

LANDIS-II Biomass Harvest v1.0 Extension User Guide

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1 Introduction

This document describes the Biomass Harvesting extension for the LANDIS-II model. Users should read the *LANDIS-II Model User's Guide* prior to reading this document.

The Biomass Harvest Extension for LANDIS-II is derived from the Base Harvest extension and therefore generally behaves the same as Base Harvest. The largest change is that Biomass Harvest supports partial thinning of cohorts; other changes are listed below. Consult the user guide for Base Harvest for further information.

Because Biomass Harvest is derived from Base Harvest, **it is required that the most recent version of Base Harvest be installed prior to using Biomass Harvest.**

1.1 Acknowledgements

Funding for the development of LANDIS-II has been provided by the Northern Research Station (Rhinelander, Wisconsin) of the U.S. Forest Service.

1.2 Extension History

This is the first public release of this extension.

2 Input Files

The inputs for this extension are identical to those listed in the Base Harvest User Guide with the exceptions listed below. The text file must comply with the general format requirements described in section 3.1 *Text Input Files* in the *LANDIS-II Model User Guide*.

2.1 LandisData

This parameter's value must be "Biomass Harvest".

2.2 Species List for Cohort Removal

When `SpeciesList` is specified for the `CohortsRemoved` parameter in a prescription, the user may specify a percentage after any individual age or range of ages. The percentage indicates what proportion of a cohort(s) will be removed. For example:

```
CohortsRemoved  SpeciesList
acersacc        1-40 (50%)  50(65%)  65-70  71-107 ( 15% )  109
```

The default percentage is 100%. If no parenthetical biomass removal values are given, the Biomass Harvest extension will behave identically to the Base Harvest extension.

2.3 BiomassMaps

The new `BiomassMaps` parameter comes after the `PrescriptionMaps` parameter. It specifies the template for the names of output maps of biomass removed from harvested sites. Like the `PrescriptionMaps` parameter, the `BiomassMaps` parameter requires that the timestep variable be used in its value:

```
BiomassMaps  harvest/biomass-removed-{timestep}.gis
```

This parameter is optional. If it is not present, then no output maps are created.