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# THE DESIGN AND ANALYSIS OF SALMONID TAGGING STUDIES IN THE COLUMBIA BASIN

VOLUME X: INSTRUCTIONAL GUIDE TO USING PROGRAM  
CAPTHIST TO CREATE SURPH FILES FOR SURVIVAL  
ANALYSIS, USING PTAGIS DATA FILES

Instructional Report



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Environment, Fish and Wildlife Division  
P.O. Box 3621  
905 N.E. 11th Avenue  
Portland, OR 97208-3621

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**THE DESIGN AND ANALYSIS OF SALMONID TAGGING  
STUDIES IN THE COLUMBIA BASIN**

**VOLUME X**

**INSTRUCTIONAL GUIDE TO USING PROGRAM CAPTHIST  
TO CREATE SURPH FILES FOR SURVIVAL ANALYSIS, USING  
PTAGIS DATA FILES**

Prepared by:

Peter Westhagen

and

John Skalski

School of Fisheries  
University of Washington  
Seattle, WA 98101

Prepared for;

U. S. Department of Energy  
Bonneville Power Administration  
Environment, Fish and Wildlife  
P.O. Box 3621  
Portland, OR 97208-3621

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## Preface

Project 89-107, Epidemiological Survival Methods, was developed to provide statistical guidance on the design and analysis of PIT-tag survival studies to the Northwest fisheries community. This study determined the statistical feasibility of conducting PIT-tag smolt survival studies, analytical capabilities for analyzing the tagging experiments, and recommendations on study design. As PIT-tag capabilities developed and research interests increased, the project has been instrumental in maintaining the statistical capabilities for designing and analyzing tagging studies as research objectives have expanded. This report describes program **CaptHist**, a data processing utility for converting raw PIT-tag data from PTAGIS into SURPH data input format for use in survival analysis.

# Executive Summary

The utility **CaptHist** was developed to act as a bridge between raw PIT tag data found in PTAGIS and the statistical survival **analysis tool SURPH**. The **raw** data from PTAGIS is in a form that must be pre-processed before it can be used as input to SURPH. A user's guide for **CaptHist** is presented here.

This work was conducted under contract **#DE-BI79-90BP02341** to Bonneville Power Administration. The goal of this task and project is to develop and disseminate the technologies needed for the design and analysis of Columbia Basin **salmonid** survival studies. This report describes an utility to streamline the conversion of PTAGIS raw data file into formats ready for survival analysis using program SURPH. 1.

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# Introduction

The SURPH program is a valuable, tool for estimating survivals and capture probabilities of fish outmigrations on-the Snake and Columbia Rivers. Using special data files, SURPH computes reach to reach statistics for any release group passing a system of detection sites.

Because the data must be recorded for individual fish, PIT tag data is best suited for use as input. However, PIT tag data as available from PTAGIS comes in a form that is not ready for use as SURPH input.

SURPH requires a capture history for each fish. A capture history consists of a series of fields, one for each detection site, that has a code for whether the fish was detected and returned to the river, detected and removed, or not detected.

The data as received from PTAGIS has one line for each detection with information such as fish **identification** (id), detection date and time, number of coil hits and detector coil id's, etc. Because an individual fish may be detected at several coils within a detection site as well as at several detections sites, each fish is often represented by multiple lines in the PTAGIS data file.

For the PTAGIS data to be usable by SURPH it must **be pre-processed**. The data must be condensed down to one line per fish with the relevant detection information from the PTAGIS file represented compactly on each line. In addition, the **PTAGIS** data file coil information must be passed through a series of logic algorithms to determine whether or not a fish is returned to the river after detection.

**Program CaptHist** was developed, to properly pre-process the PTAGIS data files for input to program SURPH. This utility **takes** PTAGIS data files as input and creates a SURPH data file as well as other output including travel **time records**, detection date records, and a data error file. **CaptHist allows** a user to, download **PTAGIS** files and easily process the data for use with SURPH.

## CaptHist User's Guide

### SURPH Input Files

SURPH input files consist of header information followed by capture histories for each-fish in the **release** group being studied. Fig. 1 shows an abbreviated example of an input file. The file has two basic parts: the header and the data. For a description of the **SURPH** data file header, consult the SURPH manual, section 5.3.

This document will focus on the data segment portion of the SURPH data file because this is where program **CaptHist** is most valuable. The data segment of the SURPH file consists of capture histories surrounded by the keywords "data" and "**enddata**".

In the example shown in Fig. 1, fish were released above Lower Granite Dam and then detected (or not detected) at detections sites on the Snake and Columbia Rivers. The capture history consists of a PIT code followed by 5 fields: the first field for the point of release, the next three fields representing recaptures at Lower Granite, Little Goose, and Lower Monumental dams, and the final field handling any other downstream recaptures. A field can have any one of several values with meanings as follows: 0 if not detected, 1 if detected and returned to the river, 2 if

detected and removed **from** the river, **and**, optionally, 3 if detected but of unknown' active status.

In the example, the fish with PIT code **222A4E2254** was released and then detected at Lower Granite and **Little Goose**, was known to be in the river after Little Goose but was not **detected** again.

---

**Figure 1: SURPH Input File**

```
Data
sample
npop 1
nper 4
ntag
9 0 0 0
full_hist
idlen 10
data
222A4E2254 1 1 1 0 0
22294C637A 1 0 0 0 0
2214636243 1 0 1 0 0
2214670160 1 0 0 1 0
221464261D 1 0 0 0 1
22142A006D 1 0 0 1 0
2214167A3B 1 0 0 1 0
22140C0306 1 0 0 1 1
221339587C 1 0 0 0 1
enddata
```

See SURPH **documentation** for explanation

**CaptHist** is especially useful in creating the capture histories in the second part of the data file. These capture histories represent the record of the fish through the river system.

---

## PTAGIS Data Files

The raw data **from** PTAGIS that corresponds to the capture history in the previous example is shown in Fig. 2. The columns in order are: tag file id, PIT code, detection site, detection date and time, number of **detector** coil reads, coil id 1, coil id 2 (XX indicates none), and travel time from release.

To translate this raw data into a SURPH data file, we must look at both the detection site

code and the coil ids. The detection site code will tell us where the fish is being detected. **In** this example we have three detection site codes for two detection sites: GOJ indicates Little Goose Dam; GRJ and GRX indicate Lower Granite Dam.

Next we look at the coil id's to determine what route the fish took through the detection site (i.e. whether or not the fish ended up back in the river). The coil id 2A at detection site GRJ **indicates** the "separator" detector, the first detector a fish would pass at Lower Granite Dam. Coil ids **3A**, 36, and 38 indicate the "diversion" detector which means the fish was diverted back to the river. The proper capture history value for this fish would be a 1, detected and returned to river.

The **GRX** detector hits are ignored. Certain coil id's associated with the GRX detection site code would indicate a removal and cause a 2 in the capture. history, but these codes indicate a return to river, which has already been indicated.

The GOJ detection site hits would be dealt with in a similar manner.

---

**Figure 2: Sample PTAGIS data**

RNI96102.AA1	222A4E2254	GOJ	15-may-1996	11:50:57	1	48	XX	32.95
RNI96102.AA1	222A4E2254	GOJ	15-may-1996	11:50:57	1	4A	XX	32.95
RNI96102.AA1	222A4E2254	GOJ	15-may-1996	11:50:58	2	4C	4E	32.95
RNI96102.AA1	222A4E2254	GOJ	15-may-1996	11:51:07	1	A6	XX	32.95
RNI96102.AA1	222A4E2254	GOJ	15-may-1996	11:51:08	2	A8	AA	32.95
RNI96102.AA1	222A4E2254	GOJ	15-may-1996	11:51:13	1	56	XX	32.95
RNI96102.AA1	222A4E2254	GOJ	15-may-1996	11:51:14	2	58	5A	32.95
RNI96102.AA1	222A4E2254	GRJ	10-may-1996	23:04:17	1	2C	XX	28.42
RNI96102.AA1	222A4E2254	GRJ	10-may-1996	23:26:01	2	36	38	28.43
RNI96102.AA1	22219432254	GRJ	10-may-1996	23:27:01	1	3A	XX	28.44
RNI96102.AA1	222A4E2254	GRX	10-may-1996	23:22:22	1	74	XX	28.43
RNI96102.AA1	222A4E2254	GRX	10-may-1996	23:22:22	1	76	Xx	28.43
RNI96102.AA1	222A4E2254	GRX	10-may-1996	23:22:26	1	80	XX	28.43
RNI96102.AA1	222A4E2254	GRX	10-may-1996	23:22:26	1	82	XX	28.43
RNI96102.AA1	222A4E2254	GRX	10-may-1996	23:22:26	1	86	XX	28.43

**This PTAGIS observation data is a detailed record of the passage of a fish through the system. The columns are tag group, PIT code, observation site, observation-date, observation time, number of coils hits, first coil, second coil, and travel time.-'**

---

## The SURPH data pre-processor CaptHist

Needless to say, repeating the above exercise for thousands of fish would become tedious which is why we developed a data handling tool to do the work, **Program CaptHist** takes data in the form of raw **PTAGIS** files and runs the necessary coil and detection site recognition algorithms to create the capture histories for use as input to **SURPH**.

In addition **CaptHist** checks for errors in the data; For instance, a fish detected downstream “before” being detected upstream. These errors are removed from the output and recorded in a separate error file.

Other outputs include a travel time file that contains travel times to each detection site for each fish, a detection date file that contains detection dates in day-of-year to each detection site for each fish, and a coil hit file that counts the number of coil hits at each site for each fish.

**Program CaptHist** was designed to work on the Snake and Lower Columbia Rivers. The program can be extended to work on other systems. All that is needed is some site **specific** information for the new system. This information is added to the program by editing the source code and so any modifications of this type must be made by the Columbia Basin Research staff.

## Using CaptHist

**Two** basic data files are needed for input to **CaptHist**: an observation (detection) file and a tag file. In addition a mortality file, if available, **allows CaptHist to** remove any fish that are **known** mortalities, and a release **date** file **allows CaptHist** to create the travel time file (Travel times given in PTAGIS data files are sometimes inconsistent due to different methods of recording). These release date and **mortality** data are readily available through **PTAGIS**.

## Observation Files

The example in Fig.2 shows the proper format for the observation **file**. This file would contain such **data for** all fish in the release. The **file** must have exactly 8 columns with dates in the proper format. Currently **CaptHist** recognizes **both** dates in the **03-feb-** 1997 format and the equivalent **02/03/97** format. Any missing coil codes should be replaced by XX.

## Tag files

Tag files give **CaptHist the** release information. Most importantly in survival calculations, the set of all fish released. This will be compared to detections to determine survivals. **Figure 3** shows the proper format for a tag file. The columns are: tag file id, PIT code, species, rearing type, length, and tag session flag (with information on special circumstance related to the fish).

---

### Figure3: Tag file format

RNI96102.AA1	2214167A3B	1	H	159	XX
RNI96102.AA1	22142A006D	1	H	138	XX
RNI96102.AA1	221464261D	1	H	114	XX
RNI96102.AA1	2214670160	1	H	118	I_I
RNI96102.AA1	2214636243	1	H	125	XX
RNI96102.AA1	22294C637A	1	H	167	XX
RNI96102.AA1	222A4E2254	1	H	178	XX

Tag file format columns as follows: tag group, PIT code, species, rearing type, length, tag session flag.

---

### Mortality files

A file that simply contains a list of PIT codes, one per line, each representing a dead fish that was not released but whose PIT code is in the tag file.

### Release files

A file with a tag file id, a release date, and a release time. An example of a release file is shown in Fig. 4.

---

### Figure 4: Release file format

RNI96096.6A1	4/06/96	14:30:00
RNI96099.6B1	4/08/96	15:28:00
RNI96100.6C1	4/09/96	11:00:00
RNI96102.6D1	4/11/96	15:00:00
RNI96102.A11	4/12/96	13:00:00

Release file has 3 columns: tag group, release date, and release time! .

---

## Downloading CaptHist

To run CaptHist you need a copy of the executable file. Go to the SURPH web page at <http://www.cqs.washington.edu/surph/> and click on utilities; This page will have further instructions.

## Running CaptHist

To test your copy, start with the following command from the DOS or UNIX prompt:

```
capthist -n
```

This should create a file called “**chrc**”, a run time configuration file that is described at length below.

If you have downloaded the complete distribution you will have some sample data files for input to CaptHist: **pw1h01.obs** and **pw1h01.tag**. Use the following command to create a surph input file:

```
capthist -Gpw1h01 -Ragr
```

This is the most basic way to run CaptHist. The **-G** flag tells CaptHist that the group prefix is **pw1h01**. CaptHist will search the data directory (defined in **chrc**, see below) for the file **pw1h01.obs** and **pw1h01.tag**. Multiple groups are possible simply by giving multiple **-G** flags. The **-R** flag lets CaptHist know where in the system the fish were released, in this case upstream of, or “above”, Lower Granite Dam (ie., agr).

The resulting capture history will have 5 fields. The first is the release field which is always one. The next field is the first detection site the fish will pass, namely Lower Granite Dam. Following are Little Goose Dam, and Lower Monumental Dam, and finally a catch-all field for any detections downstream of Lower Monumental at McNary dam, John Day dam, or Bonneville dam..

The fields in the surph file-are configured in the **chrc** file and may-include the above sites as well as the traps on the Snake River and-it’s tributaries. See the discussion below on run time configuration for more information about how to do this option.

## Run time configuration: chrc

Perhaps you do not want all five of the output files, or you would like to keep your data in one sub-directory and your output files in another. Or you would like to change-the fields that appear in the SURPH file. By editing the **chrc** file created in a previous example, you can configure CaptHist to your own needs.

The fields strings allow you to define capture history fields, the output switches allow you to switch certain files on or off, the directory configuration allows you to keep data and output separated, the file name configuration allows you to name the default mortality and release files (if used), and the file name extension section allows you to define the extension names that work best for your system.

## Fields strings

There are two sets of fields strings: **main** and last. The main fields refer to the fields in the capture history after release but before the final field. The last field is a catch all **that** will be one if the fish is detected at any one of the detection sites listed in the last fields string. These fields should be a comma separated list. For example, **gr,go,lm** specifies Lower Granite, Little Goose, and Monumental dams.

For a complete list of possible field names, see Appendix A.

## Output file switches

Determine which of the output files are written: 1 to write, 0 to not write. See the section titled “Output Files” for more information on the content of the files.

## Directory configuration

By default **CaptHist** will search for input files and write output files in the directory where it was run. To change directory for the input or output files, enter the name of the new directory here. This allows the feature of having two sub-directories of the directory that contains the **CaptHist** executable file, one for input and one for output to help keep your files organized. A ‘.’ **indicates** the current working directory.

## File name configuration

The mortality file and the release **file** are not necessary for a basic run. These files are used to check errors and to create the travel time output file. By default **CaptHist** searches the data directory (as configured above) for files of the name **all.mrt** and **all.rel**. You can change these default names here. In addition, if you wish to specify the names on the command line with a flag that is possible **too**. See the section titled “Command Line **Flags**” for more, information.

## File extension configuration

Set the filename extension for both input and output files. For instance, the SURPH data file could be **.surph**, or on the PC platform **.srh** since three letters work better there.

## Command Line Flags

Most of the **specifics** of a **run** are passed to the **CaptHist** program **via** command line flags or

**arguments. These flags tell CaptHist** what group(s) to look for, where the release was within the system, whether or not to check for errors, and other more esoteric considerations. Below is the entire list of flags along with explanations.

**-c allows single coil hits**

By default, if a fish is detected at only one coil at a given detection site, this detection record is removed because there is a good chance of it being an error. This flag allows you to include these if you desire.

**-E indicates errors should be found and removed**

Error checking is switched on with this flag .

**-Freartype:**

The rearing type. Same codes as used at **PTAGIS**.

W -> wild

H -> hatchery

u -> unknown

A->all

**-Ggroup: where group is the obs file and tag file prefix**

**The name of the group of fish to do the run on. If the group is pwl h01, by default CaptHist** will look for pwlh01.obs and **pw1h01.tag** for input and all the output will be prefixed by this group (e.g. pw **1h01.srh**, etc.) The filename extension can be changed using the “extensions” directives in the **chrc** file as explained in the section titled Run Tie **Configuration**. Multiple **-G** flags may be given.

**-ircfile: use alternate runtime configuration file**

CaptHist will read the named file for **configuration** instead of the default file “**chrc**”.

**-l include length in surph file, remove zero length fish**

**This flag causes Cap&Iii** to add length to the **SURPH** file as a **covariate** and in addition remove any fish with a zero length. **Otherwise** length is ignored,

**-Mmortfile: gives the name of the mortalities file to be searched for**

This file will be searched for in the data directory **in** place of the default file **all.mrt**.

**-n create new run time configuration file “chrc” with default settings**

Create a default **chrc** for editing and **exit**.

**-p set output prefix, used when multiple groups are given on command line**

When multiple **-G** flags are give, the output file name prefix is ambiguous. By default the output files will be named composite. \*. Use the **-p** flag to set it to something **specific**.

### **-rrelfile:**

This file will be searched for in the data directory in place of the default file **all.rel**.

### **-RreleaseSite:**

The point of release. This can either be a dam (**e.g. gr**), or upstream of a dam (e.g. agr). The following are valid release points. By placing the letter “a” in front of any **of** them, the meaning will be “above” (i.e. upstream of):

**agr** -> upstream of Lower Granite  
**gr** -> Lower -Granite Dam

If the fish are released at the dam an additional check is done so that only fish known to have entered the river below the dam (not transported or sampled) are included in the survival estimates. Otherwise the capture histories will look the same.

Not all release sites are allowable for each run. The release site must be one of the main fields as defined in the run time configuration file. If the main fields are configured as **follows**:

fields-main: **gr,go,lm**

The **accepted** release sites are agr, gr, ago, go, alm, or lm.

If the release site is not the first of main field, than the main fields up stream of the release site are not included in any of **the** output files. Program CaptHist **does use** the up stream main fields for error checking.

### **-s split input data to conserve memory.**

If data files tie so big that **CaptHist runs out** of memory, this flag can be set to conserve memory. **Data files** are split into smaller temporary files and then read in one by one, thus each data file is **small enough** (hopefully) to be loaded into memory for processing.

### **-t ignore fish found in obs tile that are not in tag file**

This flag tells **CaptHist** to ignore any fish in the-detection file that is not in **the** tag file. This is convenient when a detection file contains **observations from** many release groups and you only want to look at one of them.

### **-Species:**

Set the species. By default this is set to **chinook**. The species codes are the same as the **PTAGIS** codes. They are as follows:

0->all  
1 -> chinook  
3 -> steelhead  
4 -> sockeye

## **-U give 3 for unknown capture status**

By default a fish gets a 2 for its capture history if it is detected and then either known to have been removed or if its passage after the first detection is unknown. If the **-U** flag is given then the second group of fish with unknown fates receive a 3 in their capture history to distinguish them from the first group.

## **Output Files**

### **surph file:**

An example of a Surphfile is illustrated below for the case of a release above Lower Granite **Dam** with 4 downstream detection sites at Lower Granite, Little Goose, Lower Monumental and below Lower Monumental,

The meanings of the header fields is explained in the Surph documentation. The columns in this example are as follows:

column 1: PIT code  
column 2: release  
column 3: Lower Granite  
column 4: Little Goose  
**column 5: Lower Monumental**  
column 6: **downriver** of Lower **Monumental**

-- start file --  
Data

**pw.1.H.01**

**npop 1**  
**nper 4**

**ntag**  
1233 000

**full\_hist**

**idlen 10**

**data**  
**7F7A250404 1 0 0 0 0**  
**7F7A250402 1 1 0 0 0**  
**7F7A250361 10 0 0 0**  
**7F7A25034A 110 0 1**  
**7F7A25033A 10 10 0**  
**7F7A25032E 10 10 0**

```

...
204B543F02 10 10 0
1F6E0C7077 10 0 10
1F6E0B2D3B 10 0 0 0
1F6E0A6B7E 1 10 0 0
1F6D763F3F 10 0 0 0
enddata
-- end file --

```

**travel time file:**

A fish may be detected several times at a dam. Here we record the travel time from release to the first detection at a given site along with the travel time from release to the last detection at a given site.

The first line in the file is the header giving information about the data that follows. The header for this file looks **like**:

```

pitcode gr go go lm lm mc mc jd jd bv bv

```

From the header we **can** determine that, for instance, the second and third columns pertain to Lower Granite detections. In the travel time file as well as the detection date file, the convention is that the **first** of the two columns pertains to the first detection at the site and the second pertains to the last detection.

The columns are **determined** by the field configuration given in **chrc**. Any field listed as either a 'main field or a last field has two columns associated with it in the travel time **file** (and the detection date tile).

Here is an example:

```

-- start sample --
pitcode gr gr go go lm lm mc mc jd jd bv bv
22131C273C 0.000000 0.000000 0.000000 0.000000 9.758333
9.759027 0.000000 0.000000 14.148611 15.025000 0.000000 0.0000.00
0.000000 0.000000
22131C3661 0.000000 0.000000 0.000000 0.000000 8.398611
8.402777 0.000000 0.000000 -0.000000 0.000000 0.000000 0.000000
0.000000 0.000000
. 22131C5950 0.000000 0.000000 25.184027 25.793750 0.000000
0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000
0.000000 0.000000
22131D243F 0.000000 0.000000 0.000000 0.000000 0.000000
0.000000 19.367361 19.576389 29.576389 29.577777 0.000000
0.000000 0.000000 0.000000
2213232F29 0.000000 0.000000 x13.236805 13.242361 0.000000

```

```

0.000000 18.411111 18.744444 0.000000 0.000000 0.000000 0.000000
0.000000 0.000000
    2213261007 0.000000 0.000000 0.000000 0.000000 0.000000 0.0000.00
0.000000 0.000000 0.000000 30.296527 30.297222 0.000000 ~0.000000
0.000000 0.000000
    2213271B42 0.000000 0.000000 0.000000 0.900000 16.300000
16.302083 0.000000 0.-000000 0.000000 0.000000 0.000000 0.000000
0.000000 0.000000
    22.13293721 0.000000 0.000000 0.000000 0.000000 25.701389
25.716666 '0.000000 0.000000 0.000000 0.000000 0.000000 0.000000
0.000000 0.000000
    2213293B7D 0.000000 0.000000 0.000000 0.000000 33.355555
33.356944 0.000000 0.000000 0.000000 0.000000 0.-.000000 0.000000'
43.291666 43.291666
    22132B0C6A 0.000000 0.000000 11.331250~11.336805.15.742361
15.745139 0.000000 0.000000 0.000000 0.000000 0~000000 0.000000
0.000000 0.'000000
    -- end sample --

```

**detection date file:**

A fish may **be** detected **several** times at a detection site. Here **we** record the first detection date and 'the last detection date. All values- **are** in day-of-year with January 1 at noon being denoted 1.50 (as **opposed to** 0.50) A **0.000000** indicates no detection.

In the detection date file there are two columns for each of the fields configured-in the **chrc** file. These fields are given in the one line header of the file. The **first** column of the pair contains first detection date at the site, the second contains last detection **date** at the site.

```

-- start sample --
pitcode gr gr go go lm lm mc-mc jd jd bv bv
    22131C273C 0.000000 0.000000 0.000000 0.000000 113.300000
113.300694 0.000000 0.000000 117.690278 118.566667 0.000000
0.000000 0.000000 0.000000
    22131C3661 0.000000 0.00'0000 0.000000 0.000000 111.940278
111.944444 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000
0.000000 0.000000
    22131C5950 0.0~0000 0.000000 128.725694'129.335417 0.000000
0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000
0.000000 0.000000
    22131D243F 0.000000. 0.000000 0.000000 0.000000 0.000000
0.000000 122.909028 123.118056 133.118056 133.119444 0.000000
0.000000 0.000000 0.000000
    2213232F29 0.000000 0.000000 116.778472 116.784028 0.000000
0.000000 121.952078 0.000000 122.0286111 0.000000 0.000000

```

```

2213261007 0.00~000 0.000000 0.000'000 0.000000 0.000000
0.000000 0.000000 0.000000 133.838194 133.838889, 0.000000
0.000000 0.000000 0.000000
2213271B42 0.000000 0.000000 0.000000 0.000000 1'19.841667
119.843750 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000
0.000000 0.000000
2213293721 0.000000 0.000000 0.00'0000 0.000000 129.243056
129.258333 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000
'0.000000 0.000000
2213293B7D 0.000000 0.000000 0.000000 0.000000 136.897222
136.898611 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000
'146.833333 1461833333
22132B0C6A 0.000000 0.000000 114.872917 114.878472 119.284028
119.286806 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000
0.000000 0.000000

```

-- end sample --

error file:

**When the -E flag is given on the commandline, CaptHist checks the data for any errors and outputs whatever error information it finds to a file. If no errors are found, the file will not be created.**

**The errors are broken up into errors that are removed and errors that are recorded but not removed. The second column has either a one to indicate that the error caused removal or a zero to indicate that the error is a warning only and the fish was not removed.**

**The error file has one line for each fish and is followed by a series of either ones or zeros. A one in a given column indicates an error of that-type has occurred, a zero indicates no error. A fish is recorded in the error file only if it has at least one error.**

**Here is a brief description of the error columns:**

**column 1: PIT code**

**column 2: removal status**

**column 3: duplicate tag in tag file (not implemented)**

**column 4: negative travel time**

**column 5: length of zero**

**column 6: Fish detected upstream after downstream detection**

**column 7: Fish detected upstream of release**

**column 8: mortality in observation file**

**column 9: mortality in tag file**

**column 10: wrong species**

**column 11: wrong rearing type**

**column 12: fish not known to be returned to river**

**column 13 no release date for fish**

**Lower Granite Coil errors;**

- column 14: Detection at diverter and sample
- column 15: Detection at diverter and raceway
- column 16: Detection at raceway and downriver
- column 17:** Detection at sample and downriver

**Little Goose Coil errors:**

- column **18:** Detection at diverter and raceway
- column 19: Detection at diverter and sample
- column 20: Detection at diverter exit and raceway
- column 21:** Detection at diverter exit and sample
- column 22: Detection at raceway and downriver
- column 23: Detection at sample and downriver

**Lower Monumental Coil errors:**

- column 24: Detection at diverter and raceway
- column 25: Detection at **diverter** and sample
- column 26: Detection at diverter exit and raceway
- column 27: Detection at diverter exit and sample
- column **28:** Detection at raceway and **downriver**
- column 29: Detection at sample and downriver

```
-- begin sample --
7F7A250313 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
0 0 0
7F7A247E7D 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
000
7F7A24763E 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
0 0 0
7F7A246741 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
0 0 0
7F7A1B7A1E 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
0 0 0
7F7A1B6D3F 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
0 0 0
7F7A1A2A5F 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
0 0 0
7F7A1A2406 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
0 0 0
7F7A1A1C5F 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
00.0
7F7A1A1B27 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
0 0 0
-- end sample --
```

## coil file:

The number of coil reads at each of the detection sites is recorded in the coil file. The first line in the file is a header **line** that indicates the capture sites for the data. For instance, fish **2233445D63** was detected on 9 coils at Lower Granite Dam, 11 coils at Little Goose Dam, etc. As with the travel time file and the detection date file, the columns in this file are configured by the settings for “fields-main” and “**fields\_last**” in the **chrc** run-time configuration file.

```
-- begin sample --
```

```
pitcodegrgolmmcjdbv
```

```
2233445D63    9  11  11    0    0    0
22333C5D2E    0   8   0    0    0    0
223335137C    0   0   8    0    0    0
222A622B7D    0   0  11    0    0    0
222A5A5A76   10   0   0    0    0    0
222A4E2254    9  10   0    0    0    0
222A4D0856    0   0  11    0    0    0
222A4A3012    9   0   0    0    0    0
222A433E39    0  11   0    0    0    0
22295C470F   0-8   0    0    0    0
```

```
-- end sample --
```

## Examples

Run **CaptHist** on **pwlh01.obs** and **pw1h01.tag**, tell **CaptHist** that the fish were released above Granite and **check** for errors.

```
capthist -Gpw1h01 -Ragr -E
```

**Program CaptHist** will search the data directory (as configured in **chrc**) for the files **pw1h01.obs** and **pw1h01.tag**. If either of these files are not found, **the CaptHist will exit**. In addition, **CaptHist** will search the data directory for a mortality file and a release date file, by default **all.mrt** and **all.rel**. **Program CaptHist will complete** the run whether or not these files are found as the mortality and release date files are optional.

**The -Ragr** flag tells **CaptHist** to treat the fish as being released above Lower Granite Dam with that **being the first** detection site. The **-E** flag causes **CaptHist** to check for errors and to record them in an error iog file if any are found. For instance, if a fish is detected above the point of release, that would be an error. This would happen in the following scenario:

If the main fields are configured to be im, gr, go, and lm with the release site being agr, any detection at im would be an error. To include these fish as valid detections, the release site would have

to be aim. See the description of the error file above for a more complete description of errors.

Run **CaptHist** on a group of wild Steelhead.

```
capthist -Gpw3w01 -Ragr -E -S3 -FW
```

With the **-E** flag, fish of the wrong species are removed. The default species is chinook but with the **-S** flag with the 3 qualifier, the species has been switched to steelhead. No with the **-E** flag, any fish **that** is not a steelhead is removed. The **-F** flag with the W qualifier **tells CaptHist** that the fish are wild; by default the fish are taken to be hatchery fish (H).

Create **runtime** configuration file “**chrc**” and exit.

```
capthist -n
```

This creates the default **chrc** file which may then be edited for custom configuration.

Run **CaptHist** in split mode: splitting input data to use less memory.

```
capthist -Gpw1h01 -Ragr -E-s
```

This run will have the same results as the first example but the run will be done in a memory **minimizing** way so that files that very large **can be** handled. The **-s** flag tells **CaptHist** to break the input files into smaller files and then to work on each of these smaller tiles one at a time

**Program CaptHist** splits files so that fish are grouped by **PIT** code, taking the last n digits as the matching criterion. By default the fish are sorted by the last digit in their PIT code. This can be **modified** to create small split files by taking the last two digits of the **PIT** code by adding a qualifier to the **-s** flag: **-s2**.

## Feedback

For **additional** information, clarification or assistance in running program **CaptHist** contact Peter **Westhagen** at Columbia Basin Research, School of Fisheries, University of Washington (email: [china@u.washington.edu](mailto:china@u.washington.edu)).

# Appendix A

## Site name codes

b	v	Bonneville Dam <b>DSM1</b> Subsample
<b>cn</b>		Challis Diversion North
cl		Clearwater River Trap Juvenile
<b>gr</b>		<b>Lower Granite Dam Juvenile</b>
im		Imnaha River Trap Juvenile
<b>jd</b>		John Day Dam <b>Gatewell</b> Airlift
<b>lm</b>		Lower Monumental Dam Juvenile
mc		<b>McNary Dam Juvenile</b>
<b>pr</b>		'Prosser, Juvenile (Chandler Diversion)-
ra		<b>Redfish Lake Creek Trap A</b>
r b		<b>Redfish Lake Creek Trap B</b>
sa		Salmon River Trap Juvenile
sn		Snake River Trap Juvenile