# hatchR

Morgan Sparks<sup>1</sup> Bryan Maitland<sup>1</sup> Eli Felts<sup>2</sup> Allison Swartz<sup>3</sup> Paul Frater<sup>4</sup>

- <sup>1</sup> Rocky Mountain Research Station, US Forest Service
- <sup>2</sup> Idaho Fish and Wildlife Conservation Office, US Fish and Wildlife Service
- <sup>3</sup> College of Forestry, Oregon State University
- <sup>4</sup> Wisconsin Department of Natural Resources, Madison, WI, USA

## Introduction

**hatchR** is a software ecosystem for predicting fish developmental phenology using statistical models. It offers three primary applications:

- 1. Basic summarization, plotting, and data checks for water temperature data.
- 2. Access published parameterizations for salmonid developmental models or build your own custom parameterizations for any species with user provided data.
- 3. Predict hatch and/or emergence timing across ambient temperatures from common data sources such as HOBO loggers.

hatchR is deployed in two formats. First, it can be downloaded as a R package. Secondly, for users not familiar with R, it can also be accessed as a web app built in Shiny as a point-and-click tool.

To predict phenology, you need <u>daily average</u> <u>temperatures over incubation</u> (or data that can be <u>summarized as such</u>) and <u>spawn timing</u>.

# Using hatchR

### hatchR can be accessed two ways:

#### R package

hatchR can be downloaded from CRAN and used in the R programming language for full functionality. The hatchR website provides numerous Articles on basic to advanced use.



A toolset to predict when fish hatch and emerge in the wild



#### Shiny app

hatchR can be accessed in a point-and-click interface via Shiny. The app loads in your browser and provides much of the functionality of the R package but is less automative.



## Workflow

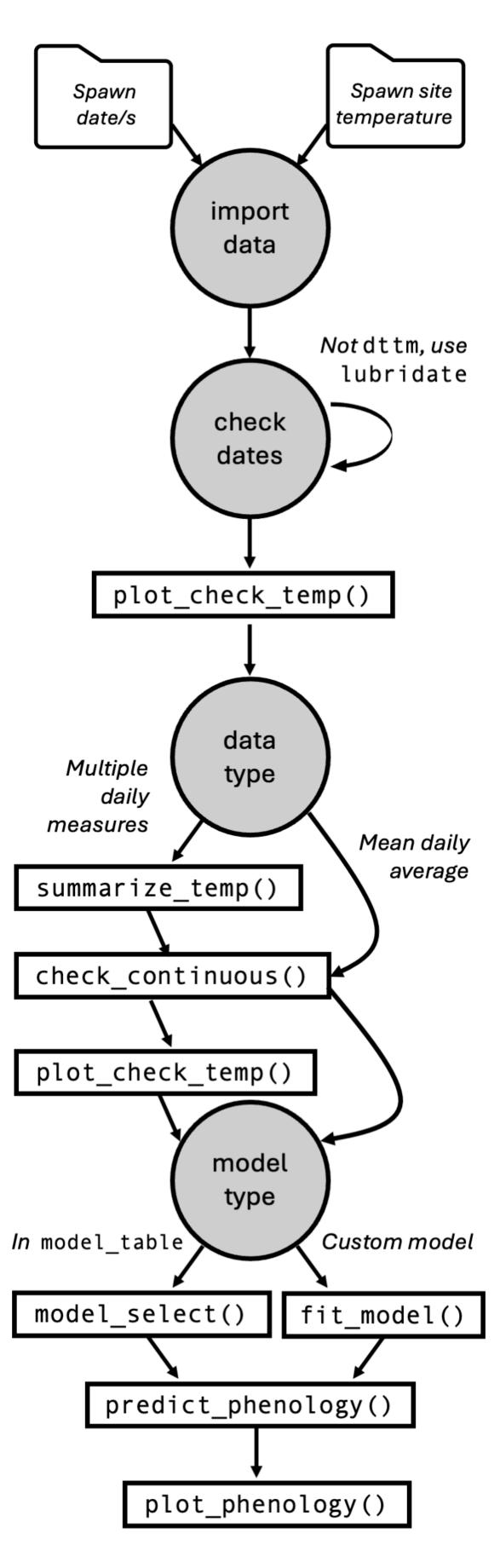


Figure 1: hatchR workflow. Processes are presented as filled circles, functions as plain text rectangles, and decisions as italicized text.