$  be independent continuous random variables with values in  $\mathbb{R}$ , denote with  $f_X$  and  $f_Y$  their probability distribution functions.*

- (1) *The distribution of the random variable  $X + Y$  is given by the probability density function*

$$f_{X+Y}(x) = \int_{\mathbb{R}} f_X(z) f_Y(z-x) dz.$$

- (2) *The distribution of the random variable  $X^2$  is given by the probability density function*

$$f_{X^2}(x) = \begin{cases} \frac{1}{2\sqrt{x}} (f_X(\sqrt{x}) + f_X(-\sqrt{x})) & x > 0 \\ 0 & x \leq 0 \end{cases}$$

- (3) *Let  $X$  only take positive values. Then the distribution of the random variable  $\sqrt{X}$  is given by the probability density function*

$$f_{\sqrt{X}}(x) = \begin{cases} 2x f_X(x^2) & x \geq 0 \\ 0 & x < 0 \end{cases}$$

*Proof.* These are well known identities. Distributions in (2) and (3) are easily verified by first considering the cumulative distribution function and subsequent derivation.  $\square$

**Theorem 3.15.** *Let  $D$  be a random variable mapping to the distance of two random points in the square  $[0, s]^2$  of side-length  $s$ . Then the distribution of  $D$  is given by the probability density function*

$$f(x) = \begin{cases} \frac{2x^4 - 8sx^3 + 2\pi s^2 x}{s^4} & x \in [0, s] \\ \frac{H(x)}{s^4} & x \in [s, s\sqrt{2}), \\ 0 & x \notin [0, s\sqrt{2}) \end{cases} \quad (3.1)$$

where

$$H(x) = 8sx\sqrt{x^2 - s^2} - 2x^3 - 4s^2x \left(1 + \arcsin\left(1 - \frac{2s^2}{x^2}\right)\right).$$

*Proof.* We follow the approach described by Moltchanov (2012). Consider two independently and uniformly distributed random points  $p_1 = (x_1, y_1)$  and  $p_2 = (x_2, y_2)$  in  $[0, s]^2$ . The distance between  $p_1$  and  $p_2$  is given as

$$\|p_1 - p_2\| = \sqrt{(x_1 - x_2)^2 + (y_1 - y_2)^2}.$$

As a first step we calculate the distribution for  $\Delta_x = x_1 - x_2$ . Denote with  $f_{\Delta_x}$  its probability density. Then, since  $f_{-x_2}(z) = f_{x_2}(-z)$ ,

$$f_{\Delta_x}(d) = f_{x_1+(-x_2)}(d) = \int_{\mathbb{R}} f_{x_1}(z) f_{x_2}(z-d) dz \quad (3.2)$$

after Lemma 3.14. Density functions for  $x_1$  and  $x_2$  are given by

$$f_{x_1}(z) = f_{x_2}(z) = \begin{cases} \frac{1}{s} & \text{for } z \in [0, s] \\ 0 & \text{otherwise.} \end{cases}$$

Thus we may only obtain non-zero values in (3.2) for  $d \in (-s, s]$ , as otherwise either one of the factors in the integrand is zero. In full we obtain the triangular distribution (Simpson 1755),

$$f_{\Delta_x}(d) = \begin{cases} 0 & d \notin (-s, s] \\ \frac{s+d}{s^2} & d \in (-s, 0] \\ \frac{s-d}{s^2} & d \in (0, s]. \end{cases}$$

Next we calculate the distribution for  $\Delta_x^2 = (x_1 - x_2)^2$ . Using Lemma 3.14 once again we obtain for  $d > 0$

$$\begin{aligned} f_{\Delta_x^2}(d) &= \frac{1}{2\sqrt{d}} \left( f_{\Delta_x}(\sqrt{d}) + f_{\Delta_x}(-\sqrt{d}) \right) \\ &= \frac{1}{2\sqrt{d}} \left( \frac{s-\sqrt{d}}{s^2} + \frac{s+\sqrt{d}}{s^2} \right) = \frac{1}{s\sqrt{d}} - \frac{1}{s^2}, \end{aligned}$$

and  $f_{\Delta_x^2}(d) = 0$  for  $d \leq 0$ . Note that of course,  $f_{\Delta_x^2} = f_{\Delta_y^2}$  and we will refer to this density function as  $f_{\Delta}^2$ . Convolution yields again the probability density function for the sum of the random variables  $\Delta_x^2$  and  $\Delta_y^2$ ,

$$f_{\Delta_x^2 + \Delta_y^2}(d) = \int_{\mathbb{R}} f_{\Delta^2}(z) f_{\Delta^2}(d-z) dz.$$

Finally Lemma 3.14 lets us compute

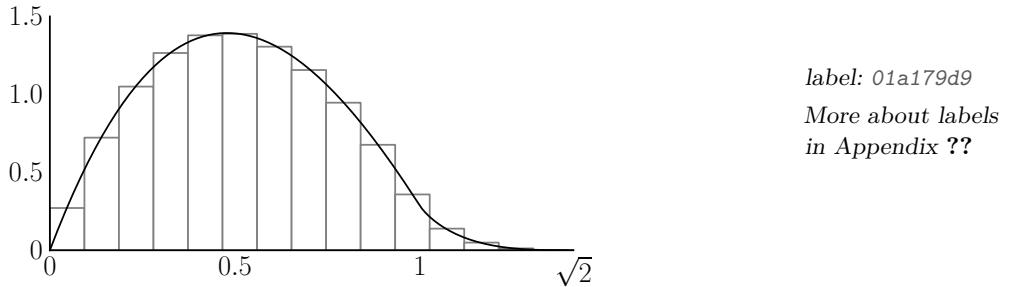
$$f_D(d) = f_{\sqrt{\Delta_x^2 + \Delta_y^2}}(d) = 2d f_{\Delta_x^2 + \Delta_y^2}(d^2),$$

for  $d \geq 0$ , yielding expression (3.1) for probability density function of  $D$  (Mathematica A.1, ??).  $\square$

The distribution for the distance between two random points in the unit square  $[0, 1]^2$ , is then obtained from (3.1) by setting  $s = 1$ . The probability density function  $f$  becomes

$$f(x) = \begin{cases} 2x^3 - 8x^2 + 2\pi x & x \in [0, 1) \\ 8x\sqrt{x^2 - 1} - 2x^3 & x \in [1, \sqrt{2}) \\ -4x - 4x \arcsin\left(1 - \frac{2}{x^2}\right) & x \in [\sqrt{2}, 1] \end{cases} \quad (3.3)$$

Plotting  $f$  in combination with the results of a numerical simulation (10000 points randomly distributed on unit square, extracted distances binned to a of width  $\sqrt{2}/15$ ) verifies our calculation:



Density functions 3.1 and 3.3 are of high importance. Here we use 3.3 to compute the probability that a random pair of vertices in the distance-dependent random graph model is connected:

**Corollary 3.16.** *Let  $v \neq w$  be vertices in an arbitrary realization of  $G_\Phi(n, C)$ . The probability  $p$  for an edge from  $v$  to  $w$  is given by*

$$p = \int_0^{\sqrt{2}} C(x)f(x) dx,$$

where  $f(x)$  is the probability density function (3.3).

**Example** Let  $C(x) = 1 - \frac{x}{\sqrt{2}}$ . Then we compute the probability to find an edge between a random vertex pair in  $G_\Phi(n, C)$  according to Theorem 3.16 as

$$p = \int_0^{\sqrt{2}} f(x) dx - \frac{1}{\sqrt{2}} \int_0^{\sqrt{2}} x f_D(x) dx = 1 - \frac{1}{\sqrt{2}} \mathbf{E}[D].$$

The expected distance between two random points on the unit square is computed as  $\mathbf{E}[D] \approx 0.521405$  ([Mathematica A.1](#)), a result confirmed by [Philip \(2007\)](#). Thus we obtain the probability for connection of  $p \approx 0.631311$



# 4

## NETWORK MODEL

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Motivated by anisotropic characteristics in connectivity in local cortical circuits found in the rat’s brain, a network model with anisotropic tissue geometry is developed. Employing both a graph theoretic definition and a numerical implementation, distance-dependent connectivity present in the model is exposed. The introduction of a rewiring algorithm and quantitative anisotropy measure lays the foundation for the analysis of structural aspects of the anisotropic network model in Chapter 5.

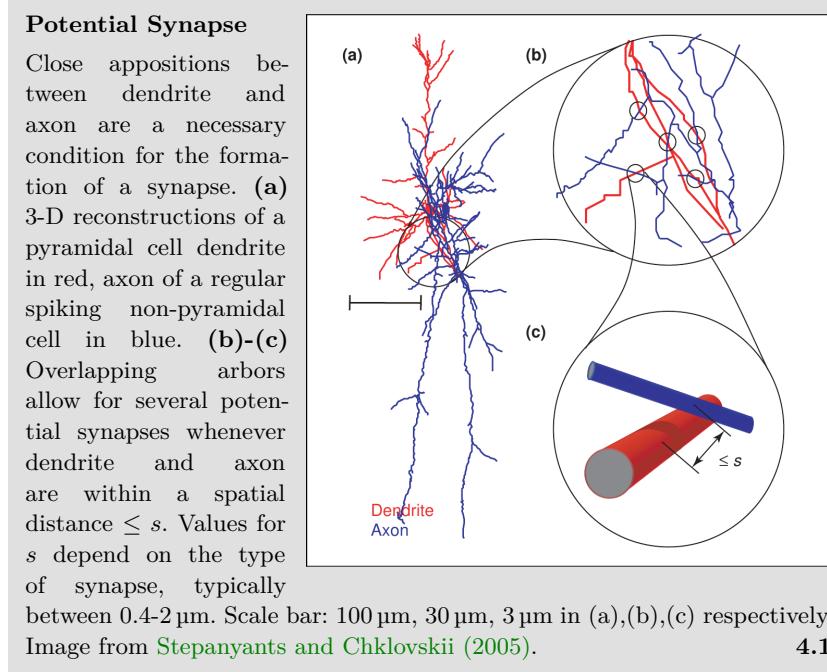
### 4.1 OVERVIEW

This chapter introduces the central object of this study, the *anisotropic network model*. Reviewing connectivity in local cortical circuits in Section 4.2, we identify anisotropy in connectivity in layer V pyramidal cells, inferred by their specific neuronal anatomy. Reducing the complex neuron morphology to characteristic axonal and dendritic profiles, we introduce the anisotropic network model in Section 4.3. While a graph theoretic definition allows for analytical considerations, a numerical implementation enables us not only to support the analytical observations but also gives access to results that go beyond. To fully harness this implementation we argue for the choice of a specific parameter set in Section 4.4, allowing us to generate a set of “sample graphs” to which we will refer through this study.

Using the analytical and numerical approach in conjunction, in our analysis of anisotropic networks we are most interested in identifying structural aspects that are due to the network’s anisotropy in connectivity and do not occur in similar, isotropic networks. To be able to make this distinction, in Section 4.5 we extract the distance-dependent connectivity of anisotropic networks, as it is imposed by the specific geometric relations present in the network. We then go on to introduce “rewiring” in Section 4.6, a method that allows us to manipulate the anisotropic networks to eventually display isotropy in connectivity. Finally then, Section 4.7 ties together the previous concepts by providing a measure for anisotropy and showing how rewiring is, in fact, providing the transition from anisotropic networks to networks with isotropy in connectivity.

## 4.2 ANISOTROPY IN NEURAL CONNECTIVITY

Neurogeometry addresses the problem of inferring synaptic connectivity from the geometric shapes of axon and dendrites. A fundamental concept in this field is that of a *potential synapse* (Stepanyants et al. 2002). Defined as the potential axonal-dendritic connection of two neurons, present whenever the axon of one neuron is within a spatial distance  $s$  of the dendrite of the other, it is a necessary, although not sufficient, condition for the formation of a synaptic connection (Figure 4.1). The existence of such close appositions solely depends on dendritic and axonal anatomy; identification of defining morphological characteristics in both axon and dendrite would therefore allow for a model of local network connectivity, assuming for example that a certain ratio  $r$  of potential synapses turn into active contacts independently. It is the hope that such a model, motivated from the geometry of a neuron's functional compartments, not only displays inherent patterns of connectivity similar to what has been observed in biological networks, but also proofs itself as a testing ground for how this connectivity may affect network dynamics.

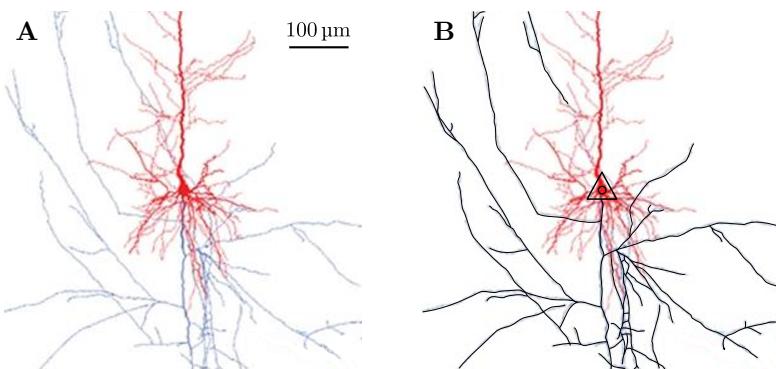


high variability in  
axonal  
morphology

Finding stereotypical anatomical characteristics however is difficult, as axonal morphology is, in general, highly diverse (Debanne 2004). Across different species, distinct regions in the central nervous system and different neuron types, axons display a wide variety of shapes characterized by morphometric parameters such as total length, branching complexity and axonal extent (Ropireddy et al. 2011). Typical exam-

ples of distinct morphology include the T-shaped axons of cerebellar granule cells branching only at a singular point (Ramon and Cajal 1911), and axons of hippocampal CA3 pyramidal cells, which, in stark contrast, may feature up to 40 branches resulting in a total length of axon collaterals of up to 12 mm (Ishizuka et al. 1990).

It is therefore imperative to confine this analysis to a specific brain region and neuron type. In this study, we set the focus on circuits of pyramidal cells in the mammalian cortex. More specifically, local circuits of thick tufted layer 5 pyramidal neurons in the rat's somatosensory cortex have been the target of advanced experimentation (Song et al. 2005; Perin et al. 2011; Romand et al. 2011; Ramaswamy et al. 2012), and will serve as a benchmark for results in neural morphology and network connectivity in this report.



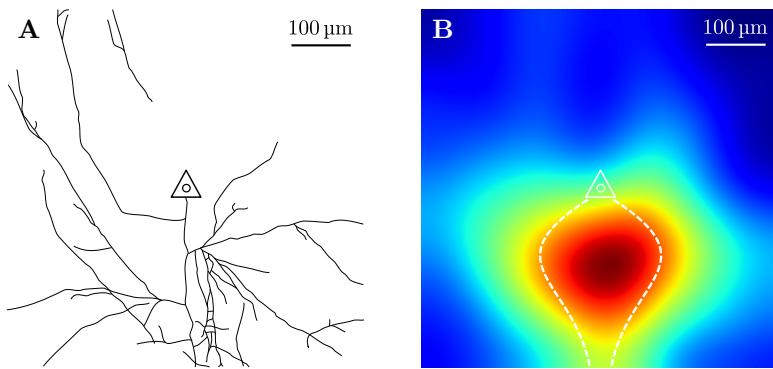
**Figure 4.2: Tracing axonal branching of a pyramidal cell** In a 3-D model reconstructed from biocytin-labeled thick-tufted layer 5 pyramidal cells in the somatosensory cortex of postnatal (day 14) Wistar rats, Romand et al. (2011) depict dendritic compartments in red, axonal compartments in blue. **A)** A 600  $\mu\text{m}$  window centered on the soma of the pyramidal cell shows the main stem of the cell's axon projecting downwards in a straight line, collaterals branching at various angles. **B)** Using image manipulation software, axon morphology was manually traced and is emphasized in black.

Axonal morphology of pyramidal cells in the cerebral cortex is well described. From the soma the single main stem of the axon originates and projects downwards, describing a trajectory closely resembling a straight line (Braitenberg and Schüz 1998). At arbitrary points along this path, collaterals branch off at various angles and constitute themselves linear paths until they further ramify or terminate. Displaying a high degree of ramification, axonal trees of cortical pyramidal cells build, in general, complex structures (Petersen et al. 2003; Ramaswamy et al. 2012). Cortical slice experiments analyzing neural anatomy are typically constrained by a slice thickness of 300  $\mu\text{m}$ . On this scale, 3-D reconstruction from labeled thick tufted layer 5 pyramidal cells reveals

cortical axons  
form straight lines,  
arborize profusely

characteristic morphology of the axonal tree (Figure 4.2). The downwards projecting, straight axon branches at several points, forming collateral branches that travel in linear path as well.

In a statistical view, this characteristic axonal morphology results in high axon branch densities along the main stem, whereas distant regions display a relatively low density (Figure 4.3). Specifically, axon collaterals do not cluster around the soma but align with the main stem's projection. As presence of an axonal branch constitutes a necessary condition for a potential synapse, a higher concentration of potential and, subsequently, realized synapses is expected in regions of high branch density. For a coherent picture of local connectivity profiles, however, dendritic morphology needs to be considered as well.

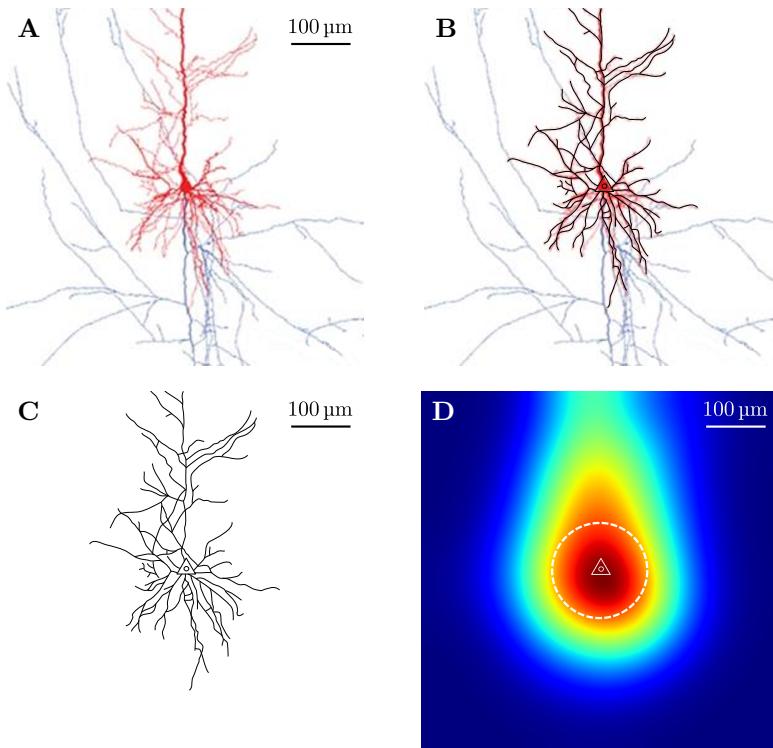


**Figure 4.3: Illustrating axonal branch density** In a sample of 5 reconstructions from thick-tufted layer 5 pyramidal cells (Romand et al. 2011), tracing axonal morphology illustrates characteristic branch density along the axon's main stem. **A)** Example of extracted axonal tree. Outline manually traced using image manipulation software. Soma indicated by triangle. Original data from Romand et al. (2011). **B)** Overlaying 5 axonal trees extracted as in A), applying a Gaussian filter and displaying high axon densities in warm colors, illustrates the characteristic higher branch densities along the axon's main stem.

*basal dendrites dominate local connectivity*

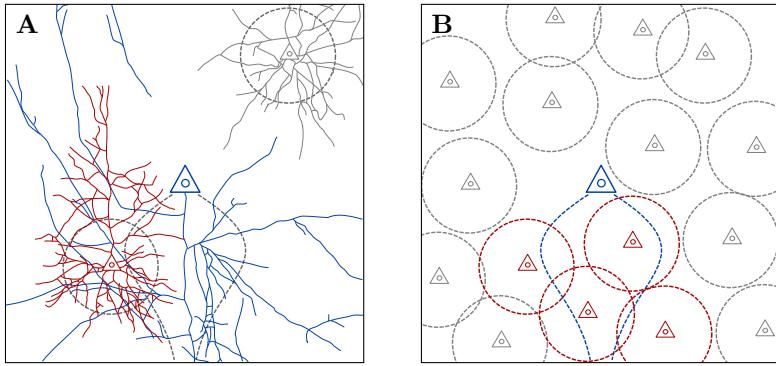
Dendritic anatomy of cortical pyramidal cells is inherently bipartite. From the soma several *basal dendrites* emerge and extend into arbitrary directions, branching profusely until they terminate. The single *apical dendrite* emerges from the apex of the pyramidal cell and ascends in a linear trajectory, forming occasional collateral branches until finally terminating into the apical tuft, where the dendrite branches several times to form a tree like structure (Feldman 1984). On the scale of typical cortical slice thickness, however, the apical dendrite is cut off and the basal dendrite dominates the dendritic morphology and potential of dendritic-axonal connections (Figure 4.4). The radial extension of dendritic branches results in a high concentration of dendritic branches

around the soma, much in the contrast to the findings of axonal branch densities before.



**Figure 4.4: Dendritic morphology and branch density** Using neuronal morphology of thick-tufted layer 5 pyramidal cells recorded by Romand et al. (2011), dendritic anatomy is traced and combined to illustrate high branch density around the soma. **A)** In a 600  $\mu\text{m}$  window centered on the soma, basal dendrites (red) are visible extending around the soma. The ascending thick apical dendrite (red) is cut off and apical tuft is not shown. **B)-C)** Manual tracing of dendritic outlines in five samples (one shown), allows for clearer identification of stereotypical morphology and later analysis. **D)** Combining 5 dendritic outlines as shown in C) and subsequent Gaussian filtering reveals the relatively high dendritic branch density around the soma.

Combining the above results of dendritic and axonal branch densities in the light of neurogeometry, a clear concept of anisotropy of neural connectivity emerges. As dendritic branches of potential post-synaptic targets extend radially from the soma and do not display a preferred direction, target neurons for outgoing synaptic contacts originating from a single pyramidal cell, cluster around the downwards projecting axon (Figure 4.5). In their in-depth study, Stepanyants and Chklovskii (2005) confirm the overrepresentation of potential synapses along the axon for pyramidal cells. Consistent with the notion that stereotypical morphology of pyramidal cells is intrinsic to the local network's connectivity



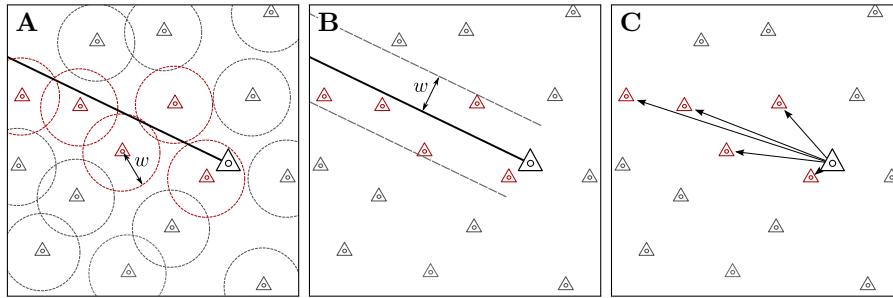
**Figure 4.5: Connected neurons of a single pyramidal cell align with axonal projection** Reducing the full axonal (blue, cf. Figure 4.2) and dendritic trees (red, gray, cf. Figure 4.4) as shown for two neurons in A) to their stereotypical axonal (blue) and dendritic profiles (red, gray) in B), demonstrates how connected neurons (red) tend to cluster around the pre-synaptic axon's profile, as spatial closeness constitutes a necessary condition for the formation of contacts. Unconnected neurons (gray) are found distant from the axon's projection, but not necessarily distant from the soma.

profile, they also find that anisotropy of this degree is *not* present in spiny stellate neurons located in lower-layer-4.

### 4.3 ANISOTROPIC GEOMETRIC NETWORK MODEL

Taking up the concept of anisotropy in neural connectivity introduced in the last section, we propose here, as basis for this study, a simple geometric network model featuring anisotropic connectivity. Constructing such a model, we're challenged with resembling the anisotropic aspects outlined in the last section as closely as possible, while at the same time basing the model on simple and abstract relations to allow for an analytical study of such anisotropic networks.

With this in mind, we propose the following model: On a square surface of side length  $s$ , a number of  $N$  point neurons are randomly, uniformly distributed. Connected neighbors are then calculated for each neuron separately and independently, by determining the randomly, uniformly distributed direction of the neuron's single axon. In this direction the axon traverses over the surface describing a straight path, terminating only when an edge of the surface is reached. Directed contacts are made with every neuron that is within a width  $w(x)$  of the axon's trajectory, where in general  $w$  depends on the axon length  $x$  at this point (Figure 4.6).



**Figure 4.6: Anisotropic geometric network model and interpretations of width parameter  $w$**  Illustrating the process of finding connections for one neuron (large triangle, black), the axon describes a linear trajectory in an arbitrary direction and until terminating on the surface's edge. Target neurons (red) are encountered along the path within a (here constant) distance  $w$ , which is in **A**) interpreted as a dendritic radius or, equivalently, in **B**) as a “bandwidth” of the axon. Connections to the encountered targets are then established as projections in **C**), consistent with the directed nature of synapses in biological networks (cf. Chapter 2).

The implementation of arbitrary axonal orientation is crucial to the model. Although cortical axons are described as consistently projecting downwards (Braitenberg and Schüz 1998, cf. Section 4.2), combining exclusively vertically aligned axons with the simplified axonal and dendritic morphological profiles would result in a “vertically staggered

random axonal  
orientation yields  
relevant  
connectivity

connectivity" - neurons could then only project to targets located below them. It is in fact not a vertical alignment of axon orientation, but the anisotropy in neural connectivity - the observation of neuronal targets aligning with the axonal projection - that we try to capture and analyze in this model.

We will refer to the model as the *anisotropic geometric network model*. Trying to provide a simple, abstract model isolating anisotropy in connectivity, in most of this study the width  $w(x)$  is assumed to be constant,  $w(x) = w$ , a notable exception being the development of tuned networks in Section 5.5. In the graph theoretic context the anisotropic network model is a random graph model, in which a realization of the random process results in a geometric directed graph with a special mode of connectivity. We can formally define such realization as:

**Definition 4.1** (Anisotropic geometric graph). Let  $n \in \mathbb{N}$  and  $w \in (0, \infty)$ . An *anisotropic geometric graph*  $G_{n,w}$  then consists of a tuple  $(G, \Phi, a)$ , of a simple directed graph  $G$  with  $|V(G)| = n$  vertices and the maps  $\Phi : V(G) \rightarrow [0, 1]^2$  and  $a : V(G) \rightarrow [0, 2\pi)$ , such that for every vertex pair  $v, v' \in V(G)$  and edge  $e \in E(G)$  with  $s(e) = v$  and  $t(e) = v'$  exists if and only if the inequalities

$$R_{-a(v)} (\Phi(v') - \Phi(v)) \hat{e}_x \geq 0 \quad \text{and} \quad \left| R_{-a(v)} (\Phi(v') - \Phi(v)) \hat{e}_y \right| \leq \frac{w}{2}$$

hold. Here  $R_\varphi$  is the rotation matrix of angle  $\varphi$  in the Cartesian plane and  $\hat{e}_x, \hat{e}_y$  are the standard unit vectors.

The anisotropic random graph model then is then giving the probability distribution over the set of anisotropic random graphs by describing a random process generating such graph.

**Definition 4.2** (Anisotropic random graph model). Let  $n \in \mathbb{N}$  and  $w > 0$ . The *anisotropic random graph model*  $G(n, w)$  is a probability space over the set of anisotropic geometric graphs with a probability distribution induced by the following process: Let  $G$  be an empty graph with  $n$  vertices. Assign randomly and uniformly to every vertex  $v \in V(G)$  a position  $\Phi(v) \in [0, 1]^2$  and axonal orientation  $0 \leq a(v) < 2\pi$ . Then add edges such that  $(G, \Phi, a)$  is an anisotropic geometric graph  $G_{n,w}$ .

As with every geometric graph model introduced, we restrict the surface to be the unit square. This does not limit the model, as only the relative width of the axon band in regard to the surface's side length is determining connectivity statistics - the expected number of connections is easily obtained by the quotient of the area covered by the axon and the surface area, making connectivity statistics in the anisotropic random graph model scale-free.

The following maybe interpreted as a study of anisotropic geometric graphs in the light of a neuroscientific context. To enable such an analysis, a few more concepts are needed. The introduction of those concepts composes the rest of the chapter. A first important step is the numerical implementation of the anisotropic network model.

#### 4.4 NUMERICAL IMPLEMENTATION

Numerical implementation of the anisotropic random graph model was achieved in Python<sup>1</sup>. Relying on NumPy as part of the scientific Python library SciPy<sup>2</sup> for the more complex mathematical computations, the implementation also uses graph-tool<sup>3</sup>, to ensure convenient and efficient handling of the created networks.

The algorithm for the generation of anisotropic networks closely resembles Definition 4.1. After randomly distributing  $N$  neurons on the square of side-length  $s$ , for every neuron a random axon orientation  $a \in [0, 2\pi)$  is chosen and an affine transformation, such that the current neuron is located at the origin and its axon projection aligns with the positive x-axis, secures a straightforward implementation of connectivity, using the the inequalities in Definition 4.1 as a rule for establishing connections.

To harness the numerical implemenation to generate networks, a set of parameters needs to be chosen. The network size  $N$  strongly influences the needed computational efforts in calculations based on the generated graphs and has thus been set to  $N = 1000$ . Choosing the surface side-length arbitrarily as  $s = 100$ , the axon width  $w$  determines connectivity in the network, the relation between width  $w$  and overall connection probability  $p$  being shown in Figure 4.7. In their analysis of connectivity of thick-tufted layer 5 pyramidal cells in neonatal rats (day 14), Song et al. (2005) report an overall connection probability of  $p = 0.116$ , consistent with prior reports of a cortical connection probability of  $p \approx 0.1$ . Choosing  $w$  to be constant, we determine the axon width such that overall connectivity matches the value report by Song et al. and obtain  $w/2 = 12.6$  (Figure 4.7).

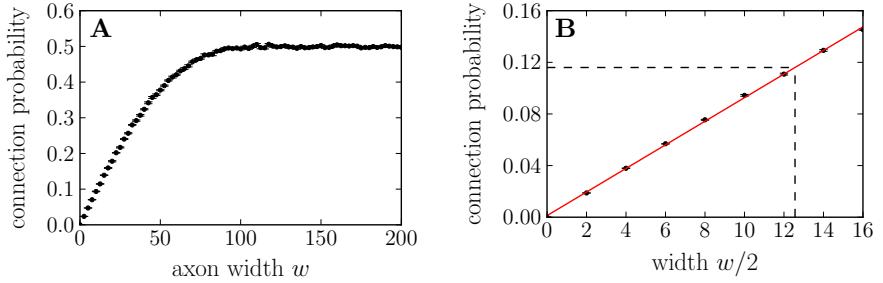
*parameter set  
chosen to resemble  
cortical circuits*

Having determined a suitable set of parameters as  $N = 1000$ ,  $s = 100$  and  $w = 25.2$ , we generate 25 graphs with this parameter set

<sup>1</sup> Python Software Foundation. Python Language Reference, version 2.7. Available at <http://www.python.org>

<sup>2</sup> Eric Jones, Travis Oliphant, Pearu Peterson and others. NumPy version 1.6.1. Available at <http://www.scipy.org>

<sup>3</sup> Tiago P. Peixoto. Efficient network analysis. Version 2.2.18. Available at <http://graph-tool.skewed.de/>



**Figure 4.7: Axon width dependent connection probability determines parameter for numerical analysis** Generating anisotropic networks with different axon widths  $w$  and extracting probability  $p$  of directed connection between two random nodes, demonstrates the dependency of  $p$  on the width parameter  $w$ . **A)** At an axon width of over  $w = 100$ , exceeding the square's side length, the connection probability saturates at  $p = 0.5$ , as axon bands are essentially “cutting” the square in a connected and unconnected half (c5b64f3e). **B)** For small  $w$  the connection probability is a linear function of  $w$ , allowing the width  $w_S/2$  at which  $p(w_S) = 0.116$  to be determined by a linear fit as  $w_S/2 = 12.6$  (585a946f).

sample graphs as  
reference for  
structural analysis

(label: N1000w\_ax126-flat\_graph0-24). This set of sample graphs will serve as a reference for the following structural analysis. Extending the set by the (partially) rewired sample graphs (see Section 4.6) and by purely distance-dependent graphs best resembling the anisotropic networks (see Section 4.5) we obtain a resourceful reference for the analysis of structural features of anisotropic geometric graphs, that we will frequently employ to obtain quantitative and qualitative results.

## 4.5 DISTANCE DEPENDENT CONNECTIVITY

In Gilbert's random graph model  $G(n, p)$ , probability of connection  $p$  is independently chosen and a fixed value for all vertex pairs. The anisotropic geometric graph model introduced in Section 4.3 is itself a random graph model - node positions as well as preferred directions of connection are uniformly at random distributed. In contrast to Gilbert's model however, neither is the probability of connection between a given vertex pair independent of the realization of other edges in the graph, nor is it a fixed value - probabilities strongly depend on internode distance in the anisotropic geometric graph model introduced.

*random graph  
models in  
Section 3.3*

Analyzing dependencies in the anisotropic model, specifically by identifying prevalent patterns of connectivity and relating these modes of non-randomness to biological findings, is the main focus of Chapter 5. However, such structural correlations may not necessarily be an inherent feature of the network's anisotropy - distance dependent connectivity alone, as imposed by the model's specific geometry, may be the cause for emerging dependencies. It is therefore a crucial initial task to map the anisotropic model's distance dependent connection probability. Inferring connection probability as a function of internode distance and comparing it with computational results, in this section we explore distance connectivity of the anisotropic network model, securing a vital component in the analysis of structural features.

**Theorem 4.3.** *Let  $(G, \Phi, a)$  represent an arbitrary realization of the anisotropic random graph model  $G(n, w)$ . Define  $C : [0, \sqrt{2}] \rightarrow [0, 1]$  as the distance-dependent connection probability profile of  $(G, \Phi)$ , that is such that  $C(x)$  is the probability that for a vertex pair  $(v, v') \in V(G)^2 \setminus \Delta_{V(G)}$  in distance  $x = \|\Phi(v) - \Phi(v')\|$  the vertex  $v$  projects to vertex  $v'$ . Then*

$$C(x) = \begin{cases} \frac{1}{2} & \text{for } x \leq w/2 \\ \frac{1}{\pi} \arcsin\left(\frac{w}{2x}\right) & \text{for } x > w/2. \end{cases}$$

*Proof.* Let  $v, v'$  be a pair of vertices in  $V(G)^2 \setminus \Delta_{V(G)}$  in Euclidean distance  $x$  of each other. The vector difference  $\Phi(v') - \Phi(v)$  may then be written as  $xe^{i\theta}$ , with  $0 \leq \theta < 2\pi$ . We have

$$R_{-\alpha(v)} xe^{i\theta} = xe^{i(\theta - \alpha(v))}.$$

Only for suitable combination of  $\theta$  and  $\alpha(v)$  an edge from  $v$  to  $v'$  exists. Assuming  $\alpha(v)$  fixed, we calculate the probability of connection depending on the random choice of  $\theta$ . We can assume  $\alpha(v) = 0$ , otherwise the same argument holds for  $\theta' = \theta - \alpha(v)$ .

From 4.1 we obtain the necessary and sufficient conditions

$$x \cos \theta \geq 0 \quad \text{and} \quad |x \sin \theta| \leq \frac{w}{2}.$$

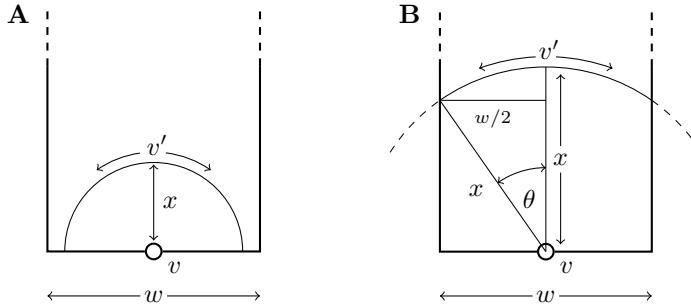
Choosing uniformly at random  $\theta$  in the range of  $[0, 2\pi)$ , the first condition is satisfied with a probability of  $\frac{1}{2}$ . Consider for the second condition  $\theta \in [0, \pi)$ . We have

$$\sin \theta \leq \frac{w}{2x},$$

and for  $x \leq \frac{w}{2}$  the inequality holds for all  $\theta$  by definition of  $\sin \theta$ . In the case of  $x > \frac{w}{2}$ , we note that for the first condition to hold  $\theta$  must already be in  $[0, \frac{\pi}{2})$  and can thus write the second condition  $\theta$  as

$$\theta \leq \arcsin \frac{w}{2x},$$

yielding  $C(x)$  by combining the considerations above and using the symmetry of sine for  $\theta$  in the third and fourth quadrant.  $\square$



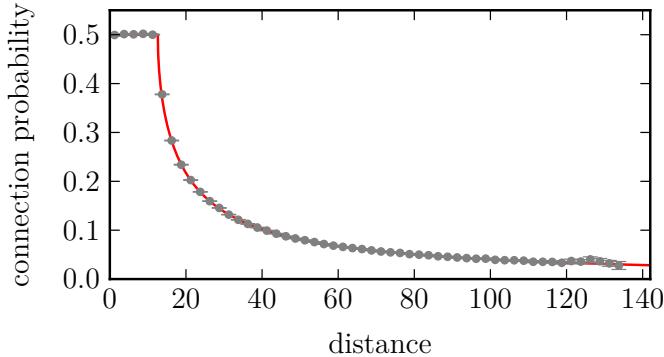
**Figure 4.8: Illustrating the proof of Theorem 4.3** Distance-dependent connectivity profile  $C(x)$  in an anisotropic geometric graph calculated from geometric dependencies. **A)** In the case of  $x \leq w/2$ , target  $v'$  may be located anywhere on the shown semicircle and therefore receives input from  $v$  with probability  $1/2$ . **B)** For  $x > w/2$ , suitable positions for target  $v'$  are dependent on  $x$ . The geometric relation  $\sin \theta = w/2x$  leads to the distance-dependent connectivity profile as described in Theorem 4.3.

distance-dependent sample graphs as reference

We can verify this result by extracting the distance-dependent connection probabilities in the sample graphs created in Section 4.4. Combining data of all 25 graphs, we find that connection probabilities perfectly match the theoretical prediction (Figure 4.9). Additionally we're able to extend the reference sample graphs by distance-dependent networks (Definition 3.13). Using Theorem 4.3 in conjunction with the sample graph parameter set ( $N = 1000$ ,  $s = 100^4$ ,  $w = 25.2$ ) we easily obtain the expected distance-dependent connectivity profile for the

<sup>4</sup> The generalization of Theorem 4.3 to allow for arbitrary side-length  $s$  is trivial and omitted here

created sample graphs and, using this profile, generate purely distance-dependent networks<sup>5</sup>. Being highly interested in structural features not explained by distance-dependent connectivity, the numerical analysis in this work will heavily rely on these networks to identify aspects that are inherent to the anisotropy in connectivity.



**Figure 4.9: Predicted distance-dependent connection probability profile is matched by numerical results** Averaging distance-dependent connection probabilities over the 25 sample graphs, we find the expected profile calculated in Theorem 4.3 is matched perfectly by the numerical results. (dbffa88e)

## 4.6 REWIRING

Distance-dependency as identified in the last section may already account for many of the structural features present in anisotropic networks. A central question of this study is: What structural aspects in the network are truly features of the anisotropy in connectivity? Although a quantitative measure for anisotropy will only be introduced in the next section, already here we are able to qualitatively observe the strong directionality in connectivity - edges originating from one node “point in the same direction”, effectively aligning with the orientation of the axonal projection of the source node (cf. Figure 4.6). To answer the question above, we need to introduce a method that eliminates this directionality, making networks essentially isotropic in connectivity. Then, structural features present in the original anisotropic networks, but not in their rewired, isotropic counterparts may be attributed to anisotropy.

eliminate  
anisotropy to find  
structures caused  
by it

Rewiring as introduced here, provides the transition from anisotropic connectivity to networks isotropic in connectivity, closely resembling purely distance-dependent networks. Applying this process only partially then allows us to analyse structural features as they change with

<sup>5</sup> label: N1000-dist\_depend-flat\_graph-00-24.xml.gz

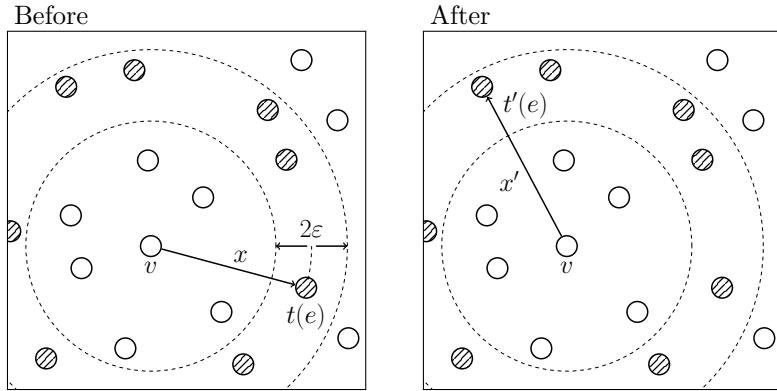
a varying degree of isotropy, asserting the importance of this process to our study. In designing the specific rewiring algorithm we identify two requirements that our implementation should satisfy:

1. elimination of anisotropy in connectivity
2. preservation of distance-dependent connectivity

The second point is especially important to us, as we want to impose isotropy on the network at “minimal cost”, that is by changing as little as possible about the other characteristics of the network’s connectivity. The following process respects both of the points above:

For every edge between vertices  $v$  and  $v'$  with inter-vertex distance  $x$ , identify neurons with distance to  $v$  in the range of  $(x - \varepsilon, x + \varepsilon)$  as potential new targets. Then pick at random one of these vertices (including  $v'$ ) as a new target for the current edge, if such an edge doesn’t already exist ([Figure 4.10](#)).

In the graph theoretic context we formally define a rewiring as follows:



**Figure 4.10: Rewiring transforms anisotropic geometric graphs to networks with isotropic connectivity** For a given edge  $e$  with a distance  $x$  from its source vertex  $v$  to its target vertex  $t(e)$ , potential new targets (striped) are found in within a distance  $(x - \varepsilon, x + \varepsilon)$  of  $v$ . The rewired edge then projects from  $v$  to a new target  $t'(e)$ , randomly chosen from the set of vertices within in this range. Inter-vertex distance between  $v$  and  $t'(e)$  differs by less than  $\varepsilon$  from  $x$ , ensuring that for small  $\varepsilon$  the original distance-dependent connectivity is preserved. (Note that all targets within range are eligible for rewiring as no other edges exist. In general this is not the case.)

**Definition 4.4.** Let  $G$  be an anisotropic geometric graph with  $|V(G)| = n$  and  $\varepsilon > 0$ . Then we define a *rewiring*  $R_\varepsilon$  of  $G$  to be probability space over  $G_\Phi^n$ , induced by the following process: For every edge  $e \in E(G)$

uniformly at random pick a potential new target  $t'(e)$  from the set  $M_e = T_e \setminus K_e$ , where  $T_e$  is the set of all vertices that differ in their distance to  $s(e)$  less than  $\varepsilon$  from the distance of  $s(e)$  to  $t(e)$ ,

$$T_e = \{v \in V(G) \setminus s(e) \mid |\text{d}(s(e), v) - \text{d}(s(e), t(e))| < \varepsilon\}$$

and  $K_e$  the set vertices that already are connected to  $s(e)$  by another rewired edge,

$$K_e = \{v \in V(G) \mid \exists e' \in E'(G) : s(e') = s(e), t(e') = v\},$$

where  $E'(G)$  is the set of all edges that have been rewired already.

Note that in the way Definition 4.4 is formulated, it is possible for  $M_e$  to be empty for some edge  $e$ . In this case no new edge is realized and the resulting, rewired network has  $|E(G)| - 1$  edges. In practice this happens negligibly seldom, out of approximately on average only 25.68 edges, with a standard deviation of 4.51 and accounting for roughly 0.02% of the rewired edges, are “lost” in this process (4afc2727).

We formulated Definition 4.4 in such a way, that distance-dependent connectivity is preserved. We verify this claim by the following estimation:

Let  $\tilde{C}(x)$  be the distance-dependent connectivity profile of a rewiring  $R_\varepsilon$  of an anisotropic graph  $G_{n,w}$ . Denote with  $C(x)$  the distance-dependent connection probability of the  $G_{n,w}$ . The expected value for  $\tilde{C}(x)$  at any inter-vertex distance  $x \in [0, \sqrt{2}]$  is given as an average over the connection probabilities  $C(x)$  of the possible sources to the new edge at distance  $x$

$$\mathbf{E} [\tilde{C}(x)] = \frac{1}{2\varepsilon} \int_{x-\varepsilon}^{x+\varepsilon} C(x') dx'.$$

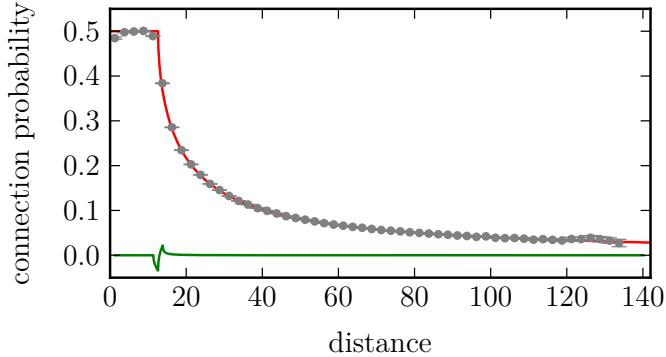
Note that for this expression to be well defined in the boundary cases, we need to extend  $C(x)$  to have  $C(x) = 0$  for  $x < 0$  and  $x > \sqrt{2}$ . We can then estimate the expected difference between  $\tilde{C}(x)$  and  $C(x)$  at any point  $x$  as

$$\begin{aligned} |\mathbf{E} [\tilde{C}(x) - C(x)]| &= \left| \frac{1}{2\varepsilon} \int_{x-\varepsilon}^{x+\varepsilon} C(x') - C(x) dx' \right| \\ &= \frac{1}{2\varepsilon} \left\{ \int_{x-\varepsilon}^x C(x') - C(x) dx - \int_x^{x+\varepsilon} C(x') - C(x) dx \right\} \\ &\leq \frac{1}{2\varepsilon}, \end{aligned} \tag{4.1}$$

The rewiring margin  $\varepsilon$  thus simultaneously governs how many new targets are available for each edge and how well distance-dependency is preserved. Setting  $\varepsilon = 1.25$  and applying the rewiring algorithm to the 25 sample graphs, we find that distance-dependent connectivity of

$d(v, w) = \|\Phi(v) - \Phi(w)\|$   
Euclidean distance  
between vertices

the original graphs is matched (Figure 4.11) while at the same time ensuring that for any edge  $e$  sufficiently many new rewiring targets are available (Figure A.1).



**Figure 4.11: Rewiring with  $\varepsilon = 1.25$  preserves distance-dependent profile in sample graphs** Comparing the distance-dependent connection probabilities of the original graph (Theorem 4.3) in red with extracted of probabilities from the rewired ( $\varepsilon = 1.25$ ) sample graphs in gray (errorbars SEM<sup>6</sup>) we verify that distance-dependent connectivity is preserved when rewiring. The green curve shows the expected difference between the original and rewired distance profiles as estimated in Equation 4.1. (4f4dfcf1)

As a generalization of Definition 4.4, we define a partial rewiring  $R_{\varepsilon,\eta}$ , finding new targets only for a fraction  $\eta$  of all edges:

**Definition 4.5.** Let  $\varepsilon > 0$  and  $0 \leq \eta \leq 1$ . A *partial rewiring*  $R_{\varepsilon,\eta}$  of an anisotropic geometric graph  $G_{n,w}$  with *rewiring factor*  $\eta$  is then a rewiring  $R_\varepsilon$  of  $G_{n,w}$ , in which every edge is rewired with a probability of  $\eta$ , otherwise it remains. To avoid the occurrence of multiple edges,  $K_e$  is then extended to include the targets of all edges originating from  $s(e)$  that will not be rewired.

Clearly, as with full rewiring, partial rewiring also preserves distance-dependent connectivity. Using the algorithm we extend our set of sample graphs once more by adding rewired versions of each graph. Choosing a rewiring margin of  $\varepsilon = 1.25$ , with fractions  $\eta = 0.25$ ,  $\eta = 0.5$ ,  $\eta = 0.75$  and  $\eta = 1$  we obtain through rewiring five stages of the sample graphs, from complete anisotropy in connectivity in the original graphs to the isotropic, fully rewired graphs<sup>7</sup>. By introducing a measure for anisotropy and applying it to the rewired and original version of the

rewiring of sample  
graphs

<sup>6</sup> standard error of the mean

<sup>7</sup> labels: N1000w\_ax126-flat\_graph0-25-rewired\_frac025, -50,-75,-100

sample graphs, we're able to solidify this notion and tie together the concepts introduced through this chapter in the next section.

#### 4.7 ANISOTROPY MEASURE

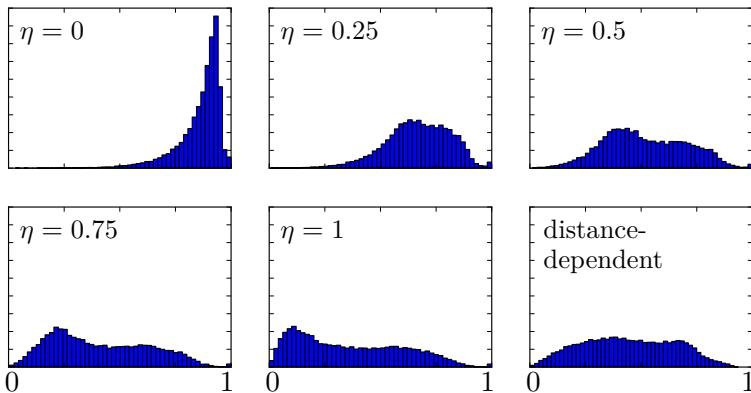
In the last section a method to rewire an anisotropic geometric graph, such that was introduced. From an . In this chapter we introduce.. capturing ..

The  $G_{n,\Phi}$  be a geometric graph. Then, for every is the *preferred direction* and its length is

Mardia and Jupp (2000)

Anisotropic model can be interpreted as a model maximizing anisotropy while keeping the cutoffs (degree distribution).

**Figure 4.12:** illustrate varying levels of anisotropy



**Figure 4.13:** Rewiring significantly reduces anisotropy In data taken from the 25 sample graphs (Section 4.4), vertex isotropy degree distribution is shown for the original set of graphs ( $\eta = 0$ ) The characteristic highly anisotropic profile found in the original is already significantly reduced by partial rewiring; anisotropy degree distribution in the fully rewired graphs resemble degree distribution of equivalent purely distance-dependent networks.

... suggesting that fully rewired anisotropic networks do not . There is however one difference in out-degree as an artifact of boundary confinement (Section 5.2).

## 4.8 SUMMARY AND DISCUSSION

has been introduced, rewiring, .

- Gilbert random graph
- distance-dependent graph
- fully rewired anisotropic graph
- anisotropic network

the claim is that distance-dependent and fully rewired are more or less equivalent (hints of non-equivalence found in ...) and so



# 5

## STRUCTURAL ASPECTS

---

Subjecting the anisotropic network model to a critical examination of its structural features, we identify prevalent patterns of connectivity and relate theoretical and computational results to findings from experiments in the rat's cortex.

### 5.1 INTRODUCTION

Investigation of the brain's connectivity is an ongoing endeavour. Concurrent collaborative efforts like the Human Connectome Project, the Open Connectome Project and the Allen Brain Atlas, intent on mapping the 'wiring' of the brain, as well as the continued development of experimental techniques and computational resources, demonstrate the great interest in advancing this field.

Research in brain connectivity spreads over the whole scale of the brain; from the mapping of fiber pathways between brain regions at the macroscopic level, to the synaptic connections of individual neurons on the microscale, researchers are trying to identify the links that enable the brain its characteristic cognitive abilities. Connections, these links are of anatomical nature. However, statistical dependencies and causal relationships between the distinct computational units in the brain are being researched with equal emphasis (Sporns 2007). Connectivity in the context of the anisotropic network model introduced in Section 4.3, refers in this chapter to structural links. So far, we have only briefly mentioned that the network's nodes should be interpreted as individual neurons; to allow for a discussion of functional relationships between nodes, we have yet to provided a physical description of a neuron's function. Here we explore the network's structural connectivity, modeling synaptic contacts between axon and dendrites of individual neurons.

In the local cortical circuits the anisotropic geometric model was derived from, synaptic connectivity is a major mode of configuration. In those networks, connectivity has been determined to be neither completely random nor exclusively specific; recurring patterns of connectivity have been identified by several reports (Sporns and Kötter 2004; Song et al. 2005; Perin et al. 2011).

HUMAN  
Connectome  
PROJECT  
[humanconnectome.org](http://humanconnectome.org)

Open Connectome  
Project  
[openconnectome-  
project.org](http://openconnectome-project.org)

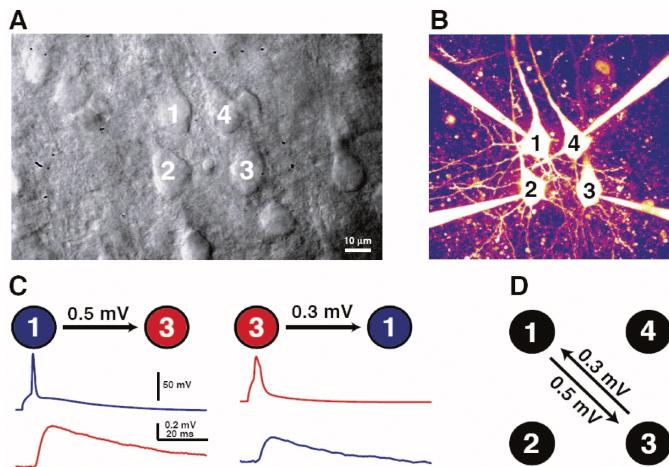
ALLEN BRAIN ATLAS  
[brain-map.org](http://brain-map.org)

synaptic  
connectivity

The impact of this structural specificity discovered in local networks is shown to be significant; while linking network structure to network dynamics remains an active field of research, several studies were able to employ computational and theoretical models to establish such a connection. A study by Zhao et al. (2011), for example, demonstrates how second order connectivity statistics affect a network's propensity to synchronize. In the same year, Alex Roxin reported on the influence of in- and out-degree distributions on dynamics of neural network (Roxin 2011). Later, Pernice et al. were able to link structural connectivity to spike train correlations in neural networks (Pernice et al. 2011).

#### *mapping synaptic connectivity in experiments*

Experimentally, paired intracellular recordings are used to determine synaptic connectivity in cortical slices. Using two electrodes, one inserted in the cell and one outside the cell, a single intracellular recording allows for measurement of a cell's membrane potential (Brette and Destexhe 2012, Chapter 3; Weckstrom 2010). Simultaneous recordings from multiple neurons are then able to infer synaptic connectivity by evoking an action potential through current injection in one neuron and observing the change of membrane potential in the other cells (Song et al. 2005).



**Figure 5.1:** Song et al. use quadruple whole-cell recordings, observing simultaneously the membrane potential of four neurons. **A)** Contrast image showing four thick-tufted L5 neurons **B)** Fluorescent image of the same cells after patching on **C)** Evoking an action potential in the presynaptic neuron causes characteristic membrane potential change in the postsynaptic neuron **D)** Inferring synaptic connectivity from the EPSP waveform observed in C). Image from Song et al. (2005).

While techniques for paired intracellular recordings are rapidly developing, their ability to capture connectivity patterns of large networks is yet very limited. To this date, the connectome of *C. Elegans* remains

the outstanding exception of a connectivity configuration that has been fully mapped (White et al. 1986). Even in the state-of-the-art experiment conducted by Perin et al., using a setup capable of recording up to twelve neurons simultaneously, the authors note that an investigation of degree distribution was not carried out, due to lack of sufficient data (Perin et al. 2011).

Working with a geometrical network model and its computational implementation, such restrictions disappear; the full information about the network, in form of its connectivity matrix, is given at point in time and can be easily queried for. Experiments that may take days to perform *in vivo*, can be completed in a matter of seconds *in silico*. As such, geometrical models lend themselves to extensive examination of their structural aspects. In trying to exploit these advantages, two approaches present themselves. One may construct a network model that extrapolates the known biological configuration; a full structural examination of these networks could possibly expose relevant patterns not yet observed. For this approach a sophisticated understanding of the biological configuration is critical. Neuron morphology, however, is difficult to describe and extract. For this analysis we suggest a reductionist approach. Having motivated an abstract model reflecting a cortical network’s anisotropy in connectivity, we distinguish emerging structural patterns, specific to anisotropic networks, from results, that only indirectly stem from the network’s anisotropy, in the hopes to be able to characterize the significance of directional heterogeneity in structural connectivity of cortical circuits.

In this chapter we subject the anisotropic network model introduced in Section 4.3 to a critical analysis of its structural aspects. General network topology, as well as specific modes and patterns of connectivity, are to be identified and laid out for comparison with findings in biological neural networks. In an effort to identify structural features that can be directly associated with the network’s anisotropy in connectivity, it is crucial to differentiate such findings from results that are only indirectly caused by the network’s anisotropy. To this end we are recruiting the different network types introduced in the previous chapter throughout this analysis. Having shown a decreasing degree of anisotropy in rewired and distance-dependent networks, both models will serve as reference to compare against for structural features found in anisotropic networks. Analyzing standard graph measures in the first two sections, we quickly move on to towards neuronal network specific connectivity and anisotropy’s role in being able to model such highly non-random patterns in the later sections.

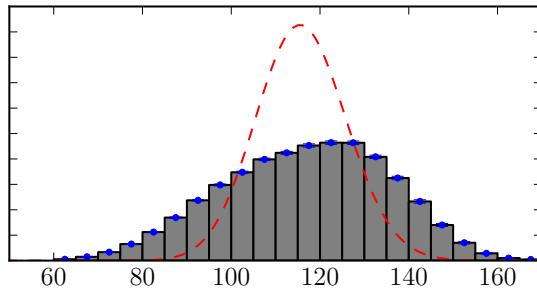
*exploiting the  
benefits of a  
geometrical model*

*rewired and  
distance-depen-  
dent networks as  
reference*

## 5.2 DEGREE DISTRIBUTION

The in- and out-degree of a vertex in a directed graph describes the number of incoming and outgoing connection from and to other vertices. *cf. Definition 3.4* As a fundamental concept in graph and network theory, the degree distribution is integral in the categorization of networks and allows for the estimation of graph properties.

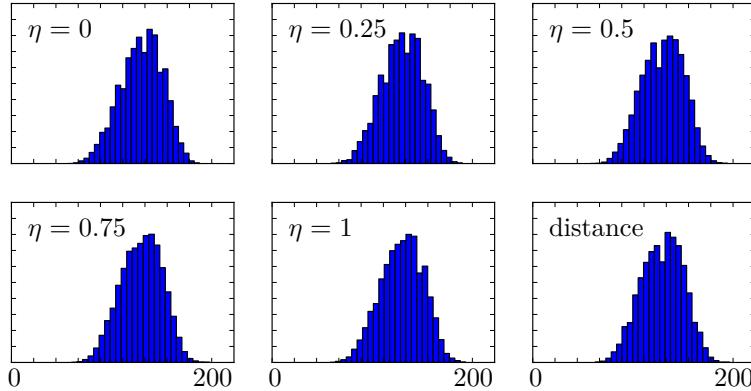
Node degree distribution was shown to have strong impact on the dynamics of neuronal networks models commonly used in computational neuroscience research (Roxin 2011). Increasing in-degree variance for example could be connected to the appearance of oscillations in the network. Extracting degree distributions from biological networks however, remains a challenge as many neurons need to be tracked simultaneously to obtain enough data to confidently estimate degree distributions.



**Figure 5.2: In-degree distribution in anisotropic networks shows comparably high variance and is skewed to the left** From 250 anisotropic networks in-degree distributions were extracted and are shown in a normed histogram plot, errorbars SEM. Comparison with the binomial degree distribution (red) of a Gilbert random graph model with matching parameter set ( $N = 1000$ ,  $p = 0.116$ ) shows higher variance of in-degrees in anisotropic networks (sample variance = 344.54, variance of binomial distribution  $Np(1 - p) = 102.44$ .) Skewness to the left of the sample is  $-0.1763$ . (9326138e)

Here we analyze in- and out-degrees in the anisotropic network model. First we find that compared to the binomial in-degree distribution of a Gilbert random graph model, in-degrees of vertices in anisotropic networks display higher variance and their distribution is skewed to the left (Figure 5.2). However, this specific in-degree profile is not an intrinsic property of anisotropy, as the distribution remains stable under manipulation of the anisotropy degree and closely matches the profile of a purely distance-dependent network (Figure 5.3). This result agrees with findings of Perin et al. (2011, Fig. S3), who were able to recreate

degree distributions from their experiment with layer 5 thick-tufted pyramidal cells in neonatal rats from the extracted distance-dependent connection profiles alone.



**Figure 5.3: In-degree distribution not affected by varying degrees of anisotropy** In-degree distributions from the 25 sample graphs and their rewiring stages are plotted in normed histograms and listed from rewiring factor  $\eta = 0$  (original anisotropic) to  $\eta = 1$  (completely rewired, maximal isotropy). Comparison shows that varying degrees of anisotropy do not influence the degree distribution, in fact in-degree distributions match with the degree distribution of an equivalent distance-dependent network shown bottom-right (77995b6b).

While the out-degree distribution of vertices in the anisotropic network also shows itself stable under rewiring, its distribution is drastically different from the out-degree distribution in a comparable distance-dependent network (Figure 5.4). The asymmetric, long-tailed distribution is identified as an artifact of the anisotropic network's spatial confinement; a neuron, closely located near a surface edge, might have an axon projection out of the square causing minimal out-degree or, projecting through the entire length of the surface, may have maximal out-degree. Approximating the expected number of outgoing connections for a vertex in an anisotropic network of size  $N$ , side-length  $s$  and axon width  $w$  as

$$N \frac{wl}{s^2},$$

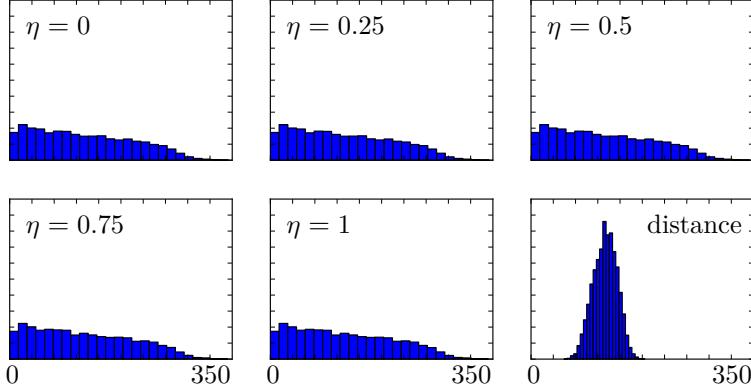
with parameters  $N = 1000$  and  $\frac{w}{s} = 0.252$ , we obtain an upper bound for the expected out-degree,

$$N \frac{wl}{s^2} \leq N \frac{w}{s} \sqrt{2} \approx 350.$$

If  $f(l)$  is the probability density function to find axon length  $l$  for a random node  $v$  in the anisotropic network model, the out-degree distribution is then approximated by

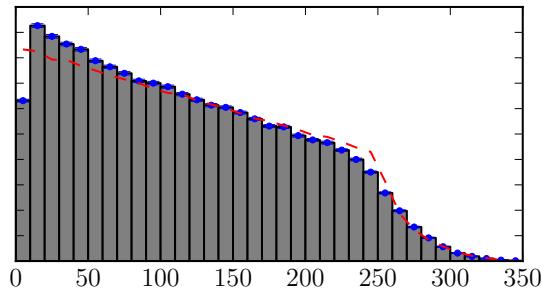
$$\mathbf{P}(d_{\text{out}}(v) = N \frac{wl}{s^2}) = f(l), \quad (5.1)$$

see also [Figure 5.5](#).



**Figure 5.4:** Out-degree distribution not affected by varying anisotropy but highly different from distance-dependent networks Out-degree distributions from the 25 sample graphs and their rewiring stages are plotted in normed histograms and listed from rewiring factor  $\eta = 0$  (original anisotropic) to  $\eta = 1$  (completely rewired, maximal isotropy). While varying degrees of anisotropy do not influence the degree distribution, the characteristic out-degree profile is drastically different from the distribution found in equivalent distance-dependent networks (77995b6b).

*Steep incline for small out-degree cut off due to binning (cf. [Figure A.2](#))*



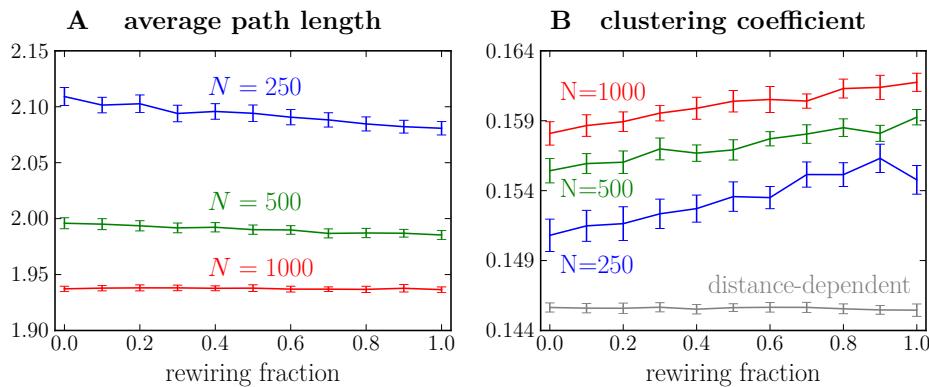
**Figure 5.5:** Characteristic out-degree distribution as an artifact of network's boundaries From 250 anisotropic networks out-degree distributions were extracted and are shown in a normed histogram plot, errorbars SEM. The characteristic distribution is identified as an artifact of the network's spatial confinement; using equation 5.1 the out-degree profile is approximated (red) by the distribution of axon lengths in the anisotropic network (019555b0).

### 5.3 SMALL WORLD PROPERTIES

Small-world networks, as described in Section 3.2, are characterized by a small average path length and comparably high clustering coefficient. In brain networks, combining advantages of sparse connectivity with mostly local and only few long-range projections, small-world properties are frequently discussed as a way of achieving high efficiency in the parallel processing of information (cf. Achard and Bullmore 2007). While most often reported on the macroscale (Sporns and Zwi 2004; Bassett and Bullmore 2006), small-worldness is also found in local cortical networks (Perin et al. 2011, SI).

Here we are interested in exploring the question whether anisotropy in connectivity influences the small-worldness of geometric networks. First we find that at a network size  $N = 1000$ , anisotropic networks display a relatively high clustering coefficient,  $c = 0.1581 \pm 0.0008$  compared with  $p = 0.116$  in random networks, and a comparable path length,  $l_{\text{aniso}} = 1.937 \pm 0.002$  and  $l_{\text{random}} = 1.8820 \pm 0.0001$ , ascertaining a small-world property in the anisotropic network model.

*in random  
networks from  
independence  
clustering = p*



**Figure 5.6: Anisotropy does not contribute to small-worldness** In increasingly rewired networks, trends show a decreasing average path length and rising clustering coefficient and thus possibly a higher degree of small-worldness in the rewired, isotropically connected networks. **A)** Average path lengths for network sizes  $N = 250$ , 500 and 1000, where vertex pairs with no existing are discarded. Individual value pairs are obtained by averaging over a trial size of 20, 15 and 5 respectively; errorbars are SEM. **B)** Network configuration as in A), additionally showing clustering coefficients for distance-dependent networks. (064f9b10)

However, is this degree of small-worldness inferred by anisotropy in connectivity? Using distance-dependent networks as a reference, we find that successively eliminating anisotropy through rewiring contributes

positively to the small-world property; with rising isotropy in the network, the characteristic path length declines in small networks and remains unchanged in larger networks, while the clustering coefficient increases regardless of network size, resulting together in rewired networks to display a higher degree of small-worldness ([Figure 5.6](#)).

In distance-dependent networks the average path length is generally smaller than in (rewired) anisotropic networks ([Figure A.3](#)), matching those of a random network as reported above. At the same time also the clustering coefficient is smaller than in anisotropic networks ([Figure 5.6](#)), resulting overall in a comparable degree of small-worldness in distance-dependent networks<sup>1</sup> and leading to the conclusion that the observed small-worldness in the anisotropic networks is due to the imposed distant-dependent connectivity rather than the anisotropy in connectivity.

#### 5.4 TWO NEURON CONNECTIONS

Connectivity in local cortical circuits exhibits a salient feature: Examining the occurrence of connections in neuron pairs, studies have repeatedly found that bidirectionally connected neuron pairs appear much more frequently than expected from the network's overall connection probability. In layer 5 of the somatosensory cortex studies from [Markram \(1997\)](#) and [Perin et al. \(2011\)](#) have found an overrepresentation of reciprocally connected pairs of thick tufted pyramidal cells, an observation that has also been reported in layer 2/3 ([Holmgren et al. 2003](#)) and layer 5 ([Song et al. 2005](#)) of the visual cortex. The overrepresentation of bidirectionally connected pairs is significant, Song et al. for example found such pairs represented four times the expected amount.

The underlying connection principle imposing this overrepresentation on the network however remains unclear. Song et al. discuss the possibility of known learning rules to explain their findings, leaving a definitive answer open to further investigation. More recent studies find overrepresentation of reciprocally connected pairs *in vitro* resulting from functional specificity ([Ko et al. 2011](#)) and *in silico* from dense neuron clustering rules ([Klinshov et al. 2014](#)), identifying specific network characteristics that may contribute to the reported overrepresentation *in vivo*.

---

<sup>1</sup> Differences in the absolute values of both path length and clustering coefficient presumably relates to difference in the out-degree distribution ([Figure 5.4](#))

Here we examine anisotropy in connectivity as a possible candidate for an underlying principle explaining the characteristic two neuron connection distribution. In random networks, the chance to encounter a specific mode of connection in a random pair of neurons can easily be computed from the overall connection probability  $p$ . For this let  $X$  be the random variable of the number of edges between two different vertices in a Gilbert graph  $G(n, p)$  with  $n \geq 2$ . As the edges are independently realized, resulting in a simple directed graph, we have

$$\begin{aligned}\mathbf{P}(X = 0) &= (1 - p)^2 && \text{unconnected pair,} \\ \mathbf{P}(X = 1) &= 2p(1 - p) && \text{single connection,} \\ \mathbf{P}(X = 2) &= p^2 && \text{reciprocal connection;}\end{aligned}\tag{5.2}$$

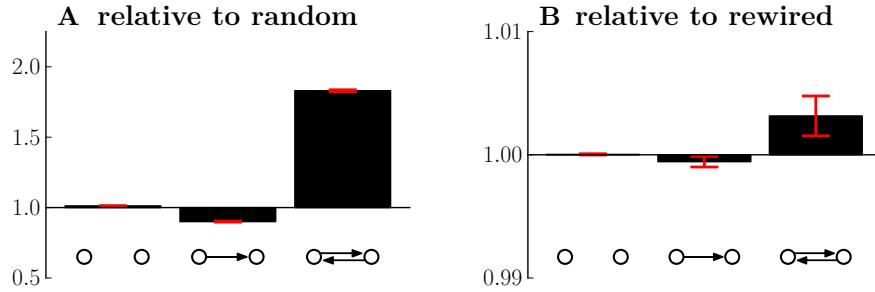
in short  $\mathbf{P}(X = k) = \mathcal{B}_{2,p}(k)$  for  $k \in \{0, 1, 2\}$  and  $\mathbf{P}(X = k) = 0$  otherwise. Using this probability distribution as the expectation for connectivity of neuron pairs in the various network types, a numeric analysis of the anisotropic sample graphs reveals that bidirectionally connected pairs appear almost twice as often as expected from the overall connection probability ( $p = 0.116$ ) and equations 5.5, similarly as reported by Song et al. (Figure 5.7 A). However, comparing the pair probabilities in anisotropic networks with the probabilities in their rewired counterparts, we find that anisotropy does not influence the occurrence of two-neuron motifs (Figure 5.7 B) In fact, expected connections in neuron pairs are identical in distance-dependent and rewired anisotropic networks (Figure A.4).

We further support this observation by computing the probability distribution for the expected number of edges between two random vertices in the anisotropic graph model. For this we assume that only the distance-dependent connection probability  $C(x)$  determines the occurrence of edges in vertex pairs in the anisotropic graph model. Then, using the probability distribution  $f(x)$  for the a random neuron pair to be at distance  $x$ , we calculate

$$\begin{aligned}\mathbf{P}(X = 0) &= \int_0^{\sqrt{2}} (1 - C(x))^2 f(x) dx, \\ \mathbf{P}(X = 1) &= \int_0^{\sqrt{2}} 2C(x)(1 - C(x))f(x) dx \quad \text{and} \\ \mathbf{P}(X = 2) &= \int_0^{\sqrt{2}} C(x)^2 f(x) dx.\end{aligned}$$

Inserting the distance-dependent connection probabilities  $C(x)$  in the anisotropic graph model as computed in Theorem 4.3 and the probability distribution  $f(x)$  from Theorem 3.15 we obtain

$$\begin{aligned}\mathbf{P}(X = 0) &= 0.791336 & 0.7907 \pm 0.0008 \\ \mathbf{P}(X = 1) &= 0.184151 & 0.1846 \pm 0.0007 \\ \mathbf{P}(X = 2) &= 0.024513 & 0.02462 \pm 0.00009,\end{aligned}$$



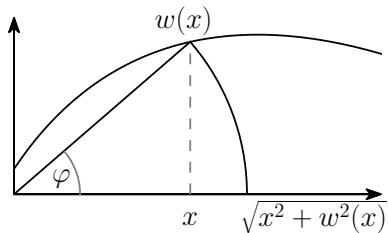
**Figure 5.7: Overrepresentation of reciprocal connections in anisotropic networks due to distance-dependent connectivity** Extracting the counts of unconnected, one-directionally and bidirectionally connected neuron pairs in the anisotropic sample graphs, overrepresentation of reciprocally connected pairs is identified as a feature of the network's distance dependency as opposed to anisotropy in connectivity. **A)** Showing the quotient of the counts for the three pair types, extracted from the set of sample graphs, with the number of expected pairs in Gilbert random graphs  $G(n, p)$ , where  $n = 1000$  and  $p = 0.116$  were matched to the sample graph parameters. While single connections appear less often than in Gilbert random graphs, reciprocal connections are significantly overrepresented. Errorbars SEM. **B)** Comparing appearance of connection pairs in the anisotropic sample graphs with their respective appearance in the rewired sample graphs, we find that eliminating anisotropy does not significantly change the counts for the connection types, indicating that anisotropy does not influence two neuron connection probabilities. Errorbars SEM. (c5f1462b)

perfectly matching the probabilities extracted from anisotropic sample graphs in the right column (error SEM, c5f1462b). Noting that distance-dependency alone is sufficient to accurately predict edge probabilities in neuron pairs in the anisotropic network model and combined with the observations in Figure 5.7, we conclude that varying degrees of anisotropy do not affect the occurrence of neuron pair motifs.

## 5.5 TUNING DISTANCE-DEPENDENCY

The discussion in the last section focused on the effect of anisotropy in connectivity on the occurrence of neuron pair motifs. Could distance-dependency itself, as imposed by the specific geometry, be a decisive factor in the distribution of edge counts in neuron pairs? Song et al. (2005), as well as Perin et al. (2011), report an overrepresentation of reciprocal connections independent from distance-dependent connectivity, opposing the observations made in the last section (Figure 5.7 A). Furthermore, the connectivity profile in the anisotropic graph model, as identified in Section 4.5, follows purely from abstract geometry rather than being motivated by connectivity found in cortical circuits. In an attempt to rectify this and to allow for a more differentiated examination of two neuron connections, in this section we step away from simplistic geometry and “tune” the anisotropic networks to display a distance-dependent connectivity as reported by Perin et al. by adjusting the width  $w(x)$  at any point  $x$  along the axon’s projection.

For this we introduce anisotropic networks tuned to reflect a given distance-dependent connection profile  $C(x)$ . We are facing the following problem: Given  $C(x) : [0, \sqrt{2}] \rightarrow [0, 1]$ , find  $w : [0, \sqrt{2}] \rightarrow [0, \infty)$  such that the probability to have a connection from  $v_1$  to  $v_2$  for arbitrary vertices  $v_1 \neq v_2$  in an anisotropic graph  $G(n, w)$  with distance  $d(v_1, v_2) = x$  is  $C(x)$ . The problem is in general highly complex when nothing can be assumed about  $C(x)$ . We find an approximate solution to the problem considering the following geometric relation:



**Figure 5.8:** Computing connection probability  $C(x)$  from non-constant  $w(x)$

From Figure 5.8 we have the relation

$$C\left(\sqrt{x^2 + w^2(x)}\right) = \frac{1}{\pi} \arctan \frac{w(x)}{x}. \quad (5.3)$$

In order to solve for  $w(x)$  we first consider a linear approximation, expanding

$$C\left(\sqrt{x^2 + w^2(x)}\right) \approx C(x) + \left(\sqrt{x^2 + w^2(x)} - x\right) C'(x).$$

The resulting transcendental equation

$$C(x) + \left( \sqrt{x^2 + w^2(x)} - x \right) C'(x) = \frac{1}{\pi} \arctan \frac{w(x)}{x}$$

is however still too complex in the context of this work. Instead we propose the approximation  $\sqrt{x^2 + w^2(x)} \approx x$ , which inserting into 5.3 yields

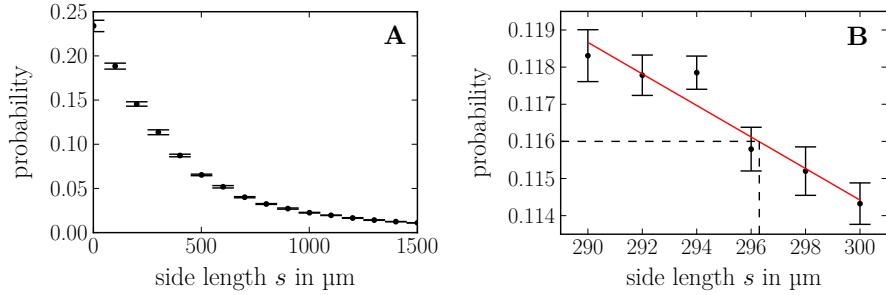
$$C(x) \approx \frac{1}{\pi} \arctan \frac{w(x)}{x}. \quad (5.4)$$

Under the assumption that  $C(x) < \frac{1}{2}$  for all  $x$  we obtain the identity

$$w(x) = x \tan(\pi C(x)), \quad (5.5)$$

being aware that it only holds as well as approximation 5.4 does.

Here we use relation 5.5 to generate anisotropic networks reflecting the distance-dependent connectivity profile as found by Perin et al. (2011). For this we finally need to adjust the before arbitrarily determined side length of the network's surface. Perin et al. mapped connectivity in layer 5 of the rat's somatosensory cortex up to a distance of 300  $\mu\text{m}$ . Using this reported distance connectivity to generate anisotropic networks via 5.5, the chosen side length  $s$  determines the networks overall connectivity (Figure 5.9 A). We determine  $s = 296 \mu\text{m}$  to match the overall connection probability of  $p = 0.116$  as used before and reported by Song et al. (Figure 5.9 B). The obtained value for  $s$  is consistent with the slice thickness of 300  $\mu\text{m}$  used in Perin et al.'s experiment.



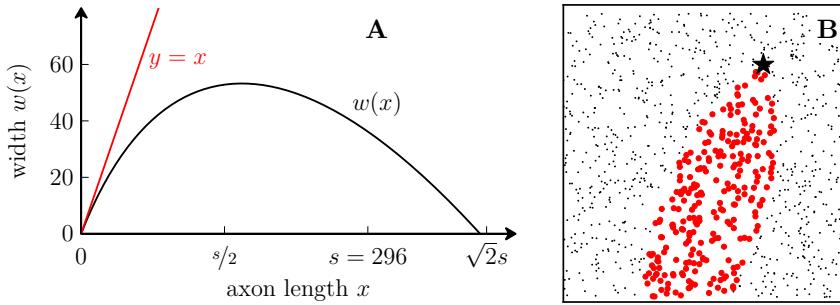
**Figure 5.9: Network side length adjusted to match overall connection probability** Side length of the network's surface determines the overall connection probability in the network when axon width function  $w(x)$  is fixed. **A)** Connection probability declines with rising side length **B)** Determining side length as  $s = 296 \mu\text{m}$  to match  $p = 0.116$  as reported by Song et al. (2005). (6154302f, ef0e785d)

Having determined the network's side length  $s$ , we're extending the quiver of generated sample networks for the numerical analysis once more by the “tuned anisotropic graphs”, in which the axon width

$w(x)$  was determined such that the networks reflect Perin's connectivity profile. Analyzing the obtained axon width function we note that  $x \gg w(x)$  holds for most  $x$ , justifying the approximation

$$\sqrt{x^2 + w^2(x)} \approx x$$

*a posteriori* (Figure 5.10). From the 25 generated networks overall connection probability is extracted as  $p = 0.1160 \pm 0.0006$  (SEM), as expected from the choice of  $s$  (f11dca65).



**Figure 5.10: Anisotropic network model with tuned axon width  $w(x)$**  **A)** Resulting axon width function  $w(x)$  from tuning to distance-dependent connection profile as reported by Perin et al. (2011), see also Figure 5.11. Note that  $x \gg w(x)$  for most  $x$ , supporting approximation 5.4. **B)** Showing for a single neuron (star) connected (red) and unconnected (gray) neurons in the tuned anisotropic network, revealing the characteristic axon shape. (d45c02e4, 8f0d65e4)

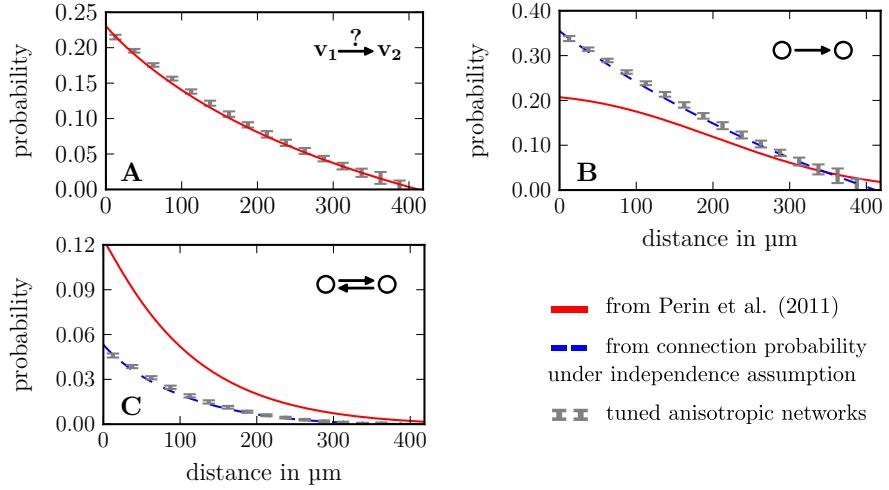
Overall distance-dependent connection probabilities in the tuned anisotropic graphs clearly match the profile of Perin et al. (Figure 5.11 A), presenting strongest the argument in support of the chosen approximation. Analyzing two neuron connections in the tuned networks, we affirm the findings of the last section. In their experiment, Perin et al. were able to show an overrepresentation of reciprocal connections at any inter-neuron distance (Figure 5.11 B-C). Rather than matching these profiles, we find that occurrences of one- and bidirectionally connected pairs in the anisotropic graphs align with probabilities obtained from the distance-dependent overall connection probability  $p(x)$  under the assumption of independence (cf. Equation 5.5),

$$\begin{aligned} \mathbf{P}_{X=1}(x) &= 2p(x)(1-p(x)) && \text{single connection}, \\ \mathbf{P}_{X=2}(x) &= p(x)^2 && \text{reciprocal connection}. \end{aligned}$$

Thus, in comparison with Perin et al.'s findings, we find that anisotropy in connectivity cannot account for the overrepresentation in reciprocal connections. While results in Section 5.4 still indicated such an overrepresentation due to distance-dependency, examining the occurrence of two neuron connections at any inter-neuron distance in anisotropic

revisiting two  
neuron  
connections

networks, tuned to a distance-dependent connection profile matching experimental findings from cortical circuits, imply complete unrelatedness of anisotropy and two-neuron connection distributions.



**Figure 5.11: Distance-independent overrepresentation of reciprocal connections** Comparison of occurrences of one- and bidirectionally connected neuron pairs in the tuned anisotropic networks (gray) with profiles found by Perin et al. (red), shows that overrepresentation of bidirectional pairs is distance-independent and not connected to anisotropy. **A)** Overall connection probability in the tuned anisotropic networks was successfully adjusted to reflect connection probability found by Perin et al. **B)-C)** Showing in blue the probabilities to obtain a neuron pair motif (single edge in B, two edges in C) calculated under independence assumption from the overall probability from A), we find that counts in the tuned anisotropic networks (gray) match the independence assumption and do *not* show the overrepresentation present in Perin et al.'s experiment. (875505b0)

## 5.6 MOTIFS

Non-random connectivity in local cortical circuits extends well beyond specific connection probability in neuron pairs: As one of their main results, [Song et al. \(2005\)](#) report a characteristic occurrence of motifs of three neurons, while [Perin et al. \(2011\)](#) find patterns in the number of connections appearing in clusters of up to 8 neurons. Motifs have been shown to have a significant influence on network dynamics, with patterns affecting dynamical correlations ([Pernice et al. 2011](#)) and network synchrony ([Zhao et al. 2011](#)).

Finding no correlation between two-neuron connection probabilities and anisotropy in connectivity prior, does anisotropy influence the occurrence of neuron motifs? In this section we analyze higher order connectivity in the different network types and observe a surprisingly strong impact of anisotropy on patterns of connected neurons, promoting the concept as an underlying principle for non-random network connectivity.

### *Three-neuron patterns*

We first investigate the occurrence of three-neuron patterns in anisotropic networks. Song et al. (2005) reported a characteristic, highly non-random motif distribution of pyramidal cells in the rat's visual cortex layer 5, a result later confirmed by Perin et al. (2011) in their experiment in the rat's somatosensory cortex layer 5. Repeating the experiment *in silico* for the different networks subject to this study, we find similar, characteristic motif distributions strongly influenced by anisotropy in connectivity.

There are  $13^2$  non-isomorphic 3-motifs in simple directed graphs. In reference to Song et al.'s result, the patterns are labeled 4 to 16,



Let  $X$  be a random variable that maps three random vertices  $v_1 \neq v_2 \neq v_3$  in a graph  $G$  to the  $n \in \{4, 5, \dots, 16\}$  labeling the isomorphism class of the full subgraph with vertex set  $\{v_1, v_2, v_3\}$  in  $G$  as above if the subgraph is connected, and let  $X$  map to  $n = 0$  otherwise. A first idea of how to compute the distribution of  $X$  is by inferring the probabilities of motif occurrence from the two-neuron connection probabilities from Section 5.4. In anisotropic networks we found that the probabilities of occurrence are

$$\begin{aligned} p_u &= 0.791336 && \text{for unconnected pairs,} \\ p_s &= 0.184151 && \text{for single connections and} \\ p_r &= 0.024513 && \text{for reciprocal connections.} \end{aligned}$$

From these we may, for example, calculate the probability of occurrence for motif 8,

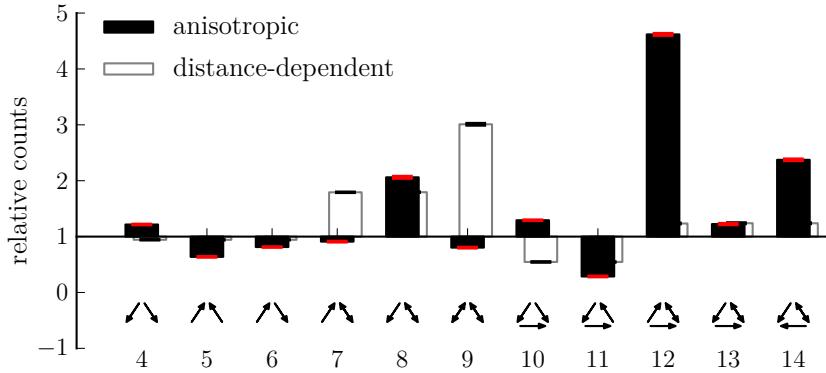
$$\mathbf{P}(X = 8) = 6 p_u p_s p_r,$$

---

<sup>2</sup> There are 16 non-isomorphic simple directed graphs with 3 nodes. Three of those graphs are unconnected (cf. Davis 1953, N. J. A. Sloane. The On-Line Encyclopedia of Integer Sequences, <http://oeis.org>. Sequence A000273).

where the factor 6 is determined by the number of different *labeled* graphs belonging to the isomorphism class. The distribution of  $X$  for the remaining motifs is given by

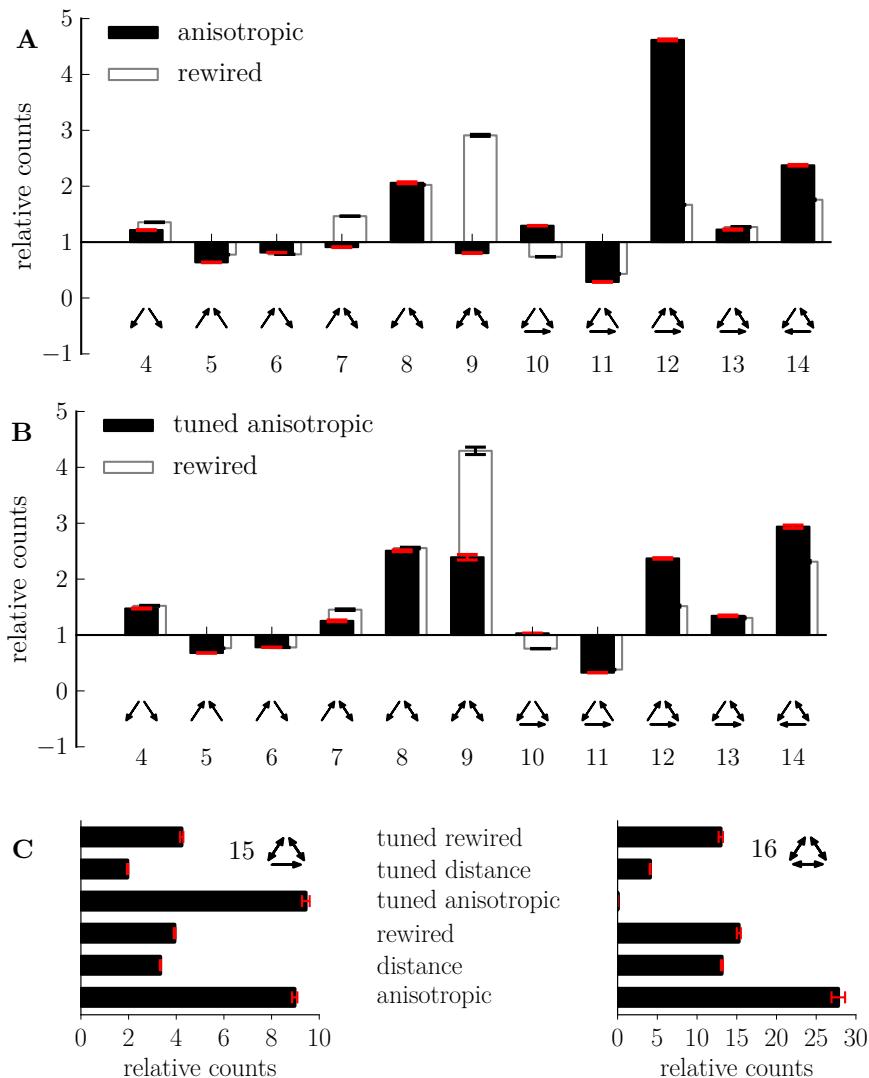
$$\begin{aligned}\mathbf{P}(X = 4) &= 3p_s^2 p_u & \mathbf{P}(X = 9) &= 3p_r^2 p_u & \mathbf{P}(X = 13) &= 6p_s^2 p_r \\ \mathbf{P}(X = 5) &= 3p_s^2 p_u & \mathbf{P}(X = 10) &= 6p_s^3 & \mathbf{P}(X = 14) &= 3p_s^2 p_r \\ \mathbf{P}(X = 6) &= 6p_s^2 p_u & \mathbf{P}(X = 11) &= 2p_s^3 & \mathbf{P}(X = 15) &= 6p_s p_r^2 \\ \mathbf{P}(X = 7) &= 6p_s p_u p_r & \mathbf{P}(X = 12) &= 3p_s^2 p_r & \mathbf{P}(X = 16) &= p_r^3.\end{aligned}$$



**Figure 5.12: Relative occurrence of three-neuron patterns** Extracting the counts of three-node motifs in anisotropic (filled bars) and distance-dependent networks (unfilled bars), the quotient of the obtained count with the number of occurrences expected from the two-neuron connection probabilities in the networks (cf. Section 5.4) shows the over- and underrepresentation of specific motifs in the network (red and black errorbars are SEM). In anisotropic networks pattern 12, for example, appears around five times more often than we would expect from the occurrence two-neuron connections. The relative counts for anisotropic networks resemble the findings of Song et al. (2005) and differ significantly from the counts in distance-dependent networks, implying that anisotropy has a strong influence on the relative occurrence of three-neuron patterns. (4839ce41)

distribution from  
neuron-pairs as  
reference

Does this distribution accurately reflect the occurrences of three-neuron motifs in anisotropic or even distance-dependent networks? Here we take the distribution determined from the two-neuron probabilities as a reference to analyze occurrences of three-neuron motifs in our sets of sample graphs. Counting the occurrences of patterns in we find that there are significant over- and underrepresentations in anisotropic as well as distance-dependent networks, relative to our expectation (Figure 5.12). We find, for example, that in anisotropic graphs pattern 12 occurs almost 5 times as often as we would have expected from the two-neuron probabilities, whereas the counts for pattern 11 only make up less than 30% of the occurrences expected.



**Figure 5.13: Three-neuron motif occurrence in different network types** **A)** Comparing counts in anisotropic sample graphs with their rewired counterparts. **B)** Three-neuron motifs occurrence in tuned anisotropic networks (cf. Section 5.5) with their rewired counterparts. For this two-neuron connection probabilities were extracted as in Section 5.4 and motif probabilities were calculated analogously to anisotropic networks. **C)** Relative counts for the high edge count motifs 15 and 16 for different network types, errorbars SEM. (4839ce41)

Comparing the relative counts for motifs in anisotropic graphs with those in comparable distance-dependent networks, we identify a strong influence of anisotropy in connectivity on three-neuron motif occurrence (Figure 5.12). In their experiments, Song et al. and Perin et al. find an overrepresentation of motifs 4, 10, 12 and 14. In anisotropic networks increased counts of motifs 4, 8, 10, 12, 13 and 14 were recorded. However, motifs 8 and 13 are overrepresented in distance-dependent

*anisotropy  
strongly affects  
3-motif occurrence*

networks as well, leaving the reported motifs 4, 10, 12 and 14 as motifs that are overrepresented due to anisotropy. To analyze this effect closer, we also compare three-neuron counts before and after rewiring in anisotropic networks ([Figure 5.13](#)).

Considering motif occurrences in anisotropic as well as tuned anisotropic networks, we once again confirm the overrepresentation of motifs 4, 10, 12 and 14. However, increased counts of pattern 4 are observed in the rewired networks as well, leading to the conclusion that increased occurrence in this motif is only implicitly affected by anisotropy. Motifs 10, 12 and 14 however show significant overrepresentation even over their rewired counterparts in anisotropic as well as in tuned anisotropic networks.

The overall motif distribution shows itself stable under changes in the distance-dependency with the notable exception of motif 9, that shows underrepresentation only in anisotropic but not in tuned anisotropic or any distance-dependent network type. Analyzing the occurrences of motifs 15 and 16 with a high edge counts ([Figure 5.13 C](#)) we find that anisotropy has strong influence on both motifs, with motif 15 being significantly overrepresented in anisotropic networks. Motif 16 shows a highly increased occurrence in anisotropic networks, however tuning causes the loss of this feature in the network connectivity.

#### *results summary*

Summarizing the above observations, we find that anisotropy in connectivity induces increased occurrence of motifs 10, 12, 14 and 15 in the network, reflecting experimental results in the rat's cortex. While over- and underrepresentation observed in local cortical circuits can be indirectly linked to anisotropy for some motifs (4, 9), it does not accurately reflect observed counts for other motifs (8) and shows instability under manipulation of distance-dependency in some patterns (9, 16).

#### *On motif occurrence in networks*

In obtaining and discussing the previous results, we closely followed the approach described by [Song et al. \(2005\)](#). Here, relying on analytical considerations, we submit these results to a critical analysis and identify potential caveats when dealing with motif distributions in distance-dependent networks.

For this consider for example the motifs 9, 15 and 16 as labeled above. In distance-dependent networks we find that each of the patterns is overrepresented, observing that the motifs occur about 3, 3 and 12 times more often than expected from the two-neuron connection probabili-

ties, respectively (Figure 5.12). Thus, in distance-dependent networks, triplets in which two of the pairs are reciprocally connected appear more often than expected, regardless of the connectivity in the third pair. This is surprising, as probabilities for the first two neuron pairs in a triplet should be independent and the probabilities of obtaining a certain connectivity should reflect the product of two-neuron probabilities. The overrepresentation found in these motifs may thus not necessarily be an inherent feature of the network connectivity but rather an artifact of triplet selection.

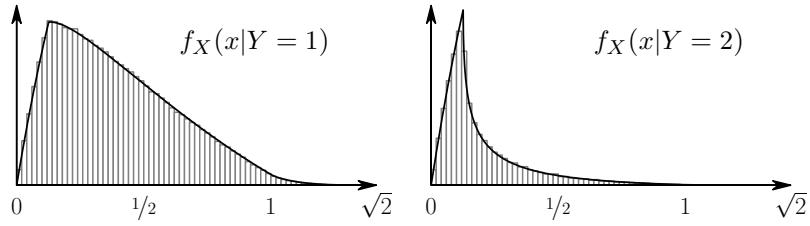
To analyze this further, we compute the relative probabilities of occurrence for motifs 9, 15 and 16. Formulated differently, given a triplet with two reciprocally connected pairs, what is probability of connection in the third pair? For this we first find the probability density function of the distance between a random neuron pair, given that we know its connectivity, meaning the existence of either 0, 1 or 2 edges between the neurons. Let  $X$  be the random variable mapping a random neuron pair of neurons to the distance between them and  $Y$  the random variable mapping to the number of edges between the pair. By the relation

$$f_X(x|Y = n) = \frac{f_{X,Y}(x,n)}{\mathbf{P}(Y = n)},$$

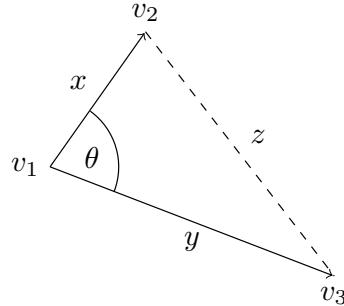
for  $n = 0, 1, 2$ , the probability density function  $f_X$  of the continuous variable  $X$  conditioned on the discrete variable  $Y$ , are then computed as the quotients

$$\begin{aligned} f_X(x|Y = 0) &= \frac{f(x)(1 - C(x))^2}{\int_0^{\sqrt{2}} f(x)(1 - C(x))^2 dx} \\ f_X(x|Y = 1) &= \frac{f(x)2C(x)(1 - C(x))}{\int_0^{\sqrt{2}} f(x)2C(x)(1 - C(x)) dx} \\ f_X(x|Y = 2) &= \frac{f(x)C(x)^2}{\int_0^{\sqrt{2}} f(x)C(x)^2 dx}, \end{aligned} \quad (5.6)$$

where  $f(x)$  is the probability density function of the distance between two random in the unit square the found in Theorem 3.15 and  $C(x)$  the distance-dependent connectivity from Theorem 4.3. Values for the denominator in Equation 5.6 have already been determined in Section 5.4. Evaluating the products in the numerator, we find that the expected probability density function is perfectly match by the distance distributions found in the simulated distance-dependent network model (38c11969):



Consider then a triplet with vertices  $v_1, v_2$  and  $v_3$ , the pairs  $(v_1, v_2)$  and  $(v_1, v_3)$  being reciprocally connected. With the density function  $f_X(x|Y = 2)$  we then have a distribution for the distances  $x = d(v_1, v_2)$  and  $y = d(v_1, v_3)$ . In the triangle spanned by the positions of the vertices, the distance  $z = d(v_2, v_3)$  is then determined by the angle  $\theta$  between  $x$  and  $y$ :



By the law of cosines, between  $x, y, z$  and  $\theta$  we have the relation

$$z = \sqrt{x^2 + y^2 - 2xy \cos \theta}. \quad (5.7)$$

From this we can, extensively using Lemma ??, calculate the probability density of  $z$  in the triplet  $(v_1, v_2, v_3)$  from the densities  $f_X(x|Y = 2)$  of  $x$  and  $y$ , similar to the proof of Theorem 3.15. Here, however, it suffices to find the expected value of  $z$ . The expected values for  $x$  and  $y$  are

$$r := \mathbf{E}[x] = \mathbf{E}[y] = \int_0^{\sqrt{2}} x f_X(x|Y = 2) dx.$$

Finding the expected value of  $z$  is then the well known problem of determining the length of a chord between two random points on the circle (cf. Weisstein 2014). For  $x, y = r$ , Equation 5.7 becomes

half-angle formula,  
Abramowitz and  
Stegun (1964)

$$z(\theta) = r\sqrt{2 - 2 \cos \theta} = 2r \left| \sin \frac{\theta}{2} \right|.$$

Using the symmetry of the problem, we then find the expected value

for  $z$  by integration,

$$\begin{aligned}\mathbf{E}[z] &= \int_0^{2\pi} \frac{1}{2\pi} z(\theta) d\theta \\ &= \int_0^{2\pi} \frac{r}{\pi} \left| \sin \frac{\theta}{2} \right| d\theta \\ &= \frac{2r}{\pi} \int_0^{\pi} \sin \frac{\theta}{2} d\theta = \frac{4r}{\pi}.\end{aligned}$$

With the expected distance  $z$  we can finally calculate the relative probabilities to obtain either motif 9, 15 or 16 from the distance-dependent connectivity profile  $C(x)$ . If  $Z$  is the random variable mapping given a triplet  $(v_1, v_2, v_3)$  with reciprocal connections between  $v_1$  and  $v_2$ , as well as  $v_1$  and  $v_3$  to the labels of the possible motifs 9, 15 or 16 describing the triplet, the distribution is (Mathematica A.2)

$$\begin{aligned}\mathbf{P}(Z = 9) &= (1 - C(\mathbf{E}[z]))^2 &= 0.697628 \\ \mathbf{P}(Z = 15) &= 2C(\mathbf{E}[z])(1 - C(\mathbf{E}[z])) &= 0.275227 \\ \mathbf{P}(Z = 16) &= C(\mathbf{E}[z])^2 &= 0.0271455.\end{aligned}\tag{5.8}$$

What relative occurrences would be expect from the two-neuron connection probabilities? For this we take the probabilities to find motifs 9, 15 and 16 under all motifs,

$$\mathbf{P}(X = 9) = 3p_r^2 p_u, \quad \mathbf{P}(X = 15) = 6p_s p_r^2, \quad \mathbf{P}(X = 16) = p_r^3,$$

and normalize the probabilities by their sum. Building the quotient of the calculated relative probabilities  $\mathbf{P}$  from (5.8) with the expected relative frequency  $\tilde{F}_2$ , inferred from two-neuron connectivity, we find the relative overrepresentation in the family of triplets with two reciprocal connections.

|    | $\tilde{F}_2$   | $\mathbf{P}/\tilde{F}_2$ | $k\mathbf{P}/\tilde{F}_2$ | $F_2$   |
|----|---|--------------------------|---------------------------|---------|
| 9  |  | 0.677625                 | 1.0295                    | 3.2408  |
| 15 |  | 0.315379                 | 0.8727                    | 2.7471  |
| 16 |  | 0.006997                 | 3.8797                    | 12.2126 |

(label: b5e4ed3e)

We note that amongst triplets with two reciprocal connections, motif 16 appears about four times as often as expected, while motifs 9 and 15 are not significantly over- or underrepresented. How do these numbers relate to the absolute overrepresentation  $F_2$  amongst all three-motifs? We find that triplets with two reciprocal connections appear  $k = 3.1478$

as often amongst all three-motifs as expected (b5e4ed3e); multiplying the quotient  $P/F_2$  with  $k$  then yields an absolute overrepresentation as found in the simulated distance-dependent networks.

#### *summary & discussion*

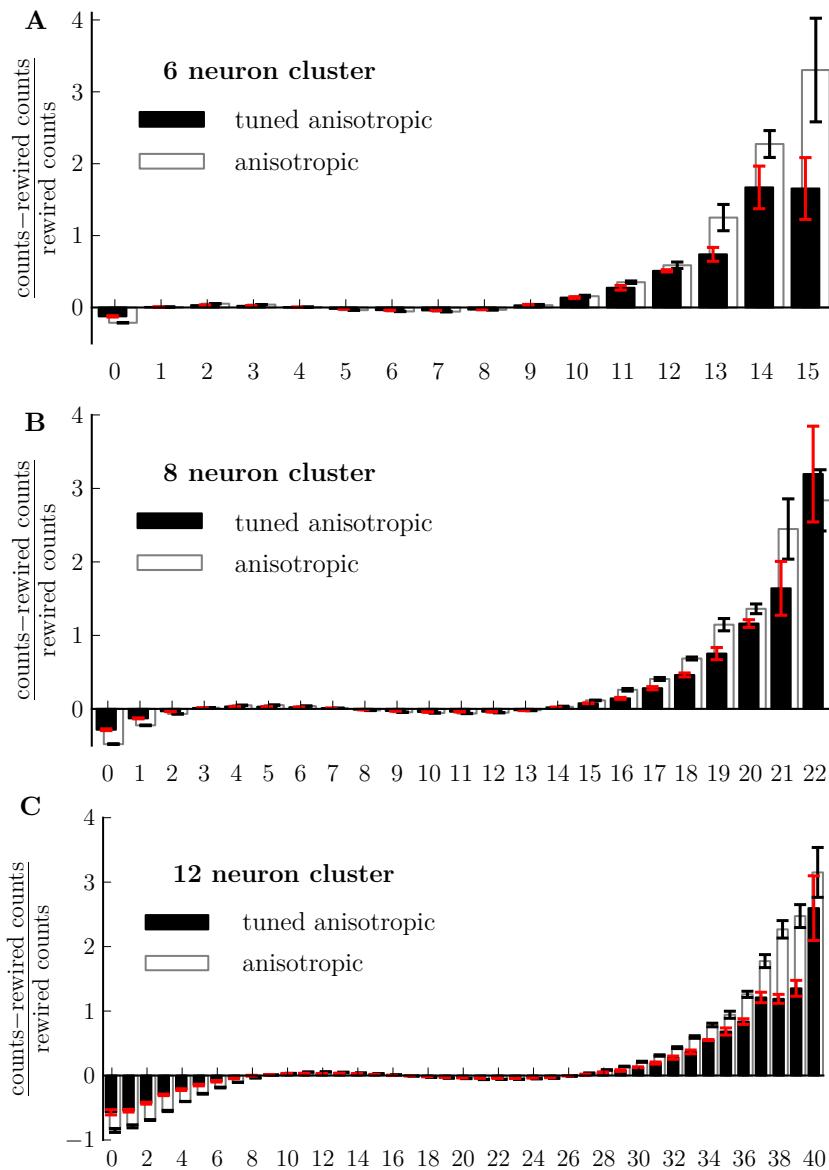
In distance-dependent networks, independence in the connectivity of the first two neuron pairs was expected. From our simulations, however, we clearly find that all triplets with at least two reciprocally occur more frequently than expected. This is due to boundary effects - neurons located close to the border or even a corner have drastically different distribution of distances to all other neurons as node located close to the center. Thus, rectifying for this bias, the “true” overrepresentation was calculated in (5.8), finding only motif 16 with an increased occurrence relative to the expectation.

Fortunately, results in anisotropic networks remain unaffected by this problem. As only statements relative towards rewired or distance-dependent networks were made, both containing this bias, overrepresentation due to anisotropy could still be accurately identified. Song et al.’s three-motif distribution also does not immediately show this problem. While, for example, motifs 9, 15 and 16 are all reported as overrepresented, it is well in the error margin of their experiment to have triplets with at least two reciprocal connections appearing overall as expected. However, the analysis shown here may well be important for further studies in brain connectivity, testing results against potential biases, or, gaining new insights on neuronal connectivity, finding without doubt that such overrepresentations do occur.

#### *Edge counts in neuron clusters*

In motifs consisting of 3 to 8 neurons, Perin et al. (2011) reported a striking statistic from their experiment with pyramidal cells in the rat’s somatosensory cortex, layer 5: Counting the number of edges appearing in a cluster of  $n$  neurons, they find that clusters with relatively high edge counts appear significantly more often than expected from the network’s distance-dependent connection probabilities alone. Do anisotropic networks exhibit a similar feature?

Recruiting the collection of sample graphs once again we analyze the occurrence of edge counts in clusters of  $n$  neurons in the different network types. In the main process, after randomly sampling  $n$  pairwise different vertices  $S_n$ , the motif  $H$  in  $G$  with vertex set  $V(H) = S_n$  is identified and its number of edges  $|E(H)|$  is recorded. Repeating this sufficiently often (order  $10^6$ ) we obtain edge counts for clusters of 6,8 and 12 neurons in the anisotropic- and tuned anisotropic networks



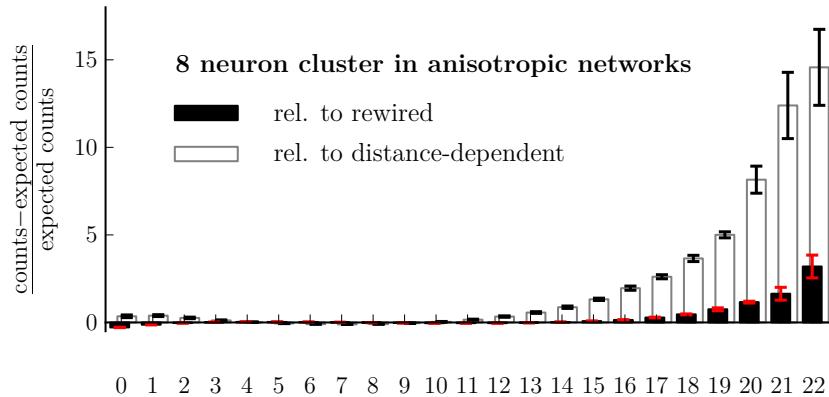
**Figure 5.14: High edge counts are overrepresented in networks with anisotropy** Normalized difference between number of edges in clusters of 6, 8 and 12 randomly selected neurons in anisotropic networks and rewired networks shows an overrepresentation of high edge counts in networks with anisotropy in connectivity. Errorbars SEM. (76cc6fa0, smtcite987992b0, 54329cf4)

as well as in their rewired counterparts. Then, showing the difference between counts in the anisotropic networks and counts in the rewired networks, normalized by the rewired counts, we identify an overrepresentation of high edge counts in the neuron similar to Perin et al. (Figure 5.14).

In anisotropic networks an overrepresentation of high edge counts is found consistently through clusters of 6, 8 and 12 neurons, with the

highest counts appearing up to four times as often as expected. Low edge counts occur less frequently than expected and an increasing number of connections shows an oscillation around the expectation. These findings are consistent with the report of Perin et al, which however indicates a higher maximal overrepresentation in clusters of 8 neurons.

Tuning the network's distance-dependency appears to not affect the overrepresentation as edge counts in anisotropic and tuned anisotropic essentially match (Figure 5.14). Anisotropy in connectivity therefore presents itself as an important factor in the occurrence of increased high edge counts in cortical networks, possibly fully reflecting the findings of Perin et al., as the overrepresentation effect is found to be even stronger when comparing edge counts to distance-dependent networks as opposed to rewired versions (Figure 5.15).



**Figure 5.15: Stronger overrepresentation of clusters with high edge counts with distance-dependent networks as reference**

Showing the occurrence of edge counts in clusters of 8 random neurons, we find that the overrepresentation of motifs with high edge counts is stronger when taking distance-dependent networks as a reference as opposed to rewired networks. (7c826e10)

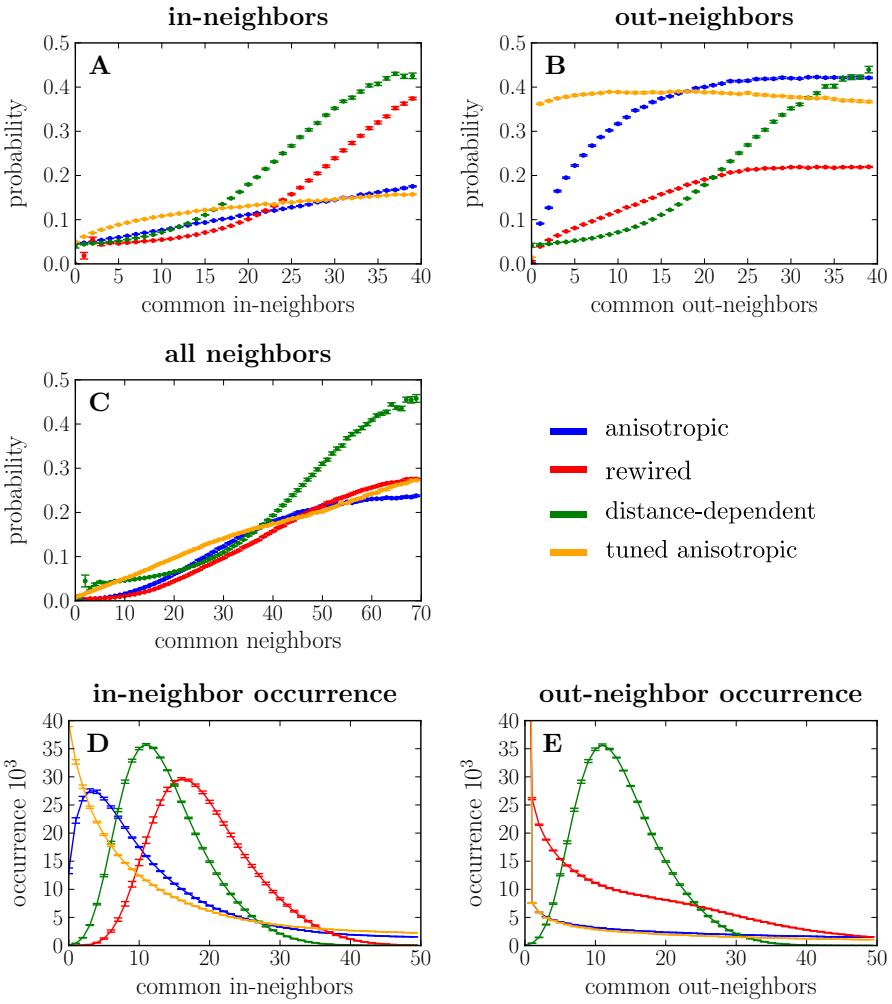
## 5.7 COMMON NEIGHBOR RULE

common neighbor rule as underlying principle?

In their study, Perin et al. follow their report of increased edge counts in neuron clusters with the observation of a “common neighbor rule”: Relying once again on their data in the rat's somatosensory cortex, Perin et al. find that not only do neuron pairs with a high number of common neighbor count appear significantly more often than expected, but also that such pairs display a higher probability of being connected. In fact, the relationship between pair connectivity and number of common neighbors appears to be linear. Perin et al. also report that this effect

is most pronounced when only considering common in-neighbors, that is other neurons that are projecting to both neurons in the pair.

Here we also investigate our networks for the existence of such a common neighbor relationship. Simultaneously recording connection probabilities and the number of common neighbors between pairs of neurons, we find inherent dependencies between the two quantities in all network types (Figure 5.16).



**Figure 5.16: Common neighbor rules in the different network types**

Showing the dependency of connection probability on the number of shared neighbors for random neuron pairs, characteristic curves for the network types (see legend) arises. High error margins (errorbars SEM) for small number of common neighbors in distance-dependent networks is induced by vanishingly low occurrence of a small common neighbor count. (5841710e)

Analyzing the results, we immediately note the sharp difference between in- and out-neighbors in their effect on connection probabili-

ties in anisotropic networks, as well as in rewired networks. Only in distance-dependent networks it appears that in- and out-neighbors can be considered equivalent in their influence on connection probabilities ([Figure 5.16 A-B](#)). Furthermore, while the distribution of the number of common neighbors is consistent in distance-dependent networks, the other network types display a characteristic distribution of common out-neighbors ([Figure 5.16 D-E](#)). While the latter observation clearly also relates to the differences in out-degree distributions found in [Section 5.2](#), finding differences between common inputs and outputs in neuron pairs is consistent with the observations of Perin et al., who report a significant difference in effect of the common neighbor rule. In trying to model the inherently asymmetric axonal-dendritic connections between pyramidal cells in cortical circuits with the anisotropic networks, finding such disparity is not only expected but gives the model further validity as an approach to obtain network connectivity going beyond the distance-dependent archetype, which here fails to produce diverging connectivity statistics for in- and outputs.

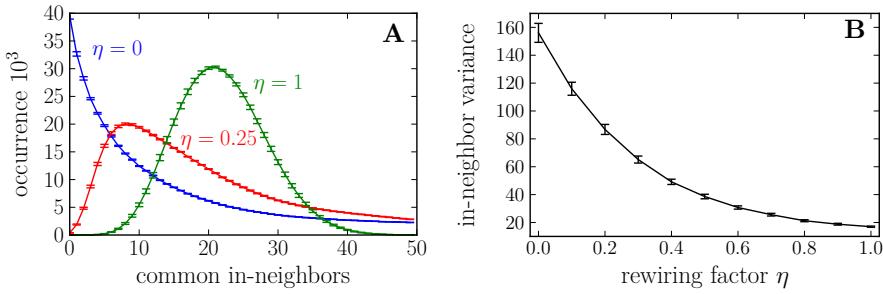
Both for in- and out-neighbors, we find characteristic curves describing their influence on connection probabilities. The in-neighbor profiles split into two categories: While networks with anisotropy in connectivity (blue, orange) display a constant increase, distance-dependent network types (red, green) show a sigmoidal shared input-connection probability curve ([Figure 5.16 A](#)). We thus find a strong influence of anisotropy on the shared input relationship, inducing a common neighbor rule characteristically different from isotropic, distance-dependent networks.

Does this anisotropy-induced rule reflect the findings in cortical networks? Perin et al. report a linear common neighbor relationship, finding a stronger effect when considering only in-neighbors. Imposing the common neighbor rule on *in silico* networks reflecting a distance-dependency as determined *in vivo*, Perin et al. were then able to reproduce the observed overrepresentation of high edge counts in neuron clusters, identifying the common neighbor effect as an underlying connection principle inducing increased high edge counts in clusters comparable to the profiles shown in [Figure 5.14](#). Showing not only the presence of such an edge count overrepresentation in anisotropic networks, but also finding that only networks featuring anisotropy display an approximately linear relationship between common inputs and connection probability, we identify anisotropy in connectivity as a candidate for an underlying connection principle motivated from neuronal morphology, to induce a common neighbor rule, that may be at the heart of many of the non-random connectivity statistics observed in local cortical networks.

*anisotropy induces  
characteristic  
common neighbor  
rule*

*anisotropy as  
underlying  
connection  
principle!*

Extending the analysis of shared inputs in the different network types, we further observe that anisotropy affects the number of common in-neighbors typically observed itself (Figure 5.16 D). We specifically find that increases anisotropy in connectivity induces an increased variance in the distribution of common inputs of a random neuron pair Figure 5.17). Such increased variance may provide an important advantage in the processing of information, allowing a heightened functional specificity in the network, where many neurons do not share many common inputs, enabling a high variety of functionality, and where few neuron pairs have a high number of shared inputs, strengthening their correlation and thus their capacity to relay related information.



**Figure 5.17: Anisotropy increases variance of common input distribution**

Recording common in-neighbor counts for random neuron pairs in tuned anisotropic networks and their rewired versions reveals increased variance in networks with a high degree of anisotropy. **A)** Common in-neighbor distribution for original tuned anisotropic networks ( $\eta = 0$ , blue) and rewired versions with 1/4 of all edges rewired ( $\eta = 0.25$ , red) and completely rewired ( $\eta = 1$ , green). (5841710e) **B)** Variance of the common in-neighbor distributions declines with increasing rewiring factor  $\eta$ ; highest variance is found in networks with the highest degree of anisotropy ( $\eta = 0$ ). Errorbars SEM. (ffcefef9b)

## 5.8 DISCUSSION

The structural analysis of anisotropic networks revealed a network topology characteristically different from comparable distance-dependent or even rewired networks. This however only, higher order connectivity motifs .

Vertex degree distributions were found to have increased variance and skew. However, as rewired and in the case of in-degrees even distance-dependent networks also show this feature, increased variance and skew are only indirectly caused by network anisotropy. Similarly, while anisotropic networks display a small-world property, the short path length and high clustering coefficient are not due to anisotropy but stem from

the network's distance-dependency. In fact, eliminating anisotropy in the network can lead to even shorter path lengths and higher clustering coefficients and thus strengthening the small-worldness of the network. Finally, anisotropy is not able to explain overrepresentation reciprocal connections reported repeatedly to be present in cortical networks.

In terms of higher order connectivity however, anisotropy yields highly interesting results. Three-neuron motifs occurrences are not only distinct, but resemble in many aspects the profiles found by [Song et al. \(2005\)](#).

In their study [Perin et al. \(2011\)](#) were able to relate increased edge-counts in neuron clusters to a *common neighbor rule* - a specific relationship between common in- or outputs in a neuron pair and the probability for the pair to be connected - identifying an underlying principle for the occurrence of non-random connectivity in local circuits. Perin et al. suggest that (fire together wire together). In this structural analysis we found non-random connectivity resembling the findings of Song et al./ and , purely from a g. Identifying a common neighbor rule as the inherent, that non-random connectivity statistics might arrise from morphological restrictions rather than plastic processes.

# A

## APPENDIX

---

### A.1 MATHEMATICA

```

In[1]:= f[d_] = Piecewise[{{1 / (s * (d)^(1/2)) - 1 / (s^2), 0 < d < s^2}, {0, d > s^2}}]

Out[1]= 
$$\begin{cases} -\frac{1}{s^2} + \frac{1}{\sqrt{d}} & 0 < d < s^2 \\ 0 & \text{True} \end{cases}$$


In[2]:= g[x_] := Convolve[f[d], f[d], d, x, Assumptions -> {d ∈ Reals, x ∈ Reals}]
Simplify[g[x], {s > 0, x ∈ Reals}]

Out[3]= 
$$\begin{cases} \frac{\pi s^2 - 4 s \sqrt{x+x}}{s^4} & x > 0 \&& s^2 \geq x \\ -\frac{2 s^2+x+\frac{4 s^3}{\sqrt{-s^2+x}}-\frac{4 s x}{\sqrt{-s^2+x}}-2 s^2 \operatorname{ArcTan}\left[\frac{s}{\sqrt{-s^2+x}}\right]+i s^2 \operatorname{Log}\left[s-i \sqrt{-s^2+x}\right]-i s^2 \operatorname{Log}\left[s+i \sqrt{-s^2+x}\right]}{s^4} & s^2 < x \&& 2 s^2 > x \\ 0 & \text{True} \end{cases}$$


In[4]:= h[x_] := g[x^2] * 2 * x

In[5]:= Simplify[h[x], {s > 0, x ∈ Reals, x > 0}]

Out[5]= 
$$2 x \begin{cases} \frac{\pi s^2-4 s x+x^2}{s^4} & s \geq x \\ -\frac{2 s^2+x^2+\frac{4 s^3}{\sqrt{-s^2+x^2}}-\frac{4 s x^2}{\sqrt{-s^2+x^2}}-2 s^2 \operatorname{ArcTan}\left[\frac{s}{\sqrt{-s^2+x^2}}\right]+2 s^2 \operatorname{ArcTan}\left[\frac{\sqrt{-s^2+x^2}}{s}\right]}{s^4} & s < x \&& \sqrt{2} > x \\ 0 & \text{True} \end{cases}$$


In[6]:= (*For s == 1, h becomes*)

In[7]:= Simplify[h[x], {s == 1, x ∈ Reals, x > 0}]

Out[7]= 
$$2 x \begin{cases} \pi + (-4+x) x & x \leq 1 \\ -2-x^2+4 \sqrt{-1+x^2}-2 \operatorname{ArcCot}\left[\frac{1}{\sqrt{-1+x^2}}\right]+2 \operatorname{ArcTan}\left[\frac{1}{\sqrt{-1+x^2}}\right] & 1 < x < \sqrt{2} \\ 0 & \text{True} \end{cases}$$


In[8]:= (*Expected Value*)
s := 1.
Integrate[x * h[x], {x, 0, Sqrt[2]}]

Out[9]= 0.521405

```

**Mathematica A.1:** Computation of probability density function for distance between two random points in square of side length  $s$  as supplement to proof of Theorem 3.15. Note that form of final result Out[7] differs from solution given in 3.15 for  $1 < x < \sqrt{2}$ . While proof of equivalence could not be achieved analytically, expressions given are numerically equivalent.

```

In[200]:= (*Expected distance between reciprocally connected pairs*)
r = NIntegrate[x * w[x] * c[x]^2 / 0.02451, {x, 0, Sqrt[2]}]
Out[200]= 0.20001

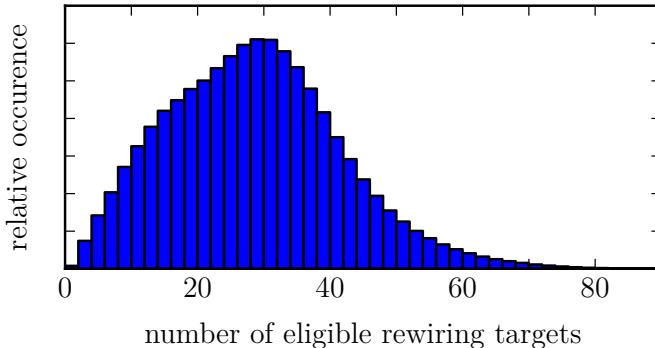
In[201]:= (*Probability of connection in third pair when other two are reciprocally connected*)
p = c[4 / Pi * r]
Out[201]= 0.164749

In[202]:= (*Probabilities for motif 9, 15 and 16 are then:*)
(1 - p)^2
2 * p * (1 - p)
p^2
Out[202]= 0.697643
Out[203]= 0.275214
Out[204]= 0.0271424

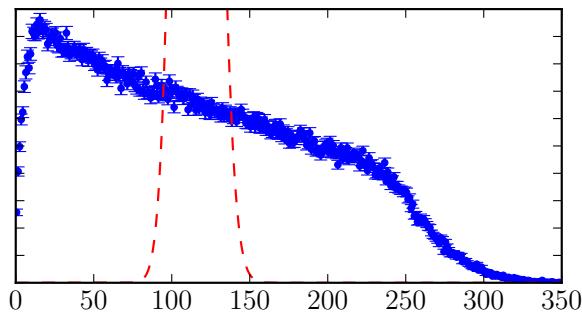
```

**Mathematica A.2:** Computation of three motifs for Section 5.6. Function  $c[x]$  is the distance-dependent probability distribution from Theorem and  $w[x]$  the probability density function for distance between two random points in a box (cf. Mathematica A.1, Moltchanov 2012).

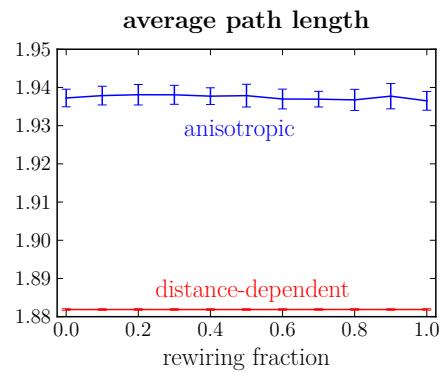
## A.2 SUPPLEMENTARY FIGURES



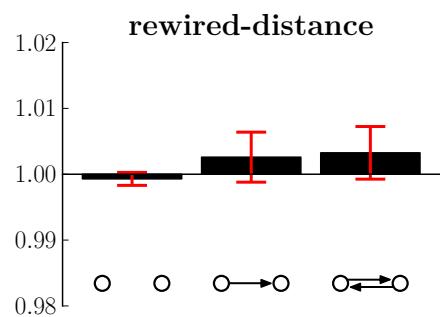
**Figure A.1:** (4afc2727)



**Figure A.2:** (c7ee86d7)



**Figure A.3:** Average path length for anisotropic and distance-dependent networks,  $N = 1000$ . (064f9b10)



**Figure A.4:** Probabilities for connections in neuron pairs are identical in distance-dependent and rewired anisotropic networks. (c5f1462b)



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## INDEX

---

- anisotropic geometric graph, 24  
random graph model, 24  
apical dendrite, 20
- basal dendrite, 20  
binomial random graph, 11
- common neighbor rule, 60
- directed graph, 5  
distance-dependent connection probability, 12  
connection profile, 12  
random graph model, 12
- edge set, 5
- geometric graph, 12  
Gilbert random graph model, 10
- in-degree, 7
- loop graph, 5
- motif, 6  
multigraph, 5
- neuro geometry, 18
- out-degree, 7
- partial rewiring, 32  
potential synapse, 18  
preferred direction, 34  
pseudograph, 5
- random graph, 10  
rewiring factor, 32
- SEM, 32  
simple graph, 5  
subgraph, 6  
full, 6
- tuned anisotropic networks, 48



## ERKLÄRUNG

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Hiermit versichere ich, dass ich diese Arbeit selbständig verfasst und nur die angegebenen Quellen und Hilfsmittel benutzt habe.

*Freiburg, 17. Juni 2014*

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Felix Hoffmann