

# Package ‘ecorest’

May 8, 2026

**Title** Conducts Analyses Informing Ecosystem Restoration Decisions

**Version** 2.0.2

**Description** Three sets of data and functions for informing ecosystem restoration decisions, particularly in the context of the U.S. Army Corps of Engineers. First, model parameters are compiled as a data set and associated metadata for over 300 habitat suitability models developed by the U.S. Fish and Wildlife Service (USFWS 1980, <<https://www.fws.gov/policy-library/870fw1>>). Second, functions for conducting habitat suitability analyses both for the models described above as well as generic user-specified model parameterizations. Third, a suite of decision support tools for conducting cost-effectiveness and incremental cost analyses (Robinson et al. 1995, IWR Report 95-R-1, U.S. Army Corps of Engineers).

**Depends** R (>= 3.5.0)

**License** CC0

**Encoding** UTF-8

**LazyData** true

**Imports** viridis, stats, graphics, grDevices

**RoxygenNote** 7.3.3

**NeedsCompilation** no

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**Repository** CRAN

**Date/Publication** 2026-03-30 19:10:02 UTC

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annualizer	<i>Time-averaged restoration project outcomes</i>
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## Description

annualizer computes time-averaged quantities based on linear interpolation.

## Usage

```
annualizer(timevec, benefits)
```

## Arguments

timevec	numeric vector of time intervals.
benefits	numeric vector of ecological output values for a single condition (e.g., future with project) to be interpolated.

## Value

A time-averaged value over the specified time horizon.

## References

Robinson R., Hansen W., and Orth K. 1995. Evaluation of environmental investments procedures manual interim: Cost effectiveness and incremental cost analyses. IWR Report 95-R-1. Institute for Water Resources, U.S. Army Corps of Engineers, Alexandria, Virginia.

## Examples

```
#Constant value through time
annualizer(c(0,50), c(100,100))
annualizer(seq(0,50), rep(100,51))

#Simple time series
annualizer(seq(0,50), seq(0,50))

#User-specified time intervals
demo.timevec <- c(0,2,20,50)
demo.ben <- c(0,100,90,80)
annualizer(demo.timevec, demo.ben)
```

---

BBfinder	<i>Identifies "best buy" actions</i>
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---

## Description

BBfinder this analysis examines the slope of the cost-effectiveness frontier to isolate how unit cost (cost/benefit) increases with increasing environmental benefit. Restoration actions with the lowest slope of unit cost are considered "best buys".

## Usage

```
BBfinder(benefit, cost, CE)
```

## Arguments

benefit	a vector of restoration benefits. Typically, these are time-averaged ecological outcomes (e.g., average annual habitat units). Often project benefits are best presented as the "lift" associated with a restoration action (i.e., the benefits of an alternative minus the benefits of a "no action" plan).
cost	a vector of restoration costs. Typically, these are monetary costs associated with a given restoration action such as project first cost or annualized economic cost. Notably, these functions are agnostic to units, so costs could also be non-monetary such as lost political capital or social costs of each alternative.
CE	numeric vector of 0's and 1's indicating whether a plan is cost-effective (1) or non-cost-effective (0). Can be derived from <code>ecorest::CEfinder</code> .

## Value

A list with summaries of all restoration actions as well as best buy plans only.

## References

Robinson R., Hansen W., and Orth K. 1995. Evaluation of environmental investments procedures manual interim: Cost effectiveness and incremental cost analyses. IWR Report 95-R-1. Institute for Water Resources, U.S. Army Corps of Engineers, Alexandria, Virginia

## Examples

```
#Identify cost-effective actions based on random vectors of benefits and costs
benefit <- runif(50,min=0,max=10)
cost <- runif(50, min=0,max=1000)
CE <- CEfinder(benefit, cost)
BBfinder(benefit, cost, CE)

#Identify cost-effective actions based on a small number of user-specified benefits and costs
restben <- c(0, 10, 5, 20, 20)
restcost <- c(0, 100, 100, 200, 150)
restCE <- CEfinder(restben, restcost)
BBfinder(restben, restcost, restCE)
```

---

CEfinder

*Finds cost-effective frontier*

---

## Description

CEfinder returns cost-effectiveness analysis for a particular set of alternatives.

## Usage

```
CEfinder(benefit, cost)
```

## Arguments

benefit	a vector of restoration benefits. Typically, these are time-averaged ecological outcomes (e.g., average annual habitat units). Often project benefits are best presented as the "lift" associated with a restoration action (i.e., the benefits of an alternative minus the benefits of a "no action" plan).
cost	a vector of restoration costs. Typically, these are monetary costs associated with a given restoration action such as project first cost or annualized economic cost. Notably, these functions are agnostic to units, so costs could also be non-monetary such as lost political capital or social costs of each alternative.

## Value

A numeric vector identifying each plan as cost-effective (1) or non-cost-effective (0). The cost-effective actions comprise the Pareto frontier of non-dominated alternatives at a given level of cost or benefit.

## References

Robinson R., Hansen W., and Orth K. 1995. Evaluation of environmental investments procedures manual interim: Cost effectiveness and incremental cost analyses. IWR Report 95-R-1. Institute for Water Resources, U.S. Army Corps of Engineers, Alexandria, Virginia

**Examples**

```
#Identify cost-effective actions based on random vectors of benefits and costs
CEfinder(runif(50,min=0,max=10), runif(50, min=0,max=1000))

#Identify cost-effective actions based on a small number of user-specified benefits and costs
restben <- c(0, 10, 5, 20, 20)
restcost <- c(0, 100, 100, 200, 150)
CEfinder(restben, restcost)
```

---

CEICApplotter

*Plots cost-effectiveness and incremental cost analysis*


---

**Description**

CEICApplotter Plots Cost-effective Incremental Cost Analysis (CEICA) in \*.jpeg format.

**Usage**

```
CEICApplotter(altnames, benefit, cost, CE, BB, figure.name)
```

**Arguments**

altnames	vector of numerics or characters as unique restoration action identifiers.
benefit	a vector of restoration benefits. Typically, these are time-averaged ecological outcomes (e.g., average annual habitat units). Often project benefits are best presented as the "lift" associated with a restoration action (i.e., the benefits of an alternative minus the benefits of a "no action" plan).
cost	a vector of restoration costs. Typically, these are monetary costs associated with a given restoration action such as project first cost or annualized economic cost. Notably, these functions are agnostic to units, so costs could also be non-monetary such as lost political capital or social costs of each alternative.
CE	numeric vector of 0's and 1's indicating whether a plan is cost-effective (1) or non-cost-effective (0). Can be derived from ecorest::CEfinder.
BB	numeric vector of 0's and 1's indicating whether a plan is a best buy (1) or not (0). Can be derived from ecorest::BBfinder.
figure.name	output figure file name structured as "filename.jpeg".

**Value**

A multi-panel \*.jpeg figure summarizing cost-effectiveness and incremental cost analyses.

**References**

Robinson R., Hansen W., and Orth K. 1995. Evaluation of environmental investments procedures manual interim: Cost effectiveness and incremental cost analyses. IWR Report 95-R-1. Institute for Water Resources, U.S. Army Corps of Engineers, Alexandria, Virginia

## Examples

```
#Identify cost-effective actions based on random vectors of benefits and costs
altnames<- paste("Alt",seq(1,50), sep="")
benefit <- runif(50,min=0,max=10)
cost <- runif(50, min=0,max=1000)
CE <- CEfinder(benefit, cost)
BB <- BBfinder(benefit, cost, CE)[[1]][,4]
CEICAplotter(altnames, benefit, cost, CE, BB, tempfile("CEICAexample",fileext=".jpeg"))
```

---

HSIarimean

*Computes Habitat Suitability Index with Arithmetic Mean*

---

## Description

HSIarimean uses arithmetic mean to combine suitability indices into an overarching habitat suitability index.

## Usage

```
HSIarimean(x)
```

## Arguments

x                    a vector of suitability indices with values ranging from 0 to 1.

## Value

A value of habitat quality from 0 to 1 ignoring NA values.

## References

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

## Examples

```
#Determine patch quality based on a vector of four suitability indices.
HSIarimean(c(0.25, 0.25, 0.25, 0.25))
```

```
#Determine patch quality based on a vector of suitability indices with an NA.
HSIarimean(c(0.25, 0.25, NA, 0.25))
```

---

HSIeqtn	<i>Computes Habitat Suitability Index based on Model-Specified Equation</i>
---------	---

---

### Description

HSIeqtn computes a habitat suitability index based on equations specified in U.S. Fish and Wildlife Service habitat suitability models contained within ecorest via HSImodels and HSImetadata. Habitat suitability indices represent an overall assessment of habitat quality from combining individual suitability indices for multiple independent variables. The function computes an overall habitat suitability index.

### Usage

```
HSIeqtn(HSImodelName, SIV, HSImetadata, exclude = NULL)
```

### Arguments

HSImodelname	a character string in quotations that must match an existing model name in HSImetadata.
SIV	a vector of suitability index values ranging from 0 to 1 used in the model specified in HSImodelName.
HSImetadata	a data frame of HSI model metadata within the ecorest package.
exclude	a vector of character strings specifying components to be excluded from calculations. Non-NULL 'exclude' inputs are only applicable to models that explicitly provide instructions for use in the note column of HSImetadata.

### Value

A numeric of the habitat suitability index ranging from 0 to 1.

### References

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

### Examples

```
#Compute patch quality for the Barred Owl model (no components)
#Allen A.W. 1982. Habitat Suitability Index Models: Barred owl. FWS/OBS 82/10.143.
#U.S. Fish and Wildlife Service. https://pubs.er.usgs.gov/publication/fwsobs82\_10\_143.
#Suitability indices relate to density of large trees, mean diameter of overstory trees,
```

```

#and percent canopy cover of overstory.
#Example suitability vectors
HSIeqtn("barredowl", c(1,1,1), HSImetadata) #c(1,1,1) should result in 1.00
HSIeqtn("barredowl", c(0.5,1,1), HSImetadata) #c(0.5,1,1) should result in 0.707
HSIeqtn("barredowl", c(0,1,1), HSImetadata) #c(0,1,1) should result in 0.00

#Compute patch quality for the Juvenile Alewife model (two components)
#Pardue, G.B. 1983. Habitat Suitability index models: alewife and blueback herring.
#U.S. Dept. Int. Fish Wildl. Serv. FWS/OBS-82/10.58. 22pp.
#Suitability indices relate to zooplankton density, salinity, and water temperature
#Example suitability vectors are c(1,1,1), c(0.5,1,1), and c(0,1,1)
HSIeqtn("alewifeJuv", c(1,1,1), HSImetadata) #c(1,1,1) should result in 1.00
HSIeqtn("alewifeJuv", c(0.5,1,1), HSImetadata) #c(0.5,1,1) should result in 0.50
HSIeqtn("alewifeJuv", c(0,1,1), HSImetadata) #c(0,1,1) should result in 0.00

#Compute patch quality for Cutthroat trout model for lacustrine habitats (7 components)
#with spawning and lacustrine habitat and with only lacustrine habitat (i.e.,
#embryo component is excluded).
#Hickman, T., and R.F. Raleigh. 1982. Habitat suitability index models:
#Cutthroat trout. U.S.D.I. Fish and Wildlife Service. FWS/OBS-82/10.5. 38 pp.
#Suitability indices relate to temperature during the warmest period of the year,
#maximum temperature during embryo development, minimum dissolved oxygen during
#the late growing season, average velocity over spawning areas, average size
#of substrate in spawning areas, annual maximal or minimal pH, and percent fines
#in the spawning area.
#Example suitability vectors are c(1,1,1,1,1,1,1), c(0.5,1,0.5,0,1,1,1) and c(1,NA,0.5,NA,NA,0.5,NA)
#c(1,1,1,1,1,1,1) should result in 1
HSIeqtn("cutthroatLacGenLtoe15C", c(1,1,1,1,1,1,1), HSImetadata)
#c(0.5,1,0.5,0,1,1,1) should result in 0
HSIeqtn("cutthroatLacGenLtoe15C", c(0.5,1,0.5,0,1,1,1), HSImetadata)
#c(1,NA,0.5,NA,NA,0.5,NA) should result in 0.63
HSIeqtn("cutthroatLacGenLtoe15C", c(1,NA,0.5,NA,NA,0.5,NA), HSImetadata, exclude=c("CE"))

```

---

HSIgeomean

*Habitat Suitability Index with Geometric Mean*


---

### Description

HSIgeomean uses geometric mean to combine suitability indices into an overarching habitat suitability index.

### Usage

```
HSIgeomean(x)
```

### Arguments

x a vector of suitability indices with values ranging from 0 to 1.

**Value**

A value of habitat quality from 0 to 1 ignoring NA values.

**References**

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

**Examples**

```
#Determine patch quality based on a vector of four suitability indices.
HSIgeomean(c(0.25, 0.25, 0.25, 0.25))
```

```
#Determine patch quality based on a vector of suitability indices with an NA.
HSIgeomean(c(0.25, 0.25, NA, 0.25))
```

```
#Determine patch quality based on a vector of suitability indices with a zero-value.
HSIgeomean(c(0.25, 0.25, 0.0, 0.25))
```

---

HSImetadata

*Habitat suitability index (HSI) model metadata*

---

**Description**

Metadata for 351 U.S. Fish and Wildlife Service Habitat suitability index (HSI) models

**Usage**

```
HSImetadata
```

**Format**

A data frame with 351 rows and 85 variables:

**model** Model name

**submodel** Model specifications

**species** Scientific nomenclature of modeled taxa

**geography** Geographic range of organism

**ecosystem** Type of habitat

**documentation** Citation of original model

**note** Conditions under which model may be applied

**website** Link to individual model source

**SIV1** Suitability index values for each organism specific condition

**SIV1B** Suitability index values for each organism specific condition

**SIV2** Suitability index values for each organism specific condition

**SIV2B** Suitability index values for each organism specific condition

**SIV3** Suitability index values for each organism specific condition

**SIV3B** Suitability index values for each organism specific condition

**SIV4** Suitability index values for each organism specific condition

**SIV4B** Suitability index values for each organism specific condition

**SIV5** Suitability index values for each organism specific condition

**SIV5B** Suitability index values for each organism specific condition

**SIV6** Suitability index values for each organism specific condition

**SIV6B** Suitability index values for each organism specific condition

**SIV7** Suitability index values for each organism specific condition

**SIV7B** Suitability index values for each organism specific condition

**SIV8** Suitability index values for each organism specific condition

**SIV8B** Suitability index values for each organism specific condition

**SIV9** Suitability index values for each organism specific condition

**SIV10** Suitability index values for each organism specific condition

**SIV11** Suitability index values for each organism specific condition

**SIV12** Suitability index values for each organism specific condition

**SIV13** Suitability index values for each organism specific condition

**SIV14** Suitability index values for each organism specific condition

**SIV15** Suitability index values for each organism specific condition

**SIV15B** Suitability index values for each organism specific condition

**SIV16** Suitability index values for each organism specific condition

**SIV16B** Suitability index values for each organism specific condition

**SIV17** Suitability index values for each organism specific condition

**SIV18** Suitability index values for each organism specific condition

**SIV19** Suitability index values for each organism specific condition

**SIV20** Suitability index values for each organism specific condition

**SIV21** Suitability index values for each organism specific condition

**SIV22** Suitability index values for each organism specific condition

**CF** Food component equation

**CRF** Food/reproduction component equation

**CRN** Roosting-nesting component equation

**CC** Cover component equation  
**CCRO** Cover roosting component equation  
**CCRF** Cover-reproduction-food component equation  
**CCF** Cover-food component equation  
**CCSF** Cover-food shrub component equation  
**CCHF** Cover-food herbaceous component equation  
**CWF** Winter food component  
**CSF** Summer food component  
**CFF** Fall food component  
**CW** Water component  
**CCB** Cover breeding component  
**CB** Brood component  
**CN** Nest component  
**CNBC** Nest-brood cover component  
**CCN** Cover nesting component  
**CP** Pair habitat component  
**CWQ** Water quality component  
**CR** Reproduction component  
**CCR** Cover reproduction component  
**CD** Disturbance component  
**COT** Other component  
**CL** Larval component  
**CEL** Embryo and larval component  
**CE** Embryo component  
**CJ** Juvenile component  
**CFr** Fry component  
**CS** Spawning component  
**CA** Adult component  
**CI** Island component  
**CIN** Interspersion component  
**CNI** Non-island component  
**CWFC** Winter cover food component  
**CFBS** Summer food brood component  
**CFSWF** Fall spring winter food component  
**CSPF** Spring food component  
**CWC** Winter cover component  
**CCFS** Fall to spring cover component  
**CSS** Substrate-suspended solids component  
**CT** Topography component  
**CTe** Temperature component  
**CJA** Juvenile adult component  
**Eqtn** HSI overarching model equation in R syntax

**Source**

<https://pubs.usgs.gov/>

---

HSImin

*Habitat Suitability Index with Minimum*

---

**Description**

HSImin uses the minimum of given suitability indices to calculate an overarching habitat suitability index.

**Usage**

HSImin(x)

**Arguments**

x                    a vector of suitability indices ranging from 0 to 1.

**Value**

A value of habitat quality from 0 to 1 ignoring NA values.

**References**

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

**Examples**

```
#Determine patch quality based on a vector of four suitability indices.  
HSImin(c(0.1, 0.25, 0.25, 0.25))
```

```
#Determine patch quality based on a vector of suitability indices with an NA.  
HSImin(c(0.1, 0.25, NA, 0.25))
```

---

HSImodels

*Habitat suitability index (HSI) models*

---

### Description

This list of data frames contains 351 U.S. Fish and Wildlife Service Habitat suitability index (HSI) models. Please note that some of the original HSI documents provide little reference data for constructing suitability curves; hence, some suitability curves are estimated using the authors' best judgement. Users should always cross-reference results with the original documentation.

### Usage

HSImodels

### Format

An object of class list of length 349.

### Details

@format A list with 351 data frames each containing an HSI model with multiple independent variables and associated habitat suitability indices (a 0 to 1 value). Data represent break points in curves with linear extrapolation between. Categorical input variables are coded as letters.

**variable1** independent variable for assessing habitat suitability

**SIV1** suitability index value relative to variable1

... additional variables and suitability indices

### Source

<https://pubs.usgs.gov/>

---

HSIplotter

*Plots habitat suitability index curves*

---

### Description

HSIplotter plots all suitability curves.

### Usage

HSIplotter(SI, figure.name)

**Arguments**

SI list, matrix, or dataframe of suitability curves ordered as parameter breakpoints and associated suitability indices for each parameter with appropriate column names. Models containing both categorical and continuous parameters must be entered as a dataframe.

figure.name output figure file name structured as "filename.jpeg".

**Value**

A multi-panel \*.jpeg figure showing all suitability curves.

**References**

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

**Examples**

```
#Build and define a matrix of the Barred Owl suitability curves
#Allen A.W. 1982. Habitat Suitability Index Models: Barred owl. FWS/OBS 82/10.143.
#U.S. Fish and Wildlife Service. https://pubs.er.usgs.gov/publication/fwsobs82_10_143.
var1 <- cbind(c(0,2,4,NA), c(0.1,1,1,NA)) #Number of trees > 51cm diameter per 0.4 ha plot
var2 <- cbind(c(0,5,20,NA), c(0,0,1,NA)) #Mean diameter of overstory trees
var3 <- cbind(c(0,20,60,100), c(0,0,1,1)) #Percent canopy cover of overstory trees
barredowl <- cbind(var1, var2, var3)
colnames(barredowl)<- c("tree.num", "tree.num.SIV",
  "avg.dbh.in", "avg.dbh.SIV", "can.cov", "can.cov.SIV")

#Create suitability curve summary plot
HSIplotter(barredowl, tempfile("BarredOwl",fileext=".jpeg"))

#Build and define a matrix of the alewifeSAEL curves
#Pardue, GB. 1983. Habitat suitability index models: alewife and blueback herring.
#U.S. Dept. Int. Fish Wildl. Serv. FWS/OBS-82/10.58. 22pp.
var1 = data.frame(subs.type.class = c("a", "b", "c", NA, NA, NA),
  subs.type.class.SIV = c(1, 0.5, 0.1, NA, NA, NA))
var2 = data.frame("avg.daily.wtr.temp.spwn.C" = c(5, 10, 15, 20, 27, 30),
  "avg.daily.wtr.temp.spwn.C.SIV" = c(0, 0, 1, 1, 0, 0))
alewifeSAEL = data.frame(var1, var2)

#Create suitability curve summary plot
HSIplotter(alewifeSAEL, tempfile("AlewifeSAEL",fileext=".jpeg"))

#Use HSIplotter with inputs from HSImodels
HSIplotter(HSImodels$barredowl, tempfile("BarredOwl",fileext=".jpeg"))
```

---

HSIwarimean

*Habitat Suitability Index with a Weighted Arithmetic Mean*

---

### Description

HSIwarimean uses a weighted arithmetic mean to combine suitability indices into an overarching habitat suitability index.

### Usage

```
HSIwarimean(x, w)
```

### Arguments

x is a vector of suitability indices ranging from 0 to 1.  
w is a vector of weights (0 to 1 values that must sum to one).

### Value

A value of habitat quality from 0 to 1 ignoring NA values.

### References

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.  
US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.  
US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

### Examples

```
#Determine patch quality based on a vector of four, equal-weight suitability indices.  
HSIwarimean(c(1, 0, 0, 0), c(0.25, 0.25, 0.25, 0.25))  
  
#Determine patch quality based on a vector of four, unequal-weight suitability indices.  
HSIwarimean(c(1, 0, 0, 0), c(1, 0, 0, 0))  
  
#Determine patch quality based on a vector of four, unequal-weight suitability indices.  
HSIwarimean(c(1, 0, 0, 0), c(0, 1, 0, 0))
```

---

 HUcalc

*Computes Habitat Quality, Quantity, and Units*


---

**Description**

HUcalc computes habitat units given a set of suitability indices, a habitat suitability index equation, and habitat quantity.

**Usage**

```
HUcalc(SI.out, habitat.quantity, HSIfunc, ...)
```

**Arguments**

SI.out	is a vector of application-specific suitability indices between 0 and 1, which can be produced from Scalc.
habitat.quantity	is a numeric of habitat size associated with these suitability indices (i.e., length, area, or volume).
HSIfunc	is a function used to combine suitability indices into a composite habitat suitability index (HSI score) (e.g., ecorest functions like HSIarimean or HSIgeomean or functions outside ecorest like max or mean)
...	optional arguments to HSIfunc.

**Value**

A vector of habitat quality, habitat quantity, and index units (quantity times quality).

**References**

US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.

US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.

US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

**Examples**

```
#Summarize habitat outcomes based on a vector of two suitability indices
#using multiple combination equations.
HUcalc(c(0.1,1), 100, HSIarimean)
HUcalc(c(0.1,1), 100, HSIgeomean)
HUcalc(c(0.1,1), 100, HSImin)
HUcalc(c(0.1,1), 100, HSIwarimean, c(1,0))
HUcalc(c(0.1,1), 100, HSIwarimean, c(0,1))
```

```
#HSIfunc can also represent functions outside of the ecorest package
HUcalc(c(0.1,1), 100, mean)
HUcalc(c(0.1,1), 100, max)
```

---

**Sicalc***Computes Suitability Indices*

---

**Description**

Sicalc computes suitability indices given a set of suitability curves and project-specific inputs. Suitability indices may be computed based on either linear interpolation (for continuous variables) or a lookup method (for categorical variables).

**Usage**

```
Sicalc(SI, input.proj)
```

**Arguments**

SI	matrix or dataframe of suitability curves ordered as parameter breakpoints and associated suitability indices for each parameter. Suitability curves that contain both continuous and categorical variables should be formatted as a dataframe rather than a matrix.
input.proj	numeric or categorical vector of application-specific input parameters associated with the suitability curve data from SI. Note that users should enter NA for excluded variables in HSI models.

**Value**

A vector of the suitability index values ranging from 0 to 1 that match given user inputs.

**References**

- US Fish and Wildlife Service. (1980). Habitat as a basis for environmental assessment. Ecological Services Manual, 101.
- US Fish and Wildlife Service. (1980). Habitat Evaluation Procedures (HEP). Ecological Services Manual, 102.
- US Fish and Wildlife Service. (1981). Standards for the Development of Habitat Suitability Index Models. Ecological Services Manual, 103.

**Examples**

```

#Build and define a matrix of the Barred Owl suitability curves
#Allen A.W. 1982. Habitat Suitability Index Models: Barred owl. FWS/OBS 82/10.143.
#U.S. Fish and Wildlife Service. https://pubs.er.usgs.gov/publication/fwsobs82_10_143.
var1 <- cbind(c(0,2,4,NA), c(0.1,1,1,NA)) #Number of trees > 51cm diameter per 0.4 ha plot
var2 <- cbind(c(0,5,20,NA), c(0,0,1,NA)) #Mean diameter of overstory trees
var3 <- cbind(c(0,20,60,100), c(0,0,1,1)) #Percent canopy cover of overstory trees
barredowl <- cbind(var1, var2, var3)
colnames(barredowl)<- c("tree.num", "tree.num.SIV",
  "avg.dbh.in", "avg.dbh.SIV", "can.cov", "can.cov.SIV")

#Set user input variables that should return (1, 0, 0)
input.demo1 <- c(2, 5, 20)
Sicalc(barredowl, input.demo1)

#Set user input variables that should return (1, 1, 1)
input.demo2 <- c(4, 20, 60)
Sicalc(barredowl, input.demo2)

#Set user input variables that should return (1, 1, 0.5)
input.demo3 <- c(4, 20, 40)
Sicalc(barredowl, input.demo3)

#Set user input variables that should return (0.1, 0.5, 0.5)
input.demo4 <- c(0, 12.5, 40)
Sicalc(barredowl, input.demo4)

#Suitability curves may also be drawn from HSImodels (data within ecorest)
#Import Barred Owl suitability curves with HSImodels$barredowl
#The input examples are repeated from above

#Set user input variables that should return (1, 0, 0)
Sicalc(HSImodels$barredowl, input.demo1)

#Set user input variables that should return (1, 1, 1)
Sicalc(HSImodels$barredowl, input.demo2)

#Set user input variables that should return (1, 1, 0.5)
Sicalc(HSImodels$barredowl, input.demo3)

#Set user input variables that should return (0.1, 0.5, 0.5)
Sicalc(HSImodels$barredowl, input.demo4)

#Import juvenile Alewife suitability curves with HSImodels$alewifeJuv
#Demonstrate how to enter NA for excluded variables in HSImodels
#Pardue, GB. 1983. Habitat suitability index models: alewife and blueback herring.
#U.S. Dept. Int. Fish Wildl. Serv. FWS/OBS-82/10.58. 22pp.

#Set user variables that should return (NA, NA, 1, 1, 0)
input.demo7 <- c(NA, NA, 125, 5, 5)
Sicalc(HSImodels$alewifeJuv, input.demo7)

```



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