# OPTIMIZING PHYLOGENETIC DIVERSITY ACROSS TWO TREES 

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#### Abstract

We present a polynomial-time algorithm for finding an optimal set of taxa that maximizes the weighted-sum of the phylogenetic diversity across two phylogenetic trees. This resolves one of the challenges proposed as part of the Phylogenetics Programme held at the Isaac Newton Institute for Mathematical Sciences (Cambridge, 2007). It also completely closes the gap between optimizing phylogenetic diversity on one tree, which is known to be in P , and optimizing phylogenetic diversity across three or more trees, which is known to be NP-hard.


## 1. Introduction

A central task in conservation biology is measuring, predicting, and preserving biological diversity as species face extinction. Dating back to Faith (1992) [2], phylogenetic diversity (PD) is a prominent tool for measuring the biodiversity of a subset of species. This measure is based on the evolutionary distance amongst the species in the subset on an underlying phylogenetic (evolutionary) tree. For a fixed integer $k$, there are polynomial-time algorithms for finding an optimal $k$-element subset of species that maximizes the PD score across one tree. However, in practice, the underlying phylogenetic tree of the species under consideration is typically unknown, or there is no 'true tree' relating the species because of evolutionary events such as recombination. Thus one usually obtains two or more different trees for the same set of species, each arising from the analysis of a different gene or section of genome, or simply from analyses that use different models of evolution. Therefore, we would ideally like to optimize PD across a (weighted) set of phylogenetic trees. It has been previously stated that across three or more trees this problem is NPhard. In this paper, we show that the problem of finding an optimal subset of species that maximizes the PD score across two trees can be solved in polynomial time.

A phylogenetic $X$-tree $T=(V, E)$ is an (unrooted) tree with no degree- 2 vertices and whose leaf set $X$ represents a set of species. Suppose the edges of $T$ have non-negative real-valued lengths $\omega: E \rightarrow \mathbb{R}^{\geq 0}$. The phylogenetic diversity (PD

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Figure 1. (a): Two phylogenetic trees $T_{1}$ and $T_{2}$ on the same set of species $\left\{x_{1}, \ldots, x_{8}\right\}$. (b): The network $D$ constructed from $T_{1}$ and $T_{2}$. All the arcs in $D$ are directed downwards. Each solid arc has the indicated cost and capacity 1 . The dotted arcs have cost 0 and capacity $k-2$.
score) of a subset $Y$ of $X$ is the sum of the edge lengths of the minimal subtree of $T$ connecting the elements in $Y$. Referring to Figure 1(a), if $Y=\left\{x_{1}, x_{2}, x_{4}\right\}$, then $P D_{T_{1}}(Y)=21$ and $P D_{T_{2}}(Y)=14$.

For one tree, the PD optimization problem is to find a subset of $X$ of a given size $k$ that maximizes the PD score amongst all subsets of $X$ of size $k$. Extending this problem to an arbitrary number of trees, we have the following family of optimization problems:

Problem: Weighted Average PD on $t$ trees $\left(\mathrm{WAPD}_{t}\right)$
Instance: A collection $\mathcal{T}=\left\{T_{1}, \ldots, T_{t}\right\}$ of phylogenetic $X$-trees whose edges have non-negative real-valued lengths, a collection $\left\{\lambda_{1}, \ldots, \lambda_{t}\right\}$ of non-negative realvalued weights, and an integer $k$.
Question: Find a subset $Y$ of $X$ of size $k$ that maximizes

$$
P D_{\mathcal{T}}(Y)=\lambda_{1} P D_{T_{1}}(Y)+\cdots+\lambda_{t} P D_{T_{t}}(Y)
$$

amongst all $k$-element subsets of $X$.
The value $P D_{\mathcal{T}}(Y)$ is the phylogenetic diversity (PD score) of $Y$ across the trees in $\mathcal{T}$. To allow a weighting scheme on the individual trees, we have additionally included the weights $\lambda_{1}, \ldots, \lambda_{t}$. Of course, by multiplying all edge lengths of $T_{i}$ by $\lambda_{i}$ to obtain $T_{i}^{\prime}$ for all $i$, the PD score of $Y$ across $\mathcal{T}$ is $P D_{T_{1}^{\prime}}(Y)+\cdots+P D_{T_{t}^{\prime}}(Y)$. Thus for computational purposes, no generality is lost by assuming that $\lambda_{i}=1$ for all $i$. We make this assumption in the rest of the paper.

The problem $\mathrm{WAPD}_{1}$ can be solved by a greedy approach $[8,5]$ in polynomial time. An implementation of this approach with run time $O(n \log k)$ is available [3], where $n=|X|$. Furthermore, it is even possible to solve $\mathrm{WAPD}_{1}$ in $O(n)$ time [7].

The problem $\mathrm{WAPD}_{t}$ for $t \geq 2$ appears to have first been raised in [3]. For $t=3$, Spillner et al. [7] noted without proof that $\mathrm{WAPD}_{3}$, and therefore $\mathrm{WAPD}_{t}$ for all $t \geq 3$, is NP-hard using a reduction from 3 -dimensional matching. This left open the problem of determining the computational complexity of $\mathrm{WAPD}_{2}$, explicitly stated in [7] and subsequently asked again in [9] where a prize was offered for resolving the problem. In this paper, we show that $\mathrm{WAPD}_{2}$ can be solved by a polynomial-time algorithm by reformulating the problem as a set of minimum-cost flow problems. Furthermore, for completeness, we explicitly show that $\mathrm{WAPD}_{3}$ is NP-hard using a reduction from vertex cover on cubic graphs.

## 2. A Polynomial-Time Algorithm for $\mathrm{WAPD}_{2}$

In this section, we show that $\mathrm{WAPD}_{2}$ is solvable in polynomial time. To do this, we initially show that a restricted version of $\mathrm{WAPD}_{2}$ is solvable in polynomial time. In this restriction, we are additionally given a distinguished element, $x$ say, of $X$ in the instance and are asked to find a subset of $Y$ of $X$ of size $k$ that contains $x$ and maximizes

$$
P D_{\mathcal{T}}(Y)=P D_{T_{1}}(Y)+\cdots+P D_{T_{t}}(Y)
$$

It will then immediately follow that $\mathrm{WAPD}_{2}$ is solvable in polynomial time.
To show that this restricted version of $\mathrm{WAPD}_{2}$ is solvable in polynomial time, we reformulate it into a network flow problem. Let $X=\left\{x_{1}, x_{2}, \ldots, x_{n}\right\}$, and suppose that the edges of $T_{1}$ and $T_{2}$ are assigned non-negative real-valued lengths $\omega$ : $E\left(T_{1}\right) \cup E\left(T_{2}\right) \rightarrow \mathbb{R}^{\geq 0}$. Without loss of generality, choose $x_{1}$ to be the distinguished element of $X$. For the purposes of the reformulation, we distinguish between the vertices of $T_{1}$ and $T_{2}$ that share a common label in $X$ by relabelling $x_{i}$ with $x_{i 1}$ in $T_{1}$ and relabelling $x_{i}$ with $x_{i 2}$ in $T_{2}$ for all $i \in\{1,2, \ldots, n\}$. Furthermore, we view the edges of $T_{1}$ (resp. $T_{2}$ ) as arcs directed away from $x_{11}$ (resp. $x_{12}$ ).

Now we construct a network $D$ from $T_{1}$ and $T_{2}$, where the source and $\operatorname{sink}$ of $D$ will be $x_{11}$ and $x_{12}$, respectively. The vertex set $V$ of $D$ is $V\left(T_{1}\right) \cup V\left(T_{2}\right)$ and the arc set $A$ of $D$ is constructed in the following way:
(i) For each $\operatorname{arc}(u, v)$ in $T_{1}$, add two $\operatorname{arcs}(u, v)_{1}$ and $(u, v)_{2}$ in parallel directed from $u$ to $v$ with $(u, v)_{1}$ having capacity 1 and $\operatorname{cost}-\omega(u, v)$, and $(u, v)_{2}$ having capacity $k-2$ and cost 0 .
(ii) For each $\operatorname{arc}(u, v) \in T_{2}$, add two $\operatorname{arcs}(v, u)_{1}$ and $(v, u)_{2}$ in parallel directed from $v$ to $u$ with $(v, u)_{1}$ having capacity 1 and $\operatorname{cost}-\omega(u, v)$, and $(v, u)_{2}$ having capacity $k-2$ and cost 0 .
(iii) For each $i \in\{2,3, \ldots, n\}$, add the $\operatorname{arc}\left(x_{i 1}, x_{i 2}\right)$ with capacity 1 and cost 0 .

To illustrate this construction, consider the two phylogenetic trees $T_{1}$ and $T_{2}$ in Fig. 1(a). The flow network for these two trees with $x_{11}$ and $x_{12}$ as the source and sink, respectively, is shown in Fig. 1(b).

Noting that the network resulting from the above construction $D$ has a feasible flow of $k-1$ units, the following lemma states the key property of $D$.

Lemma 2.1. Let $f$ be an integer-valued minimum-cost flow of $k-1$ units from $x_{11}$ to $x_{12}$ in $D$. Then

$$
Y_{f}=\left\{x_{1}\right\} \cup\left\{x_{i} \in X-x_{1}: f\left(x_{i 1}, x_{i 2}\right)>0\right\}
$$

is an optimal solution to the restricted version of $\mathrm{WAPD}_{2}$ in which $x_{1}$ is the distinguished element of $X$.

Proof. Let $Y_{\text {opt }}$ be an optimal solution for the restricted version of $\mathrm{WAPD}_{2}$ in which $x_{1}$ is the distinguished element of $X$. The goal is to show that

$$
P D_{T_{1}}\left(Y_{f}\right)+P D_{T_{2}}\left(Y_{f}\right)=P D_{T_{1}}\left(Y_{\mathrm{opt}}\right)+P D_{T_{2}}\left(Y_{\mathrm{opt}}\right) .
$$

For each arc $a=(u, v)$ of $T_{1}$ (resp. $T_{2}$ ), let $l_{a}$ denote the number of leaves in $Y_{\text {opt }}$ that can be reached by a directed path in $T_{1}\left(\right.$ resp. $\left.T_{2}\right)$ starting at $v$. Let $f_{Y_{\text {opt }}}$ be the flow of $k-1$ units on $D$ that is defined as follows:
(i) For each arc $a=(u, v) \in T_{1}$, set

$$
f_{Y_{\text {opt }}}\left((u, v)_{1}\right)=\min \left\{1, l_{a}\right\} \text { and } f_{Y_{\text {opt }}}\left((u, v)_{2}\right)=\max \left\{0, l_{a}-1\right\} .
$$

(ii) For each arc $a=(u, v) \in T_{2}$, set

$$
f_{Y_{\text {opt }}}\left((v, u)_{1}\right)=\min \left\{1, l_{a}\right\} \text { and } f_{Y_{\text {opt }}}\left((v, u)_{2}\right)=\max \left\{0, l_{a}-1\right\} .
$$

(iii) For each $x_{i} \in X-x_{1}$, set $f_{\text {opt }}\left(x_{i 1}, x_{i 2}\right)=1$ if $x_{i} \in Y_{\mathrm{opt}}$; otherwise set $f_{Y_{\text {opt }}}\left(x_{i 1}, x_{i 2}\right)=0$.

It is easily checked that $f_{Y_{\text {opt }}}$ is indeed a flow of $k-1$ units on $D$ and, moreover, the cost of this flow is $\operatorname{cost}\left(f_{Y_{\text {opt }}}\right)=-\left(P D_{T_{1}}\left(Y_{\mathrm{opt}}\right)+P D_{T_{2}}\left(Y_{\mathrm{opt}}\right)\right)$.

We next show that the cost of $f$ is

$$
\begin{equation*}
\operatorname{cost}(f)=-\left(P D_{T_{1}}\left(Y_{f}\right)+P D_{T_{2}}\left(Y_{f}\right)\right) . \tag{1}
\end{equation*}
$$

To establish (1), it suffices to show that the set of arcs used by $f$ of the form $(u, v)_{1}$ (derived from $T_{1}$ ) or the form $(v, u)_{1}$ (derived from $T_{2}$ ) are precisely the union of the arcs in the minimal subtree of $T_{1}$ connecting the elements in $Y_{f}$ and the minimal subtree of $T_{2}$ connecting the elements in $Y_{f}$. Note that if $f$ uses an arc from $u$ to $v$ and both arcs from $u$ to $v$ have cost 0 (i.e. a tree edge has length 0 ), we may assume that $f$ uses the arc of the form $(u, v)_{1}$. If an arc of $D$ of the form $(u, v)_{1}$ (derived from $T_{1}$ ) is used by $f$, then it is clear that there exists an element in $Y_{f}-x_{1}$ in the subtree of $T_{1}$ below $(u, v)$. Thus, as $x_{1} \in Y_{f}$, the arc $(u, v)$ is in the minimal subtree of $T_{1}$ connecting the elements in $Y_{f}$. Similarly, if an arc of $D$ of the form $(v, u)_{1}$ (derived from $T_{2}$ ) is used by $f$, then $(u, v)$ is in the minimal subtree of $T_{2}$ connecting the elements in $Y_{f}$. Now assume that $(u, v)$ is in the minimal subtree of $T_{1}$ connecting the elements in $Y_{f}$. Then $(u, v)$ is on the directed path from $x_{1}$ to an element, $x_{j}$ say, in $Y_{f}$. Since $x_{j}$ is an element of $Y_{f}$ and since, up to parallel edges, there is a unique directed path from $x_{11}$ to $x_{1 j}$ in $D$, the minimum-cost flow $f$ must use $(u, v)_{1}$. An analogous argument also holds in the case that $(u, v)$ is in
the minimal subtree of $T_{2}$ connecting the elements in $Y_{f}$. This establishes (1). It now follows that

$$
\begin{aligned}
\operatorname{cost}\left(f_{Y_{\mathrm{opt}}}\right) & =-\left(P D_{T_{1}}\left(Y_{\mathrm{opt}}\right)+P D_{T_{2}}\left(Y_{\mathrm{opt}}\right)\right) \\
& \leq-\left(P D_{T_{1}}\left(Y_{f}\right)+P D_{T_{2}}\left(Y_{f}\right)\right)=\operatorname{cost}(f)
\end{aligned}
$$

Hence, as $f$ and $f_{Y_{\text {opt }}}$ are both flows of $k-1$ units on $D$, but $f$ has minimum cost, we have

$$
P D_{T_{1}}\left(Y_{\mathrm{opt}}\right)+P D_{T_{2}}\left(Y_{\mathrm{opt}}\right)=P D_{T_{1}}\left(Y_{f}\right)+P D_{T_{2}}\left(Y_{f}\right)
$$

This completes the proof of the lemma.
Lemma 2.2. The restricted version of $W A P D_{2}$ can be solved in $O\left(n^{2} \log ^{2} n\right)$ time, where $n=|X|$.

Proof. The construction above can certainly be done within time $O\left(n^{2} \log ^{2} n\right)$. Finding a minimum-cost flow in such a network is an old and well-studied problem, and can be solved using an algorithm that runs in time $O((|A| \log |V|)(|A|+$ $|V| \log |V|)$ ) (see, for example, [1]). Since $D$ has $O(n)$ vertices and arcs this yields a run time of $O\left(n^{2} \log ^{2} n\right)$. Furthermore, as all the capacities in $D$ and the target flow of $k-1$ units are integral, there is an integral minimum-cost flow, and this is found by the above algorithm [1].
Theorem 2.3. The problem $\mathrm{WAPD}_{2}$ (without restriction) can be solved in $O\left(n^{3} \log ^{2} n\right)$ time, where $n=|X|$.

Proof. Suppose $Y_{\text {opt }}$ is an optimal solution to $\mathrm{WAPD}_{2}$ and let $x$ be an element of $Y_{\mathrm{opt}}$. Then $Y_{\mathrm{opt}}$ is an optimal solution to the restricted version of $\mathrm{WAPD}_{2}$ in which $x$ is the distinguished element of $X$. Consequently, by solving the restricted version of $\mathrm{WAPD}_{2}$ for each element of $X$-thus running the method described above $n$ times - and choosing the solution that maximizes $P D_{T_{1}}(Y)+P D_{T_{2}}(Y)$, we obtain an optimal solution to $\mathrm{WAPD}_{2}$. The theorem now follows from Lemma 2.2

## 3. NP-Hardness of $\mathrm{WAPD}_{3}$

In this section, we explicitly show that $\mathrm{WAPD}_{3}$ is NP-hard. The reduction is from a restricted version of the following classic NP-complete problem:

## Problem: VertexCover

Instance: A graph $G=(V, E)$ and an integer $k$.
Question: Does there exist a subset $C \subseteq V$ such that $|C|=k$ and, for every edge $\{u, v\} \in E$, the intersection $\{u, v\} \cap C$ is non-empty.

VertexCover remains NP-complete even if the input graph is restricted to be a 3 -connected cubic planar graph [11]. The reduction proceeds as follows. Take a 3 -connected cubic planar instance $G$. Colour the edges of $G$ with three colours $\{1,2,3\}$ such that no two adjacent edges receive the same colour. Due to a classic construction of Tait [10], this is equivalent to four-colouring the faces of a planar drawing of $G$ which can be done in quadratic time [6]. For each colour $c \in\{1,2,3\}$, let $T_{c}$ be the phylogenetic $V$-tree that consists of a (central) vertex $z_{c}$ of degree
$|V| / 2$, where the $|V| / 2$ neighbours of $z_{c}$ each have degree 3 and the $|V|$ leaves are arranged so that, for each edge $\{u, v\}$ of $G$ coloured $c$, the vertices $u$ and $v$ are adjacent to the same degree-3 vertex. As $G$ is a cubic graph, $T_{c}$ is well-defined for all $c$.

For each of $T_{1}, T_{2}$, and $T_{3}$, assign length 1 to all edges. It now follows that the PD score of an optimal solution to $\mathrm{WAPD}_{3}$ across $T_{1}, T_{2}$, and $T_{3}$ is equal to $|E|+3 k$ if and only if $G$ has a vertex cover of size $k$, where $k \geq 3$. To see this, consider a vertex cover $C$ of size $k$. Each element in $C$ appears as a leaf in each tree, so the edges incident with these elements contribute $3 k$ to the overall PD score. Since $C$ covers every edge of $G$, each edge of $G$ corresponds to a unique edge incident with one of $z_{1}, z_{2}, z_{3}$, and $k \geq 3$, these latter edges contribute exactly $|E|$ to the overall PD score. Thus the PD score of $C$ across $T_{1}, T_{2}$, and $T_{3}$ is $|E|+3 k$. The converse is similar.

We remark here that, in the above reduction, $T_{1}, T_{2}$, and $T_{3}$ could have been made binary (that is, each internal node has degree 3) if desired by refining $z_{1}$, $z_{2}$, and $z_{3}$ and assigning length $\epsilon$ to each new edge such that $\epsilon$ is smaller than $3 /(|V| / 2)$.

## 4. Concluding Remarks

We end the paper with two remarks. First, the problem $\mathrm{WAPD}_{1}$ can be considered as a special case of $\mathrm{WAPD}_{2}$ when one of the initial trees is degenerated to a tree with a single internal vertex. Hence, our algorithm for solving $\mathrm{WAPD}_{2}$ also applies to $\mathrm{WAPD}_{1}$. However, the greedy approach for the latter problem mentioned in the introduction, yields a much better asymptotic run time. Note that the greedy approach used to solve $\mathrm{WAPD}_{1}$ can fail to produce optimal solutions for $\mathrm{WAPD}_{2}$ - this follows from the discussion in [4, Section 7].

Second, a rooted phylogenetic $X$-tree $T$ is a rooted tree with leaf set $X$ and whose root has degree at least 2 and all other internal vertices have degree at least 3. Assuming that the edges of $T$ are assigned non-negative real-valued lengths, the PD score of a subset $Y$ of $X$ on $T$ is the sum of the lengths of the edges in the minimal subtree of $T$ connecting the elements in $Y \cup \rho$, where $\rho$ is the root of $T$. The family of problems $\mathrm{WAPD}_{t}$ can be interpreted for rooted phylogenetic trees in the obvious way. Indeed, for $t=1, t=2$, and for all $t \geq 3$, the analogous unrooted computational results hold. In particular, $\mathrm{WAPD}_{2}$ for rooted phylogenetic trees can be solved in polynomial time. It can be directly interpreted as the restricted version of $\mathrm{WAPD}_{2}$ described in Section 2, where the distinguished element is $\rho$, the common root label of the two rooted trees.

## References

[1] R. Ahuja, T. Magnanti, and J. Orlin. Network Flows: Theory, Algorithms, and Applications. Prentice Hall, London, 1993.
[2] D. P. Faith. Conservation evaluation and phylogenetic diversity. Biological Conservation, 61:1-10, 1992.
[3] B. Q. Minh, S. Klaere, and A. von Haeseler. Phylogenetic diversity within seconds. Systematic Biology, 55:769-773, 2006.
[4] V. Moulton, C. Semple, and M. Steel. Optimizing phylogenetic diversity under constraints. Journal of Theoretical Biology, 246:186-194, 2007.
[5] F. Pardi and N. Goldman. Species choice for comparative genomics: being greedy works. PLoS Genetics, 1(6), 2005.
[6] N. Robertson, D. Sanders, P. Seymour, and R. Thomas. A new proof of the four-colour theorem. Electron. Res. Announc. Amer. Math. Soc., 2:17-25, 1996.
[7] A. Spillner, B. T. Nguyen, and V. Moulton. Computing phylogenetic diversity for split systems, 2007. submitted.
[8] M. Steel. Phylogenetic diversity and the greedy algorithm. Systematic Biology, 54:527-529, 2005.
[9] M. Steel. Phylognetics: Challenges and conjectures. In Newton Institute Programme on Phylogenetics, 2007.
[10] P.G. Tait. On the colouring of maps. In Proceedings of the Royal Society of Edinburgh, Section $A$, volume 10, pages 501-503, 1878-1880.
[11] R. Uehara. NP-complete problems on a 3-connected cubic planar graph and their applications. Technical Report TWCU-M-0004, Tokyo Woman's Christian University, 1996.

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