

# Package: PEcAn.settings (via r-universe)

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**Title** PEcAn Settings package

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**LazyData** FALSE

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**Description** Contains functions to read PEcAn settings files.

**Depends** methods

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---

addSecrets	<i>Add Users secrets</i>
------------	--------------------------

---

## Description

Add secret information from ~/.pecan.xml

## Usage

```
addSecrets(settings, force = FALSE)
```

## Arguments

settings	settings file
force	Logical: add secrets even if they have been added previously?

**Details**

Copies certain sections from ~/.pecan.xml to the settings. This allows a user to have their own unique parameters also when sharing the pecan.xml file we don't expose these secrets. Currently this will copy the database and browndog sections

**Value**

will return the updated settings values

**Author(s)**

Rob Kooper

---

check.bety.version      *Check BETY Version*

---

**Description**

check to make sure BETY is up to date

**Usage**

check.bety.version(dbcon)

**Arguments**

dbcon                  database connection object

---

check.database          *Check Database*

---

**Description**

Check Database

**Usage**

check.database(database)

**Arguments**

database                  settings list to check. You'll probably use settings\$database

check.database.settings

*Check Database Settings*

---

### **Description**

Check Database Settings

### **Usage**

check.database.settings(settings)

### **Arguments**

settings          settings file

---

check.ensemble.settings

*Check ensemble Settings*

---

### **Description**

Check ensemble Settings

### **Usage**

check.ensemble.settings(settings)

### **Arguments**

settings          settings file

---

check.inputs

*Check Inputs*

---

### **Description**

check to see if inputs are specified - this should be part of the model code

### **Usage**

check.inputs(settings)

### **Arguments**

settings          settings file

---

check.model.settings    *Check Model Settings*

---

**Description**

Check Model Settings

**Usage**

check.model.settings(settings, dbcon = NULL)

**Arguments**

settings	settings file
dbcon	database connection.

---

check.run.settings    *Check Run Settings*

---

**Description**

Check Run Settings

**Usage**

check.run.settings(settings, dbcon = NULL)

**Arguments**

settings	settings file
dbcon	database connection.

check.settings            *Check Settings*

---

**Description**

Sanity checks. Checks the settings file to make sure expected fields exist. It will try to use default values for any missing values, or stop the execution if no defaults are possible.

**Usage**

```
check.settings(settings, force = FALSE)
```

**Arguments**

settings	settings file
force	Logical: Rerun check even if these settings have been checked previously?

**Details**

Expected fields in settings file are:

- pfts with at least one pft defined

**Value**

will return the updated settings values with defaults set.

**Author(s)**

Rob Kooper, David LeBauer

---

check.workflow.settings  
*Check Workflow Settings*

---

**Description**

Check Workflow Settings

**Usage**

```
check.workflow.settings(settings, dbcon = NULL)
```

**Arguments**

settings	settings file
----------	---------------

---

clean.settings	<i>Cleans PEcAn settings file</i>
----------------	-----------------------------------

---

**Description**

This will try and clean the settings file so it is ready for a new run. This will remove all run specific information and set the outdir to be 'pecan' for the next run.

**Usage**

```
clean.settings(inputfile = "pecan.xml", outputfile = "pecan.xml", write = TRUE)
```

**Arguments**

inputfile	the PEcAn settings file to be used.
outputfile	the name of file to which the settings will be written inside the outputdir.
write	Indicates whether to write the modified settings to a file.

**Value**

list of all settings as saved to the XML file(s)

**Author(s)**

Rob Kooper

**Examples**

```
## Not run:  
clean.settings('output/PEcAn_1/pecan.xml', 'pecan.xml')  
  
## End(Not run)
```

---

createMultiSiteSettings	<i>Transform Settings into multi-site MultiSettings</i>
-------------------------	---

---

**Description**

Create a MultiSettings object containing (identical) run blocks for multiple different sites

**Usage**

```
createMultiSiteSettings(templateSettings, siteIds)
```

**Arguments**

templateSettings      A [Settings](#) object that will be the template for the resulting MultiSettings.

siteIds                The site IDs to be used in the resulting MultiSettings

**Details**

Starts with a template settings object, and duplicates the run\$site block once for each specified site ID. The resulting MultiSettings is thus identical to the input, except ready to run for each site in the vector of site IDs.

**Value**

A MultiSettings object with the same settings as templateSettings but replicated run\$site blocks, one for each specified site ID.

**Author(s)**

Ryan Kelly

**Examples**

```
dontrun <- function() { # Added by Alexey Shiklomanov
  # so this doesn't run and break the build

# This isn't necessarily a fully working settings object.
# Enough to get the idea though.
# Note it has a $run block with settings that will be shared across all sites

template <- Settings(list(
  info = structure(list(
    notes = NULL, userid = "1000000005", username = "Ryan Kelly",
    date = "2016/07/13 13:23:46 -0400"),
    .Names = c("notes", "userid", "username", "date")),
  database = structure(list(
    bety = structure(
      list(user = "bety", password = "bety", host = "psql-pecan.bu.edu",
      dbname = "bety", driver = "PostgreSQL", write = "TRUE"),
      .Names = c("user", "password", "host", "dbname", "driver", "write")),
    fia = structure(
      list(user = "bety", password = "bety", host = "psql-pecan.bu.edu",
      dbname = "fia5", driver = "PostgreSQL", write = "true"),
      .Names = c("user", "password", "host", "dbname", "driver", "write"))),
    .Names = c("bety", "fia")),
  pfts = structure(list(
    pft = structure(
      list(comment = NULL, name = "temperate.Evergreen_Hardwood",
      constants = structure(list(num = "1"), .Names = "num")),
      .Names = c("comment", "name", "constants")),
    pft = structure(
      list(name = "temperate.Hydric",
```



```

        constants = structure(list(num = "2"), .Names = "num"),
        .Names = c("name", "constants")),
    .Names = c("pft", "pft")),
meta.analysis = structure(list(
  iter = "3000", random.effects = list(on = FALSE, use_ghs = TRUE),
  update = "AUTO", threshold = "1.2"),
  .Names = c("iter", "random.effects", "update", "threshold")),
ensemble = structure(list(size = "1", variable = "NPP"),
  .Names = c("size", "variable")),
model = structure(list(id = "2000000005",
  edin = "/home/rykelly/pecan/RK_files/ED2IN/ED2IN.rgit.mandifore_04",
  config.header = structure(list(
    radiation = structure(list(lai_min = "0.01"), .Names = "lai_min"),
    ed_misc = structure(list(output_month = "12"),
      .Names = "output_month")),
    .Names = c("radiation", "ed_misc")),
  phenol.scheme = "0", prerun = "module load hdf5/1.8.11",
  binary = "/usr2/postdoc/rykelly/ED2/ED/build/ed_2.1-opt"),
  .Names = c("id", "edin", "config.header", "phenol.scheme", "prerun",
    "binary")),
host = structure(list(name = "geo.bu.edu", user = "rykelly",
  folder = "/projectnb/dietzelab/pecan.data/output/rykelly",
  qsub = "qsub -V -N @NAME@ -o @STDOUT@ -e @STDERR@ -S /bin/bash",
  qsub.jobid = "Your job ([0-9]+) .*",
  qstat = "qstat -j @JOBID@ || echo DONE",
  prerun = "module load udunits R/R-3.0.0-gnu-4.4.6",
  dbfiles = "/projectnb/dietzelab/pecan.data/input",
  modellauncher = structure(list(
    binary = "/usr2/postdoc/rykelly/pecan/utils/modellauncher/modellauncher",
    qsub.extra = "-pe omp 20"),
    .Names = c("binary", "qsub.extra")),
  .Names = c("name", "user", "folder", "qsub", "qsub.jobid", "qstat",
    "prerun", "dbfiles", "modellauncher")),
run = structure(list(
  inputs = structure(list(
    met = structure(list(source = "NARR", output = "ED2"),
      .Names = c("source", "output")),
    lu = structure(list(id = "294",
      path = "/projectnb/dietzelab/EDI/ed_inputs/glu/"),
      .Names = c("id", "path")),
    soil = structure(list(id = "297",
      path = "/projectnb/dietzelab/EDI/faoOLD/FAO_"),
      .Names = c("id", "path")),
    thsum = structure(list(id = "295",
      path = "/projectnb/dietzelab/EDI/ed_inputs/"),
      .Names = c("id", "path")),
    veg = structure(list(id = "296",
      path = "/projectnb/dietzelab/EDI/oge2OLD/OG2_"),
      .Names = c("id", "path")),
    pss = structure(list(source = "FIA"), .Names = "source")),
    .Names = c("met", "lu", "soil", "thsum", "veg", "pss")),
  start.date = "2004/01/01",
  end.date = "2004/01/31"),

```

```

    .Names = c("inputs", "start.date", "end.date"))
  ))

  sitegroupId <- 1000000002
  startDate <- "2000/01/01"
  endDate <- "2015/12/31"
  nSite <- 10
  outDir <- "~/multisite_setup_test"

  template <- setDates(template, startDate = startDate, endDate = endDate)
  template <- setOutDir(template, outDir)

  multiRunSettings <- createSitegroupMultiSettings(
    template,
    sitegroupId = sitegroupId,
    nSite = nSite)

  dir.create(outDir, showWarnings = FALSE)
  write.settings(multiRunSettings, outputfile = "pecan.xml")

} # dontrun

```

---

```

createSitegroupMultiSettings
      Create Sitegroup MultiSettings

```

---

## Description

Helps to create a MultiSettings object to run some or all sites in a Sitegroup.

## Usage

```

createSitegroupMultiSettings(
  templateSettings,
  sitegroupId,
  nSite,
  con = NULL,
  params = templateSettings$database$bety
)

```

## Arguments

templateSettings	A <a href="#">Settings</a> object that will be the template for the resulting MultiSettings.
sitegroupId	The Bety ID of the sitegroup to draw from
nSite	The number of sites to randomly select (without replacement) from the site-Group. Omit to use all sites in the group.
con, params	Bety DB connection or parameters. Passed directly to <a href="#">db.query</a>

**Details**

Starts with a template settings object, and fills in the run block with site info sampled from the sitegroup. The template could be fully set up except for the site info, or more or less empty if you plan to fill in the other settings later. A `MultiSettings` is created from `templateSettings`, `nSite` sites (or all of them, if `nSite` is unset) are selected from Bety, and their info is dropped into the `MultiSettings`.

**Value**

A `MultiSettings` object with the same settings as `templateSettings` but site information for the selected sites

**Author(s)**

Ryan Kelly

**Examples**

```
dontrun <- function() { # Added by Alexey Shiklomanov
  # so this doesn't run and break the build

# This isn't necessarily a fully working settings object.
# Enough to get the idea though.
# Note it has a $run block with settings that will be shared across all sites

template <- Settings(list(
  info = structure(list(
    notes = NULL, userid = "1000000005", username = "Ryan Kelly",
    date = "2016/07/13 13:23:46 -0400"),
    .Names = c("notes", "userid", "username", "date")),
  database = structure(list(
    bety = structure(
      list(user = "bety", password = "bety", host = "psql-pecan.bu.edu",
      dbname = "bety", driver = "PostgreSQL", write = "TRUE"),
      .Names = c("user", "password", "host", "dbname", "driver", "write")),
    fia = structure(
      list(user = "bety", password = "bety", host = "psql-pecan.bu.edu",
      dbname = "fia5", driver = "PostgreSQL", write = "true"),
      .Names = c("user", "password", "host", "dbname", "driver", "write")),
    .Names = c("bety", "fia")),
  pfts = structure(list(
    pft = structure(
      list(comment = NULL, name = "temperate.Evergreen_Hardwood",
      constants = structure(list(num = "1"), .Names = "num")),
      .Names = c("comment", "name", "constants")),
    pft = structure(
      list(name = "temperate.Hydric",
      constants = structure(list(num = "2"), .Names = "num")),
      .Names = c("name", "constants")),
    .Names = c("pft", "pft")),
  meta.analysis = structure(list(
```

```

    iter = "3000", random.effects = list(on = FALSE, use_ghs = TRUE),
    update = "AUTO", threshold = "1.2"),
  .Names = c("iter", "random.effects", "update", "threshold")),
ensemble = structure(list(size = "1", variable = "NPP"),
  .Names = c("size", "variable")),
model = structure(list(id = "2000000005",
  edin = "/home/rykelly/pecan/RK_files/ED2IN/ED2IN.rgit.mandifore_04",
  config.header = structure(list(
    radiation = structure(list(lai_min = "0.01"), .Names = "lai_min"),
    ed_misc = structure(list(output_month = "12"),
      .Names = "output_month")),
    .Names = c("radiation", "ed_misc")),
  phenol.scheme = "0", prerun = "module load hdf5/1.8.11",
  binary = "/usr2/postdoc/rykelly/ED2/ED/build/ed_2.1-opt"),
  .Names = c("id", "edin", "config.header", "phenol.scheme", "prerun",
    "binary")),
host = structure(list(name = "geo.bu.edu", user = "rykelly",
  folder = "/projectnb/dietzelab/pecan.data/output/rykelly",
  qsub = "qsub -V -N @NAME@ -o @STDOUT@ -e @STDERR@ -S /bin/bash",
  qsub.jobid = "Your job ([0-9]+) .*",
  qstat = "qstat -j @JOBID@ || echo DONE",
  prerun = "module load udunits R/R-3.0.0-gnu-4.4.6",
  dbfiles = "/projectnb/dietzelab/pecan.data/input",
  modellauncher = structure(list(
    binary = "/usr2/postdoc/rykelly/pecan/utis/modellauncher/modellauncher",
    qsub.extra = "-pe omp 20"),
    .Names = c("binary", "qsub.extra")),
  .Names = c("name", "user", "folder", "qsub", "qsub.jobid", "qstat",
    "prerun", "dbfiles", "modellauncher")),
run = structure(list(
  inputs = structure(list(
    met = structure(list(source = "NARR", output = "ED2"),
      .Names = c("source", "output")),
    lu = structure(list(id = "294",
      path = "/projectnb/dietzelab/EDI/ed_inputs/glu/"),
      .Names = c("id", "path")),
    soil = structure(list(id = "297",
      path = "/projectnb/dietzelab/EDI/faoOLD/FAO_"),
      .Names = c("id", "path")),
    thsum = structure(list(id = "295",
      path = "/projectnb/dietzelab/EDI/ed_inputs/"),
      .Names = c("id", "path")),
    veg = structure(list(id = "296",
      path = "/projectnb/dietzelab/EDI/oge2OLD/OG2_"),
      .Names = c("id", "path")),
    pss = structure(list(source = "FIA"), .Names = "source")),
    .Names = c("met", "lu", "soil", "thsum", "veg", "pss")),
  start.date = "2004/01/01",
  end.date = "2004/01/31"),
  .Names = c("inputs", "start.date", "end.date"))
))

```

```
sitegroupId <- 1000000002
startDate <- "2000/01/01"
endDate <- "2015/12/31"
nSite <- 10
outDir <- "~/multisite_setup_test"

template <- setDates(template, startDate = startDate, endDate = endDate)
template <- setOutDir(template, outDir)

multiRunSettings <- createSitegroupMultiSettings(
  template,
  sitegroupId = sitegroupId,
  nSite = nSite)

dir.create(outDir, showWarnings = FALSE)
write.settings(multiRunSettings, outputfile = "pecan.xml")

} # dontrun
```

---

expandMultiSettings    *generic function for expanding multi-settings.*

---

### Description

generic function for expanding multi-settings.

### Usage

```
expandMultiSettings(x)
```

### Arguments

x                    object to be expanded.

---

fix.deprecated.settings

*Fix Deprecated Settings*

---

### Description

Checks for and attempts to fix deprecated settings structure

### Usage

```
fix.deprecated.settings(settings, force = FALSE)
```

**Arguments**

settings	settings list
force	Logical: re-run fixing of deprecated settings even if it has been done previously?

**Value**

updated settings list

**Author(s)**

Ryan Kelly

---

getRunSettings	<i>Build run MultiSettings for a single site id</i>
----------------	---

---

**Description**

Processes one site from the siteIds argument of createMultiSiteSettings. You probably don't need to call it directly.

**Usage**

```
getRunSettings(templateSettings, siteId)
```

**Arguments**

templateSettings	A <a href="#">Settings</a> object that will be the template for the resulting MultiSettings.
siteId	site to process. See createMultiSiteSettings

---

get_args	<i>Get Args</i>
----------	-----------------

---

**Description**

Used in web/workflow.R to parse command line arguments. See also <https://github.com/PecanProject/pecan/pull/2626>.

**Usage**

```
get_args()
```

**Value**

list generated by [parse\\_args](#); see there for details.

**Examples**

```
## Not run: ./web/workflow.R -h
```

---

listToXml	<i>A generic function to convert list to XML</i>
-----------	--

---

**Description**

A generic function to convert list to XML

**Usage**

```
listToXml(x, ...)
```

**Arguments**

x	list to be converted
...	arguments passed to methods

---

listToXml.default	<i>List to XML</i>
-------------------	--------------------

---

**Description**

Convert List to XML

**Usage**

```
## Default S3 method:  
listToXml(x, ...)
```

**Arguments**

x	object to be converted. Despite the function name, need not actually be a list
...	further arguments. Used to set the element name of the created XML object, which is taken from an argument named tag if present, or otherwise from the first element of ...

**Details**

Can convert list or other object to an xml object using xmlNode

**Value**

xmlNode

**Author(s)**

David LeBauer, Carl Davidson, Rob Kooper

---

loadPath.sitePFT	<i>Title loadPath.sitePFT</i>
------------------	-------------------------------

---

**Description**

The csv or the text file needs to have a header and be separated using comma. Under the first column in the text file, one needs to specify the site id and in the second column there has to be the name of the PFT.

**Usage**

```
loadPath.sitePFT(settings, Path)
```

**Arguments**

settings	pecan setting list.
Path	Character of file name with extension. The path will be generated using the outdir tag in pecan settings.

**Value**

a dataframe of two columns of site and pft

---

MultiSettings	<i>Create a PEcAn MultiSettings object</i>
---------------	--

---

**Description**

Create a PEcAn MultiSettings object

**Usage**

```
MultiSettings(...)
as.MultiSettings(x)
is.MultiSettings(x)
```

**Arguments**

...	Settings objects to concatenate
x	object to test or coerce

**Value**

list with class "Multisettings"



**Functions**

- `as.MultiSettings()`: coerce an existing object to `MultiSettings`
- `is.MultiSettings()`: test if an object is a `MultiSettings`

**Author(s)**

Ryan Kelly

---

papply

*Apply functions to PEcAn MultiSettings*

---

**Description**

Works like `lapply()`, but for PEcAn `Settings` and `MultiSettings` objects

**Usage**

```
papply(settings, fn, ..., stop.on.error = FALSE)
```

**Arguments**

<code>settings</code>	A <code>MultiSettings</code> , <code>Settings</code> , or <code>list</code> to operate on
<code>fn</code>	The function to apply to <code>settings</code>
<code>...</code>	additional arguments to <code>fn</code>
<code>stop.on.error</code>	Whether to halt execution if a single element in <code>settings</code> results in error. See <code>Details</code> .

**Details**

`papply` is mainly used to call a function on each `Settings` object in a `MultiSettings` object, and returning the results in a list. It has some additional features, however:

- If the result of `fn` is a `Settings` object, then `papply` will coerce the returned list into a new `MultiSettings`.
- If `settings` is a `Settings` object, then `papply` knows to call `fn` on it directly.
- If `settings` is a generic list, then `papply` coerces it to a `Settings` object and then calls `fn` on it directly. This is meant for backwards compatibility with old-fashioned PEcAn settings lists, but could have unintended consequences
- By default, `papply` will proceed even if `fn` throws an error for one or more of the elements in `settings`. Note that if this option is used, the returned results list will have entries for *only* those elements that did not result in an error.

**Value**

A single `fn` return value, or a list of such values (coerced to `MultiSettings` if appropriate; *see Details*)

**Author(s)**

Ryan Kelly

**Examples**

```
f = function(settings, ...) {
  # Here's how I envisioned a typical use case within a standard PEcAn function
  if(is.MultiSettings(settings)) {
    return(papply(settings, f, ...))
  }

  # Don't worry about the below, it's just some guts to make the function do something we can see
  l <- list(...)
  for(i in seq_along(l)) {
    ind <- length(settings) + 1
    settings[[ind]] <- l[[i]]
    if(!is.null(names(l))) {
      names(settings)[ind] <- names(l)[i]
    }
  }
  return(settings)
}

# Example
settings1 <- Settings(list(a="aa", b=1:3, c="NA"))
settings2 <- Settings(list(a="A", b=4:5, c=paste))
multiSettings <- MultiSettings(settings1, settings2)

# The function should add element $d = D to either a Settings, or each entry in a MultiSettings
f(settings1, d="D")
print(f(multiSettings, d="D"), TRUE)
```

---

```
prepare.settings
```

```
Prepare Settings
```

---

**Description**

Update, set defaults for, and otherwise prepare a PEcAn Settings object

**Usage**

```
prepare.settings(settings, force = FALSE)
```

**Arguments**

settings	settings list
force	Whether to force the function to run even if it determines it has been run on these settings already.

**Details**

Performs various checks, fixes deprecated constructs, and assigns missing values where possible.

**Author(s)**

Ryan Kelly  
Betsy Cowdery

---

printAll	<i>generic function for printing contents of objects.</i>
----------	---

---

**Description**

generic function for printing contents of objects.

**Usage**

```
printAll(x)
```

**Arguments**

x	object to be printed.
---	-----------------------

---

read.settings	<i>Loads PEcAn settings file</i>
---------------	----------------------------------

---

**Description**

This will try and find the PEcAn settings file in the following order:

1. --settings <file> passed as command line argument using --settings
2. inputfile passed as argument to function
3. PECAN\_SETTINGS environment variable PECAN\_SETTINGS pointing to a specific file
4. ./pecan.xml pecan.xml in the current folder

**Usage**

```
read.settings(inputfile = "pecan.xml")
```

**Arguments**

inputfile	the PEcAn settings file to be used.
-----------	-------------------------------------

**Details**

Once the function finds a valid file, it will not look further. Thus, if `inputfile` is supplied, `PECAN_SETTINGS` will be ignored. Even if a file argument is passed, it will be ignored if a file is passed through a higher priority method.

**Value**

list of all settings as loaded from the XML file(s)

**Author(s)**

Shawn Serbin  
Rob Kooper  
David LeBauer  
Ryan Kelly  
Betsy Cowdery

**Examples**

```
## Not run:  
## bash shell:  
## example workflow.R and pecan.xml files in pecan/tests  
R --vanilla -- --settings path/to/mypecan.xml < workflow.R  
  
## R:  
  
settings <- read.settings()  
settings <- read.settings(file="willowcreek.xml")  
test.settings.file <- system.file("tests/test.xml", package = "PEcAn.all")  
settings <- read.settings(test.settings.file)  
  
## End(Not run)
```

---

SafeList

*Create a SafeList object*

---

**Description**

SafeList is a wrapper class for the normal R list. It should behave identically, except for the `$` operator being overridden to require exact matches.

**Usage**

```
SafeList(...)  
  
as.SafeList(x)  
  
is.SafeList(x)
```



**Details**

Sets the start/end dates in `settings$run` to the specified dates, and sets the corresponding years for `settings$ensemble` and `settings$sensitivity.analysis`. Either date can be omitted to leave it unchanged.

**Value**

The original Settings object with updated dates

**Author(s)**

Ryan Kelly

---

setOutDir	<i>Set the Output Directories of PEcAn Settings</i>
-----------	---

---

**Description**

Sets the main output directory and nulls out the others

**Usage**

```
setOutDir(settings, outDir)
```

**Arguments**

settings	A <a href="#">Settings</a> object
outDir	The desired output directory

**Details**

Sets the main output directory (`settings$outdir`) to `outDir`, and sets numerous others (`settings$modeloutdir`, `settings$host$rundir`, `settings$host$outdir`, `settings$host$modeloutdir`) to NULL so they will revert to defaults when `check.settings` is run.

**Value**

The original Settings object with updated output directories

**Author(s)**

Ryan Kelly

---

settingNames	<i>function that can retrieve or update the names of multi-settings.</i>
--------------	--

---

**Description**

function that can retrieve or update the names of multi-settings.

**Usage**

```
settingNames(multiSettings, settingNames)
```

**Arguments**

multiSettings	object for which to retrieve or set the names.
settingNames	names to be set for the multi-settings object.

---

Settings	<i>Create a PEcAn Settings object</i>
----------	---------------------------------------

---

**Description**

Create a PEcAn Settings object

**Usage**

```
Settings(...)  
as.Settings(x)  
is.Settings(x)
```

**Arguments**

...	objects to concatenate
x	object to test or coerce

**Value**

a list containing all objects in ..., with class c("Settings", "SafeList", "list").

**Functions**

- `as.Settings()`: coerce an object to Settings
- `is.Settings()`: test if object is already a Settings

**Author(s)**

Ryan Kelly

---

```
site.pft.link.settings
      site.pft.link.settings
```

---

**Description**

This function reads in a pecan setting and check for the pft.site xml tag under run>inputs. If a path or a ID for the input is defined then, it will be used for linking sites with the pfts.

**Usage**

```
site.pft.link.settings(settings)
```

**Arguments**

```
settings      settings list
```

**Value**

```
pecan xml setting file
```

---

```
site.pft.linkage      site.pft.linkage
```

---

**Description**

This function creates the required tags inside pecan.xml to link sites with pfts given a look up table. If the required tags are already defined in the pecan xml then they will be updated. If there are multiple pfts that they need to be used for a site, each pft needs to have a separate row in the lookup table, resulting multiple rows for a site.

**Usage**

```
site.pft.linkage(settings, site.pft.links)
```

**Arguments**

```
settings      pecan settings list.
site.pft.links dataframe. Your look up table should have two columns of site and pft with site
ids under site column and pft names under pft column.
```



**Value**

pecan setting list

**Examples**

```
## Not run:
#setting up the Look up tables
site.pft.links <-tribble(
  ~site, ~pft,
  "1000025731", "temperate.broadleaf.deciduous1",
  "1000025731", "temperate.needleleaf.evergreen",
  "1000000048", "temperate.broadleaf.deciduous2",
  "772", "temperate.broadleaf.deciduous3",
  "763", "temperate.broadleaf.deciduous4"
)

# sending a multi- setting xml file to the function
site.pft.linkage(settings,site.pft.links)

## End(Not run)
```

---

update.settings

*Update Settings*

---

**Description**

Updates a pecan.xml file to match new layout. This will take care of the conversion to the latest pecan.xml file.

**Usage**

```
## S3 method for class 'settings'
update(settings, force = FALSE)
```

**Arguments**

settings            settings file  
force               Logical: update even if settings have previously been updated?.

**Value**

will return the updated settings values

**Author(s)**

Rob Kooper

---

write.settings	<i>Write settings</i>
----------------	-----------------------

---

**Description**

Takes in a settings object, performs a series of checks, fixes & updates settings and produces pecan.CHECKED.xml

**Usage**

```
write.settings(settings, outputfile, outputdir = settings$outdir)
```

**Arguments**

settings	settings list
outputfile	the file name to write to
outputdir	the directory to write to

**Author(s)**

Ryan Kelly  
Betsy Cowdery

---

\$.SafeList	<i>Extract SafeList component by name</i>
-------------	---

---

**Description**

Extract SafeList component by name

**Usage**

```
## S3 method for class 'SafeList'  
x$name
```

**Arguments**

x	the SafeList object
name	the name of the component

**Details**

Overrides \$.list, and works just like it except forces exact match (i.e., makes x\$name behave exactly like x[[name, exact=T]])

**Value**

The specified component

**Author(s)**

Ryan Kelly

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