

# Package ‘FAIBCompiler’

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SIndexR,

FAIBOracle,

FAIBBase

**Suggests** knitr

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ageByForester

Site age by experienced forester

---

**Description**

This function derives the site age based on experienced forester. It is equivalent to vri\_age macro in original SAS compiler.

**Usage**

ageByForester(projectID, sampleNumber, sampleTypeCode)

**Arguments**

- projectID      character, Project ID.
- sampleNumber    character, Sample number.
- sampleTypeCode   character, Sample type code.

**Value**

Age provided by experienced forester

**Note**

Contact Bob Krahn for details

**Author(s)**

Yong Luo

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ageRangeClassifier

Derive age range code

---

**Description**

This function derives age range code based on age, species and FIZ. The returned age range code includes: 1-young(immature), 2-older(immature), 3-mature and 4-overmature. This function is equivalent to age\_rng.sas macro.

**Usage**

ageRangeClassifier(age, species, FIZ)

**Arguments**

age	numeric, Usually layer mean age. what does this mean?
species	character, Tree basic species code, which is SP0 in VRI original data.
FIZ	character, BC forest inventory zone.

**Value**

DWB age range code

**Author(s)**

Yong Luo

---

applyDWB	<i>Apply decay, waste and breakage percentage to gross merchantable volume-VRI specific</i>
----------	---

---

**Description**

This function calculates merchantable volume after removing decay, waste and breakage in VRI compiler. The function is part of cp\_vegi\_2017.sas to derive tree\_ms7.

**Usage**

```
applyDWB(treeMS)
```

**Arguments**

treeMS	data.table, Compiled full and enhanced trees with percentage of decay, waste and breakage.
--------	--

**Value**

A data table that contains VOL\_NTWB (net volume that waste 2 wood and breakage), VOL\_D (merchantable volume after removing decay), VOL\_DW (merchantable volume after removing decay and waste) and VOL\_DWB (merchantable volume after removing decay, waste and breakage).

**Author(s)**

Yong Luo

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auxiTreeCompiler	<i>Derive volume components for H-enhanced and non-enhanced trees using external coeff and ratio-VRI specific</i>
------------------	---

---

## Description

Estimates volume components for H-enhanced and non-enhanced trees using regression and ratio methods. For H-enhanced trees, the whole stem volume and gross merchantable volume are already calculated directly using taper equations; and rest of volume components will be calculated using ratio method in this function. For non-enhanced trees, the whole stem volume is derived using regression equation between basal area and whole stem volume and the rest of volume components will be computed using ratio method in this function.

## Usage

```
auxiTreeCompiler(
  fullMeasuredTrees,
  auxiTrees,
  clusterPlotHeader,
  fixedCoeff,
  randomCoeff,
  ratios
)
```

## Arguments

fullMeasuredTrees	Compiled tree-level data in vi_c, which contains full measured trees, enhanced trees and H-enhanced trees. This data is output of <a href="#">DWBCompiler</a>
auxiTrees	data.table, Non-enhanced trees in auxillary plots, however, it may have enhanced trees and H-enhanced trees. An output from <a href="#">VRIInit_auxTree</a> .
clusterPlotHeader	data.table, Cluster and plot-level information. An output of <a href="#">VRIInit_clusterplot</a> .
fixedCoeff	data.table, Specifies the WSV-BA equations by project group PRJ_GRP, live and dead status LV_D, stand and falling status SF_COMPILE and species code SP0. The table can be generated using <a href="#">WSV_BARegression</a> .
randomCoeff	data.table, Specifies the WSV-BA equations by project group PRJ_GRP, live and dead status LV_D, stand and falling status SF_COMPILE and species code SP0. The table can be generated using <a href="#">WSV_BARegression</a> .
ratios	data.table, Specifies toWSV ratio by project group PRJ_GRP, live and dead status LV_D, stand and falling status SF_COMPILE and species code SP0. The table can be generated using <a href="#">toWSVRatio</a> .

## Value

A list of four tables: 1. fullenhancedtrees: full and enhanced trees; 2. HnonenhancedTrees: Height enhanced and non-enhanced trees;

**Note**

The data selection procedure for regression has been standardized as following:

1. Start from vi\_c, which has all trees have minimum information of DBH and Height;
2. Select all the full, enhanced and H-enhanced trees;
3. Remove observations in Audit plots and have zero whole stem volume;
4. Select the latest observation for each tree by live\_dead status. A tree's identity is considered same when it is from same proj\_id, samp\_no and plot.

The data selection for ratio has been standardized as following:

1. Start from the that used for regression;
2. Select all the full and enhanced trees;
3. Select the trees with DBH  $\geq 10$ cm

**Author(s)**

Yong Luo

---

boredAgeCalculator\_Bore

*Derive bored age using office and field bored age*

---

**Description**

This function is to derive bore age based on either office bored age (officeBoredAge) or field bored age (fieldBoredAge). When both bore age information are available, the function takes officeBoredAge as priority. The function is one of the four functions that derive bored age using different method. The rests are [boredAgeCalculator\\_Total](#), [boredAgeCalculator\\_Phys](#) and [boredAgeCalculator\\_Prorated](#).

**Usage**

```
boredAgeCalculator_Bore(officeBoredAge, fieldBoredAge)
```

```
## S4 method for signature 'numeric,numeric'
boredAgeCalculator_Bore(officeBoredAge, fieldBoredAge)
```

```
## S4 method for signature 'numeric,missing'
boredAgeCalculator_Bore(officeBoredAge)
```

```
## S4 method for signature 'missing,numeric'
boredAgeCalculator_Bore(fieldBoredAge)
```

**Arguments**

officeBoredAge    numeric, Office bored age, which is measured in lab by professionals.

fieldBoredAge    numeric, Field bored age, estimated in field by field crew.

**Value**

bored age

**Author(s)**

Yong Luo

**See Also**

[boredAgeCalculator\\_Total](#) [boredAgeCalculator\\_Phys](#) [boredAgeCalculator\\_Prorated](#)

---

boredAgeCalculator\_Crted

*Derive age at breast height*

---

**Description**

This function uses site tools to derive age at breast height for the bored age that is not taken at breast height, i.e., 1.3 m. The process was documented in BC VRI Sample Data Compilation Process.

**Usage**

```
boredAgeCalculator_Crted(boredAge, boredHeight, treeHeight, species, FIZ)
```

```
## S4 method for signature 'numeric,numeric,numeric,character,character'  
boredAgeCalculator_Crted(boredAge, boredHeight, treeHeight, species, FIZ)
```

**Arguments**

boredAge	numeric, Age at bored height
boredHeight	numeric, Height at the bore core.
treeHeight	numeric, Total tree height
species	character, Species code, must be consistent with the species code in site tools. Can be derived using <a href="#">siteToolsSpeciesConvertor</a> .
FIZ	character, Forest inventory zone.

**Value**

Tree age at breast height (1.3 m). For the estimated age  $\leq 0.1$ , the bored age will be used as breast age and a warning message will be given.

**Author(s)**

Yong Luo

---

boredAgeCalculator\_Phys

*Derive bored age using physiological age*


---

### Description

This function is to derive bore age based on physiological age (physAge). The function is one of the four functions that derive bored age using different method. The rests are [boredAgeCalculator\\_Bore](#), [boredAgeCalculator\\_Total](#) and [boredAgeCalculator\\_Prorated](#).

### Usage

```
boredAgeCalculator_Phys(physAge)
```

```
## S4 method for signature 'numeric'
boredAgeCalculator_Phys(physAge)
```

### Arguments

physAge                  numeric, Pysiological age.

### Value

bored age

### Author(s)

Yong Luo

### See Also

[boredAgeCalculator\\_Total](#) [boredAgeCalculator\\_Phys](#) [boredAgeCalculator\\_Prorated](#)

---

boredAgeCalculator\_Prorated

*Derive bored age using pro-rated age*


---

### Description

This function is to derive bore age based on diameter at bore (boreDiameter), bark thickness (barkThickness), pro-rated ring length (ringLength\_prorated) and pro-rated ring count (ringCount\_prorated). The function is one of the four functions that derive bored age using different method. The rests are [boredAgeCalculator\\_Bore](#), [boredAgeCalculator\\_Total](#) and [boredAgeCalculator\\_Phys](#).



**Usage**

```
boredAgeCalculator_Prorated(  
  ringLength_prorated,  
  ringCount_prorated,  
  boreDiameter,  
  barkThickness  
)  
  
## S4 method for signature 'numeric,numeric,numeric,numeric'  
boredAgeCalculator_Prorated(  
  ringLength_prorated,  
  ringCount_prorated,  
  boreDiameter,  
  barkThickness  
)  
  
## S4 method for signature 'numeric,numeric,numeric,missing'  
boredAgeCalculator_Prorated(  
  ringLength_prorated,  
  ringCount_prorated,  
  boreDiameter  
)
```

**Arguments**

ringLength_prorated	numeric, Pro-rated ring length in cm
ringCount_prorated	numeric, Pro-rated ring count
boreDiameter	numeric, Diameter at bore in cm
barkThickness	numeric, Bark thickness in mm. If missing, 0.05 is used in the function.

**Value**

bored age

**Author(s)**

Yong Luo

**See Also**

[boredAgeCalculator\\_Total](#) [boredAgeCalculator\\_Phys](#) [boredAgeCalculator\\_Prorated](#)

---

boredAgeCalculator\_Total

*Derive bored age using total age*


---

### Description

This function is to derive bore age based on total age (totalAge). The function is one of the four functions that derive bored age using different method. The rests are [boredAgeCalculator\\_Bore](#), [boredAgeCalculator\\_Phys](#) and [boredAgeCalculator\\_Prorted](#).

### Usage

```
boredAgeCalculator_Total(totalAge)
```

```
## S4 method for signature 'numeric'
boredAgeCalculator_Total(totalAge)
```

### Arguments

totalAge            numeric, Total tree age, ie., age at height of 0.

### Value

bored age

### Author(s)

Yong Luo

### See Also

[boredAgeCalculator\\_Total](#) [boredAgeCalculator\\_Phys](#) [boredAgeCalculator\\_Prorted](#)

---

checkMaps

*Load maps from map source*


---

### Description

This function is to load maps from mapSourcePath and save them to mapPath. For TSA and BEC, the maps are direct from bcmeps package.

### Usage

```
checkMaps(mapPath)
```

### Arguments

mapPath            character, The path to save all the maps. Note that all the saved maps have time stamps, which suggest when the files have been modified in mapSourcePath.

**Value**

no value is returned from this function.

**Note**

all the maps relies on the bcdata package.

**Author(s)**

Yong Luo

---

compilerPathSetup	<i>Setup output paths of the compiler</i>
-------------------	---

---

**Description**

This function does two things: 1. create folders that will store compiled data; 2. return paths that directs the compiled folder.

**Usage**

```
compilerPathSetup(compilationPath = ".", compilationDate)
```

**Arguments**

compilationPath

character, Specifies a path to store whole compilation process. If missing, the current work directory will be used.

compilationDate

character, Specifies a compilation date. It should be in format of YYYYMMDD. It will be used for archive the compilation outputs.

**Value**

Seven paths will be returned as following:

- raw\_from\_oracle Path to save all data that read from both oracle and txt database without merging.
- compilation\_sa Path to save merged data for key tables (i.e., vi\_a to vi.g) from both oracle and txt database.
- compilation\_db Path to save compiled outputs.
- compilation\_archive Path to archive all compilation process.
- compilation\_report Path to report compilation process.
- compilation\_map Path to archive all maps for compilation process.
- compilation\_coeff Path to archive all coefficients for compilation process.

compilation\_last Path to archive last compilation process.

**Note**

Could overwrite the existing output folder, depending on user's choice, i.e., yes or no.

**Author(s)**

Yong Luo

DWBCompiler

*Compile decay, waste and breakage for standard tables-VRI specific***Description**

This function compiles decay, waste and breakage for standard tables in VRI compiler. The function is equivalent to `dwb_vri_2017.sas`.

**Usage**

```
DWBCompiler(treeMS, siteAge, treeLossFactors, equation)

## S4 method for signature 'data.table,data.table,data.table,character'
DWBCompiler(treeMS, siteAge, treeLossFactors, equation)

## S4 method for signature 'data.table,data.table,data.table,missing'
DWBCompiler(treeMS, siteAge, treeLossFactors)
```

**Arguments**

<code>treeMS</code>	<code>data.table</code> , Tree-level data that has been compiled whole stem volume and gross merchantable volume for full and enhanced trees.
<code>siteAge</code>	<code>data.table</code> , Cluster-level summaries of age and height. This table is an output from <a href="#">siteAgeSummary</a>
<code>treeLossFactors</code>	<code>data.table</code> , The tree loss factor data, an output of <a href="#">VRIInit_lossFactor</a> . In this function, this table provides loss indicator.
<code>equation</code>	character, Specifies whether the compiler is based on KFIZ or KBEC. Default is set as KBEC.

**Value**

A compiled volume after removing decay, waste and breakage; a log file

**Author(s)**

Yong Luo

---

DWBGenerator_BEC	<i>Collect decay, waste and breakage factor in BEC routine</i>
------------------	--

---

### Description

This function is to collect the dead, waste and breakage factor from lookup table and join them into tree data. Instead of reading the lookup table from disk, the function uses hard-coded the lookup table. This function is equivalent to dwb\_v3.sas macro. For FIZ routine, the decay, waste and breakage are collected using funtion [DWBGenerator\\_FIZ](#)

### Usage

```
DWBGenerator_BEC(DBH, height, species, meanAge, BEC, riskGroup, adjustID)
```

```
## S4 method for signature
## 'numeric,numeric,character,numeric,character,character,character'
DWBGenerator_BEC(DBH, height, species, meanAge, BEC, riskGroup, adjustID)
```

### Arguments

DBH	numeric, Tree DBH.
height	numeric, Tree height.
species	character, Tree basic species code, which is SP0 in VRI original data.
meanAge	numeric, Mean site age.
BEC	character, BC BEC zone.
riskGroup	character, Specifies the risk group. It must be one of 1, 2 or 3. It can be derived from <a href="#">riskGroupDeriver</a> .
adjustID	character, Adjustment identifier. Blank is no adjustment; QCI is queen charlottes; WET is wetbelt and GLD_NW golden ?.

### Value

A list of decay, waste and breakage percentage.

### Author(s)

Yong Luo

---

DWBGenerator_FIZ	<i>Collect decay, waste and breakage factor in FIZ routine</i>
------------------	--

---

### Description

This function is to collect the dead, waste and breakage factor from lookup table and join them into tree data. Instead of reading the lookup table from disk, the function uses hard-coded the lookup table. This function is equivalent to second part of dwb\_fct.sas macro. For BEC routine, the decay, waste and breakage are collected using funtion [DWBGenerator\\_BEC](#)

**Usage**

```
DWBGenerator_FIZ(DBHClass, tabNumber, riskGroup)

## S4 method for signature 'numeric,character,character'
DWBGenerator_FIZ(DBHClass, tabNumber, riskGroup)
```

**Arguments**

DBHClass	numeric, Tree DBH class. Currently, The function is capatable to the DBH class with 5 cm interval. It can be derived using <a href="#">DBHClassifier</a> .
tabNumber	character, This character consist of 4 numbers. The first two number is the species number, and the last is the series number.
riskGroup	character, Specifies the risk group. It must be one of 1, 2 or 3. It can be derived from <a href="#">riskGroupDeriver</a> .

**Value**

A list of decay, waste and breakage percentage.

**Author(s)**

Yong Luo

---

getDWBSeries	<i>Get local DWB series</i>
--------------	-----------------------------

---

**Description**

Join local DWB (old/imperial factors) by species, PSYUB, ageRangeClass and FIZ. This function is the first part of dwb\_fct.sas.

**Usage**

```
getDWBSeries(species, ageRangeClass, PSYUB, FIZ, source)

## S4 method for signature 'character,character,character,character,character'
getDWBSeries(species, ageRangeClass, PSYUB, FIZ, source)

## S4 method for signature 'character,character,character,missing,character'
getDWBSeries(species, ageRangeClass, PSYUB, source)

## S4 method for signature 'character,character,missing,character,character'
getDWBSeries(species, ageRangeClass, FIZ, source)
```

**Arguments**

species	character, Tree basic species code, which is SP0 in VRI original data.
ageRangeClass	character, The classified age range into 1 to 4. Output from <a href="#">ageRangeClassifier</a> .
PSYUB	character, Unique PSYU+PSYU_BLK code.
FIZ	character, BC forest inventory zone.

source character, Series source, must one of local, zonal and reversingZonal. local is based on species, PSYUB and ageRangeClass; zonal is based on FIZ, species and ageRangeClass; and reversingZonal is based on reversing FIZ zones, i.e., changing interior to coastal and changing coastal to interior.

### Value

DWB series, a two number character.

### Author(s)

Yong Luo

---

heightEstimate\_byHeightModel

*Estimate height using height-DBH curves*

---

### Description

Estimate height using height-DBH curves by bec subzone and species

### Usage

```
heightEstimate_byHeightModel(beczone, subzone, species, DBH, heightModels)
```

### Arguments

beczone	character, BEC zone.
subzone	character, BEC subzone.
species	character, Species.
DBH	numeric, Diameter at breast height.
heightModels	data.table, Specifies the best model and coefficients by each BEC subzone and species.

### Value

projected total height

### Author(s)

Yong Luo

---

heightSmry_byC	<i>Summarize mean and lorry's height by cluster-VRI specific</i>
----------------	--

---

### Description

Summarizes mean and lorry's height by cluster for standing trees, standing + live trees, and standing + live + non-broken top trees. The function is improved version to calculate mean height in vol\_ha\_2017.sas by outputting lorey's height. For both fixed and variable area plots, the function computes mean height by using plot weight (PLOT\_WT) weighted height. For lorey's height computation, the function treats variable and fixed area plots differently. Specifically, the function uses the mean height as lorey's height for variable plots, while uses height that weighted both by plot weight (PLOT\_WT) and basal area (BA\_TREE) for fixed area plots.

### Usage

```
heightSmry_byC(treeMC)

## S4 method for signature 'data.table'
heightSmry_byC(treeMC)
```

### Arguments

treeMC	data.table, Compiled tree-level data that contains both measured trees and counted trees.
--------	---

### Value

A table contains computed mean height MN or MEAN and lorey's height LRY for all standing trees ALL, standing and live trees 1 and standing and non-broken top trees 2. The output is equivalent to height table in original compiler.

### Author(s)

Yong Luo

---

ISMCCompiler	<i>ISMCC compiler - Adapted from VRI compiler to compile data converted from VGIS database</i>
--------------	--

---

### Description

This compiler is adapted version of original VRI compiler. It loads data from ISMC, pipes data into compilation processes and outputs compiled results at both tree and stand levels.



**Usage**

```
ISMCCompiler(
  oracleUserName,
  oraclePassword,
  oracleEnv = "INT",

  compilationPath = "//albers/gis_tib/VRI/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/FromRCompiler",
  equation = "KBEC",
  walkThru = TRUE,
  logMinLength = 0.1,
  stumpHeight = 0.3,
  breastHeight = 1.3,
  UTOPDIB = 10,
  utilLevel = 4,
  weirdUtil = "No"
)
```

**Arguments**

<code>oracleUserName</code>	character, User name to access to ISMC database.
<code>oraclePassword</code>	character, Password to access to ISMC database.
<code>oracleEnv</code>	character, Specify which environment of ISMC database the data download from. Currently, it supports 1) INT for intergration environment; 2) TST for test environment; 3) PROD for final production environment.
<code>compilationPath</code>	character, Specifies the path that stores all the data/processes. By specifying this, four folders will be created to record all the data/processes. Specifically, <code>raw_from_oracle</code> stores the data just after oracle and ascii without editing; <code>compilation_sa</code> stores key data (not all) that after editing and before volume and age compilation; <code>compilation_db</code> stores compiled results for volume and age compilation at both tree level and cluater level; <code>Archive_YYYYMMDD</code> achives all the data mentioned above for the future use or reference. By default, this path is set as <code>//albers/gis_tib/VRI/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/RCompilation</code> , which is consistent with our <code>rdw</code> system.
<code>equation</code>	character, Specifies the taper equation that is used for compiler. Currently supports BEC-based (KBEC) and FIZ-based (KFIZ).
<code>walkThru</code>	logical, Speciefies whether the data had been collected using work through method. Default is TRUE, if it is not specified.
<code>logMinLength</code>	numeric, Specifies minimum length of log when doing log length adjustment, see <a href="#">logMatrixAdjustment</a> for details. If missing 0.1 is used.
<code>stumpHeight</code>	numeric, Stump height. If missing 0.3 is used.
<code>breastHeight</code>	numeric, Breast height. If missing 1.3 is used.
<code>UTOPDIB</code>	numeric, Threshold inside-bark diameter for merchantable volume. If missing, UTOPDIB is 10.
<code>utilLevel</code>	numeric, Specifies utilization level in summrizing tree volumes at cluster and species level. Default is 4.
<code>weirdUtil</code>	character, Specifies weird utilization in summarizing tree volumes at cluster and species level. Default is no, if missing. Otherwise, a number should be provided.

**Value**

This function compiles data and save outputs in `compilationPaths$compilation_db` and no file is returned.

**Note**

Improvements include:

1. specifies trees in B plots as height enhanced trees
2. removes the sas-dependent `sindex` functions
3. introduce the `SIndexR` package

Currently, the compiler supports compilation for the below sample types:

- Q: Regular VRI sample with five point clusters design
- T: ??, sample protocol and sample design are same as Q samples
- B: ??, same plot layout as Q sample, with height is measured for all trees and no call grading information
- M: CMI sample, with all trees have call grading information in the field
- L: LiDAR project, same plot layout and same design but without call grading information
- Y: YSM plots, population between 15 and 50 years
- F: NFI plots, trees measured all DBH, height and call grading
- N: NVAF plots
- A: VRI audit plots

**Author(s)**

Yong Luo

**References**

VRI compiler manual

---

ISMC_VGISTranslator	<i>Translate ismc data formats into vgis formats</i>
---------------------	--

---

**Description**

This function is to translate ismc data formats into vgis formats

**Usage**

```
ISMC_VGISTranslator(inputPath, outputPath, coeffPath)
```

**Arguments**

inputPath	character, Specifies the path that stores data from oracle data base.
outputPath	character, Specifies the path to save your outputs. If missing, the current working directory will be choose.
coeffPath	character, Specifies the path where the crosswalk table is stored.

**Value**

no item returned

**Author(s)**

Yong Luo

---

loadASCII

*Load the data from ASCII files*


---

**Description**

This function is to read the data from ASCII files (in txt format).

**Usage**

```
loadASCII(txtLocation, saveThem = FALSE, savePath)
```

**Arguments**

txtLocation	character, Specifies the location of ASCII files.
saveThem	logical, Specifies whether the loaded data should be saved or returned. The default value is FALSE, which means the function will not save files for you.
savePath	character, Specifies the path that directs to the VRI original data source, i.e., //Mayhem/GIS_TIB/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/vri_sa.

**Value**

no files

**Author(s)**

Yong Luo

**Examples**

```
## Not run:
loadASCII(txtLocation = "Q:/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/vri_raw",
          saveThem = TRUE,
          savePath = "F:/vricompilertest/OracleTests")

## End(Not run)
```

---

logAdjustment	<i>Adjust log length - VRI specific</i>
---------------	---

---

### Description

The function is to adjust the log length to fit actual height. This function is equivalent to `log_adj_new` macro in original VRI compiler

### Usage

```
logAdjustment(treeData, stumpHeight)

## S4 method for signature 'data.table,numeric'
logAdjustment(treeData, stumpHeight)

## S4 method for signature 'data.table,missing'
logAdjustment(treeData)
```

### Arguments

treeData	data.table, Must have tree data information. The table is an output of <a href="#">VRIInit_measuredTree</a> .
stumpHeight	numeric, Length of stump. As default, this argument is set as 0.3 m.

### Value

Data table that contains the adjusted log length

### Note

Please see Bob for details about input files

### Author(s)

Yong Luo

---

logAttributesLongForm	<i>Transpose wide form table to long form table-VRI specific</i>
-----------------------	--

---

### Description

This function transposes wide form outputs to long form outputs. This function is not included in the VRI compiler anymore.

### Usage

```
logAttributesLongForm(treeData, maximumLogNO)

## S4 method for signature 'data.table,numeric'
logAttributesLongForm(treeData, maximumLogNO)

## S4 method for signature 'data.table,missing'
logAttributesLongForm(treeData, maximumLogNO)
```

**Arguments**

- treeData** data.table, an output from logValueCalculator function, i.e., tree\_ms6. This table currently has top diameter (LOG\_D\_X), length (LOG\_L\_X), volume (LOG\_V\_X), merchantable volume (LOG\_VM\_X), grade (LOG\_G\_X), sound percentage (LOG\_S\_X) and value (LOG\_c\_x). X is log number from 1 to maximum log number.
- maximumLogNO** numeric, determine the maximum number of logs. In VRI compiler, it is 9. Therefore, 9 is default.

**Value**

A data table and a log file

**Author(s)**

Yong Luo

---

logFileProducer	<i>To display a table to log file</i>
-----------------	---------------------------------------

---

**Description**

Simple tool to display a table to log file.

**Usage**

```
logFileProducer(
  reason,
  action,
  displayTable,
  displayColumn,
  changedVariable,
  fromTo
)

## S4 method for signature
## 'character,character,data.table,character,character,character'
logFileProducer(
  reason,
  action,
  displayTable,
  displayColumn,
  changedVariable,
  fromTo
)

## S4 method for signature
## 'character,character,data.table,missing,character,character'
logFileProducer(reason, action, displayTable, changedVariable, fromTo)

## S4 method for signature
```

```
## 'character,character,data.table,character,missing,missing'
logFileProducer(reason, action, displayTable, displayColumn)

## S4 method for signature
## 'character,character,data.table,missing,missing,missing'
logFileProducer(reason, action, displayTable)
```

### Arguments

reason	character, Reason to trigger an action.
action	character, Specifies action from one of removed, no and changed.
displayTable	data.table, A table of interest
displayColumn	character, Specifies which column(s) will be displayed in the log file.
changedVariable,	
	charcater, Specifies the variable that has been modified, must be present if action is changed.
fromTo	character, This is two vectors character. Specifies columns that before and after alteration. Must be present when action is set as changed.

### Value

A tring of text

### Author(s)

Yong Luo

---

logMatrixAdjustment	<i>Adjust log length matrix</i>
---------------------	---------------------------------

---

### Description

This function is to adjust log length matrix based on tree height, minimum log length and default log length. This function is equivalent to vol\_tree\_log\_validation macro in original sas compiler.

### Usage

```
logMatrixAdjustment(
  logLengthMatrix,
  height,
  stumpHeight,
  logMinLength,
  logDefaultLength
)

## S4 method for signature 'data.table,numeric,numeric,numeric,numeric'
logMatrixAdjustment(
  logLengthMatrix,
  height,
  stumpHeight,
```

```

    logMinLength,
    logDefaultLength
)

## S4 method for signature 'data.table,numeric,missing,numeric,numeric'
logMatrixAdjustment(logLengthMatrix, height, logMinLength, logDefaultLength)

## S4 method for signature 'data.table,numeric,numeric,missing,numeric'
logMatrixAdjustment(logLengthMatrix, height, stumpHeight, logDefaultLength)

## S4 method for signature 'data.table,numeric,numeric,numeric,missing'
logMatrixAdjustment(logLengthMatrix, height, stumpHeight, logMinLength)

## S4 method for signature 'data.table,numeric,missing,missing,numeric'
logMatrixAdjustment(logLengthMatrix, height, logDefaultLength)

## S4 method for signature 'data.table,numeric,missing,numeric,missing'
logMatrixAdjustment(logLengthMatrix, height, logMinLength)

## S4 method for signature 'data.table,numeric,missing,missing,missing'
logMatrixAdjustment(logLengthMatrix, height)

```

### Arguments

logLengthMatrix	data.table, A matrix of log length for each tree, NA is accepted in the matrix. The order of log from bottom to top must be presented from left to right in the table
height	numeric, Tree height
stumpHeight,	numeric, Stump height. If missing, 0.3 m is used.
logMinLength	numeric, Minimum log length. If missing, 3 m is used.
logDefaultLength	numeric, Default log length. If missing, 5 m is used.

### Value

A data.table that contains the matrix of adjusted log length

### Author(s)

Yong Luo

---

lookupCheck

*Test whether the lookup table is updated*


---

### Description

Reports whether a lookup table is updated. This function is highly recommended before proceed the VRI compiler.

**Usage**

```
lookupCheck(lookupName, lookupPath)

## S4 method for signature 'character,character'
lookupCheck(lookupName, lookupPath)

## S4 method for signature 'character,missing'
lookupCheck(lookupName)
```

**Arguments**

lookupName	character, Specifies the name of lookup table.
lookupPath	character, Path that directs to lookup tables.

**Value**

Not value returned. A warning message is given if the lookup table is changed.

**Author(s)**

Yong Luo

---

measNumCorrect_PSP	<i>Measurement number correction for raw PSP data</i>
--------------------	---

---

**Description**

This function is to correct the measurement number for the remeasured PSP data. The remeasured PSP data is defined as PSP that sits outside of GYS oracle database. In those remeasurement data, some measurement number has not been correctly assigned, and conflicts the measurement number for the same samples in GYS oracle database.

**Usage**

```
measNumCorrect_PSP(sampleID, measureNumber)
```

**Arguments**

sampleID	character, Specifies how the PSP sample ID.
measureNumber	numeric, Specifies measurement number before correction.

**Value**

corrected measurement number

**Note**

with confirmation by Rene

**Author(s)**

Yong Luo



---

mergeAllVolTrees	<i>Merge all volume trees-VRI specific</i>
------------------	--

---

### Description

Merge all the volume trees, including full trees (fully-measure trees in IPC), enhanced trees (fully-measured trees in auxi plots), H-enhanced trees (Height measured in auxi plots) and non-enhanced trees (only DBH measured in auxi plots). The function is part of `vol_ha_2017.sas` and modified dramatically in R compiler.

### Usage

```
mergeAllVolTrees(treeMS, treeAX)

## S4 method for signature 'data.table,data.table'
mergeAllVolTrees(treeMS, treeAX)
```

### Arguments

treeMS	data.table, Compiled full, enhanced and H-enhanced trees. This data should be listed in <code>vi_c</code> table. This data is an output of <a href="#">DWBCompiler</a>
treeAX	data.table, Non-enhanced trees in auxilary plots ( <code>vi_i</code> ). Supposedly, the table only contains non-enhanced tree list. However, some enhanced and H-enhanced trees also been stored in this dataset. An output from <a href="#">VRIInit_auxTree</a> .

### Value

A data table that contains all volume trees without duplicates. Equivalent to `tree_vb` table.

### Author(s)

Yong Luo

---

mergeOaData	<i>Merge oracle and ascii data for <code>vi_(a, ..i)</code></i>
-------------	---

---

### Description

This function is to merge oracle and ascii data before the VRI compiler

### Usage

```
mergeOaData(
  oracleSourcePath,
  asciiSourcePath,
  coeffPath,
  fizmapPath,
  fizmapName,
  fizmapFormat,
  outputPath
)
```

**Arguments**

oracleSourcePath	character, Specifies the path that stores data from oracle data base. In VRI compiler, this should be the savePath for <a href="#">loadVGIS</a> .
asciiSourcePath	character, Specifies the path that stores data from ascii data base. In VRI compiler, this should be the savePath for <a href="#">loadASCII</a> .
coeffPath	character, Specifies the path that stores coefficients and spatial lookup tables.
fizmapPath	character, Specifies the path to forest inventory zone map. By default, it is set to //spatialfiles2.bcgov/work/for/vic/hts/dam/workarea/data/infrastructure, which is maintained by FAIB employee.
fizmapName	character, Specifies the name of forest inventory zone map. By default, it is set to FIZ_REG_COMPARTMENT, which is maintained by FAIB employee.
fizmapFormat	character, Specifies the format of forest inventory zone map. Currently, it can be specified as gdb for geodatabase format and shp for shapefile format. By default, it is set to gdb, which is maintained by FAIB employee.
outputPath	character, Specifies the path to save your outputs. If missing, the current working directory will be choosed.

**Value**

no item returned

**Author(s)**

Yong Luo

---

netVolumeCalculator	<i>Calcualtes total net volume and merchantable volume-VRI specific</i>
---------------------	---

---

**Description**

This function calculates total net volume and net merchantable volume for each tree based on ground called sound percentage. From the second column to the last column, the grossVolMatrix table should have same dimensions (i.e., number of rows and columns) of netFactorMatrix and grossMerchVolMatrix if they are provided. Furthermore, be aware of the correspondingness among the matrix. This function is part of log\_valu\_2017.sas.

**Usage**

```
netVolumeCalculator(grossVolMatrix, grossMerchVolMatrix, netFactorMatrix)

## S4 method for signature 'data.table,data.table,data.table'
netVolumeCalculator(grossVolMatrix, grossMerchVolMatrix, netFactorMatrix)

## S4 method for signature 'missing,data.table,data.table'
netVolumeCalculator(grossMerchVolMatrix, netFactorMatrix)

## S4 method for signature 'data.table,missing,data.table'
```

```
netVolumeCalculator(grossVolMatrix, netFactorMatrix)

## S4 method for signature 'data.table,data.table,missing'
netVolumeCalculator(grossVolMatrix, grossMerchVolMatrix)
```

### Arguments

**grossVolMatrix** data.table, Calculated gross volume for each log. The first column of this table is the volume for the stump. If missing, the function calculates the total net merchantable volume.

**grossMerchVolMatrix** data.table, Calculated gross merchantable volume for each log. If missing, all the merchantable volume is assigned as 0.

**netFactorMatrix** data.table, Ground call for sound percentage. If missing, the net factoring will be assigned as 100.

### Value

Data table that contains total net volume (VOL\_NET) and total net merchantable volume (VOL\_NETM) for each tree.

### Author(s)

Yong Luo

---

prj\_ID2BEC

---

Assign BEC based on project ID

---

### Description

This function takes lookup table that connects project id to BEC zone and joins BEC zone by project id. The function uses a hardcoded lookup table `vri_bec`. The function is equivalent to `group_bec.sas`.

### Usage

```
prj_ID2BEC(projectID)

## S4 method for signature 'character'
prj_ID2BEC(projectID)
```

### Arguments

**projectID** character, Specifies project ID.

### Value

BEC, Unknown will be return if project id does have any match in lookup table.

### Author(s)

Yong Luo

---

prj_ID2Grp	<i>Group project ID into project group</i>
------------	--

---

### Description

This function takes lookup table that connect project id to project group and joins project group by project id. The function uses hardcoded lookup table vri\_grp. The function is equivalent to group\_prj.sas.

### Usage

```
prj_ID2Grp(projectID)

## S4 method for signature 'character'
prj_ID2Grp(projectID)
```

### Arguments

projectID      character, Specifies project ID.

### Value

Project group, Unknown will be return if project id does have any match in lookup table.

### Author(s)

Yong Luo

---

readHandHeld	<i>load json data from handheld and save tables into target folder</i>
--------------	--

---

### Description

This function is to load JSON file data from FAIB new handheld.

### Usage

```
readHandHeld(
  fileName,
  savePath = getwd(),
  saveFormat = "rdata",
  saveName = "myHandHeldFile",
  overWrite = FALSE
)
```

**Arguments**

fileName	character, JSON file name.
savePath	character, A folder path user wants to save all the outputs. Default is the current work directory.
saveFormat	character, Specifies the format user wants to save. Currently, the function supports <code>xlsx</code> and <code>rdata</code> . Default is <code>rdata</code> .
saveName	character, Specifies the save name.
overWrite	logical, Determine if the file with same name as user specifies will be overwritten. Default is <code>FALSE</code> .

**Value**

No value will be returned. All the outputs will be saved in a target folder.

**Author(s)**

Yong Luo

**Examples**

```
## Not run:
## given, in the current work directory, there is a ismc-file-transfer.json file
icmcjsonFile <- "ismc-file-transfer.json"
## create a folder in current work directory and save the outputs into it
dir.create("ismc_test_folder")

readHandHeld(fileName = icmcjsonFile,
              savePath = file.path(getwd(), "ismc_test_folder"))

## End(Not run)
```

---

readHandHeld_mod	<i>load json data from handheld and save tables into target folder</i>
------------------	--

---

**Description**

This function is to load JSON file data from FAIB new handheld.

**Usage**

```
readHandHeld_mod(
  fileName,
  savePath = getwd(),
  saveFormat = "rdata",
  saveName = "myHandHeldFile",
  overWrite = FALSE
)
```

**Arguments**

fileName	character, JSON file name.
savePath	character, A folder path user wants to save all the outputs. Default is the current work directory.
saveFormat	character, Specifies the format user wants to save. Currently, the function supports xlsx and rdata. Default is rdata.
saveName	character, Specifies the save name.
overWrite	logical, Determine if the file with same name as user specifies will be overwritten. Default is FALSE.

**Value**

No value will be returned. All the outputs will be saved in a target folder.

**Author(s)**

Yong Luo

**Examples**

```
## Not run:
## given, in the current work directory, there is a ismc-file-transfer.json file
icmcjsonFile <- "ismc-file-transfer.json"
## create a folder in current work directory and save the outputs into it
dir.create("ismc_test_folder")

readHandHeld_mod(fileName = icmcjsonFile,
                  savePath = file.path(getwd(), "ismc_test_folder"))

## End(Not run)
```

---

regBA\_WSV

---

*To fit basal area and whole stem volume regression using mixed effect model*


---

**Description**

This function takes the selected data for fitting regression model between basal area and whole stem volume using mixed effect model. The models are fitted based on strata of BEC+SP0+LV\_D

**Usage**

```
regBA_WSV(inputData, needCombs)
```

**Arguments**

inputData	data.table, The data for fitting regression.
needCombs	data.table, The combinations of BEC+SP0+LV_D are needed to derive coefficients.

**Value**

coefficient table

**Author(s)**

Yong Luo

---

regRatioDataSelect	<i>Select the data for regression and ratio</i>
--------------------	---

---

**Description**

This function selects the data for fitting regression model between basal area and whole stem volume, and for ratios to whole stem volume in VRI compiler. The regression method is used to derive whole stem volume for the trees that just have DBH information. The ratio method is used to derive netted merchantable volume for the trees that do not have call grading information. The data selection should be done annually at the end of every March to allow newer and higher quality data enter the data.

**Usage**

```
regRatioDataSelect(sampledata, alltreedata, usage)
```

**Arguments**

sampledata	data.table, The data contains the sample level information.
alltreedata	data.table, The is the data pool which contains the full/enhanced trees and H-enhanced trees.
usage	character, Specifies compiler name. It supports ismc and vgis.

**Value**

Selected data for regression and ratio.

**Author(s)**

Yong Luo

---

riskGroupDeriver	<i>Derive risk group for standard sample compilation/data</i>
------------------	---

---

### Description

will refine. This function is equivalent to risk\_grp.sas macro.

### Usage

```
riskGroupDeriver(species, pathIndex, series, height, method)

## S4 method for signature 'character,character,character,numeric,character'
riskGroupDeriver(species, pathIndex, series, height, method)

## S4 method for signature 'character,character,missing,missing,character'
riskGroupDeriver(species, pathIndex, series, height, method)
```

### Arguments

species	character, Tree basic species code, which is SP0 in VRI original data.
pathIndex	character, A character with length of 8, consists of 0 or 1.
series	character, DWB series. It is a length of 2 number character and can be derived using <a href="#">getDWBSeries</a> function.
height	numeric, Total tree height.
method	character, Specifies the method between FIZ and KBEC to categorize the risk group. The FIZ method derives risk group by species, pathIndex, series and height. KBEC method derives the risk group using species and pathIndex.

### Value

Risk group, which is character

### Author(s)

Yong Luo

---

siteAgeCompiler	<i>Compile breast age, total age, and site index where possible-VRI specific</i>
-----------------	--

---

### Description

This function takes site age tree data ie., vi\_h, an output of [VRIInit\\_siteTree](#) to compute the breast height age, total age, and site index where possible. This function is equivalent to site\_age.sas. The function heavily depends on site tools program.



**Usage**

```
siteAgeCompiler(siteAgeData)

## S4 method for signature 'data.table'
siteAgeCompiler(siteAgeData)
```

**Arguments**

siteAgeData      data.table, Site age data with plot header information. An output from [VRIInit\\_siteTree](#) function.

**Value**

A data table and a log file.

**Author(s)**

Yong Luo

---

siteAgeSummary	<i>Summarize site age data by cluster and cluster/species-VRI specific</i>
----------------	--

---

**Description**

This function takes compiled site age tree data, an output of [siteAgeCompiler](#), to derive mean age and height results. The compiled data must have breast height age, total age, and site index. This function is equivalent to mean\_ht1.sas.

**Usage**

```
siteAgeSummary(cp1dSiteAgeData)

## S4 method for signature 'data.table'
siteAgeSummary(cp1dSiteAgeData)
```

**Arguments**

cp1dSiteAgeData      data.table, Compiled site age tree data, an output of [siteAgeCompiler](#).

**Value**

Two data tables: cl\_ah is the age/height summary at cluster level and spc\_ah is the age/height summary at cluster and species level

**Author(s)**

Yong Luo

---

```
siteToolsSpeciesConvertor
```

```
Convert VRI species code to site tools species code
```

---

### Description

This function converts BC VRI species code to site tools species code. The conversion is based on a hardcoded lookup table spv\_frd.

### Usage

```
siteToolsSpeciesConvertor(species)
```

```
## S4 method for signature 'character'
siteToolsSpeciesConvertor(species)
```

### Arguments

```
species          character, Species code in VRI data sets.
```

### Value

Converted species codes that can be recognized by site tools program. NA is the species that failed to be converted.

### Author(s)

```
Yong Luo
```

---

```
SiteTools_HTBoredAge2SI
```

```
Calculate site index using site tools
```

---

### Description

This function calculates site index based on bored age (boredAge), tree height (height), species (species) and region (ICRegion) using site tools program. This function is equivalent to sin-dex\_httoage.sas.

### Usage

```
SiteTools_HTBoredAge2SI(boredAge, height, species, ICRegion, ageType, estType)
```

```
## S4 method for signature
## 'numeric,numeric,character,character,numeric,numeric'
SiteTools_HTBoredAge2SI(boredAge, height, species, ICRegion, ageType, estType)
```

Arguments

boredAge	numeric, Age at bored height.
height	numeric, Total tree height.
species	character, Species code, must be consistent with the species code in site tools, which can be converted from the original species code by using <a href="#">siteToolsSpeciesConvertor</a> .
ICRegion	character, Must be either I (interior) and C (coastal). IC regions can be derived using <a href="#">BEC2IC</a> .
ageType	numeric, Must be either 0 or 1. 0 stands for total age, for which site index is calculated for 50 years of total tree age. While 1 stands for breast height age, for which site index is calculated for 50 year old at breast height.
estType	numeric, Defines how the site tools estimate site index. Valued as 0 and 1, 0 is interactive and while 1 is directive. Default is 1, which is directive.

Value

Site index

Author(s)

Yong Luo

---

SiteTools_Y2BH	<i>Derive years to breast height using site tools</i>
----------------	---

---

Description

Derive years to breast height based on species (species), region (ICRegion) and site index (siteIndex) using site tools. This function is equivalent to `sindex_httoage.sas`.

Usage

```
SiteTools_Y2BH(species, ICRegion, siteIndex)

## S4 method for signature 'character,character,numeric'
SiteTools_Y2BH(species, ICRegion, siteIndex)
```

Arguments

species	character, Species code, must be consistent with the species code in site tools, which can be converted from the original species code by using <a href="#">siteToolsSpeciesConvertor</a> .
ICRegion	character, Must be either I (interior) and C (coastal). IC regions can be derived using <a href="#">BEC2IC</a> function.
siteIndex	numeric, Site index. Defined as tree height at 50 years old.

Value

Years to breast height

Author(s)

Yong Luo

---

smallTreeVolSmry	<i>Summarizes the volume for small trees - VRI specific</i>
------------------	---

---

### Description

Calculates and summarizes volume of small trees at both cluster/species level and cluster. This function is equivalent to `sml_tree.sas` in original compiler.

### Usage

```
smallTreeVolSmry(smallTreeData, smallTreePlotHeader)

## S4 method for signature 'data.table,data.table'
smallTreeVolSmry(smallTreeData, smallTreePlotHeader)
```

### Arguments

`smallTreeData` data.table, Small tree data. This data is from card f, i.e., `vi_f`.  
`smallTreePlotHeader` data.table, Plot header data for stump and small tree data. The data is from card e, i.e., `vi_e`.

### Value

Two tables: `stmp_c` is summarized volume at cluster level; `stmp_cs` is summarized volume at cluster/species level.

### Author(s)

Yong Luo

---

speciesCode2speciesType	<i>Group species into deciduous and coniferous species group</i>
-------------------------	--

---

### Description

This function is to group species into deciduous and coniferous species types based on BC species code and `sp_type` lookup table. This function uses hardcoded `sp_type` lookup table.

### Usage

```
speciesCode2speciesType(species)

## S4 method for signature 'character'
speciesCode2speciesType(species)
```

### Arguments

`species` character, Tree basic species code, which is SPECIES in VRI original data.

**Value**

Species type: D-deciduous species and C-coniferous species. NA, with a warning message, is given if a species fails to be grouped.

**Author(s)**

Yong Luo

---

speciesComp_byC	<i>Calculates species composition based on cluster/species summary</i>
-----------------	--

---

**Description**

Calculates species composition at cluster level based on cluster/species summary. The cluster/species-level summaries is an output of [volSmry\\_byCS](#) function. This function is equivalent to `sp_comp.sas` in original compiler.

**Usage**

```
speciesComp_byC(CSSmryTable, basedOn, speciesMaxNO, smallTreeCompile)
```

```
## S4 method for signature 'data.table,character,numeric,logical'
speciesComp_byC(CSSmryTable, basedOn, speciesMaxNO, smallTreeCompile)
```

```
## S4 method for signature 'data.table,character,numeric,missing'
speciesComp_byC(CSSmryTable, basedOn, speciesMaxNO)
```

**Arguments**

CSSmryTable	data.table, Summarized volume components for both measured and counted trees at cluster and species level. See <a href="#">volSmry_byCS</a> for details.
basedOn	character, Specifies which component is used for species composition summary.
speciesMaxNO	numeric, Maximum number of species entries to calculate.
smallTreeCompile	logical, Defines whether the function is used for calculate species composition for small trees. If missing, FALSE is used.

**Value**

A data table that contains species composition at cluster level.

**Author(s)**

Yong Luo

---

speciesCorrection	<i>Species correction, based on Rene and Dan's correction routine. See emails on May 12th, 2021.</i>
-------------------	--

---

### Description

This function is to correct species codes

### Usage

```
speciesCorrection(species, BEC, BEC_subzone)
```

### Arguments

species	character, Specifies the original species.
BEC	character, BEC zone.
BEC_subzone	character, BEC subzone.

### Value

corrected species code

### Author(s)

Yong Luo

---

stumpVolSmry	<i>Summarizes the volume for stumps - VRI specific</i>
--------------	--

---

### Description

Calculates stump volume at cluster/species level and cluster. This function is equivalent to `stmpvol.sas` in original compiler.

### Usage

```
stumpVolSmry(stumpData, stumpPlotHeader)

## S4 method for signature 'data.table,data.table'
stumpVolSmry(stumpData, stumpPlotHeader)
```

### Arguments

stumpData	data.table, Stump data. This data is from card g, i.e., vi_g.
stumpPlotHeader	data.table, Plot header data for stump and small tree data. The data is from card e, i.e., vi_e.

Value

Two tables: stmp\_c is summarized volume at cluster level; stmp\_cs is summarized volume at cluster/species level.

Author(s)

Yong Luo

---

toWSVRatio	<i>To toWSV ratio for each of strata</i>
------------	--

---

Description

This function takes the selected data for derive ratio for each of strata by BEC+SP0+LV\_D

Usage

```
toWSVRatio(inputData, needCombs, minDBH = 10, minObs = 30)
```

Arguments

inputData	data.table, The data for deriving ratios, that must contain full and enhanced trees.
needCombs	data.table, The combinations of BEC+SP0+LV_D are needed to derive coefficients.
minDBH	numeric, The minimum DBH for selecting trees to derive ratios. If missing, 10 cm will be used.
minObs	numeric, The minimum samples size for a stratum. If missing, 30 observations will be used.

Value

ratio table

Author(s)

Yong Luo

---

toWSVRatio_curve	<i>To toWSV ratio for each of strata</i>
------------------	--

---

### Description

This function takes the selected data for derive ratio for each of strata by BEC+SP0+LV\_D

### Usage

```
toWSVRatio_curve(inputData, needCombs, minDBH = 10, minObs = 30)
```

### Arguments

inputData	data.table, The data for deriving ratios, that must contain full and enhanced trees.
needCombs	data.table, The combinations of BEC+SP0+LV_D are needed to derive coefficients.
minDBH	numeric, The minimum DBH for selecting trees to derive ratios. If missing, 10 cm will be used.
minObs	numeric, The minimum samples size for a stratum. If missing, 30 observations will be used.

### Value

ratio table

### Author(s)

Yong Luo

---

treeVolEst_RegRatio	<i>Estimate volume for H-enhanced and non-enhanced trees-VRI specific</i>
---------------------	---

---

### Description

This function estimates the volumes for JH-enhanced and non-enhanced trees using BA-WSV equation and toWSV ratio methods. For H-enhanced trees, the whole stem volume and gross merchantable volume are already calculated directly using taper equations; and rest of volume components will be calculated using ratio method in this function. For non-enhanced trees, the whole stem volume is derived using regression equation between basal area and whole stem volume and the rest of volume components will be computed using ratio method in this function.

### Usage

```
treeVolEst_RegRatio(nonVolTrees, fixedCoeffTable, randomCoeffTable, ratioTable)
```



Arguments

nonVolTrees	data.table, H-enhanced trees and non-enhanced trees.
fixedCoeffTable	data.table, Specifies the WSV-BA equations by project group PRJ_GRP, live and dead status LV_D, stand and falling status SF_COMPILE and species code SP0. The table can be generated using <a href="#">WSV_BARegression</a> .
randomCoeffTable	data.table, Specifies the WSV-BA equations by project group PRJ_GRP, live and dead status LV_D, stand and falling status SF_COMPILE and species code SP0. The table can be generated using <a href="#">WSV_BARegression</a> .
ratioTable	data.table, Specifies toWSV ratio by project group PRJ_GRP, live and dead status LV_D, stand and falling status SF_COMPILE and species code SP0. The table can be generated using <a href="#">toWSVRatio</a> .

Value

A data table that has compiled non volume trees.

Author(s)

Yong Luo

---

updateSpatial	<i>Used for updating spatial attributes based on locations</i>
---------------	--

---

Description

Used for updating spatial attributes based on locations.

Usage

```
updateSpatial(samplesites, mapPath)
```

Arguments

samplesites	data.table A table that contains spatial loctions. Currently it is designed for accepting UTM coordinates. Therefore, zone, northing and easting must be provided.
mapPath	character The path to fiz map.

Value

a table that contains spatial attributes of bc albers, longitude/latitude, TSA, BEC, FIZ, TFL and OWNERSHIP.

Author(s)

Yong Luo

---

valueCalculator	<i>Calcualte tree value-VRI specific</i>
-----------------	--

---

### Description

This function calculates tree value for each tree based on ground called grade. From the second column to the last column, the grossVolMatrix table should have same dimensions (i.e., number of rows and columns) of callGradeMatrix and grossMerchVolMatrix if they are provided. Furthermore, be aware of the correspondingness among the matrix. In the function, two lookup table are hardcoded (i.e., spv\_spc and sp\_cost). This function is part of the log\_valu\_2017.sas.

### Usage

```
valueCalculator(species, grossVolMatrix, grossMerchVolMatrix, callGradeMatrix)

## S4 method for signature 'character,data.table,data.table,data.table'
valueCalculator(species, grossVolMatrix, grossMerchVolMatrix, callGradeMatrix)
```

### Arguments

species	character, Species codes in BC inventory system.
grossVolMatrix	data.table, Calculated gross volume for each log. The first column of this table is the volume for the stump. If missing, the function calculates the total net merchantable volume.
grossMerchVolMatrix	data.table, Calculated gross merchantable volume for each log. If missing, all the merchantable volume is assigned as 0.
callGradeMatrix	data.table, Ground call grading table.

### Value

Data table that contains net value (VAL\_NET) and net merchantable value (VAL\_MER).

### Author(s)

Yong Luo

---

volSmry_byC	<i>Summarize volume components at cluster level-VRI specific</i>
-------------	--

---

### Description

This function summarizes the cluster-level volume components using cluster/species-level summaries. The cluster/species-level summaries is an output of [volSmry\\_byCS](#) function.

**Usage**

```
volSmry_byC(volSmryByCS)

## S4 method for signature 'data.table'
volSmry_byC(volSmryByCS)
```

**Arguments**

volSmryByCS      data.table, Summarized volume components for both measured and counted trees. See [volSmry\\_byCS](#) for details.

**Value**

A data table

**Author(s)**

Yong Luo

---

volSmry_byCS	<i>Summarize volume components per hectare by cluster and species-VRI specific</i>
--------------	--

---

**Description**

Summarizes volume components per hectare by cluster and species. The function is last part of vol\_ha\_2017.sas.

**Usage**

```
volSmry_byCS(treeMC, utilLevel, weirdUtil, equation)

## S4 method for signature 'data.table,numeric,character,character'
volSmry_byCS(treeMC, utilLevel, weirdUtil, equation)

## S4 method for signature 'data.table,missing,character,character'
volSmry_byCS(treeMC, weirdUtil, equation)

## S4 method for signature 'data.table,numeric,missing,character'
volSmry_byCS(treeMC, utilLevel, equation)

## S4 method for signature 'data.table,numeric,numeric,character'
volSmry_byCS(treeMC, utilLevel, weirdUtil, equation)

## S4 method for signature 'data.table,numeric,character,missing'
volSmry_byCS(treeMC, utilLevel, weirdUtil)

## S4 method for signature 'data.table,missing,missing,missing'
volSmry_byCS(treeMC)
```

Arguments

treeMC	data.table, Tree-level compiled data for all volume trees.
utilLevel	numeric, Utilization levels. Default is 4.
weirdUtil	character, Weird util. Default is No. Otherwise need to be specified as a number.
equation	character, Specifies whether the compiler is based on KBEC or KFIZ. Default is KBEC.

Value

A data table summarizes volume components by cluster and species. Equevalent to smy\_cs.

Author(s)

Yong Luo

---

VRICompiler	<i>VRI compiler - VRI specific</i>
-------------	------------------------------------

---

Description

This function compiles VRI data by calling specific VRI functions. Unlike the original compiler (i.e., SAS compiler), the R version compiler hardcodes all the lookup tables in the compilation process. Please refer the descriptions for lookup table to see whether they are same as the original lookup table.

Usage

```
VRICompiler(  
  oracleUserName,  
  oraclePassword,  
  asciiTxtPath = "//albers/gis_tib/VRI/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/vri_raw",  
  
  compilationPath = "//albers/gis_tib/VRI/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/FromRCompiler",  
  
  coeffPath = "//albers/gis_tib/VRI/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/FromRCompiler/Coeffs",  
  
  fizmapPath = "//spatialfiles2.bcgov/work/for/vic/hts/dam/workarea/data/infrastructure",  
  fizmapName = "FIZ_REG_COMPARTMENT",  
  fizmapFormat = "gdb",  
  equation = "KBEC",  
  walkThru = TRUE,  
  logMinLength = 0.1,  
  stumpHeight = 0.3,  
  breastHeight = 1.3,  
  UTOPDIB = 10,  
  utilLevel = 4,  
  weirdUtil = "No"  
)
```

**Arguments**

oracleUserName	character, User name to access to oracle database.
oraclePassword	character, Password to access to oracle database.
asciiTxtPath	character, Path to ascii txt files. By default, the argument is specified to //albers/gis_tib/VR/IRDW. However, user can modify.
compilationPath	character, Specifies the path that stores all the data/processes. By specifying this, four folders will be created to record all the data/processes. Specifically, raw_from_oracle stores the data just after oracle and ascii without editing; compilation_sa stores key data (not all) that after editing and before volume and age compilation; compilation_db stores compiled results for volume and age compilation at both tree level and cluster level; Archive_YYYYMMDD archives all the data mentioned above for the future use or reference. By default, this path is set as //albers/gis_tib/VR/IRDW/IRDW_Data2/Work_Areas/VR_ASCII_PROD/RCompilation, which is consistent with our rdw system.
coeffPath	character, Specifies the path that stores coefficients and spatial lookup tables.
fizmapPath	character, Specifies the path to forest inventory zone map. By default, it is set to //spatialfiles2.bcgov/work/for/vic/hts/dam/workarea/data/infrastructure, which is maintained by FAIB employee.
fizmapName	character, Specifies the name of forest inventory zone map. By default, it is set to FIZ_REG_COMPARTMENT, which is maintained by FAIB employee.
fizmapFormat	character, Specifies the format of forest inventory zone map. Currently, it can be specified as gdb for geodatabase format and shp for shapefile format. By default, it is set to gdb, which is maintained by FAIB employee.
equation	character, Specifies the taper equation that is used for compiler. Currently supports BEC-based (KBEC) and FIZ-based (KFIZ).
walkThru	logical, Specifies whether the data had been collected using work through method. Default is TRUE, if it is not specified.
logMinLength	numeric, Specifies minimum length of log when doing log length adjustment, see <a href="#">logMatrixAdjustment</a> for details. If missing 0.1 is used.
stumpHeight	numeric, Stump height. If missing 0.3 is used.
breastHeight	numeric, Breast height. If missing 1.3 is used.
UTOPDIB	numeric, Threshold inside-bark diameter for merchantable volume. If missing, UTOPDIB is 10.
utilLevel	numeric, Specifies utilization level in summarizing tree volumes at cluster and species level. Default is 4.
weirdUtil	character, Specifies weird utilization in summarizing tree volumes at cluster and species level. Default is no, if missing. Otherwise, a number should be provided.

**Value**

This function compiles data and save outputs in compilationPaths\$compilation\_db and no file is returned.

**Note**

Improvements include:

1. specifies trees in B plots as height enhanced trees
2. removes the sas-dependent sindex functions
3. introduce the SIndexR package

Currently, the compiler supports compilation for the below sample types:

- Q: Regular VRI sample with five point clusters design
- T: ??, sample protocol and sample design are same as Q samples
- B: ??, same plot layout as Q sample, with height is measured for all trees and no call grading information
- M: CMI sample, with all trees have call grading information in the field
- L: LiDAR project, same plot layout and same design but without call grading information
- Y: YSM plots, population between 15 and 50 years
- F: NFI plots, trees measured all DBH, height and call grading
- N: NVAF plots
- A: VRI audit plots

#### Author(s)

Yong Luo

#### References

VRI compiler manual

---

VRIInit_auxTree	<i>Load and select auxiliary plot trees-VRI specific</i>
-----------------	--

---

#### Description

This function loads and selects auxiliary data (vi\_i, cardi) based on cluster/plot header.

#### Usage

```
VRIInit_auxTree(clusterplotHeader, dataSourcePath)
```

#### Arguments

clusterplotHeader	data.table, Cluster and plot level attributes, an output from <a href="#">VRIInit_clusterplot</a> .
dataSourcePath	character, Specifies the path that directs to the VRI original data source, i.e., //Mayhem/GIS_TIB/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/vri_sa.

#### Value

A data table that contains auxiliary plot tree data.

#### Author(s)

Yong Luo

---

VRIInit\_clusterplot      *Load and select cluster and plot level data- VRI specific*


---

**Description**

This function prepares the cluster/plot-level inputs for VRI compiler. Specifically, it standardizes names for the variables; reports and removes the duplicate observations at cluster, cluster/plot.

**Usage**

```
VRIInit_clusterplot(dataSourcePath)
```

**Arguments**

`dataSourcePath` character, Specifies the path that directs to the VRI original data source, i.e.,  
//Mayhem/GIS\_TIB/RDW/RDW\_Data2/Work\_Areas/VRI\_ASCII\_PROD/vri\_sa.

**Value**

A data table that contains key information at cluster/plot level and compiler log file.

**Author(s)**

Yong Luo

---

VRIInit\_lossFactor      *Load and select trees that have loss factor information-VRI specific*


---

**Description**

This function loads and selects trees that have loss factor information (`vi_d`, `cardd`) based on selected trees from `vi_c`.

**Usage**

```
VRIInit_lossFactor(fullMeasuredTrees, dataSourcePath)
```

**Arguments**

`fullMeasuredTrees`  
data.table, Selected trees in `vi_c`, which includes full, enhanced and H-enhanced trees. An output of [VRIInit\\_measuredTree](#).

`dataSourcePath` character, Specifies the path that directs to the VRI original data source, i.e.,  
//Mayhem/GIS\_TIB/RDW/RDW\_Data2/Work\_Areas/VRI\_ASCII\_PROD/vri\_sa.

**Value**

A data table that contains loss factor data. A log file documents the detailed process

**Author(s)**

Yong Luo

---

VRIInit\_measuredTree     *Load and select fully measured tree data-VRI specific*


---

### Description

This function selects the tree-level data from vi\_c (cardc) based on selected cluster/plot headers. Additionally, the function calculates basal area and tree per ha factor.

### Usage

```
VRIInit_measuredTree(clusterplotHeader, dataSourcePath, walkThru = TRUE)
```

### Arguments

clusterplotHeader	data.table, Cluster and plot-level attributes, an output from <a href="#">VRIInit_clusterplot</a> .
dataSourcePath	character, Specifies the path that directs to the VRI original data source, i.e., //Mayhem/GIS_TIB/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/vri_sa.
walkThru	logical, Indicates whether walkthrough sampling protocol is used, Tree weight is determined by walkthrough method. In walkthrough method, a tree is identified as NA (no walkthrough applied), 0 for out tree (not counted), and W for double counted tree.

### Value

A data table that contains tree-level information. A log file that describes the detailed process.

### Author(s)

Yong Luo

---

VRIInit\_siteTree     *Load and select site trees-VRI specific*


---

### Description

This function connects site tree data (vi\_h, cardh) to selected cluster/plot-level data. Site tree data is located in //Mayhem/GIS\_TIB/RDW/RDW\_Data2/Work\_Areas/VRI\_ASCII\_PROD/vri\_sa

### Usage

```
VRIInit_siteTree(clusterplotHeader, dataSourcePath)
```

### Arguments

clusterplotHeader	data.table, contains cluster/plot-level attributes, an output from <a href="#">VRIInit_clusterplot</a> .
dataSourcePath	character, Specifies the path that directs to the VRI original data source, i.e., //Mayhem/GIS_TIB/RDW/RDW_Data2/Work_Areas/VRI_ASCII_PROD/vri_sa.



**Value**

A data table that contains site tree data information. A log file documents the detailed process

**Note**

VRI specific

**Author(s)**

Yong Luo

---

VRISummaries	<i>Summarize the tree-level data at cluster or cluster/species level-VRI specific</i>
--------------	---

---

**Description**

Summarizes the compiled tree data (including both enhanced tree data and non-enhanced tree data) at cluster level. This function is equivalent to the summary part in sas compiler in cp\_vegi\_2017.sas. Different from the original compiler, this function outputs the summaries by summarized components, rather than putting all together.

**Usage**

```
VRISummaries(allVolumeTrees, clusterPlotHeader, utilLevel, weirdUtil, equation)

## S4 method for signature 'data.table,data.table,numeric,character,character'
VRISummaries(allVolumeTrees, clusterPlotHeader, utilLevel, weirdUtil, equation)
```

**Arguments**

- allVolumeTrees data.table, All tree data from vi\_c and vi\_i that have been compiled with tree volume.
- clusterPlotHeader data.table, Cluster and plot-level information. An output of [VRIInit\\_clusterplot](#).
- utilLevel numeric, Utilization levels.
- weirdUtil character, Weird util. Default is No. Otherwise need to be specified as a number.
- equation character, Specifies whether the compiler is based on KBEC or KFIZ.

**Value**

Cluster and species-level volume summaries; cluster-level volume summaries; cluster-level height summaries; cluster-level species composition summaries and log file.

**Author(s)**

Yong Luo

VRIVolTree

*Calcualte tree volume-VRI specific***Description**

This function use BEC(or FIZ) and species-specific taper equation to calculate tree volume. Before calculation, the function adjusts height for broken top trees: scenario 1 (D scenario): availability of DBH, DIB at broken height; scenario 2 (H scenario): availability of projected tree height in the field. This functions also assigns the volume multiplier adjustment. The function is equivalent to `vir_vol_tree_2011` macro in original SAS compiler.

**Usage**

```
VRIVolTree(
  treeData,
  equation,
  logMinLength,
  stumpHeight,
  breastHeight,
  UTOPDIB,
  bestHeightModels,
  HTBTOPModel
)
```

**Arguments**

<code>treeData</code>	data.table, An output from <a href="#">VRIInit_measuredTree</a> function, i.e., <code>vi_c</code> data.
<code>equation</code>	character, Specifies which taper equation form will be used to calculate diameter inside bark for a given height. Must be either KBEC or KFIZ3. If missing, default is KBEC
<code>logMinLength</code>	numeric, Specifies a minimum length for a log.
<code>stumpHeight</code>	numeric, Specifies stump height. If missing, 0.3 m will be used.
<code>breastHeight</code>	numeric, Specifies breast height. 1.3 m will be used when this argument is missing.
<code>UTOPDIB</code>	numeric, Specifies minimum merchantable inside bark diameter. 10 cm is used as a default.
<code>bestHeightModels</code>	data.table, External table that contains the best height/DBH model and coefficients by becsubzone and species.
<code>HTBTOPModel</code>	character, Specifies whether the height estimate for broken top trees either from taper or from height.

**Value**

A data table

**Author(s)**

Yong Luo

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