

VERSION 2

OPEN BACCESS

DOI:

dx.doi.org/10.17504/protocol s.io.rm7vzbryrvx1/v2

Protocol Citation: Vidya S Vuruputoor, karlfetter, victoriaburton, Jill Wegrzyn 2023. Collecting needle and branch samples for terpenoid and expression analysis.

protocols.io

https://dx.doi.org/10.17504/p rotocols.io.rm7vzbryrvx1/v2V ersion created by Vidya S Vuruputoor

License: This is an open access protocol distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working We use this protocol and it's working

Created: May 25, 2023

Last Modified: Jun 30, 2023

PROTOCOL integer ID:

82480

Collecting needle and branch samples for terpenoid and expression analysis V.2

Vidya S Vuruputoor¹, karlfetter¹, victoriaburton¹, Jill Wegrzyn¹

¹UConn

PlantCompGenomics



Vidya S Vuruputoor

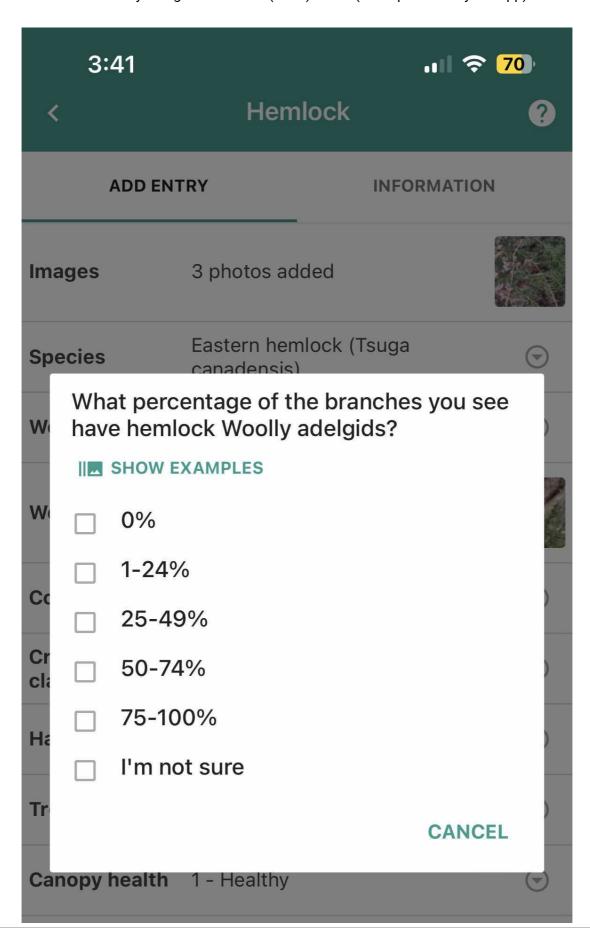
ABSTRACT

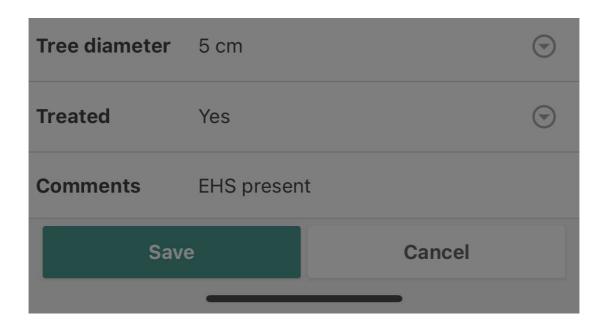
This protocol is used by the PCG group for sampling needle and branch tissue from hemlock conifers for terpenoid analysis, as well as RNA expression analysis.

IMAGE ATTRIBUTION

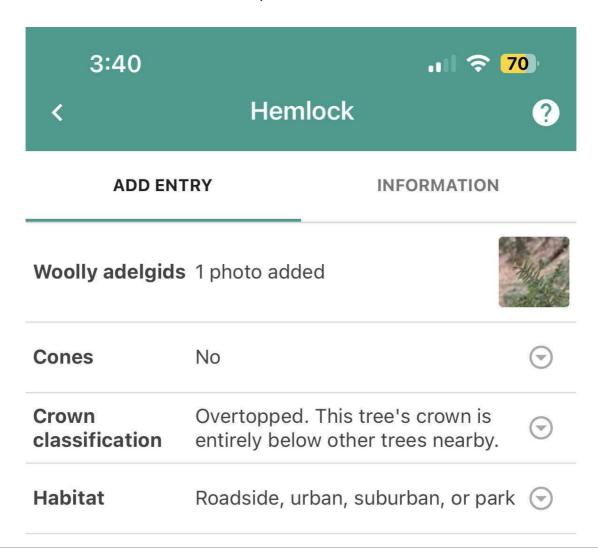
The image was taken by Dr. Karl Fetter

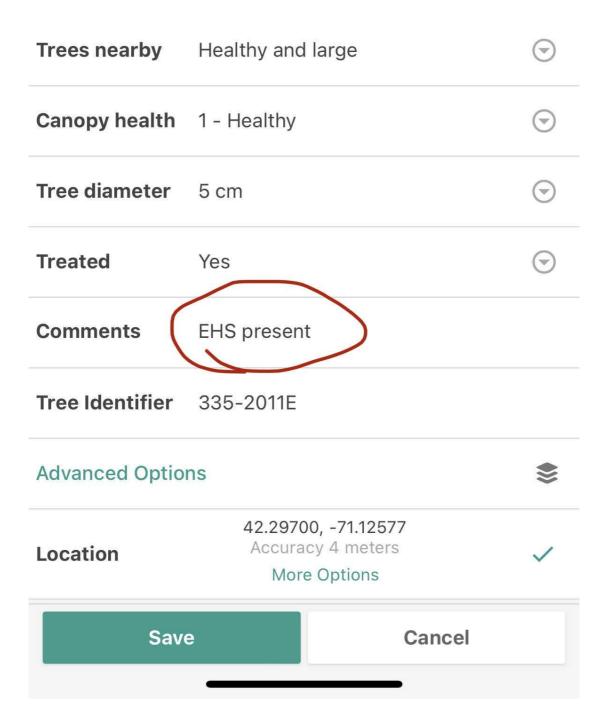
1 Open Treesnap (https://treesnap.org/), record the accession number on the tree and fill in notes for hemlock woolly adelgid infestation (HWA) levels (scale provided by the app).





1.1 Use the comment field to enter: *EHS present* OR *EHS absent* to record the hemlock scale.





- 1.2 Take at least 3 photos with TreeSnap. This includes at least one of the trees itself, one photo of the tree tag if that is present, and one photo of the collection vial with the label as well.
- **1.3** Geolocation will be recorded automatically but TreeSnap can also connect with a more precise external GPS device.

1.4	Note DBH in appropriate column
1.5	Note any other observation you may have- including plant condition (Excellent/Good/Poor), and placement of individual (lower canopy, full sun, etc.)
2	If gathering samples for pooled data, choose four random areas around the tree to collect needle and branch samples
	Note
	For RNA collections- please use gloves, and spray RNase before collections (separate vial)
3	Look for the second branch from the shoot tip (stick to the right first) and snip off 5 cm to place in 10 ml cryovial
4	Repeat for the other 3 directions (check appropriate column in the sheet provided). Place about 3-4 stems in each 10 ml cryovial.
5	Close cryovial, and using the appropriate gloves to handle liquid nitrogen (LN), place cryovial in LN dewar.
	Safety information
	Use appropriate wear while handling liquid nitrogen

5.1

Cap vial tightly to prevent opening in the LN.

5.2 Depending on dewer configuration, you can tie tubes with string to a 'halo' to help keep vials from falling to the bottom. Fully submerged tubes can open.

Note

You can hold upto 12 cryovials in a 6L dewer

6 Clean trimmers between trees with ethanol (spray bottle).

Note

For collection for RNA- spray clippers with RNase as well

- 7 Use clipboard to manually record notes on a pre-printed spreadsheet with sample names.
- **8** After returning from the field, ensure that notes are consistent with TreeSnap entries. These entries can be amended after collection as needed.