ORIGINAL ARTICLE

The Space of Tree-Based Phylogenetic Networks

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Abstract

Phylogenetic networks are generalizations of phylogenetic trees that allow the representation of reticulation events such as horizontal gene transfer or hybridization, and can also represent uncertainty in inference. A subclass of these, *tree-based* phylogenetic networks, have been introduced to capture the extent to which reticulate evolution nevertheless broadly follows tree-like patterns. Several important operations that change a general phylogenetic network have been developed in recent years and are important for allowing algorithms to move around spaces of networks; a vital ingredient in finding an optimal network given some biological data. A key such operation is the nearest neighbour interchange, or NNI. While it is already known that the space of unrooted phylogenetic networks is connected under NNI, it has been unclear whether this also holds for the subspace of tree-based networks. In this paper, we show that the space of unrooted tree-based phylogenetic networks is indeed connected under the NNI operation. We do so by explicitly showing how to get from one such network to another one without losing tree-basedness along the way. Moreover, we introduce some new concepts, for instance "shoat networks", and derive some interesting aspects concerning tree-basedness. Last, we use our results to derive an upper bound on the size of the space of tree-based networks.

1 Introduction

Phylogenetic networks have become widely studied structures in the mathematics of evolution, because they capture a realistic range of evolutionary events beyond speciation, which itself is encapsulated elegantly via phylogenetic trees. In particular, phylogenetic networks are able to represent explicit events such as horizontal

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gene transfer and hybridization, as well as the presence of uncertainty. Phylogenetic networks, like trees, have also been widely studied using geometric approaches that consider the set of networks as a space in which one may move between the objects by operations that change a feature of the graph. Such operations, prominent examples of which use the *nearest neighbour interchange* (NNI), *subtree prune and regraft* (SPR), and *tree bisection and reconnection* (TBR), generalized from trees, are valuable in computational applications because it is often necessary to search the space of networks to find one that is optimal with respect to some specific criterion (Robinso[n](#page-16-0) [1971;](#page-16-0) Allen and Stee[l](#page-16-1) [2001;](#page-16-1) Huber et al[.](#page-16-2) [2016](#page-16-2)). Furthermore, such operations may be used to define proximity measures to tree-based networks, extending the notion of tree-basedness to the full class of phylogenetic networks (Fischer and Franci[s](#page-16-3) [2020\)](#page-16-3).

At the same time, motivated by various biological and mathematical considerations, various classes of phylogenetic network have been intensively studied. The focus of this paper is the class of *tree-based networks*, that are essentially "trees with additional edges" that pass between the tree edges (Francis and Stee[l](#page-16-4) [2015;](#page-16-4) Francis et al[.](#page-16-5) [2018](#page-16-5); Jetten and van Ierse[l](#page-16-6) [2016](#page-16-6); Hendrikse[n](#page-16-7) [2018\)](#page-16-7). While the space of phylogenetic networks of a given "tier" (defined below) is connected under NNI (Huber et al[.](#page-16-2) [2016\)](#page-16-2), we show in this paper that the space of tree-based unrooted phylogenetic networks is also connected under NNI moves. That is, it forms a connected subspace of the space of phylogenetic networks. Likewise, the diameter of the space of unrooted phylogenetic networks has been bounded in Francis et al[.](#page-16-8) [\(2018](#page-16-8)); here, we show that for a given tier, the diameter of the space of unrooted tree-based phylogenetic networks is of order $O(n^2)$, where *n* is the number of leaves.

We begin by setting out the necessary definitions, including a new class of (treebased) phylogenetic networks that we call *shoat* networks. In Sect. [3,](#page-4-0) we describe circumstances in which an NNI move on a tree-based network will produce another tree-based network (not all do) and show that it is possible to move between various classes of tree-based network using NNI moves, while staying tree-based. Our main result, the connectedness of the space of tree-based networks, is proved in Sect. [4;](#page-10-0) we finish with a discussion and further questions in Sect. [5.](#page-13-0)

While this manuscript was in preparation, a paper containing a statement similar to one of our main results appeared on the arXiv (Janssen and Klawitte[r](#page-16-9) [2019,](#page-16-9) Theorem 4.10). The result in that paper relates to unrooted networks that differ from ours in that they permit parallel edges, i.e. they are multigraphs. However, in the context of phylogenetic networks, it is more common (and arguably biologically more natural) to consider simple graphs (Pons et al[.](#page-16-10) [2018;](#page-16-10) Fischer et al[.](#page-16-11) [2018](#page-16-11); Francis et al[.](#page-16-5) [2018](#page-16-5); Stee[l](#page-16-12) [2016](#page-16-12); Gambette et al[.](#page-16-13) [2012](#page-16-13)). In contrast, our work takes a different approach to the question, and it is not clear whether the approach in Janssen and Klawitte[r](#page-16-9) [\(2019\)](#page-16-9) can be adapted to deal with networks as we define them here, i.e. without parallel edges.

2 Definitions and Background

A binary unrooted phylogenetic network on a set *X* (typically a set of species or taxa) is a connected simple graph whose vertices are degree 1 or 3, and whose degree 1 vertices

(leaves) are bijectively labelled by the elements of *X*. Some of our results extend to the non-binary case, in which non-leaf vertices may have degree greater than 3, and we will remark on that where it arises. Note that, as is common in the phylogenetics literature, non-binary networks are networks that are *not necessarily* binary, but may be; that is, the class of non-binary networks also (somewhat counter-intuitively) contains the binary networks. In the following, whenever there is no ambiguity, we use the term network to refer to an unrooted phylogenetic network. Note that the special case of an acyclic phylogenetic network is called a phylogenetic tree.

Throughout this manuscript, we will assume that $|X| > 2$, and that N is "proper". A proper network is one for which all components obtained by removing a cut edge or cut vertex contain at least one element of *X*.

In the following, we denote by $V^1(N)$ the set of degree 1 vertices in N, i.e. the set of leaves $V^1(N) = X$, and $\mathring{V} = V \setminus V^1$ denotes the set of inner vertices of N. If *k* is minimal such that the deletion of *k* edges of *N* would turn *N* into a tree (i.e. a connected acyclic graph), we say that *N* has *tier k*. Note that the tier does not depend on *N* being a phylogenetic network—in fact, the tier of a connected graph can be defined analogously (in which case it is often referred to as *cyclomatic number* or *cycle rank* in graph theory), and for technical reasons, we need this later on in this manuscript. We denote the set of tier-*k* phylogenetic networks on *n* leaves by *N*(*n*, *k*).

The *triangle operation*, introduced in Huber et al[.](#page-16-2) [\(2016](#page-16-2)), allows the replacement of a vertex with a triangle (a cycle of length three) to go up a tier and the reverse to go down a tier.

More formally, the ''blow-up'' triangle operation Δ^+ (that raises the tier of a network by 1) replaces an internal, degree 3 vertex v, and its incident edges $\{v, w_1\}$, $\{v, w_2\}$, $\{v, w_3\}$, by three vertices v_1, v_2, v_3 and six new edges: $\{v_1, w_1\}$, $\{v_2, w_2\}$, $\{v_3, w_3\}$, $\{v_1, v_2\}, \{v_2, v_3\}, \{v_3, v_1\}.$ Inversely, the "collapse" triangle operation Δ^- (lowering the tier by 1) replaces a triangle (a 3-cycle) and the edges outside the cycle but incident to its vertices by a single vertex and three incident edges. That is, given a 3-cycle of vertices $\{v_1, v_2, v_3\}$ and edges $\{v_1, v_2\}$, $\{v_2, v_3\}$, $\{v_3, v_1\}$, and three incident edges $\{v_1, w_1\}, \{v_2, w_2\}, \{v_3, w_3\}$, replace these by a single vertex v and three edges $\{v, w_1\}$, $\{v, w_2\}$, $\{v, w_3\}$. We denote these operations $\Delta^+(N, v)$ and $\Delta^-(N, \{v_1, v_2, v_3\})$, for v an internal vertex and $\{v_1, v_2, v_3\}$ a 3-cycle in the network N.

A*blob* of a network (or, more generally, of a graph) is a maximal connected subgraph that has no cut edge (if such a blob consists of only one vertex, it is called *trivial*). Note that in a binary phylogenetic network, blobs cannot contain any cut vertices (as all cut vertices in a binary network are incident to a cut edge (Fischer et al[.](#page-16-11) [2018,](#page-16-11) Lemma 8). A phylogenetic network is called *simple* if it contains at most one non-trivial blob.

A *support tree T* of a network *N* is a spanning tree of *N* satisfying $V^1(T) =$ $V^1(N) = X$, that is, whose leaf set coincides with the leaf set *X* of *N*. If *N* contains such a support tree *T* , it is called *tree-based*. Note that a support tree *T* of *N* is not necessarily a phylogenetic tree as it may contain degree-2 vertices. The space of tree-based networks in tier *k* on *n* leaves is denoted *TBN*(*n*, *k*).

Another concept that we need in the following, recently introduced in Fischer et al[.](#page-16-14) [\(2020\)](#page-16-14), is the *leaf cut graph* $LCUT(N)$ of a proper network $N \in N(n, k)$, with $|V(N)| > 3$, which is the graph *G* obtained from *N* by deleting all leaves and their incident edges. Note that this may result in some vertices of degree 2 and—e.g. if *N* is a tree—even new leaves not labelled by *X*, which we do *not* remove.

Our main results require the notions of *lineal* tree-based networks and *shoat* networks.

A tree-based network is called *lineal* if it has a support tree consisting of a single path *p* between two leaves, with paths from *p* to other leaves of length one. For instance, a tree-based network that is *pseudo-Hamiltonian* (in the sense of Francis et al[.](#page-16-8) [\(2018\)](#page-16-8): a network whose *LCUT* (*N*) graph (Fischer and Franci[s](#page-16-3) [2020](#page-16-3)) has a Hamiltonian cycle, cf. also Fischer et al[.](#page-16-14) [\(2020](#page-16-14))) will be lineal if there are two vertices that are connected to leaves and that are adjacent on the Hamiltonian cycle.

The second new family of graphs that we define are called *shoat networks*. [1](#page-3-0) These are a subclass of the pseudo-Hamiltonian, tree-based phylogenetic networks, defined below.

Definition 2.1 A binary tier $k \ge 1$ phylogenetic network $N \in \text{TBN}(n, k)$ is a *shoat network* if it has a Hamiltonian $\mathcal{LCUT}(N)$ graph with the properties that:

- (1) there are two leaves *x*, $y \in X$ whose adjacent interior vertices *a* and *b* are adjacent to each other, and such that there is a path *p* from *x* to *y* that visits all interior vertices of *N*;
- (2) the interior vertices of *N* are partitioned into sets V_L , V_M , V_R and $\{a, b\}$, with properties:
	- the path *p* consists (in order) of *x*, *a*, elements of V_L , elements of V_M , elements of *VR*, *b*, *y*;
	- $|V_L|, |V_R| = k 1, |V_M| = n 2;$
	- each vertex in V_L is adjacent to its neighbouring vertices on p and one vertex in V_R ;
	- each vertex in V_R is adjacent to its neighbouring vertices on p and one vertex in V_I ;
	- each vertex in V_M is adjacent its neighbouring vertices on p , and a leaf in $X \setminus \{x, y\}.$

Note, shoat networks can be defined in the non-binary setting by allowing $|V_L|, |V_R| \le$ $k - 1$ and $|V_M| \leq n - 2$, and allowing vertices in V_L and V_R to also be adjacent to one or more leaves. Note also that it is possible for the sets V_L , V_R or V_M to be empty.

An illustration of this definition is shown in Fig. [1.](#page-4-1)

The *nearest neighbour interchange (NNI)* is a replacement of a path in the network with an alternative path and was defined for unrooted phylogenetic networks in Huber et al[.](#page-16-2) [\(2016\)](#page-16-2):

Definition 2.2 (NNI) Let *N* be a phylogenetic network in which (a, b, c, d) is a path for which neither $\{a, c\}$ nor $\{b, d\}$ is an edge. The NNI operation on this path replaces it with the path (a, c, b, d) : the edges $\{a, b\}$ and $\{c, d\}$ are deleted, and edges $\{a, c\}$ and {*b*, *d*} are added.

¹ Because of their close resemblance to the juvenile boars that frequent the streets of northern Germany.

3 NNI Moves Among Tree-Based Networks

Any NNI move will change the underlying properties of a phylogenetic network, and in particular, can change whether a network is tree-based or not. Examples of NNI moves that make a tree-based network not tree-based are plentiful. For instance, there are exactly two level-5 non-tree-based binary networks (Fischer et al[.](#page-16-11) [2018\)](#page-16-11), and so there are many NNI moves from each of them that lead to tree-based networks. Consequently, each of those NNI moves in reverse makes a tree-based network nontree-based. However, as the main result of this paper (Theorem [4.1\)](#page-10-1) will show, in order to get from one tree-based network to another one, there must be at least one NNI path using only tree-based networks. In other words, tree-based networks are connected under NNI moves.

In this section, we prove some preliminary results about the effect of NNI moves on tree-based networks, which will all be needed for the proof of this main theorem, including showing that some different subclasses of tree-based networks are connected. We begin by proving that certain NNI moves preserve tree-basedness.

Lemma 3.1 *If N is a tree-based network with support tree T , then an NNI move on the path* (*a*, *b*, *c*, *d*) *will produce another tree-based network if either:*

- *(1)* the edges $\{a, b\}$, $\{b, c\}$, and $\{c, d\}$ are all in T;
- *(2)* {*b*, *c*} *is in T but* {*a*, *b*} *and* {*c*, *d*} *are not; or*
- (3) $\{b, c\}$, $\{c, d\}$, and $\{c, d'\}$ are in T for some other vertex d' , but $\{a, b\}$ is not.
- *Proof* (1) Suppose *T* is a support tree for the network *N*. If all edges in the path (a, b, c, d) are in T, shown as bold dashed and solid lines in Fig. [2,](#page-5-0) then the effect of the NNI produces a new support tree \tilde{T} that has edges $\{a, c\}$ and $\{b, d\}$ instead of $\{a, b\}$ and $\{c, d\}$, as shown in the right side of the figure. It is clear that \tilde{T} is still a support tree for *N* because it still covers all vertices in *N*; it has no additional leaves beyond those of *T* ; and it has not generated any cycles (there are still unique paths between all vertices a, b, c, d in \overline{T} , as there were in T).
- (2) If on the other hand neither $\{a, b\}$ nor $\{c, d\}$ are in the support tree T, but $\{b, c\}$ is, then there must be another edge incident to b that is in T (clearly, b is not a leaf in *N* because it is contained in the path (a, b, c, d) , i.e. it has degree > 1 in *N* and thus also cannot be a leaf in *T*). Likewise there is an edge in *T* incident to *c* other than $\{b, c\}$, for the same reasons. Without loss of generality suppose the edges $\{a', b\}$ and $\{c, d'\}$ are in *T*, shown in Fig. [2](#page-5-0) as solid lines. Then, the NNI on the path (a, b, c, d) has no effect at all on the spanning tree T, but simply

Fig. 2 An NNI move, such as this one on the path (*a*, *b*, *c*, *d*), can preserve tree-basedness in certain contexts (Lemma [3.1\)](#page-4-2)

shifts around the attachment edges that pass between vertices of the support tree *T* . Thus, *T* remains a support tree for *N*.

(3) Finally, if (b, c, d) is a path in *T* as described, and $\{c, d'\}$ another edge in *T*, then since *b* is not a leaf in *N*, there is another edge in *T* incident to it other than $\{a, b\}$ (because $\{a, b\}$ is not contained in T by assumption in this case, but *b* cannot be a leaf of *T* as it is not a leaf in *N*, either): say without loss of generality that it is $\{a', b\}$. Then, the effect of the NNI move on (a, b, c, d) is to shift the points that *a* and *d* connect to the path (a', b, c, d') in *T* from *b* to *c* and from *c* to *b*, respectively. These shifts do not cause any vertex to become uncovered by the tree, and cannot create cycles.

Thus, in all cases, the NNI move sending (a, b, c, d) to (a, c, b, d) changes the tree-based network *N* to another tree-based network.

Note, other scenarios may or may not preserve tree-basedness. For instance, if the path (a, b, c, d') in Fig. [2](#page-5-0) was in a support tree for *N* but the edges $\{a', b\}$ and $\{c, d\}$ were not, then the NNI move shown in the figure could turn *b* into a leaf of the spanning tree, which would then contain edges $\{a, c\}$, $\{c, d'\}$, and $\{b, c\}$. Whether it did in fact or not would depend on other features of the network outside the local frame of this NNI move.

In the remainder of this section, we prove several lemmas that show that a treebased network can be transformed to any shoat network by NNI moves that stay within *TBN*(*n*, *k*). It is important to note that all these NNI moves can of course be reverted i.e. the same arguments can be used to go back from said shoat network to the original tree-based network while staying within *TBN*(*n*, *k*).

Lemma 3.2 *A network in TBN*(*n*, *k*) *can be transformed to a lineal tree-based network using only NNI moves within TBN*(*n*, *k*)*.*

Proof Let *N* be a tree-based network. Of all support trees of *N*, choose a tree *T* which has the longest maximal length path *p*. Let this maximal length path consist of vertices v_1, \ldots, v_t in *T*. Suppose there is a vertex v_i on the path for which there is a path in *T* \ *p* to a leaf of length greater than 1 (note $i \neq 1$ or *t*, by maximality of *p*, as this implies that both v_1 and v_t must be leaves and thus have no neighbours outside the path). As long as N is non-lineal, such a vertex v_i must exist. We argue that there is an NNI move that extends the length of the maximal path and reduces the length of the path from v_i to a leaf, as follows.

Let w_1 and w_2 be the first two vertices along the path of length greater than 1 from v_i in *T*, with w_1 adjacent to v_i . Note that this necessarily implies that

Fig. 3 An NNI move that incorporates an edge from v_i into a maximal path (..., $v_{i-1}, v_i, v_{i+1}, \ldots$) within the support tree *T* of a tree-based network. Edges in *N* that are not in *T* are not shown

 w_1 cannot be adjacent to either v_{i-1} or v_{i+1} in *N*, because otherwise, the path v_1, \ldots, v_t could be extended by replacing (v_{i-1}, v_i, v_{i+1}) by $(v_{i-1}, w_1, v_i, v_{i+1})$ (or by $(v_{i-1}, v_i, w_1, v_{i+1})$, respectively). This would turn *T* into a support tree *T'* of *N* with a longer maximal path, contradicting the choice of *T* .

So as w_1 is not adjacent to either v_{i-1} or v_{i+1} in N, we can perform a legal NNI move on the path (v_{i-1}, v_i, w_1, w_2) , to create the path (v_{i-1}, w_1, v_i, w_2) , as shown in Fig. [3.](#page-6-0) Then, the modified tree *T* has a new maximal path $(v_1, \ldots, v_{i-1}, w_1, v_i, \ldots, v_t)$, and the length of the other path from v_i to a leaf has been reduced in length by 1. Furthermore, it is still tree-based, by Lemma [3.1\(](#page-4-2)i).

This process can be continued until the support tree *T* of *N* takes the form of a single long path with edges from it connecting directly to leaves of *N*—that is, until the resulting tree-based network is lineal.

Lemma 3.3 *A lineal phylogenetic network in TBN*(n , k), for $k \geq 1$, can be transformed *into a shoat network in TBN*(*n*, *k*) *by NNI moves within TBN*(*n*, *k*)*.*

Proof Let *N* be a lineal phylogenetic network in $TBN(n, k)$ with $k \geq 1$.

Choose a maximal length path *p* in *N*. Because *N* is lineal, this induces also a support tree *T* for *N*. Label the leaves at each end of the path *x* and *y*, and the vertices along *p* by $x = v_1, v_2, \ldots, v_m = y$. Because *N* is lineal, the edges connected to the other vertices along the path are either incident to leaves (if $n > 2$), or other vertices along the path.

Let

$$
L = \{v_i \mid \{v_i, v_j\} \in E(N), i + 1 < j\}
$$
\n
$$
R = \{v'_j \mid \{v'_i, v'_j\} \in E(N), i + 1 < j\},
$$

so that *L* (resp. *R*) is the set of vertices at the left (resp. right) end of edges between internal vertices that are not on the path *p* (we assume without loss of generality that "left" refers to vertices closer to x and "right" refers to vertices closer to y). These sets are disjoint, since each vertex has degree 3, and non-empty, since $k > 1$. There may also be vertices in *L* to the right of some (but not all) vertices in *R*; let

Fig. 4 A lineal network with $L = \{v_2, v_3, v_4, v_9\}$, $R = \{v_7, v_8, v_{10}, v_{12}\}$. This network has overlap set $\Theta = \{(v_9, v_7), (v_9, v_8)\}\$ and $||\Theta|| = 2 + 1 = 3$

Fig. 5 An NNI move reducing $||\Theta||$ by moving a vertex in *L* to the left of a vertex in *R*

$$
\Theta := \{(v_i, v_j) \in L \times R \mid i > j\}
$$

be the "overlap set": the pairs of vertices (v_i, v_j) for which v_i is an element of *L* but is nevertheless to the right of $v_j \in R$, as in Fig. [4.](#page-7-0) Define the *extent* of the overlap to be the non-negative integer:

$$
||\Theta|| := \sum_{(v_i, v_j) \in \Theta} (i - j).
$$

We will describe NNI moves that change the network to strictly decrease $||\Theta||$.

Suppose $\Theta \neq \emptyset$, and let *i* be minimal such that $(v_i, v_j) \in \Theta$. Because *i* is minimal, the vertex v_{i-1} to the left of v_i in *p* must either be (i) in *R* or (ii) connected to a leaf. We now consider these two cases.

(i) Suppose v_{i-1} ∈ *R*, and note that this means *i* > 4 since there must be a vertex to the left of v_{i-1} that is connected to it by an edge not in *p*, $v_1 = x$ is a leaf, and *N* has no parallel edges. We perform the NNI move

$$
v_{i-2}, v_{i-1}, v_i, v_{i+1} \rightarrow v_{i-2}, v_i, v_{i-1}, v_{i+1},
$$

shown in Fig. [5.](#page-7-1) The result is still tree-based by Lemma [3.1\(](#page-4-2)i).

Now consider the effect on Θ of this NNI move. Before the move, the elements of Θ that involve v_{i-1} or v_i are precisely the elements of the disjoint union:

$$
\{(v_i, v_{i-1})\} \cup \{(v_j, v_{i-1}) \mid \forall j > i, v_j \in L\} \cup \{(v_i, v_k) \mid \forall k < i-1, v_k \in R\}.
$$

After the NNI move, the elements of Θ that involve v_i _{−1} or v_i are precisely

$$
\{(v_j, v_i) \mid \forall j > i, v_j \in L\} \cup \{(v_{i-1}, v_k) \mid \forall k < i-1, v_k \in R\}.
$$

Note that elements that do not involve v_{i-1} or v_i are unaffected by the move, and that the cardinalities of the corresponding sets have not changed, but each corresponding

Fig. 6 An NNI move reducing $||\Theta||$ by moving a vertex in L to the left of a vertex attached to a leaf

term in $||\Theta||$ has decreased by 1. Therefore, the value of $||\Theta||$ has been decreased by the move by

$$
1 + |\{v_j \in L \mid j > i\}| + |\{v_k \in R \mid k < i - 1\}|.
$$

In particular, $||\Theta||$ has strictly decreased as a result of the move.

(ii) If v_{i-1} is not in *R* but connects to a leaf *z* (so that in particular, $n \ge 3$), there is at least one vertex v_j further left of v_{i-1} that is in *R*. Perform the NNI move shown in Fig. 6 :

$$
v_{i-2}, v_{i-1}, v_i, v_{i+1} \rightarrow v_{i-2}, v_i, v_{i-1}, v_{i+1}.
$$

As before, this still results in a tree-based network by Lemma [3.1\(](#page-4-2)i). After the operation and relabelling vertices along the path, *i* is no longer minimal such that (v_i, v_j) ∈ Θ , because (v_{i-1}, v_j) is now in Θ .

The effect on Θ of this move is that an ordered pair $(v_i, v_j) \in L \times R$ with $j < i$ is replaced in Θ by the pair (v_{i-1}, v_i) . As before, this has the effect of strictly reducing $||\Theta||$, as required.

Since such moves are always possible while $||\Theta|| > 0$, and they each strictly reduce $||\Theta||$, it follows that successive application of these moves can reduce $||\Theta||$ to zero. At that point, we must have $\Theta = \emptyset$.

Having reduced the size of the overlap set $||\Theta||$ to zero, we need to move any leaves (apart from *x* and *y*) to the middle.

Suppose there are leaves attached to vertices v_j in p , for $j \neq 1, m$, other than $x = v_1$ and $y = v_m$. We now want to bring those leaves to the middle of the path, between the vertices in *L* and those in *R*.

Suppose a leaf $z \neq x$ is attached to a vertex v_{i-1} which has a vertex $v_i \in L$, to its right. Perform the NNI move

$$
v_{j-2}, v_{j-1}, v_j, v_{j+1} \rightarrow v_{j-2}, v_j, v_{j-1}, v_{j+1},
$$

which moves the attachment of the leaf *z* one position towards the right along the path *p* (the same move shown in Fig. [6\)](#page-8-0). The new network is still tree-based, by Lemma [3.1\(](#page-4-2)i). As long as there is a leaf attached to the left of a vertex from *L*, this can be repeated until there are none left. Similarly, the reverse can be performed for leaves that are attached to a vertex that has a vertex in *R* to their left, until there are no such vertices.

The result is that the leaves (apart from x and y) are attached to vertices in p with all vertices in *L* to their left, and all vertices in *R* to their right.

It is now elementary to make the network shoat. Take the leftmost vertex in *L*, which must be v_2 . It is connected by an edge $\{v_2, v_k\}$ to a vertex $v_k \in R$. If $k \neq m-1$ (so that v_k is not the rightmost vertex on the path before the leaf *y*), the vertex v_{k+1} is also in *R* and is connected to an edge $\{v_{\ell}, v_{k+1}\}$ for some $v_{\ell} \in L$. Perform the NNI

$$
v_2, v_k, v_{k+1}, v_{\ell} \rightarrow v_2, v_{k+1}, v_k, v_{\ell},
$$

so that v_2 is now connected by an edge to a vertex one further to the right. This is again tree-based by Lemma [3.1\(](#page-4-2)ii). This can be continued until there is an edge $\{v_2, v_{m-1}\}\$, at which point the network is shoat at which point the network is shoat.

Lemma 3.4 *Shoat networks are a connected subspace of TBN*(*n*, *k*)*. That is, we can use NNI moves between tree-based networks to move from one shoat network S*¹ *on leaf set X with tier k to another such shoat network S*2*.*

Proof Note that both *S*¹ and *S*² have two distinguished leaves (corresponding to *x* and *y* in Fig. [1\)](#page-4-1), and let x_1 and y_1 be the distinguished leaves in S_1 , with x_2 and y_2 the two distinguished leaves in S_2 . We will show how to transform S_1 into S_2 .

If the pairs $\{x_1, y_1\}$ and $\{x_2, y_2\}$ coincide, we do not have to bring the distinguished leaves into their positions: they already are. However, if, say, x_1 is not a distinguished leaf in *S*2, we have to use NNI moves to bring it to its correct position according to S_2 (left of the "left" vertices), and we do this along the maximal length path from x_1 to y_1 in S_1 . Note that all these NNI moves are necessarily legal because x_1 is moved left past all the "left" vertices in *S*1, meaning at no point the main rule for NNI (that the first and third or second and fourth vertices are adjacent) is violated. They also preserve tree-basedness because they are operations of the first type in Lemma [3.1.](#page-4-2)

Once we are done with bringing the first distinguished leaf of S_1 which is not a distinguished leaf of S_2 into position, we bring the corresponding distinguished leaf of *S*2, that is misplaced in *S*1, to the former position of *x*¹ by NNI moves. Again, these NNI moves are necessarily legal and preserve tree-basedness by similar arguments to the above.

We repeat the same procedure with y_1 if applicable.

When both distinguished leaves are in the correct position (as induced by S_2), we use NNI moves to arrange the other $n - 2$ leaves correctly. Again note that all these NNI moves are necessarily legal and preserve tree-basedness by Lemma [3.1.](#page-4-2)

Last, we re-arrange the at most $k - 1$ "left" vertices between x_2 and the nondistinguished leaves of S_2 as well as the at most $k-1$ "right" vertices between y_2 and the non-distinguished leaves of *S*² by NNI moves (it is sufficient to rearrange just either the left or the right sides). Again note that all these NNI moves are necessarily legal and preserve tree-basedness by Lemma [3.1.](#page-4-2)

4 Connectedness of *TBN(n, k)*

We are now in a position to prove the main result of this manuscript, Theorem [4.1.](#page-10-1) Its proof is informed by the idea of the proof of Francis et al[.](#page-16-8) [\(2018\)](#page-16-8), Theorem 3, which describes a particular path through the space of all phylogenetic networks. However, in contrast to Francis et al[.](#page-16-8) [\(2018\)](#page-16-8), Theorem 3, we are considering only tree-based networks, and so the results of the previous section laying out how NNI moves can preserve tree-basedness are essential: our path through the space is essentially laid out by the lemmas in Sect. [3.](#page-4-0) In particular, we show that one can move from any tree-based network to any other tree-based network of the same tier and on the same taxon set by NNI moves only between tree-based networks.

Theorem 4.1 *The space of tree-based unrooted phylogenetic networks on leaf set X of tier k is connected under NNI moves.*

Proof The case $k = 0$, i[n](#page-16-0) which networks are trees, has been established in Robinson [\(1971\)](#page-16-0), so we may assume $k \ge 1$ (in fact, since all such networks are tree-based for $k \leq 4$ (Francis et al[.](#page-16-5) [2018\)](#page-16-5), the result is also immediate for $k \leq 4$ because the space of networks is connected (Huber et al[.](#page-16-2) [2016\)](#page-16-2)).

Lemma [3.2](#page-5-1) shows that a tree-based network can be transformed into a lineal network while remaining inside $TBN(n, k)$. Lemma [3.3](#page-6-1) shows that a lineal tree-based network can be transformed into a shoat network, and Lemma [3.4](#page-9-0) shows that any two shoat networks are connected by NNI moves within $TBN(n, k)$ (both these results are restricted to $k \ge 1$ because shoat networks are only defined for $k \ge 1$). Noting that all NNI moves can always be reversed, this proves the theorem (because you can go from any tree-based network N to any other tree-based network N' by first modifying *N* into a shoat network, possibly modifying this shoat network to give another shoat network, and then continuing from this second shoat network to *N'*—all the time only using NNI moves through tree-based networks).

In the following, we use the fact that the number of vertices in an unrooted phylogenetic network of tier *k* is exactly² 2($n + k - 1$).

Corollary 4.2 *The diameter of the space of tree-based networks with n leaves and tier k under NNI is at most*

$$
f(n,k) = {k-1 \choose 2} + {n-2 \choose 2} + 4nk + 4k^2 + 4n - 4k - 17.
$$

That is, the diameter is in $O((n + k)^2)$ *, and for fixed tier k, it is in* $O(n^2)$ *.*

Proof Transforming one tree-based network to another takes the steps described in Theorem [4.1:](#page-10-1)

(1) make the networks lineal;

² Note that a binary phylogenetic tree with *ⁿ* leaves has precisely 2*ⁿ* [−] 2 vertices (Semple and Stee[l](#page-16-15) [2003](#page-16-15)). Any of the extra *k* edges added to such a tree will induce two new vertices. In total, this is $2n - 2 + 2k =$ $2(n + k - 1)$ vertices.

- (2) make the lineal networks shoat graphs; and
- (3) transform between shoat graphs.

Note that as the NNI moves are reversible, the total pathlength between two networks will be at most twice the sum of the maxima of the first two steps, plus the maximum of the third.

The algorithm for making a network lineal, given in Lemma [3.2,](#page-5-1) takes a support tree with a maximal path, and incorporates other edges of the tree into that path while remaining tree-based. Each edge is incorporated in a single NNI move, so the number of moves required to make the network lineal is at most the number of edges in a spanning tree minus the number of leaves (because leaf edges are not incorporated into the path), minus the length of the initial maximal path.

We need to bound two numbers: the number of edges in a support tree, and the length of a maximal path. The number of edges in a support tree is the number of vertices minus one, and for a binary network the number of vertices² is $2(n + k - 1)$. Therefore, the number of edges in the support tree is at most $2(n + k - 1) - 1$. The number of edges in a maximal path in a support tree would be 2 if the network was a star tree, so we can use this as a very weak lower bound on this.

That is, there are at most $(2(n + k - 1) - 1) - n - 2 = n + 2k - 5$ moves required.

network \rightarrow lineal: $n + 2k - 5$.

To transform a lineal network into a shoat graph using the procedure described in Lemma [3.3,](#page-6-1) we need to shift the "left" vertices to the left, and the vertices attached to leaves, to the middle. Beginning with the "left", each must pass by at most $n - 2$ vertices attached to leaves, and $2k - 2$ other vertices attached to edges (since there must be at least one "right" vertex to its right, and one other, since it cannot be adjacent to its partner. This gives a total of at most $k(n-2+2k-2) = k(n+2k-4)$ moves. To move the $n - 2$ leaf-adjacent vertices to the middle, they each have to pass at most *k* left or *k* right vertices, for a total of $k(n-2)$ moves. Thus, there is a total of $k(2n + 2k - 6)$ moves to make the network shoat.

$$
linear \rightarrow \text{shot: } k(2n + 2k - 6).
$$

Finally, a shoat network N_1 with distinguished leaves x_1 and y_1 can be arranged into another (along the lines of Theorem [4.1\)](#page-10-1), in $\binom{k-1}{2} + \binom{n-2}{2} + 2n + 4k - 7$ NNI moves:

• If one or both distinguished leaves are distinguished in both shoat networks, then the symmetry of the graph allows us to flip N_1 in the vertical axis so that these are aligned. Thus, either one or both distinguished leaves will need to be moved into position from the set of leaves in the middle (between the vertices in *L* and *R*). This will take $(n - 1) + (k - 1)$ for the first (moving past possibly $n - 1$ leaves and $k - 1$ vertices in say *L*), and $(n - 2) + (k - 1)$ for the second, since there is one fewer leaf available to move past. The leaves that were previously distinguished then move back to the middle in $k - 1$ moves each, for a total of $(n-1) + (k-1) + (n-2) + (k-1) + 2(k-1) = 2n + 4k - 7$ moves;

- at most $\binom{n-2}{2}$ moves to arrange the leaves in the leaf zone correctly (the diameter of the symmetric group on $n - 2$ objects, under the operation of adjacent transpositions); and
- at most $\binom{k-1}{2}$ moves to arrange the $k-1$ endpoints on the right or left to agree with N_2 .

$$
\text{shoat} \rightarrow \text{shoat:} \binom{k-1}{2} + \binom{n-2}{2} + 2n + 4k - 7.
$$

Therefore, the total maximal distance between any two networks is at most

$$
\binom{k-1}{2} + \binom{n-2}{2} + (2n+4k-7) + 2((n+2k-5) + k(2n+2k-6))
$$

=
$$
\binom{k-1}{2} + \binom{n-2}{2} + 4nk + 4k^2 + 4n - 4k - 17,
$$

as required. This completes the proof.

We are now able to show the connectivity of the entire space of tree-based networks on a fixed set of leaves X, using the triangle operations Δ^+ (replacing an inner degree-3 vertex by a triangle) and Δ^- (collapsing a triangle to a single vertex) as defined in Sect. [2.](#page-1-0)

First, it is straightforward to show that either triangle operation on a tree-based network is still tree-based, as follows:

Lemma 4.3 *If N is a tree-based network in TBN*(*n*, *k*)*, then*

(1) for
$$
v \in V(N) \setminus X
$$
, $\Delta^+(N, v) \in TBN(n, k + 1)$; and
(2) for $\{v_1, v_2, v_3\}$ a 3-cycle in N, $\Delta^-(N, \{v_1, v_2, v_3\}) \in TBN(n, k - 1)$.

Proof If *N* is tree-based and $v \in V(N)$ has degree 3, then any support tree for *N* includes v and at least two of the three edges incident to it, $\{v, w_1\}$, $\{v, w_2\}$, $\{v, w_3\}$, because the support tree is a spanning tree of *N* whose only leaves are elements of *X*, and hence not v.

We first show (1). Without loss of generality, suppose there is a support tree *T* for *N* containing $\{v, w_1\}$ and $\{v, w_2\}$. Then, replacing the edges $\{v, w_1\}$ and $\{v, w_2\}$ in *T* by the edges $\{w_1, v_1\}, \{v_1, v_3\}, \{v_3, v_2\}, \{v_2, w_2\}$ creates a tree (no cycles are generated) that is a support tree for $\Delta^+(N,\,v)$ (it is connected and has no additional leaves), which is also now in tier $k + 1$, so in summary, we have $\Delta^+(N, v) \in TBN(n, k + 1)$.

Likewise, for (2), if *N* is tree-based and has a 3-cycle $\{v_1, v_2, v_3\}$, with additional edges $\{v_1, w_1\}, \{v_2, w_2\}, \{v_3, w_3\}$, then it has a support tree that includes all of $\{v_1, v_2, v_3\}$, and at least two of the edges $\{v_1, w_1\}$, $\{v_2, w_2\}$, $\{v_3, w_3\}$. Replacing these by a single vertex in both the network and the support tree does not disconnect the tree, create cycles, or make any vertex uncovered by the collapsed tree.

Theorem 4.4 *The space of binary tree-based networks on X is connected under NNI moves and triangle moves.*

$$
\underline{\textcircled{\tiny 2}}
$$
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Proof This is immediate from Theorem [4.1](#page-10-1) and Lemma [4.3](#page-12-0) as long as we can show that it is always possible to move to a tree-based network with a 3-cycle in it, in order to move down a tier.

If the network does not have a triangle already, we can produce one using a series of NNI moves that stays within *TBN*(*n*, *k*) as follows.

Let *T* be a support tree for *N*, and find a cycle in *N* that is made up of edges in *T* except for one. Such a cycle always exists, just by considering an edge $\{v_1, v_2\}$ in $N \setminus T$ and a path within *T* between its endpoints. Denote the vertices of this path (v_3,\ldots,v_i) , with $i > 3$, so that the cycle in *N* is $(v_1, v_2,\ldots,v_i, v_1)$. Since all vertices in the cycle have degree three in *N*, there is an edge $\{v_3, v'_3\}$ that is not in the cycle.

Perform an NNI on the path (v_1, v_2, v_3, v'_3) , which is legal because the edges $\{v_1, v_3\}$ and $\{v_2, v'_3\}$ are not in *N* (if they were *N* would have a triangle). The result is still a tree-based network because if $\{v_3, v_3'\}$ is not in *T*, then Lemma [3.1\(](#page-4-2)ii) applies, whereas if $\{v_3, v'_3\} \in T$, then Lemma [3.1\(](#page-4-2)iii) applies. This network has a cycle $(v_1, v_3, v_4, \ldots, v_i, v_1)$, which is of strictly shorter length than the original cycle (because it does not contain v_2 anymore).

In this way, we can use NNI to shorten any cycle in *N* of length greater than 3, and so eventually obtain a triangle. Deleting that triangle via a triangle move remains tree-based, which completes the proof.

We end by noting that while the diameter of the full space of tree-based networks on *n* leaves is infinite, since the tier is unbounded, the distance between a pair of networks in different tiers is also $O(n^2)$.

Corollary 4.5 *The distance between networks* $N_1 \in TBN(n, k_1)$ *and* $N_2 \in TBN(n, k_2)$ *, for fixed* k_1 *and* k_2 *such that* $k_1 \leq k_2$ *, is in* $\mathcal{O}(n^2)$ *.*

Proof Given $N_1 \in \text{TBN}(n, k_1)$, it can be transformed into a network in $\text{TBN}(n, k_2)$ by successively inserting triangles at vertices $k_2 - k_1$ times. Once in tier k_2 , it is at most $f(n, k_2)$ moves from N_2 , and so the distance between N_1 and N_2 is at most *f* (*n*, *k*₂) + (*k*₂ − *k*₁). Since *f* (*n*, *k*₂) is in $\mathcal{O}(n^2)$ for fixed *k*₂, the result follows. \Box

5 Discussion

In this paper, we have shown that the space of unrooted binary tree-based networks within a given tier is connected under NNI moves, and that if the triangle moves are included the whole space of unrooted binary tree-based networks is connected (Theorems [4.1](#page-10-1) and [4.4\)](#page-12-1). This connectivity gives rise to several possible applications and interesting additional questions.

The connectivity of the space means that the class of tree-based networks can be searched by use of NNI moves. This has potential benefits for the sampling of the space using MCMC approaches, as it already does for phylogenetic trees (Lakner et al[.](#page-16-16) [2008;](#page-16-16) Höhna and Drummon[d](#page-16-17) [2011](#page-16-17); Whidden and Matsen I[V](#page-16-18) [2015\)](#page-16-18) and in future most likely also will for general phylogenetic networks (Huson et al[.](#page-16-19) [2010\)](#page-16-19). To further develop this application, it would be interesting to obtain estimates of the size of the tree-based neighbourhood of a given network in *TBN*(*n*, *k*).

Fig. 7 The only two binary, non-tree-based phylogenetic networks of tier 5 (Fischer et al[.](#page-16-11) [2018,](#page-16-11) Fig. 7). These are both rank −1, that is, for both of them it takes only one NNI move to make them tree-based. However, no single NNI move can convert one of these networks into the other one

The alert reader may note that the upper bound on the diameter of the space of all tier-*k* unrooted phylogenetic networks *N*(*n*, *k*) obtained in Francis et al[.](#page-16-8) [\(2018\)](#page-16-8) is of a similar order to the upper bound on the diameter for the space of tier-*k* tree-based networks $TBN(n, k)$ obtained here. While the density of $TBN(n, k)$ in $N(n, k)$ is 1 for $n \leq 3$, because the sets are equal (Fischer et al[.](#page-16-11) [2018](#page-16-11)), it is likely that the density decreases markedly as *n* grows. That said, this is unknown and would be interesting to establish. The similarity in the diameters of the spaces could be due to the fact that two tree-based networks might be close in $N(n, k)$, but distant in $TBN(n, k)$, because the distance in the latter requires a path through networks that remain treebased. Or it could be due to the subspace of tree-based networks making up the majority of the space of networks, which seems unlikely but has not been ruled out. Finally, it could be simply that there are tree-based networks at the "extremities" of the space of all networks, and it just happens that they are dispersed widely within the space.

Another relevant question for future research might be to establish a lower bound on the diameter of the space of all tier- k tree-based networks *TBN* (n, k) . However, the methods used to establish a lower bound on the diameter of $N(n, k)$ obtained in Francis et al[.](#page-16-8) [\(2018](#page-16-8)) do not apply in the tree-based case in any obvious way, which is why this is a particularly interesting open problem that presumably requires a new approach.

Moreover, there are questions relating to the proximity measures introduced in Fischer and Franci[s](#page-16-3) [\(2020](#page-16-3)) (generalizing those for rooted networks in Francis et al[.](#page-16-20) [\(2018\)](#page-16-20)). In that paper, networks are given ranks according to their distance from the boundary of the space of tree-based networks. The boundary of $TBN(n, k)$ is the subset of tree-based networks that can be made non-tree-based with a single NNI move. Networks in the boundary are said to have *tree-based rank* 0. Tree-based networks that are at most *i* NNI moves from the boundary are rank *i*, and non-tree-based networks that are at most *i* NNI moves from the boundary are rank −*i*. For instance, the two non-tree-based tier-5 networks in Fig. [7](#page-14-0) (taken from Fischer et al[.](#page-16-11) [2018](#page-16-11), Fig. [7\)](#page-14-0) are rank -1 .

These two are not connected by a single NNI move, 3 so the set of networks of treebased rank −1 is not connected. Is this also the case for other negative ranks? And especially, what about networks of positive tree-based rank, in the "core", as it were, of the space of tree-based networks? While it is certainly of interest to use the results of (Fischer and Franci[s](#page-16-3) [2020](#page-16-3)) to analyse the structure of the space of binary tree-based networks more in-depth, note that in (Fischer and Franci[s](#page-16-3) [2020\)](#page-16-3), the authors discuss general tree-based networks, that is, non-binary ones. On the other hand, the results of the present manuscript are limited to the binary case. This is not merely due to the fact that binary networks play a fundamental role in mathematical phylogenetics. On the contrary: It can be easily seen that the space of *non-binary* tree-based networks is *not* connected under NNI. To see this, note that NNI does not change the total degree of the underlying graph, which is why an NNI move cannot connect networks whose total degrees differ.

A direct conclusion from the present manuscript is the fact that Theorem [4.1](#page-10-1) shows that $TBN(n, k)$ is also connected under the related SPR and TBR moves, which have recently been defined for phylogenetic networks, because all NNI moves are also SPR and TBR moves (Francis et al[.](#page-16-8) [2018](#page-16-8)), Lemma 7. But we have left open the question of the diameter of the space under these moves, except to note that clearly it is also within $O(n^2)$, and so may be smaller.

However, there are two more open questions arising from this manuscript: First, our approach of using NNI to go from a binary unrooted network to a lineal network, then to a shoat network and then modify the shoat network, go back to a lineal network and finally arrive at the target network does probably not lead to a shortest path. The approach is constructive as it shows how to construct a valid tree-based path from any binary tree-based network *N* to any other one *N*2, but there may be shorter paths from *N*¹ to *N*2. In particular, it might be possible to avoid shoat networks altogether and directly move from one lineal network to another one. If this was possible, it would reduce the complexity of our construction.

Second, it should be noted that NNI moves on phylogenetic networks have been introduced in the *rooted* setting (Gambette et al[.](#page-16-21) [2017](#page-16-21)), and of course this is also the original setting for the introduction of tree-based phylogenetic networks (Francis and Stee[l](#page-16-4) [2015\)](#page-16-4). While there are concrete connections between rooted and unrooted tree-based networks (e.g. Fischer and Franci[s](#page-16-3) [2020](#page-16-3), Sect. 6), it is not clear whether the results in the present paper will lift to the rooted context.

³ We used the co[m](#page-16-22)puter algebra system Mathematica (Wolfram [2017\)](#page-16-22) to verify that the shortest path from network (i) in Fig. [7](#page-14-0) to a network isomorphic to (ii) requires 5 NNI moves. While this is tricky to see, it is combinatorially rather easy to see that network (i) has 24 1-step NNI neighbours, which can be divided into two classes (i.e. the NNI neighbourhood of network (i) contains only two non-isomorphic networks): those isomorphic to (i), and those isomorphic to a specific different network, which is in fact tree-based. So network (ii) cannot be in the 1-step neighbourhood of (i).

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