ENCODING AND CONSTRUCTING 1-NESTED PHYLOGENETIC NETWORKS WITH TRINETS

K. T. HUBER AND V. MOULTON.

ABSTRACT. Phylogenetic networks are a generalization of phylogenetic trees that are used in biology to represent reticulate or non-treelike evolution. Recently, several algorithms have been developed which aim to construct phylogenetic networks from biological data using *triplets*, i.e. binary phylogenetic trees on 3-element subsets of a given set of species. However, a fundamental problem with this approach is that the triplets displayed by a phylogenetic network do not necessary uniquely determine or encode the network. Here we propose an alternative approach to encoding and constructing phylogenetic networks, which uses phylogenetic networks on 3-element subsets of a set, or *trinets*, rather than triplets. More specifically, we show that for a special, well-studied type of phylogenetic network called a 1-nested network, the trinets displayed by a 1-nested network always encode the network. We also present an efficient algorithm for deciding whether a *dense* set of trinets (i.e. one that contains a trinet on every 3-element subset of a set) can be displayed by a 1-nested network or not and, if so, constructs that network. In addition, we discuss some potential new directions that this new approach opens up for constructing and comparing phylogenetic networks.

Keywords phylogenetic network, triplets, trinets, reticulate evolution

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1. INTRODUCTION

Phylogenetic networks are a generalization of phylogenetic trees that are used in biology to represent reticulate or non-treelike evolution (cf. [12, 23] for recent overviews). There are various types of phylogenetic networks, but in this paper we shall focus on phylogenetic networks that *explicitly* represent the evolution of a given set of species. Such networks (whose formal definition is presented in Section 2) can be essentially regarded as directed acyclic graphs having a single root,

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whose internal vertices represent ancestral species and whose leaves represent the set species (see e.g. Fig. 1). They have been used, just to name a few examples, to represent the evolution of viruses [28], bacteria [25], plants [22], and fish [20].

Recently, several algorithms have been developed which aim to construct phylogenetic networks (cf. [12, 23]). However, as stated in [12, p.xi], "While there is a great need for practical and reliable computational methods for inferring rooted phylogenetic networks to *explicitly* describe evolutionary scenarios involving reticulate events, generally speaking, such methods do not yet exist, or have not yet matured enough to become standard tools".

Probably one of the main reasons for this is that we do not yet have a very good understanding of how to build up complex phylogenetic networks from simpler structures. An important case in point is the construction of phylogenetic networks from phylogenetic trees. Even though there has been a great deal of recent work on this problem (cf. [12, Chapter 11], [23, Section 2]), especially concerning the construction of networks from *triplets* (i.e. binary phylogenetic trees with three leaves) [10, 11, 13, 14, 16, 17, 30]), there is a fundamental obstacle to this approach: The trees displayed by a phylogenetic network do not necessarily determine or *encode* the network [10] (even on 3 species – see e.g. Fig. 1) and, in fact, we do not even know when a phylogenetic network is uniquely determined by *all* of the trees that it displays [32].

As an alternative approach to tackling the problem of constructing phylogenetic networks, in this paper we shall investigate the following strategy: Instead of constructing phylogenetic networks from trees, try to build them up from (simpler) phylogenetic networks. More specifically, we investigate how to construct phylogenetic networks from *trinets*, that is, phylogenetic networks having just three leaves (see, for example, the networks N_1 and N_2 in Fig. 1).



FIGURE 1. Two distinct phylogenetic networks N_1 and N_2 with leaf set $\{x, y, z\}$ that display the same set $\{T_1, T_2\}$ of phylogenetic trees. In particular, neither of these two networks is encoded by this set of trees.

One of the main difficulties that we had to overcome before being able to put this strategy into practice was to find an appropriate definition for the set of trinets that is displayed by a phylogenetic network (see Definition 3.1). However, with this definition in hand, we are able to show that any *1-nested network* – a quite simple and well-studied type of phylogenetic network [7] – is always encoded by the set of trinets it displays (Theorem 6.3). Moreover, using this fact, we provide a polynomial-time algorithm for deciding whether a given *dense* set of trinets (i.e. one that contains a trinet on every 3-element subset of a set) can be displayed by a 1-nested network or not and, if so, constructs that network (see Fig. 10 and Theorem 7.3).

We now describe the contents of the rest of the paper. In Section 2 we introduce some relevant, basic terminology concerning phylogenetic networks. In Section 3 we define the rather natural concept of a recoverable network, and show that, although a phylogenetic network need not be recoverable in general, a 1-nested network always is. In the following section, we show that a recoverable phylogenetic network is 1-nested if and only if all of its displayed trinets are 1-nested (Theorem 4.3). Using this fact and certain operations on 1-nested networks that are closely related to those presented in [7] and that are presented in Section 5, we then establish Theorem 6.3 in Section 6. As a corollary, we obtain a new (and efficiently computable) proper metric on the set of 1-nested networks all having the same leaf set (see Corollary 6.4). In Section 7 we present our main algorithm for checking whether or not a dense set of trinets is displayed by a 1-nested network. We conclude in Section 8 with a discussion on some possible future directions, including some ideas about how trinets might be used in practical applications.

2. Preliminaries

For the rest of this paper, X is a non-empty, finite set (which will usually correspond to a set of species or organisms). For consistency, we follow the notation presented in [7] where appropriate.

An $rDAG \ N = (V, A)$ is a directed acyclic graph (DAG) with nonempty vertex set V = V(N), non-empty arc set A = A(N) (with no multiple arcs) and single root $\rho = \rho_N$ (i.e. a DAG with precisely one source ρ). We let \langle_N denote the usual partial order on V induced by N. The underlying graph of N is denoted \underline{N} . A cycle in \underline{N} is a subset $C = \{v_1, v_2, \ldots, v_n\} \subseteq V(\underline{N}), n \geq 3$, such that $\{v_i, v_{i+1}\} \in E(\underline{N})$ for all $1 \leq i \leq n-1$ and $\{v_1, v_n\} \in E(\underline{N})$. If C is some cycle in \underline{N} and there is some $v \neq w \in V$ so that the union of all of the arcs in Nhaving both vertices in C is the union of two directed paths in N that both start at v and end at w, then v(w) is called the *split (end)* vertex of C. We denote an arc $a \in A$ with tail x (= tail(a)), and head y by (x, y). We call (x, y) a *cut arc (of* N) if the removal of the edge $\{x, y\}$ from $E(\underline{N})$ disconnects \underline{N} . A vertex $v \in V$ is a called a *leaf* of N if *indegree*(v) = 1 and *outdegree*(v) = 0. We denote the set of leaves of N by L(N). Every vertex of N that is neither the root ρ_N nor has outdegree 0 is called an *interior* vertex of N. A tree vertex $v \in V$ is an interior vertex of N with *indegree*(v) = 1, and a *hybrid vertex* $v \in V$ is an interior vertex with *indegree* $(v) \geq 2$. Note that neither the root ρ_N nor a leaf of N is a tree vertex and that a hybrid vertex of N cannot be a leaf.

Now, an X-rDAG is an rDAG N = (V, A) with leaves uniquely labeled by the elements in X (i.e. there is a map $\phi_N : X \to V$ such that ϕ maps X bijectively onto L(N)). We will usually just assume L(N) =X in case the labeling map is clear from the context. A phylogenetic network N = (V, A) (on X) is an X-rDAG such that every tree vertex has outdegree at least 2 and every hybrid vertex has outdegree at least 1. If N is such a network and N' = (V', A') is a phylogenetic network on a non-empty finite set Y, then N is isomorphic to N' if there is a bijection $\xi : X \to Y$ and a directed graph isomorphism $\iota : V \to V'$ between N and N' such that $\phi_{N'} = \iota \circ \phi_N \circ \xi^{-1}$. In particular, in case Y = X we consider X as being a subset of both V and V', and hence N is isomorphic to N' if and only if ι restricted to X is the identity map on X.

A phylogenetic network N = (V, A) on X is

- a bush (on X) if it is isomorphic to the phylogenetic network with vertex set $V = X \cup \{v\}, v \notin X$, and arc set $A = \{(v, x) : x \in X\}$,
- a two-leafed network (on X) if $X = \{x, y\}$, and N is isomorphic to the phylogenetic network on X with vertex set $V = \{u, v, w, x, y\}$ and arc set $A = \{(u, w), (u, v), (v, w), (v, x), (w, y)\}$,
- *binary* if all of its hybrid vertices have indegree 2 and outdegree 1 and all of its tree vertices have outdegree 2,
- 1-nested if every pair of cycles in \underline{N} intersect in at most 1 vertex¹,
- a galled tree if every pair of cycles in \underline{N} is disjoint,
- a (rooted) phylogenetic tree if \underline{N} is a tree, and
- a trinet if |L(N)| = |X| = 3.

¹Note that in [7], 1-nested networks are defined in such a way that every hybrid vertex has indegree 2 - we do not make this assumption, but we will use the same name rather than introducing another term.



FIGURE 2. The fourteen possible non-isomorphic, 1nested trinets on the set $\{x, y, z\}$. Directions on arcs are omitted for clarity; internal vertices indicated with a dot are all hybrid vertices. Leaves that are at the bottom of a trinet are indicated with large dots and vertices hanging off the side of a trinet with a square.

Note that a 1-nested network N on X with |X| = 1 is a bush with arc set consisting of precisely one arc, and if |X| = 2 then N is isomorphic to either a two-leafed network or a phylogenetic tree with 2 leaves.

In Fig. 2 we picture the set of all possible non-isomorphic 1-nested trinets on $\{x, y, z\}$. If N is a 1-nested trinet on X, |X| = 3, that is not isomorphic to a phylogenetic tree on X, then we say that $t \in X$ is at the bottom of N if it corresponds to one of the vertices represented by larger dots in Fig. 2, and we say that t hangs off the side of N if it corresponds to one of the vertices represented by a square in that figure (note that, in particular, there may be more than one element at the bottom of a trinet).

Finally, let \mathcal{T} denote a non-empty set of trinets such that $L(T) \in {X \choose 3}$ for all $T \in \mathcal{T}$ (which we shall also call a *trinet set* (on X) for short). If $Y \subseteq X$, $|Y| \ge 3$, we let \mathcal{T}_Y be the subset of \mathcal{T} consisting of those trinets $T \in \mathcal{T}$ with $L(T) \subseteq Y$. In addition, we call \mathcal{T} dense (on X) if ${X \choose 3} = \{L(N) : N \in \mathcal{T}\}$ and $|\mathcal{T}| = {|X| \choose 3}$.

3. TRINETS AND RECOVERABLE NETWORKS

In this section, we investigate networks that display only 1-nested trinets. In particular, we show that even if every trinet displayed by a network N is 1-nested, it does not necessarily follow that N is 1-nested.

In addition, we shall introduce a rather natural condition on N (that it is a 'recoverable network') for which this statement does in fact hold (see Theorem 4.3 in the next section).

Suppose N = (V, A) is a phylogenetic network on X, $|X| \ge 3$, and Y is a non-empty subset of $V - \{\rho_N\}$. Let v(Y) to be the last vertex in V - Y that lies on all paths in N from ρ_N to every $y \in Y$. Note that if Y consists of a single vertex y, then $v(\{y\})$ is known as the *immediate dominator of* y [21] (see also [12, p. 143] where it is called the *lowest stable ancestor of* y)).

We now present a key definition (see also Fig. 3):

Definition 3.1. Given a phylogenetic network N on X and some $Y \in {\binom{X}{3}}$, we define the trinet on Y displayed by N to be the trinet N_Y with leaf set Y which is obtained from N by first taking the network \tilde{N} consisting of the union of all directed paths in N starting at v(Y) and ending at some element in Y, and then repeatedly first (i) suppressing all vertices v with indegree(v) = outdegree(v) = 1, and then (ii) suppressing all multiple arcs that might result, until a trinet on Y is obtained. Put $Tr(N) = \{N_Y : Y \in {\binom{X}{3}}\}$.

Given a phylogenetic network N on X, we say that a trinet set \mathcal{T} on X is *displayed* by N if $\mathcal{T} \subseteq Tr(N)$. Moreover, we say that \mathcal{T} encodes N if $\mathcal{T} \subseteq Tr(N)$ and, if N' is any other phylogenetic network on X with $\mathcal{T} \subseteq Tr(N')$, then N' is isomorphic to N.

Note that in Definition 3.1 it is necessary to consider (at least) 3element subsets of X, since if 'binets' are defined in a similar way for 2-element subsets, then the resulting set would not in general encode the network (even if the network is a tree). Also note that we do not define a trinet on Y displayed by N to be the network consisting of the union of all directed paths in N to the elements of Y as this can result in networks with vertices having in- and outdegree 1, that is, networks that are not phylogenetic networks.

The proof of the following lemma is straight-forward and is omitted:

Lemma 3.2. Suppose that N is a 1-nested network on X, $|X| \ge 3$. Then any element in Tr(N) is isomorphic to one of the fourteen trinets on $\{x, y, z\}$ presented in Fig. 2.

Remark 3.3. If N is a 1-nested network on X, $|X| \ge 3$, then N is binary if every element in Tr(N) is isomorphic to either $T_1(x, y, z)$ or one of $N_i(x, y, z)$, $1 \le i \le 7$. Moreover, binary level-1 networks and galled trees (as defined in [7]) can be characterized in a similar manner.

Now, suppose that N is a phylogenetic network on X such that every trinet in Tr(N) is isomorphic to one of the fourteen trinets presented



FIGURE 3. (a) A phylogenetic network N on $X = \{x_1, \ldots, x_n, z, y_1, \ldots, y_n\}, n \ge 1$. (b) The subnetwork N' obtained by taking the union of the directed paths from v(Y) to every element in $Y = \{x_2, y_n, z\}$. (c) The subnetwork N'' obtained from N' by suppressing all multiple arcs of N'. (d) The trinet obtained from N'' by suppressing all vertices $v \in V(N'')$ with indegree(v) = outdegree(v) = 1. Directions of arcs are omitted when clear.



FIGURE 4. A phylogenetic network N on $\{x, y, z\}$ for which Tr(N) consists of precisely the trinet $T_1(x, y, z)$ but N is not a phylogenetic tree on $\{x, y, z\}$. As before, directions are omitted for clarity when clear. Also only the vertices that are leaves are marked by a dot.

in Fig. 2. It is tempting to think that this should imply that N is 1-nested. However, this is not the case. For example, even if N is a phylogenetic network such that every trinet in Tr(N) is isomorphic to either $T_1(x, y, z)$ or $T_2(x, y, z)$ in Fig. 2, then N is not necessarily isomorphic to a phylogenetic tree (see e.g. Fig. 4). Even so, as we shall show in the next section (see Theorem 4.3), the aforementioned statement is almost correct.

To this end, we now introduce a special class of networks. Suppose that N is a phylogenetic network on X with $|X| \ge 3$. We say that a vertex $v \in V(N)$ is reachable from a vertex $w \in V(N) - v$ if there exists a directed path in N starting at w and ending in v. In addition, if $z \in V(N)$ is a vertex of N that lies on that path then we say that v is reachable from w by crossing z. We denote by $v_N^* \in V(N)$ the (necessarily unique) vertex in N for which there exist some distinct $x, y \in X$ with $v_N^* = v(\{x, y\})$ and, for all $\{u, v\} \in \binom{V(N)}{2} - \{x, y\}$, either $v_N^* = v(\{u, v\})$ holds or $v(\{u, v\})$ is reachable from v_N^* .

Now, we say that N is recoverable if $\rho_N = v_N^*$. We use the term recoverable, since for biological data it would not be possible to infer the structure of the network above v_N^* in case N is not recoverable, as there would be no way to 'detect' vertices above v_N^* using any pair of elements in X. As an illustration, the vertex v in the phylogenetic network N on $\{x, y, z\}$ pictured in Fig. 4 is the vertex $v_N^* = v(\{x, z\})$. Since $v_N^* \neq \rho_N$, N is not recoverable.

We now characterize recoverable networks N on X, $|X| \ge 3$, in terms of a special type of vertex. A vertex $v \in V(N)$ is a *cut vertex* of Nif the deletion of v (plus its incident edges) from <u>N</u> disconnects <u>N</u>. We denote the resulting graph by $\underline{N}\backslash v$. If, in addition, there exists a connected component K of $\underline{N}\backslash v$ such that $V(K) \cap L(N) = \emptyset$ then we call v a separating vertex of N. For example, in Fig. 4 v is a separating vertex of N whereas vertex w is a cut vertex of N.

Proposition 3.4. Suppose N is a phylogenetic network on X, $|X| \ge 3$. Then the following statements hold:

- (i) If N is not recoverable then v_N^* is a cut vertex of N.
- (ii) N is recoverable if and only if v_N^{*} is not a separating vertex of N.

Proof. (i) Note first that $\rho_N \neq v_N^*$ as N is not recoverable. Let $x, y \in X$ distinct such that $v_N^* = v(\{x, y\})$, and assume for contradiction that v_N^* is not a cut vertex of N. Then there must exist some leaf $l \in L(N)$ of N that is reachable from ρ_N without crossing v_N^* . Hence, there exists some $z \in \{x, y\}$ such that $v_N^* \neq v(\{l, z\})$ and v_N^* is reachable from $v(\{l, z\})$; a contradiction. Thus, v_N^* is a cut vertex of N.

(ii) We prove the contrapositive of the statement i. e. we show that N is not recoverable if and only if v_N^* is a separating vertex of N. Suppose $\{x, y\} \in \binom{X}{2}$ such that $v_N^* = v(\{x, y\})$. Assume first that N is not recoverable. Then $\rho_N \neq v_N^*$ and, by (i), v_N^* is a cut vertex of N. Hence, for every leaf $l \in L(N)$ of N, every directed path from ρ_N to l must cross v_N^* . Let K_{ρ_N} denote the connected component of $\underline{N} \setminus v_N^*$ that contains ρ_N in its vertex set. Then $V(K_{\rho_N}) \cap L(N) = \emptyset$. Thus v_N^* is a separating vertex of N. Conversely, suppose that v_N^* is a separating vertex of N. Then v_N^* is a cut vertex of N and so every directed path from ρ_N to a leaf $z \in L(N)$ of N must cross v_N^* . If N were recoverable then $\rho_N = v_N^*$ would follow, implying that every vertex in N must lie on a directed path from v_N^* to a leaf of N. But then $V(K) \cap L(N) \neq \emptyset$ for every connected component K in $\underline{N} \setminus v_N^*$; a contradiction. Thus, N cannot be recoverable. \Box

It immediately follows that 1-nested networks are always recoverable:

Corollary 3.5. Suppose N is a phylogenetic network on X, $|X| \ge 3$. If N is 1-nested, then N is recoverable.

Proof. Suppose for contradiction that there exists a 1-nested network N on X that is not recoverable, that is, $\rho_N \neq v_N^*$. Then, by Proposition 3.4(i), v_N^* is a cut vertex of N. Hence, for every leaf $l \in L(N)$ of N, every directed path from $\rho_N \neq v_N^*$ to l must cross v_N^* . Since $outdegree(\rho_N) \geq 2$ and N cannot have multiple arcs, it follows that there exist (at least) 3 distinct directed paths in N from ρ_N to v_N^* . But then there must exist two cycles in \underline{N} which intersect in at least 2 vertices; a contradiction.

4. 1-NESTED TRINETS IMPLY 1-NESTED NETWORKS

In the last section, we proved that if N is a 1-nested phylogenetic network on X, $|X| \ge 3$, then N is recoverable. We shall now prove that if all of the trinets displayed by a recoverable network are 1-nested, then the network is 1-nested (Theorem 4.3).

To this end, suppose that N is a phylogenetic network on $X, |X| \ge 3$, and that C is a cycle of <u>N</u>. Put

$$Z(C) = \{ v \in C : \text{ there exist } \{a, a'\} \in \binom{A(C)}{2} \text{ with } tail(a) = tail(a') = v \}.$$

Clearly, $Z(C) \neq \emptyset$.

Now, suppose $l \in L(N)$ is a leaf of N that is reachable from a hybrid vertex of N. We denote by p(l) the number of distinct directed paths in N from ρ_N to l. Clearly $p(l) \geq 2$. Moreover, we denote by w(l) the unique vertex of N distinct from l that simultaneously lies on every directed path from ρ_N to l such that (i) w(l) is a hybrid vertex of N, and (ii) there is a unique directed path from w(l) to l such that every interior vertex of N on this path is a tree vertex of N. To illustrate these definitions, consider the network N on $\{x, y, z\}$ depicted in Fig. 4. Then w(y) is the unique hybridization vertex of N and p(y) = 3.

We now prove some useful, but somewhat technical, results concerning the set Z(C).



FIGURE 5. The situation considered in the proof of Proposition 4.1. The vertices in C which have two of their incoming (outgoing) arcs contained in A(C) are marked with squares (triangles). The leaves of N plus the vertex $v(\{z_1, \ldots, z_m\}$ are marked by dots. For clarity all other vertices are not marked. The directed lines represent directed paths rather than arcs.

Proposition 4.1. Suppose N is a recoverable phylogenetic network on $X, |X| \ge 3$, such that every trinet in Tr(N) is isomorphic to one of the fourteen trinets on $\{x, y, z\}$ depicted in Fig. 2. Then |Z(C)| = 1, for all cycles C in <u>N</u>.

Proof. Suppose for contradiction that \underline{N} contains a cycle C with $m := |Z(C)| \geq 2$. Put $Z(C) = \{z_1, \ldots, z_m\}$. Since C is a cycle in \underline{N} there must exist distinct vertices $h_i \in C$, $1 \leq i \leq m$, such that, for all $1 \leq i \leq m$, two of the incoming arcs of h_i are contained in A(C) and h_i can be reached from z_i and from z_{i+1} , $1 \leq i \leq m$, where we define $z_{m+1} := z_1$. Moreover for each such vertex h_i there must exist a leaf $l_i \in L(N)$ of N that is reachable from h_i . Note that some of the leaves l_i might be the same (see Fig. 5 for a representation of the generic situation in which all leaves l_i , $1 \leq i \leq m$, are distinct).

Choose some $i \in \{1, \ldots, m\}$, say i = 1, and let σ be the ordering l_1, l_2, \ldots, l_m of the leaves $l_j, 1 \leq j \leq m$ induced by C via the vertices $h_i, 1 \leq i \leq m$. If there exist at least three distinct leaves in that ordering, then let $l_{i_1}, l_{i_2}, l_{i_3}$ denote the first three distinct leaves in σ . Note that $l_1 = l_{i_1}$ and each of l_{i_1}, l_{i_2} , and l_{i_3} is reachable from $v(\{z_1, z_2, \ldots, z_p\}) \in V(N)$, where $p \in \{1, \ldots, m\}$ is such that for all $i_3 \leq q \leq p$ we have $l_q = l_{i_3}$. But then $p \geq 3$ and the trinet N' on $\{l_{i_1}, l_{i_2}, l_{i_3}\}$ displayed by N contains $v(\{z_2, \ldots, z_p\})$ in its vertex set if $p \neq m$ and, otherwise, the vertex $v(\{z_1, z_2, \ldots, z_p = z_m\})$. Hence, if $p \neq m$ then $z_j \in V(N'), 2 \leq j \leq p-1$, and otherwise, $z_j \in V(N')$ with $j = 1, \ldots, m$. Consequently, $\underline{N'}$ contains two cycles that intersect in a path of length 1 or more in each case. Since, by construction, each cycle is the union of two directed

paths in N that have the same start and end vertex this implies that N' is not of the specified form, a contradiction.

Now, if there exist just two leaves l_{i_1} and l_{i_2} in σ that are distinct, then choose some $l \in L(N) - \{l_{i_1}, l_{i_2}\}$, which must exist as $|X| \ge 3$. Since each of l_{i_1} and l_{i_2} is reachable from $v(\{z_1, z_2, \ldots, z_m\}) \in V(N)$ it follows that $v(\{z_1, z_2, \ldots, z_m\})$ is a vertex in the trinet N' on $\{l, l_{i_1}, l_{i_2}\}$ displayed by N. But then $z_j \in V(N')$, $1 \le j \le m$ which implies that $\underline{N'}$ contains two cycles that intersect in a path of length at least 1. As before, this yields a contradiction.

So suppose that $l_i = l_j$ for all $i, j \in \{1, \ldots, m\}$. Let $L_{w(l_1)} \subseteq L(N)$ denote the set of leaves of N that are reachable from $w(l_1)$. We claim that $w(l_1)$ is not a cut vertex of N. Suppose for contradiction that $w(l_1)$ is a cut vertex of N. Then, since N is recoverable, there must exist a leaf $l \in L(N) - L_{w(l_1)}$ that is reachable from ρ_N without crossing $w(l_1)$. Choose some $l' \in L(N) - \{l_1, l\}$, which must exist as $|X| \ge 3$. Since l_1 is reachable from each of $z_j, 1 \le j \le m, v(\{z_1, z_2, \ldots, z_m\}) \in V(N)$ must be a vertex in the trinet N' on $\{l, l', l_1\}$ displayed by N. But then $z_i \in V(N'), 1 \le i \le m$, and so we obtain a contradiction as before. Thus, $w(l_1)$ cannot be a cut vertex of N, as claimed.

Thus, there must exist some leaf $l \in L_{w(l_1)}$ that is reachable from ρ_N without crossing $w(l_1)$. But then $l_1 \neq l$, by the definition of $w(l_1)$. Arguments similar to the ones used in the previous case can be now used to obtain a final contradiction. Thus, |Z(C)| = 1 must hold for every cycle C of \underline{N} .

To establish Theorem 4.3 we will use one further result that follows from the last proposition. Suppose N is a phylogenetic network on X, $|X| \ge 3$, and C is a cycle in <u>N</u> with |Z(C)| = 1. Then we denote the unique vertex in C that has two of its incoming arcs contained in A(C)by h_C .

Corollary 4.2. Let N be a recoverable phylogenetic network on X, $|X| \ge 3$, such that every trinet in Tr(N) is isomorphic to one of the fourteen trinets on $\{x, y, z\}$ depicted in Fig. 2. Let C_1 and C_2 denote two distinct cycles of <u>N</u> for which $A(C_1) \cap A(C_2) \neq \emptyset$ holds, and let $l \in L(N)$ denote a leaf of N that is reachable from both h_{C_1} and h_{C_2} . Then w(l) is not a cut vertex of N.

Proof. Suppose for contradiction that this is not the case, that is, there exists a recoverable phylogenetic network N on X, two distinct cycles C_1 and C_2 in \underline{N} with $A(C_1) \cap A(C_2) \neq \emptyset$, and a leaf $l \in L(N)$ of N that is reachable from h_{C_1} and from h_{C_2} but that w(l) is a cut vertex

of N. Since N is recoverable, there must exist a leaf $l' \in L(N) - \{l\}$ of N that is reachable from ρ_N without crossing w(l).

Now let $z_i \in V(N)$ denote the unique vertex in $Z(C_i)$ i = 1, 2. Note that $z_1 = z_2$ might hold. Since l is clearly also reachable from both z_1 and z_2 , there must exist a directed path from ρ_N to $v(\{z_1, z_2\})$ that crosses $v(\{l, l'\})$. Choose some $l'' \in L(N) - \{l, l'\}$ which must exist as $|X| \geq 3$. Then the trinet N' on $\{l, l', l''\}$ displayed by N contains the vertex $v(\{z_1, z_2\})$ and thus every arc in $A(C_1) \cup A(C_2)$. Since $A(C_1) \cap A(C_2) \neq \emptyset$ it follows that N' is not of the specified form which is impossible. \Box

We now prove the main result of this section:

Theorem 4.3. Suppose that N is a recoverable phylogenetic network on X, $|X| \ge 3$. Then N is 1-nested if and only if every trinet in Tr(N)is isomorphic to one of the fourteen trinets depicted in Fig. 2.

Proof. If N is 1-nested then, by Lemma 3.2, the trinets in Tr(N) are of the specified form.

Conversely, suppose that the trinets in Tr(N) are of the specified form. Assume for contradiction that N is not 1-nested. Then there must exist two cycles C_1 and C_2 in <u>N</u> which intersect in more than one vertex. Moreover, amongst all such pairs of cycles, there must exist a pair C_1 and C_2 for which the following holds: There is a path P with $V(P) \subseteq C_1 \cap C_2$ which has an end vertex $x_2 \in V(P)$ such that the edge $\{x_1, x_2\} \in E(P)$ is the arc (x_1, x_2) in A(N) and $\{y, x_2\} \notin$ $E(C_1) \cap E(C_2)$, for all $y \in (C_1 \cap C_2) - \{x_1, x_2\}$. Choose some $z_i \in Z(C_i)$, i = 1, 2 and note that, by Proposition 4.1, $|Z(C_i)| = 1$. However note that $z_1 = z_2$ might hold.

Let $l_i \in L(N)$ denote a leaf of N that is reachable from $h_i = h_{C_i}$, i = 1, 2. Then one of the three generic cases (a) - (c) pictured in Fig. 6 must hold. Note that in the case of (b) and (c) we can choose l_1 to equal l_2 since in case of (b) we have $x_2 = h_2$ and in case of (c) we have $x_2 = h_2 = h_1$.

Suppose first that Case (a) holds. We begin by considering the case $l_1 = l_2$. Since N is recoverable, Corollary 4.2 implies that $w(l_1)$ is a not a cut vertex of N. Let $L_{w(l_1)} \subseteq L(N)$ denote the set of leaves of N that are reachable from $w(l_1)$. Then there must exist a leaf $l \in L_{w(l_1)}$ of N that is reachable from ρ_N without crossing $w(l_1)$. By the definition of $w(l_1), l_1 \neq l$. Since l is reachable from z_1 and from z_2 , there must exist a directed path from ρ_N to $v(\{z_1, z_2\})$ that crosses $v(\{l_1, l\})$. Choose some $l' \in L(N) - \{l_1, l\}$, which must exist as $|X| \geq 3$. Then the trinet N' on $\{l_1, l, l'\}$ displayed by N contains the vertex $v(\{z_1, z_2\})$. Thus,

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FIGURE 6. The three generic cases considered in the proof of Theorem 4.3. The vertices in C_i , i = 1, 2 which have two of their incoming (outgoing) arcs contained in $A(C_i)$, i = 1, 2 are marked with squares (triangles). The leaves of N plus the vertices ρ_N , x_1 , x_2 , $v(\{z_1, z_2\})$ and $w(l_i)$, i = 1, 2 are marked by dots. For clarity all other vertices are not marked. The directed lines represent directed paths rather than arcs.

<u>N'</u> contains two cycles that intersect in the edge $\{x_1, x_2\}$. Since each cycle is the union of two directed paths in N that have the same start vertex and the same end vertex, it follows that N' is not of the specified form, a contradiction. Thus $l_1 \neq l_2$ must hold.

Since $|X| \geq 3$, we may choose some $l \in L(N) - \{l_1, l_2\}$. But then similar arguments applied to the trinet N' on $\{l_1, l_2, l\}$ displayed by N yields a contradiction.

Similar arguments can be used to show that Case (b) and Case (c) lead to a contradiction. But this implies that there cannot exist two distinct cycles of \underline{N} that intersect in more than one vertex. Thus, N must be 1-nested.

As a corollary we see that if all of the trinets displayed by a recoverable phylogenetic network are trees then the network must be a tree.

Corollary 4.4. Suppose N is a recoverable phylogenetic network on $X, |X| \ge 3$. Then N is a phylogenetic tree on X if and only if every trinet in Tr(N) is isomorphic to either the trinet $T_1(x, y, z)$ or the trinet $T_2(x, y, z)$ on $\{x, y, z\}$.

Proof. This is an immediate consequence of Theorem 4.3 and the fact that if N is recoverable 1-nested network on X then \underline{N} contains a cycle if and only if there exists a trinet $N' \in Tr(N)$ such that $\underline{N'}$ contains a cycle.

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5. Cherries, cactuses and reductions

In the next section, we shall show that the set of trinets displayed by a phylogenetic network encode the network. To do this, we will use some operations that can be performed on 1-nested networks to produce new 1-nested networks which we shall now introduce. These operations are very closely related to the "R, T and G-operations" presented in [7, Section 4]. In consequence, we shall omit the proofs of the results that we state concerning our operations, instead citing the related results in [7, Section 4] which have very similar proofs.

Suppose N = (V, A) is a 1-nested network on X, $|X| \ge 2$. We call a subset $S \subseteq X$ a cherry of N if $|S| \ge 2$ and there is some $v_S \in V$ such that $(v_S, x) \in A$ for all $x \in S$ and $(v_S, x) \notin A$ for all $x \in X - S$ (see Fig. 7(a)). Moreover, we shall call such a cherry *isolated* if *outdegree* $(v_S) = |S|$ and *indegree* $(v_S) = 1$ (see Fig. 7(b)). Note that if S is a cherry of N and S = X, then N is isomorphic to a bush on X. We now define a related concept. If $|X| \ge 2$, we call a tuple $H = (a_1, a_2, \ldots, a_p : b_1, b_2, \ldots, b_q : z)$ of distinct elements of X with $p \ge 1$, $q \ge 0$ a *cactus* of N (with support $S = \{a_1, a_2, \ldots, a_p, b_1, b_2, \ldots, b_q, z\}$) if there is cycle C_H in \underline{N} with split vertex v_H such that the network induced by N on $C_H \cup S$ is as pictured in Fig. 7(c) (note that if q = 0, we take the tuple to be $H = (a_1, a_2, \ldots, a_p : \emptyset : z)$). Moreover, such a cactus H is called *isolated* if *indegree* $(v_H) = 1$ and *outdegree* $(v_H) = 2$ (see Fig. 7(d)). Note that a two-leafed network on a set of size two is a cactus.

Now, suppose that N is 1-nested network on X, $|X| \ge 2$. In case there is a non-isolated cherry S of N and $z \in S$, then we define a



FIGURE 7. (a) A cherry $S = \{x_1, x_2, \ldots, x_m\}, m \ge 2$, (b) an isolated cherry $S = \{x_1, x_2, \ldots, x_m\}, m \ge 2$, (c) a cactus $H = (a_1, a_2, \ldots, a_p : b_1, b_2, \ldots, b_q : z), p \ge 1, q \ge 0$, and (d) an isolated cactus $H = (a_1, a_2, \ldots, a_p : b_1, b_2, \ldots, b_q : z), p \ge 1, q \ge 0$. Note that the arcs ending at v_S and v_H in (a) and (c) do not necessarily exist.

cherry reduction $C = C_{z:S}$ on N to be the network $C_{z:S}(N)$ which is obtained by removing all leaves in S except z from N, together with their incident arcs. In addition, if S is an isolated cherry of N and $z \in S$, then we define an *isolated cherry reduction* $\overline{C} = \overline{C}_{z:S}$ on N to be the network $\overline{C}_{z:S}(N)$ which is obtained by removing all leaves in Sfrom N, together with their incident arcs, and replacing the vertex v_S by z, which now becomes a leaf of the new network.

Similarly, suppose there is a cactus $H = (a_1, a_2, \ldots, a_p : b_1, b_2, \ldots, b_q : z)$ of N with support S. If H is not isolated, then we define a *cactus reduction* $H = H_{z:S} = H_{a_1,a_2,\ldots,a_p:b_1,b_2,\ldots,b_q:z}$ on N to be the network $H_{a_1,a_2,\ldots,a_p:b_1,b_2,\ldots,b_q:z}(N)$ which is obtained by removing the vertices $(C_H - \{v_H\}) \cup (S - \{z\})$, together with their induced arcs plus the two outgoing arcs of v_H contained in A(C), from N and then adding in the new arc (v_H, z) . In addition, if H is isolated, then we define an *isolated cactus reduction* $\overline{H} = \overline{H}_{z:S} = \overline{H}_{a_1,a_2,\ldots,a_p:b_1,b_2,\ldots,b_q:z}$ on N to be the network $\overline{H}_{a_1,a_2,\ldots,a_p:b_1,b_2,\ldots,b_q:z}(N)$ which is obtained by removing the vertices $(C_H - \{v_H\}) \cup (S - \{z\})$, together with their induced arcs plus the two outgoing arcs of v_H , from N and replacing v_H with z.

It is clear that the networks $C_{z:S}(N)$, $\overline{C}_{z:S}(N)$, $H_{a_1,a_2,\ldots,a_p:b_1,b_2,\ldots,b_q:z}(N)$ and $\overline{H}_{a_1,a_2,\ldots,a_p:b_1,b_2,\ldots,b_q:z}(N)$ are all 1-nested networks on the set $X - (S - \{z\})$ and that they all have |S| - 1 less leaves than N. Moreover we have:

Proposition 5.1. [7, Proposition 2] Suppose that N is a 1-nested network on X, $|X| \ge 1$. If $|X| \ge 2$, then at least one of the reductions C, \overline{C} , H, \overline{H} may be applied to N. Moreover, if none of the reductions C, \overline{C} , H, \overline{H} may be applied to N, then |X| = 1 and N is the bush on X.

We can also define 'inverses' of $C^{-1}, \overline{C}^{-1}, H^{-1}, \overline{H}^{-1}$ of the reductions $C, \overline{C}, H, \overline{H}$ as follows. Given a 1-nested network N on $X, |X| \ge 1$, a leaf $z \in X$ of N, and a set finite S with $|S| \ge 2$ and $S \cap X = \{z\}$, we define the *cherry expansion* $C_{z:S}^{-1}$ of N to be the network $C_{z:S}^{-1}(N)$ obtained by replacing leaf z by a new vertex v, and adding in new arcs (v, s) for all $s \in S$. Clearly $C_{z:S}^{-1}(N)$ is a 1-nested network on $X \cup S$. Isolated cherry, cactus and isolated cactus expansions $\overline{C}^{-1}, H^{-1}, \overline{H}^{-1}$, corresponding to \overline{C}, H and \overline{H} , are defined in a similar way.

It is straight-forward to see that a reduction and its corresponding expansion are mutual inverses, in that when one is applied to a 1nested network N on X and then its inverse, we obtain a network that is isomorphic to N. Moreover, we have:

Lemma 5.2. [7, Lemma 4] Let N and N' be two 1-nested networks on $X, |X| \ge 3$. If N and N' are isomorphic, then if one of the reductions

 $C, \overline{C}, H, \overline{H}$ (respectively, expansions $C^{-1}, \overline{C}^{-1}, H^{-1}, \overline{H}^{-1}$) may be applied to N, then the same one may also be applied to N' and the two resulting 1-nested networks are isomorphic.

6. Encoding 1-nested networks with trinets

In this section we show that the set of (necessarily 1-nested) trinets displayed by a 1-nested network N on X encodes N (see Theorem 6.3).

We begin by describing how to characterize cherries and cactuses in a 1-nested network in terms of their trinets, starting with cherries. To this end, we associate to a trinet set \mathcal{T} on X and a non-empty subset $S \subseteq X$ the trinet set

$$\mathcal{T}|_S := \{ N \in \mathcal{T} : S \cap L(N) \neq \emptyset \}.$$

Lemma 6.1. Suppose N = (V, A) is a 1-nested network on $X, |X| \ge 3$, and let $S \subseteq X$ with |S| > 2. Let \mathcal{T} be a non-empty subset of Tr(N). Then S is a cherry of N with $\mathcal{T} = Tr(N)|_S$ if and only if \mathcal{T} satisfies the following properties:

- (C1) $L(N') \cap S \neq \emptyset$, for all $N' \in \mathcal{T}$ (or equivalently, $\mathcal{T}|_S = \mathcal{T}$).
- (C2) For all $\{x, y\} \in {S \choose 2}$ and all $z \in X S$, either $T_1(x, y, z)$, $T_2(x, y, z), N_3(z, x, y), N_4(x, y, z), N_9(x, y, z) \text{ or } N_{10}(z, x, y) \text{ is}$ in \mathcal{T} .
- (C3) For all $\{x, y, z\} \in {S \choose 3}$, $T_2(x, y, z) \in \mathcal{T}$. (C4) There is no $S' \subseteq X$ such that $S \subset S'$ and \mathcal{T} satisfies (C2) and (C3) with S replaced by S'.

Moreover, if this is the case and $S \neq X$ (or, equivalently, $|X - S| \geq 1$), then S is isolated if and only if \mathcal{T} also satisfies:

(C5) For all $\{x, y\} \in {S \choose 2}$ and all $z \in X - S$, either $T_1(x, y, z)$, $N_3(z, x, y)$ or $N_4(x, y, z)$ is contained in \mathcal{T} .

Proof. Suppose $\mathcal{T} = Tr(N)|_S$ holds for some cherry S of N. Then it is straight-forward to check that \mathcal{T} satisfies (C1)–(C4).

Conversely, suppose \mathcal{T} satisfies (C1)–(C4). Let v = v(S). Note that $v(\{x,y\}) = v$ for all $\{x,y\} \in {S \choose 2}$, since otherwise there would exist some $z \in S$ such that $T_2(x, y, z) \notin \mathcal{T}$, in contradiction to (C3). Moreover, suppose there were some $z \in X - S$, $x \in S$ with $v(\{z, x\}) >_N$ v. Let $y \in S - \{x\}$ (which exists since $|S| \geq 2$). Then none of the trinets $T_1(x, y, z)$, $T_2(x, y, z)$, $N_3(z, x, y)$, $N_4(x, y, z)$ $N_9(x, y, z)$ or $N_{10}(z, x, y)$ could be contained in \mathcal{T} , in contradiction to (C2). Thus, for all $z \in X - S$ and all $x \in S$, we have $v(\{z, x\}) <_N v$ with possibly equality holding. It follows that $(v, x) \in A$ for all $x \in S$.

Now, suppose there is some $r \in X - S$ with $(v, r) \in A$. Let $S' = S \cup \{r\}$. Then it is straight-forward to check that, for all $x \in S'$ and all $z \in X - S'$, either $T_1(x, r, z)$, $T_2(x, r, z)$, $N_3(z, x, r)$, $N_4(x, r, z)$, $N_9(x, r, z)$ or $N_{10}(z, x, r)$ is in \mathcal{T} , and that $T_2(x, y, r) \in \mathcal{T}$ for all $\{x, y\} \in \binom{S'}{2}$. This implies that S' satisfies (C2) and (C3) with S replaced by S', which contradicts (C4). In particular, it follows that S is a cherry of N.

To see that $\mathcal{T} = Tr(N)|_S$ holds note first that $\mathcal{T} \subseteq Tr(N)|_S$ is a consequence of (C1). To see that $Tr(N)|_S \subseteq \mathcal{T}$ suppose $N' \in Tr(N)|_S$. Then $L(N') \cap S \neq \emptyset$ and so $N' \in \mathcal{T}$ follows from considering the size of the intersection $L(N') \cap S$ in conjunction with Properties (C2) and (C3).

To complete the proof, suppose that $X \neq S$. First note that if Sis an isolated cherry of N, then (C5) clearly holds. Conversely, if \mathcal{T} satisfies (C5), then let $v \in V$ be the vertex with $(v, x) \in A$ for all $x \in S$ and $(v, x) \notin A$ for all $x \in X - S$ (which exists since S is a cherry by (C2)–(C4)). Then *outdegree*(v) = |S|, since otherwise there would exist some $\{x, y\} \in {S \choose 2}$ and $z \in X - S$ with $z >_N v$ such that either $T_2(x, y, z)$ or $N_{10}(z, x, y) \in \mathcal{T}$, in contradiction to (C5).

Now, since $|X - S| \ge 1$, $indegree(v) \ge 1$. Suppose indegree(v) > 1. Then there must exist some $z \in X - S$ and $\{x, y\} \in {S \choose 2}$ such that $N_9(x, y, z) \in \mathcal{T}$, which contradicts (C5). Therefore indegree(v) = 1, which completes the proof.

We now present a similar result for cactuses.

Lemma 6.2. Let N be a 1-nested network on X, $|X| \ge 3$, and let $H = (a_1, \ldots, a_p : b_1, \ldots, b_q : z)$ be a tuple of distinct elements in X with $p \ge 1$ and $q \ge 0$. Put $S = \{a_1, \ldots, a_p, b_1, \ldots, b_q, z\}$ and let \mathcal{T} be a non-empty subset of Tr(N). Then H is a cactus of N with support S and $\mathcal{T} = Tr(N)|_S$ if and only if, with $A = \{a_1, \ldots, a_p\}$ and $B = \{b_1, \ldots, b_q\}$, \mathcal{T} satisfies the following properties:

- (H1) $L(N') \cap S \neq \emptyset$ for all $N' \in \mathcal{T}$ (or, equivalently $\mathcal{T}|_S = \mathcal{T}$).
- (H2) $N_1(x, z, y) \in \mathcal{T}$ for all $x \in A, y \in B$.
- (H3) $N_2(z, x, x') \in \mathcal{T}$ for all $x = a_i, x' = a_j, 1 \le i < j \le p$, or $x = b_r, x' = b_s, 1 \le r < s \le q$.
- (H4) $T_1(x, x', x'') \in \mathcal{T}$ for all $x = a_i, x' = a_j, x'' = a_k, 1 \le i < j < k \le p$, or $x = b_r, x' = b_s, x'' = b_t, 1 \le r < s < t \le q$.
- (H5) For all $w \in X-S$ either $N_5(z, x, w)$, or $N_6(z, x, w)$, or $N_7(w, x, z)$, or $N_8(z, x, w)$, or $N_{11}(z, x, w)$, or $N_{12}(w, x, z)$ is contained in \mathcal{T} , for all $x \in A$ or $x \in B$.

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- (H6) For all $w \in X S$, either $T_1(x, x', w)$, or $N_3(w, x, x')$, or $N_4(x, x', w)$ is contained in \mathcal{T} , for all $x \neq x' \in A$ or $x \neq x' \in B$.
- (H7) For all $w \in X S$, one of $T_1(x, y, w)$, $T_1(x, w, y)$, $T_1(y, w, x)$, $T_2(x, y, w)$, $N_4(x, y, w)$, and $N_1(x, w, y)$ is contained in \mathcal{T} , for all $x \in A$ and $y \in B$.
- (H8) There exists no tuple $H = (c_1, \ldots, c_t : d_1 \ldots, d_s : z)$ of distinct elements in $X, t \ge 1$ and $s \ge 0$, with $S \subsetneq S' := \{c_1, \ldots, c_t, d_1 \ldots, d_s, z\}$ such that \mathcal{T} satisfies (H2)-(H7) for S'.

Moreover, if this is the case and $S \neq X$, then H is isolated if and only if \mathcal{T} also satisfies:

(H9) For all $w \in X - S$, $T_2(x, y, w) \notin \mathcal{T}$, for all $x \in A$, $y \in B$, and $N_8(z, x, w) \notin \mathcal{T}$, $N_{11}(z, x, w) \notin \mathcal{T}$ and $N_{12}(w, x, z) \notin \mathcal{T}$ for all $x \in A$ or $x \in B$.

Proof. Suppose H is a cactus of N with support S and $\mathcal{T} = Tr(N)|_S$. Then it is straight-forward to see that \mathcal{T} must satisfy (H1)–(H8).

Conversely, suppose \mathcal{T} satisfies (H1)–(H8) with A and B as specified. We claim that H is a cactus of N with support S. We prove the claim for q = 0 and remark that the proof for $q \ge 1$ is similar. Let $x \in A$. If |S| = 2 then choose some $w \in X - S$. By (H5), one of the trinets $N_5(z, x, w)$, $N_6(z, x, w)$, $N_7(w, x, z)$, $N_8(z, x, w)$, $N_{11}(z, x, w)$, $N_{12}(w, x, z)$ must be contained in \mathcal{T} . But then H must clearly be a cactus of N (with support S).

Assume that $|S| \geq 3$. Then $|A| \geq 2$ and, by (H3), $N_2(z, a_i, a_j) \in \mathcal{T}$ or $N_2(z, a_j, a_i) \in \mathcal{T}$ holds for all $\{i, j\} \in \binom{\{1, \dots, p\}}{2}$. Since $\mathcal{T} \subseteq Tr(N)$, there must exist a cycle $C_{i,j}$ in \underline{N} with split vertex $v_{i,j} := v_{C_{i,j}}$ and end vertex $b_{i,j} := b_{C_{i,j}}$ that gives rise to that trinet on $\{z, a_i, a_j\}, \{i, j\} \in \binom{\{1, \dots, p\}}{2}$. We show that $C_{i,j} = C_{k,l}$ holds for all $\{i, j\}, \{k, l\} \in \binom{\{1, \dots, p\}}{2}$. To see this it suffices to show that $C_{i,j} = C_{i,l}$ holds for all $i \in \{1, \dots, p\}$ and all $\{k, l\} \in \binom{\{1, \dots, p\}-\{i\}}{2}$. So assume for contradiction that there exists some $i \in \{1, \dots, p\}$ and some $\{j, l\} \in \binom{\{1, \dots, p\}-\{i\}}{2}$ with $C_{i,j} \neq C_{i,l}$. Without loss of generality assume that i = 1.

Note that since N is 1-nested there must exist, for all $t \in \{2, \ldots, p\}$ and all $x \in \{a_1, a_t, z\}$, a unique last vertex $v_x^{1,t}$ in $C_{1,t}$ that lies on every path from ρ_N to x. Clearly, $v_z^{1,t}$ is the end vertex of $C_{1,t}$ and $v_{a_1}^{1,t}$ is neither the end vertex nor the split vertex of $C_{1,t}$, $t \in \{2, \ldots, p\}$. Put $v = v(\{v_{1,j}, v_{1,l}\})$.

We first show that $b_{1,j} = b_{1,l}$. Suppose for contradiction that $b_{1,j} \neq b_{1,l}$. Then since indegree(z) = 1, there must exist a vertex y_z distinct from z that lies simultaneously on any path in N from $b_{1,j} = v_z^{1,j}$ to z and on any path in N from $b_{1,l} = v_z^{1,l}$ to z. Without loss of generality,

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we may assume that y_z is as close to z as possible. So there must exist a cycle C in \underline{N} with $\{v, v_{1,j}, b_{1,j}, y_z, b_{1,l}, v_{1,l}\} \subseteq C$ with possibly $v = v_{1,j}$ or $v = v_{1,l}$ or $v = v_{1,j} = v_{1,l}$ or $b_{1,j} = y_z$ or $b_{1,l} = y_z$ holding. Since $\{v_{1,j}, b_{1,j}\} \subseteq C \cap C_{1,j}$ and N is 1-nested this is impossible. Thus $b_{1,j} = b_{1,l}$, as required.

Similar arguments with z replaced by a_1 in the definition of y_z also imply that $v_{a_1}^{1,j} = v_{a_1}^{1,l}$ must hold. But then $C_{1,j}$ and $C_{1,l}$ intersect in more than one vertex which is impossible as N is 1-nested. Thus $C_{1,j} = C_{1,l}$ must hold for all $j, l \in \{2, \ldots, p\}$. Moreover, by (H4), $v_{a_r}^{1,2} \neq v_{a_s}^{1,2}$ for all $\{r, s\} \in \binom{\{1, \ldots, p\}}{2}$. Thus there exists a directed path P from $v_{1,2}$ to $b_{1,2}$ that crosses the vertices $v_{a_1}^{1,2}, v_{a_2}^{1,2}, \ldots, v_{a_p}^{1,2}$ in that order.

To finish the proof of the claim that H is a cactus of N with support S, we next establish that $V(P) = Y := \{v_{a_1}^{1,2}, v_{a_2}^{1,2}, \dots, v_{a_p}^{1,2}, v_{1,2}, b_{1,2}\}$. Suppose for contradiction that this is not the case and that there exists some $u \in V(P) - Y$. Without loss of generality, we may assume that $(u, v_{a_1}^{1,2}) \in A(P)$. Since N is 1-nested, there exists some leaf $w \in L(N) - S$ that is reachable from u without crossing any further vertex in $C_{1,2}$. We distinguish the cases that X = S and that $X \neq S$. If X = S then this is impossible and so V(P) = Y, as required. Since $C_{1,2}$ is a cycle in \underline{N} and N is 1-nested it follows that $(v_{1,2}, b_{1,2})$ is an arc in N. But this implies that H is a cactus of N (with support S).

So assume that $S \neq X$. Then (H6) applied to a_1, a_2 , and w, combined with the fact that N is 1-nested, implies that the trinet $T_1(a_1, a_2, w)$ is contained in \mathcal{T} . But then \mathcal{T} satisfies (H2)–(H7) for the support $S \cup \{w\}$ of the tuple $H' = (w, a_1, \ldots, a_p : \emptyset : z)$. In view of (H8), this is impossible. Thus, V(P) = Y, as required.

We now show that $(v_{1,2}, b_{1,2}) \in A(C_{1,2})$. Suppose this is not the case and there exists some $u \in C_{1,2} - V(P)$. Without loss of generality we may assume $(v_{1,2}, u) \in A(C_{1,2})$. Then there exists a leaf $w \in L(N) - S$ such that u is the last vertex in $C_{1,2}$ on any path from ρ_N to w. But then the trinet on $\{w, a_1, z\}$ is not as specified in (H5) which is impossible. Thus, $(v_{1,2}, b_{1,2}) \in A(C_{1,2})$, as required. It follows that H must be a cactus of N (with support S) in this case, too.

To see that $Tr(N)|_S = \mathcal{T}$, let $N' \in Tr(N)|_S$. Then $L(N') \cap S \neq \emptyset$. By distinguishing the cases that $|L(N') \cap S| = 1, 2, \text{ or } 3$, it is straight forward to show that $N' \in \mathcal{T}$ using Properties (H2)–(H8). Also $\mathcal{T} \subseteq Tr(N)|_S$ holds by Property (H1).

It remains to show that if H is a cactus of N with support S and $S \neq X$ then H is isolated if and only if \mathcal{T} satisfies (H9). Assume that H is a cactus of N with support S and that $S \neq X$. Then it is straight forward to check that \mathcal{T} satisfies (H9).

Conversely, assume that \mathcal{T} satisfies (H9). We need to show that $outdegree(v_H) = 2$ and that $indegree(v_H) = 1$. We again prove the case q = 0 and remark that the arguments for $q \ge 1$ are similar. Since H is a cactus of N we clearly have $outdegree(v_H) \ge 2$. Assume for contradiction that $outdegree(v_H) > 2$. Then since N is 1-nested and $X \ne S$ there must exist some $w \in X - S$ that is reachable from v_H without crossing a vertex in $C_H - \{v_H\}$, where C_H is the cycle in \underline{N} corresponding to H. But then there exists some $x \in A$ such that $N_8(z, x, w)$ or $N_{12}(w, x, z)$ is contained in \mathcal{T} contradicting (H9). Thus, $outdegree(v_H) = 2$, as required. But then $indegree(v_H) \ge 1$ as $S \ne X$. Assume for contradiction that $indegree(v_H) > 1$. Then since $S \ne X$ there must exist some $w \in X - S$ such that $N_{11}(z, a, w) \in \mathcal{T}$ for some $a \in A$ contradicting again (H9). Thus, $indegree(v_H) = 1$. This completes the proof of Lemma 6.2.

Now, let $R_{z:S}$ $(R_{z:S}^{-1})$ denote any of the four reductions $C, \overline{C}, H, \overline{H}$ with z and S as specified in the definition of the reductions. Then it is straight-forward to check that if N is a 1-nested network on X, $|X| \geq 3$, and

$$\mathcal{T}_{z:S} = \{ N' \in Tr(N) : S \cap L(N') \neq \emptyset \text{ and } S \cap L(N') \neq \{z\} \},\$$

then

(1)
$$Tr(N) = Tr(R_{z:S}(N)) \amalg \mathcal{T}_{z:S},$$

or, in other words, $Tr(R_{z:S}(N)) = Tr(N) - \mathcal{T}_{z:S}$.

Theorem 6.3. Suppose that N and N' are both 1-nested networks on $X, |X| \ge 3$. Then Tr(N) = Tr(N') if and only if N is isomorphic to N'.

Proof. Suppose first that N is isomorphic to N'. Then Tr(N) = Tr(N') follows immediately by using induction on |X|, Lemma 5.2 and (1).

To prove the converse we also use induction on |X|. If |X| = 3, then the converse obviously holds. So, suppose that, for all $1 \le |X| \le m$, $m \ge 3$, if Tr(N) = Tr(N') then N is isomorphic to N'.

Let |X| = m + 1, and suppose that N and N' are 1-nested networks on X with Tr(N) = Tr(N'). By Proposition 5.1 we can apply at least one of the reductions $R = \overline{C}, C, H, \overline{H}$ to N. Therefore, since Tr(N) =Tr(N'), by Lemmas 6.1 and 6.2, we may also apply the same reduction R to N'. Moreover, by (1) we have Tr(R(N)) = Tr(R(N')). So, by induction, R(N) is isomorphic to R(N'). Therefore, by Lemma 5.2, $R^{-1}(R(N))$ is isomorphic to $R^{-1}(R(N'))$, i.e. N is isomorphic to N', as required. There has been some interest in the literature in defining metrics on networks [12, page 172], and various metrics have been defined for different types of phylogenetic networks including 1-nested networks [4, 5, 6, 7, 8, 10]. Thus the following result could be of interest. For X with $|X| \ge 3$, let $\mathcal{N}_1(X)$ denote the set of 1-nested networks on X. In addition, define the map

$$d: \mathcal{N}_1(X) \times \mathcal{N}_1(X) \to \mathbb{R}; (N, N') \mapsto d(N, N') := |Tr(N)\Delta Tr(N')|,$$

for all $N, N' \in \mathcal{N}_1(X)$. Then the last theorem immediately implies:

Corollary 6.4. For X with $|X| \ge 3$, the map d is a (proper) metric on $\mathcal{N}_1(X)$.

Note that the metric d can be efficiently computed since, for $N \in \mathcal{N}_1$, it is possible to compute every trinet in Tr(N) efficiently (essentially because for any $Y \in {X \choose 3}$ the vertex v(Y) can be computed efficiently using, e.g. the algorithm presented in [21]).

7. Constructing 1-nested networks from dense sets of trinets

In this section, we present an efficient algorithm which, given a dense set \mathcal{T} of trinets, can decide whether or not it is displayed by a 1-nested network, and if this is the case, constructs the network displaying \mathcal{T} (see Fig. 10).

We begin by describing efficient algorithms for detecting cherries and cactuses. Given a dense set \mathcal{T} of trinets on X, we say that $S \subseteq X$, $|S| \ge 2$ is a *cherry of* \mathcal{T} if the set $\mathcal{T}|_S$ satisfies conditions (C2)–(C4) (note that it necessarily satisfies (C1)), and that it is *isolated* if it also satisfies (C5). We now show that cherries can be found in polynomial time in a dense set of trinets using the algorithm presented in Fig. 8.

Lemma 7.1. Given a dense set \mathcal{T} of trinets on X, $|X| \ge 3$, algorithm FINDCHERRY is correct and has run-time that is polynomial in |X|.

Proof. It is straight-forward to see that algorithm FINDCHERRY has run-time that is polynomial in |X|.

To see that algorithm FINDCHERRY is correct, first note that it will clearly terminate. Now, suppose that the algorithm outputs a (nonempty) set S. Then, in view of line 7, $\mathcal{T}|_S$ must satisfy (C2) and (C3). Moreover, in view of the while loop (lines 6–10) $\mathcal{T}|_S$ must satisfy (C4), So S must be a cherry of \mathcal{T} . Moreover, if the output indicates that S is isolated (i.e. that $S \neq X$ and that $\mathcal{T}|_S$ satisfies (C5)), then this must be the case in view of line 8. Now, suppose that algorithm FINDCHERRY outputs "No cherry of \mathcal{T} exists", and that, for the purposes of contradiction, a cherry S of \mathcal{T} does exist. Then, as any cherry has cardinality at least 2, if a cherry exists then at some stage the while loop in lines 2–12 must encounter some $\{x, y\} \in \binom{X}{2}$ with $\{x, y\} \subseteq S$. Clearly, the algorithm will then have to output S, a contradiction. Thus the algorithm FINDCHERRY is correct.

FINDCHERRY (X, \mathcal{T})

Inpu Out	it: A put: A if e	a set X , $ X \ge 3$, and a dense set \mathcal{T} of trinets on X . A cherry S of \mathcal{T} , and a boolean variable $I \in \{T, F\}$, with $I = T$ S is isolated and $I = F$ else, or the statement "No cherry of \mathcal{T} xists".
1.	Let $S =$	$\emptyset, I = F, G = {X \choose 2}.$
2.	While the	here is some $\{x, y\} \in G$ do
3.	If T_1	$(x, y, z), T_2(x, y, z), N_3(z, x, y), N_4(x, y, z), N_9(x, y, z)$
4.	or N	$T_{10}(z, x, y)$ is contained in \mathcal{T} for all $z \in X - \{x, y\}$ then do
5.	Let ,	$S = \{x, y\}, G = \emptyset \text{ and } U = X - \{x, y\}.$
6.	Whi	e there is some $u \in U$ do
7.	It	$\mathcal{T} _{S \cup \{u\}}$ satisfies (C2) and (C3), then let $S = S \cup \{u\}$.
8.	It	$U = \{u\}, S \neq X$, and S satisfies (C5), then let $I = T$.
9.	I	$et U = U - \{u\}.$
10.	end	"do (line 6)"
11.	else	let $G = G - \{\{x, y\}\}.$
12.	end "do	(line 2)"
13.	If $S = \emptyset$	then output "No cherry of \mathcal{T} exists" else output S and I.

FIGURE 8. Pseudo-code for an algorithm that either finds a cherry of a dense trinet set \mathcal{T} and also checks whether it is isolated or not or determines that no cherry of \mathcal{T} exists.

Now, given a dense trinet set \mathcal{T} on X, we say that a tuple $H = (a_1, a_2, \ldots, a_p : b_1, b_2, \ldots, b_q : z)$ of distinct elements of $X, p \ge 1, q \ge 0$ is a *cactus of* \mathcal{T} (with support $S = A \cup B \cup \{z\}, A = \{a_1, \ldots, a_p\}$ and $B = \{b_1, \ldots, b_q\}$) if $\mathcal{T}|_S$ satisfies conditions (H2)–(H8) of Lemma 6.2 (note that $\mathcal{T}|_S$ necessarily satisfies (H1)). Moreover, such an H is *isolated* if $S \ne X$ and $\mathcal{T}|_S$ also satisfies condition (H9) of Lemma 6.2.

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Note that if $H = (a_1, a_2, \ldots, a_p : b_1, b_2, \ldots, b_q : z)$ is a cactus of \mathcal{T} , then the relation $\sim_{\mathcal{T}}$ defined on the set $Y = S - \{z\} = A \cup B$ by putting $y \sim_{\mathcal{T}} y'$ if and only if y = y' or $N_2(z, y, y')$ or $N_2(z, y', y) \in \mathcal{T}$, for all $y, y' \in Y$, is an equivalence relation on Y with (at most two) equivalence classes A, B. Moreover, the relation $<_{\mathcal{T}}$ defined on Y by $y <_{\mathcal{T}} y'$ if and only if $N_2(z, y, y') \in \mathcal{T}$, for all $y, y' \in Y$, is a strict partial order on Y, which restricts to a strict linear order on A and also on B.

Using these observations, we now show that the algorithm presented in Fig. 9 can be used to detect cactuses in a dense set of trinets in polynomial time.

Lemma 7.2. Given a dense set \mathcal{T} of trinets on X, $|X| \ge 3$, algorithm FINDCACTUS is correct and has run-time that is polynomial in |X|.

Proof. First note that the algorithm will clearly terminate. Moreover, if it does output a tuple then in view of lines 12 and 13 this must be a cactus of \mathcal{T} and it will be isolated only if I = T. In addition, if the algorithm outputs "No cactus of \mathcal{T} exists", then this must be the case. Otherwise, suppose there is some cactus $K = (a_1, \ldots, a_p : b_1, \ldots, b_q : z)$ of $\mathcal{T}, p \ge 1, q \ge 0$. Setting $A = \{a_1, ..., a_p\}$ and $B = \{b_1, ..., b_q\}$ it follows that $S = A \cup B \cup \{z\}$ is the support of K and that z must be at the bottom of some trinet in \mathcal{T} . Thus the while loop (lines 2–20) would eventually find z at line 3. Since K is a cactus of \mathcal{T} , for each element $y \in Y := A \cup B$, there exists some $N \in \mathcal{T}$ such that y hangs off the side of N and z is at the bottom of N. Moreover, A and B(in case $B \neq \emptyset$) are the equivalence classes of the relation $\sim_{\mathcal{T}}$ defined on Y and the elements in A and B (again in case $B \neq \emptyset$) are strictly linearly ordered by $<_{\mathcal{T}}$. Thus, the algorithm would form the tuple $F = (a_1, \ldots, a_p : b_1, \ldots, b_q : z)$ (lines 10 and 11). Clearly, the support of F is S. Since $\mathcal{T}|_S$ satisfies (H2)–(H8) it follows that F is returned by the algorithm. However since F = K, this is impossible.

Finally, to see that algorithm FINDCACTUS is polynomial in |X|, it is sufficient to note that lines 6–7, 8–9 and 12–13 can all clearly be executed in time that is polynomial in |X|.

We now use the algorithms FINDCHERRY and FINDCACTUS to show that it can be decided in polynomial time whether or not a dense set of trinets is displayed by a 1-nested network using the algorithm presented in Fig. 10.

Theorem 7.3. For X with $|X| \ge 3$ and \mathcal{T} a dense set of trinets on X, algorithm BUILDNET has run-time that is polynomial in |X| and is correct.

FINDCACTUS (X, \mathcal{T})

Inpu Out	at: A set $X, X \ge 3$, and a dense set \mathcal{T} of trinets on X . put: A cactus H of \mathcal{T} and a boolean variable $I \in \{T, F\}$, with $I = T$ if H is isolated and $I = F$ else, or the statement "No cactus of \mathcal{T} exists".
1.	Put $H = \emptyset$, $I = F$, $G = X$.
2.	While there is some $z \in G$ do
3.	If there is a trinet $N \in \mathcal{T}$ such that z is at the bottom of N, then do
4.	Let Y be the set of $y \in X - \{z\}$ such that y hangs off the side of
5.	some $N \in \mathcal{T}$ for which z is at the bottom of N.
6.	If the relation $\sim_{\mathcal{T}}$ is an equivalence relation on Y
7.	that has at most two equivalence classes E, E' , then do
8.	If the relation $<_{\mathcal{T}}$ on Y is a partial order on Y that also restricts
9.	to give a strict linear order on E and on E' then do
10.	Let $F = (a_1,, a_p : b_1,, b_q : z)$ and $S = Y \cup \{z\}$, where
11.	$E = \{a_1, \ldots, a_p\}$ and $E' = \{b_1, \ldots, b_q\}$ are ordered relative to $<_{\mathcal{T}}$.
12.	If $\mathcal{T} _S$ satisfies (H2)–(H8), then let $H = F$ and $G = \emptyset$ and, if
13.	$\mathcal{T} _S$ also satisfies (H9) then let $I = T$, else let $G = G - \{z\}$.
14.	end "do (line 11)"
15.	else let $G = G - \{z\}.$
16.	end "do (line 8)"
17.	else let $G = G - \{z\}.$
18.	end "do (line 3)"
19.	else let $G = G - \{z\}.$
20.	end "do (line 2)"
21.	If $H = \emptyset$ then output "No cactus of \mathcal{T} exists" else output H and I .

FIGURE 9. Pseudo-code for an algorithm that either finds a cactus of a dense trinet set \mathcal{T} and also decides whether it is isolated or not or determines that no cactus of \mathcal{T} exists.

Proof. Algorithm BUILDNET has run-time that is polynomial in |X| since the check required in line 2 can be executed in time that is polynomial in |X| by Lemmas 7.1 and 7.2

Now, if algorithm BUILDNET outputs "There is no 1-nested network displaying \mathcal{T} ", then by Proposition 5.1, Lemma 6.1 and Lemma 6.2,

BUILDNET(\mathcal{T})

Inpu Outj	t: A set $X, X \ge 3$, and a dense set \mathcal{T} of trinets on X . out: A 1-nested network N on X with $Tr(N) = \mathcal{T}$, or the statement "There is no 1-nested network displaying \mathcal{T} ".
1.	$Stack = \emptyset, G = X$
2.	While there is some cherry S in \mathcal{T} with $z \in S$ or some cactus
3.	$H = (a_1, \dots, a_p : b_1, \dots, b_q : z)$ with support $S = \{a_1, \dots, a_p, b_1, \dots, b_q, z\}$
4.	in \mathcal{T} do
5.	Put the symbol $R_{z:S}$ on the top of <i>Stack</i> .
6.	If $ G - (S - \{z\}) \le 2$, then let N be either the bush on G or
7.	the two-leafed network on G , depending on \mathcal{T} .
8.	Let $\mathcal{T} = \mathcal{T} - \mathcal{T}_{z:S}, G = G - (S - \{z\}).$
9.	end "do (line 2)"
10.	If $ G \geq 3$, then output "There is no 1-nested network displaying \mathcal{T} "
11.	else do
12.	While there is some $R_{z:S}$ on the top of $Stack$, do $N = R_{z:S}^{-1}(N)$.
13.	Output N
13.	end "do (line 12)"

FIGURE 10. Pseudo-code for an algorithm to construct a 1-nested network from a dense set of trinets, or decide that such a network does not exist.

there is no 1-nested network N on X with $Tr(N) = \mathcal{T}$. Moreover, if BUILDNET outputs a network N, then N is clearly 1-nested, and $Tr(N) = \mathcal{T}$ by (1). This completes the proof.

Remark 7.4. Although we have shown that algorithm BUILDNET has run-time that is polynomial in |X|, it could be of interest to see if faster, more sophisticated algorithms can be developed.

8. DISCUSSION

In this paper, we have shown that we can recover a 1-nested network from 'perfect data', viz. the dense set of 1-nested trinets that is displayed by the network. In practice, we will not usually have access to such information for biological datasets. Even so, it should be quite straight-forward to at least compute a dense set of trinets for any given biological dataset using existing phylogenetic network methods.



FIGURE 11. The 1-nested network N on $\{w, x, y, z\}$ depicted in (a) is uniquely determined by the two trinets pictured in (b). As before, directions are omitted for clarity when clear. Also only the vertices that are leaves are marked by a dot.

For example, given a multiple sequence alignment, one could compute the most parsimonious or most likely trinet for every sub-alignment of 3 sequences (using, e.g. methods described in [18, 19]), which would be feasible as there are a bounded number of 1-nested trinets. Note that this would have the advantage that no 'breakpoints' would need to be computed for the multiple alignment, which is a first (and sometimes quite difficult) step that is usually required when constructing phylogenetic networks from phylogenetic trees (cf. e.g. [12, Chapter 11], [23, Section 2]).

Given that computing dense sets of trinets is feasible for biological data, it could be reasonable to develop methods for finding 1-nested networks displaying as many trinets as possible from a dense set of trinets. Similar techniques have been developed for triplets e.g. [11, 13, 31, although it is worth noting that it is NP-hard to find a tree displaying a maximum number of rooted triplets from an arbitrary set of triplets [2, 15, 33] (even if the set is dense [3]). Alternatively, it might be of interest to investigate if there might be an 'Aho-type' algorithm [1] to determine if an arbitrary subset of 1-nested trinets encodes a 1nested network, and, if so, adapt this to give 'Min-Cut' type algorithms for building 1-nested networks from sets of trinets (cf. [24, 26, 27]). A first step in this direction could be to determine whether or not it is an NP-complete problem to decide if an arbitrary subset of 1-nested trinets encodes a 1-nested network (in particular, note that there are non-dense sets of 1-nested trinets that encode 1-nested networks – e.g. the 1-nested network N on $\{w, x, y, z\}$ pictured in Fig. 11(a) is the only 1-nested network on $\{w, x, y, z\}$ displaying the two trinets presented in Fig. 11(b)).

In another direction, clearly we can ask for results along the lines of those presented above for *level-k networks* [9], $k \geq 2$, phylogenetic networks that have a bounded level of complexity depending on k



FIGURE 12. A level-*n* phylogenetic network *N* on $\{x_1, x_2, \ldots, x_n\}, n \ge 4$, for which every trinet in Tr(N) is of level-3. For clarity arc directions are omitted when clear.

(and also, of course, 'k-nested' networks). Note that there are nonrecoverable level-2 networks (e.g. Fig. 4), and so this could be rather more technical. Moreover, it should be noted that, for $k \ge 3$, there are level-k networks that are not of level-(k - 1) all of whose trinets have fixed level (see Fig. 12). Thus, the levels of the trinets displayed by a network do not necessarily determine the level of a network. For practical purposes, it might also be of interest to determine a way to enumerate the level-k trinets, $k \ge 2$.

Another avenue worth exploring, could be to try generalizing the above results to 'r-nets', $r \ge 4$, i.e. phylogenetic networks with r-leaves (note that in case r = 4 quartet trees are commonly used to build phylogenetic trees, e.g. [29]). Note that it is straight-forward to extend Definition 3.1 to obtain a set of r-nets displayed by a phylogenetic network. This could be quite useful in practice since it might be possible to obtain more accurate estimates for r-nets than trinets (at least for r = 4) before we try to piece them together, although, technically speaking, this could be very challenging.

Finally, we conclude with what we consider to be a rather bold conjecture:

Conjecture 8.1. If N is a recoverable phylogenetic network on X, then Tr(N) encodes N, that is, if N' a recoverable phylogenetic network on X such that Tr(N) = Tr(N') then N is isomorphic to N'.

A first (and probably quite instructive!) 'exercise' could be to try and show that this conjecture at least holds for level-2 networks. Note that if this conjecture were true, then as in Corollary 6.4, we would immediately obtain a new proper metric on the set of recoverable phylogenetic networks on X. Acknowledgements VM thanks the Royal Society for supporting his visit to New Zealand. Both authors thank Mike Hendy, David Penny, Charles Semple, Peter Stadler and Mike Steel for hosting them during their sabbatical, during which this work was conceived and undertaken.

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