

# Package: kwb.epanet (via r-universe)

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**Title** R Package for Interfacing EPANET

**Version** 0.2.0

**Description** Functions enabling the reading and writing of EPANET  
(<http://www.epa.gov/nrmrl/wswrd/dw/epanet.html>) input files and  
reading of output files.

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**URL** <https://github.com/KWB-R/kwb.epanet>

**BugReports** <https://github.com/KWB-R/kwb.epanet/issues>

**Imports** gtools, kwb.plot, kwb.utils, lattice, plot3D

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**Repository** <https://kwb-r.r-universe.dev>

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<code>availableSections</code>	<i>Available Sections</i>
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**Description**

Names of sections available in EPANET input file

**Usage**

`availableSections(inpfile)`

**Arguments**

`inpfile`            full path to EPANET input file

**Value**

character vector containing the names of the sections contained in the EPANET input file (without brackets)

**See Also**

[readEpanetInputFile](#)

---

<code>calculateSpecificEnergyDemand</code>	<i>Calculate Specific Energy Demand</i>
--	---

---

**Description**

Calculate Specific Energy Demand

**Usage**

```
calculateSpecificEnergyDemand(
    waterDemand,
    totalEnergy,
    COLNAMES = list(Q = "Q.m3.per.hour.sum", E = "Kw.hr.per.m3.avg", Eff =
        "Average.Efficiency.avg")
)
```

**Arguments**

waterDemand	minimum constraint for water demand
totalEnergy	list element "energyTotal", as retrieved by runOptimisationStrategy
COLNAMES	list with elements $Q$ , $E$ , $Eff$ , corresponding to discharge, energy demand and efficiency, respectively. Default: list(Q = "Q.m3.per.hour.sum", E = "Kw.hr.per.m3.avg", Eff = "Average.Efficiency.avg")

**Value**

list with elements  $Q$ ,  $E$ ,  $Eff$ , holding the column names of *totalEnergy*, corresponding to discharge, energy demand and efficiency, respectively. Default: list(Q = "Q.m3.per.hour.sum", E = "Kw.hr.per.m3.avg", Eff = "Average.Efficiency.avg")

---

calibrateModel	<i>Calibrate Model</i>
----------------	------------------------

---

**Description**

Calibrate Model

**Usage**

```
calibrateModel(
  configuration,
  pipeIDs = NULL,
  measured,
  pumpsToCalibrate = NULL,
  parameterName = "Diameter",
  parameterRange,
  showLivePlot = TRUE,
  ...
)
```

**Arguments**

configuration	EPANET parameterisation, e.g. as retrieved by readEpanetInputFile()
pipeIDs	regular expression or name of pipeID(s) to be used for calibration
measured	measurement data for all pumps as data.frame e.g. data.frame(pumpNames=c("pmpW1", "pmpW2"), measuredQ=c(140,190))
pumpsToCalibrate	regular expression or name of pumps to be used for calibration: e.g. "pmpW1"
parameterName	name of ONE EPANET pipe parameters to be calibrated: e.g. Diameter or Roughness
parameterRange	min/max range of possible calibration parameter values: 0-1; with parameterRange*cun = newParameterValue, e.g. parameter range = 0.5 -> 50% reduction of initial value of parameterName for all pipeIDs defined in calibrateModel()

showLivePlot    current calibration status is plotted if showLivePlot=TRUE. Default: TRUE  
...            additional parameters to be passed to fitnessAdaptedModelConfiguration()

---

checkReportFileForErrors

*Check Report File For Errors*

---

### **Description**

Check Report File For Errors

### **Usage**

```
checkReportFileForErrors(reportFile)
```

### **Arguments**

reportFile    full path to report file

---

createOptimisationResultsTable

*Create Optimisation Results Table*

---

### **Description**

Create Optimisation Results Table

### **Usage**

```
createOptimisationResultsTable(  
  optimisationStrategies,  
  averageWaterDemand,  
  currentEnergyDemand,  
  onlyBestSolutions = FALSE  
)
```

### **Arguments**

optimisationStrategies    average daily water demand in m3/h to be satisfied  
averageWaterDemand        current specific energy demand  
currentEnergyDemand        should only the best solutions be written to data.frame? Default: FALSE  
onlyBestSolutions         return only best solutions? (default: FALSE)

---

curvesToText      *Curves To Text*

---

**Description**

Curves To Text

**Usage**

curvesToText(curves)

**Arguments**

curves      data frame representing curves, with columns *ID*, *X\_Value*, *Y\_Value*, as returned by [readEpanetInputFile](#) in list element *CURVES*

---

defaultReportVariables      *Default Report Variables*

---

**Description**

Default Report Variables

**Usage**

defaultReportVariables()

**Value**

list of report variable definitions as returned by [reportVariable](#)

---

epanetInputFileLines      *Epanet Input File Lines*

---

**Description**

Epanet Input File Lines

**Usage**

epanetInputFileLines(inpdat, dbg = FALSE)

**Arguments**

inpdat      input data to be saved in EPANET's input file format  
 dbg      if TRUE, debug messages are shown. Default: FALSE

---

exampleInputFiles	<i>Example Input Files</i>
-------------------	----------------------------

---

**Description**

Example Input Files

**Usage**

```
exampleInputFiles()
```

**Value**

full path(s) to EPANET example input file(s)

---

extdata_file	<i>Get Path to File in This Package</i>
--------------	---

---

**Description**

Get Path to File in This Package

**Usage**

```
extdata_file(..., must_exist = TRUE)
```

**Arguments**

...	parts of the file path to be passed to <a href="#">system.file</a>
must_exist	if TRUE (the default) and the specified file does not exist, the program stops with an error message

---

getEpanetInstallationPath	<i>Get Epanet Installation Path</i>
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---

**Description**

Get Epanet Installation Path

**Usage**

```
getEpanetInstallationPath()
```

---

getLinkResults      *Get Link Results*

---

**Description**

Get Link Results

**Usage**

```
getLinkResults(
  outdat,
  links,
  vars = c("q", "v", "hl", "wq", "sta", "set", "rr", "ff")
)
```

**Arguments**

outdat	output data read from EPANET output file, as returned by <a href="#">readEpanetOutputFile</a>
links	names of links to be included in the returned data frame. You may use <a href="#">getNamesOfPipes</a> , <a href="#">getNamesOfPumps</a> , <a href="#">getNamesOfValves</a> in order to get the names of available links
vars	acronyms of variables to be included in the returned data frame. "Q" = flow, "v" = velocity, "hl" = headloss, "wq" = avg. water quality, "sta" = status, "set" = setting, "rr" = reaction rate, "ff" = friction factor

---

getLinkTimeseriesFromOutputData  
*Get Link Timeseries From Output Data*

---

**Description**

Get Link Timeseries From Output Data

**Usage**

```
getLinkTimeseriesFromOutputData(outdat)
```

**Arguments**

outdat	data structure read from EPANET output file, as returned by <a href="#">readEpanetOutputFile</a> .
--------	--



---

getNamesOfCurves	<i>Get Names Of Curves</i>
------------------	----------------------------

---

**Description**

Get Names Of Curves

**Usage**

```
getNamesOfCurves(inpdat, pattern = ".*")
```

**Arguments**

inpdat	imported EPANET file (as retrieved by readEpanetInputFile)
pattern	optional filter pattern (default: ".*")

---

getNamesOfJunctions	<i>Get Names Of Junctions</i>
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---

**Description**

Get Names Of Junctions

**Usage**

```
getNamesOfJunctions(inpdat, pattern = ".*")
```

**Arguments**

inpdat	imported EPANET file (as retrieved by readEpanetInputFile)
pattern	optional filter pattern (default: ".*")

---

getNamesOfPipes	<i>Get Names Of Pipes</i>
-----------------	---------------------------

---

**Description**

Get Names Of Pipes

**Usage**

```
getNamesOfPipes(inpdat, pattern = ".*")
```

**Arguments**

inpdat	imported EPANET file (as retrieved by readEpanetInputFile)
pattern	optional filter pattern (default: ".*")

---

getNamesOfPumps      *Get Names Of Pumps*

---

**Description**

Get Names Of Pumps

**Usage**

```
getNamesOfPumps(inpdat, pattern = ".*")
```

**Arguments**

inpdat	imported EPANET file (as retrieved by readEpanetInputFile)
pattern	optional filter pattern (default: ".*")

---

getNamesOfReservoirs      *Get Names Of Reservoirs*

---

**Description**

Get Names Of Reservoirs

**Usage**

```
getNamesOfReservoirs(inpdat, pattern = ".*")
```

**Arguments**

inpdat	imported EPANET file (as retrieved by readEpanetInputFile)
pattern	optional filter pattern (default: ".*")

---

getNamesOfTanks      *Get Names Of Tanks*

---

**Description**

Get Names Of Tanks

**Usage**

```
getNamesOfTanks(inpdat, pattern = ".*")
```

**Arguments**

inpdat	imported EPANET file (as retrieved by readEpanetInputFile)
pattern	optional filter pattern (default: ".*")

---

getNamesOfValves	<i>Get Names Of Valves</i>
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---

**Description**

Get Names Of Valves

**Usage**

```
getNamesOfValves(inpdat, pattern = ".*")
```

**Arguments**

inpdat	imported EPANET file (as retrieved by readEpanetInputFile)
pattern	optional filter pattern (default: ".*")

---

getNodeResults	<i>Get Node Results</i>
----------------	-------------------------

---

**Description**

Get Node Results

**Usage**

```
getNodeResults(outdat, nodes, vars = c("d", "h", "p", "wq"))
```

**Arguments**

outdat	output data read from EPANET output file, as returned by <a href="#">readEpanetOutputFile</a>
nodes	names of nodes to be included in the returned data frame. You may use <a href="#">getNamesOfJunctions</a> , <a href="#">getNamesOfReservoirs</a> , <a href="#">getNamesOfTanks</a> in order to get the names of available nodes
vars	acronyms of variables to be included in the returned data frame. "d" = demand, "h" = head, "p" = pressure, "wq" = water quality

---

getNodeTimeseriesFromOutputData  
*Get Node Timeseries From Output Data*

---

**Description**

Get Node Timeseries From Output Data

**Usage**

getNodeTimeseriesFromOutputData(outdat)

**Arguments**

outdat            data structure read from EPANET output file, as returned by [readEpanetOutputFile](#).

---

getNumberOfPeriods    *Number of Simulation Periods*

---

**Description**

number of simulation periods, calculated from duration and hydraulic time step both of which must be given in the [TIMES] section of the EPANET configuration

**Usage**

getNumberOfPeriods(configuration)

**Arguments**

configuration    EPANET configuration, as retrieved by [readEpanetInputFile](#)

---

getPipeCoordinates    *Get Pipe Coordinates*

---

**Description**

Get Pipe Coordinates

**Usage**

getPipeCoordinates(inpdat)

**Arguments**

inpdat            imported EPANET file (as retrieved by [readEpanetInputFile](#))

---

getPumpInfo	<i>Get Pump Info</i>
-------------	----------------------

---

**Description**

Get Pump Info

**Usage**

```
getPumpInfo(inpdat)
```

**Arguments**

inpdat	imported EPANET file (as retrieved by readEpanetInputFile)
--------	--

---

getPumpPerformance	<i>Get Pump Performance</i>
--------------------	-----------------------------

---

**Description**

Get time series of pump performance from EPANET result using head curves and efficiency curves as contained in EPANET input file

**Usage**

```
getPumpPerformance(inpdata, outdata, pumppnames)
```

**Arguments**

inpdata	data structure read from EPANET input file, as returned by <a href="#">readEpanetInputFile</a> .
outdata	data structure read from EPANET output file, as returned by <a href="#">readEpanetOutputFile</a> .
pumppnames	vector with names of pumps for which the pump performance should be evaluated

**Value**

data frame with columns  $Q$  (discharge),  $H$  (head),  $Eff$  (efficiency),  $specEn$  (specific efficiency),  $En$  (energy)

**See Also**

[plotPumpPerformance](#)

---

getSection	<i>Get Section</i>
------------	--------------------

---

**Description**

Get section from EPANET input file

**Usage**

```
getSection(inpfile, sectionName)
```

**Arguments**

inpfile	full path to EPANET input file
sectionName	name of section to be read, for possible section names see the documentation of the EPANET Toolkit

**Value**

data frame representing the content of the section in the input file. If possible, column names are read from the section's header line

---

outputFileSize	<i>Size of Binary Output File</i>
----------------	-----------------------------------

---

**Description**

Size of binary output file in Bytes, kB (rounded), MB (rounded)

**Usage**

```
outputFileSize(configuration)
```

**Arguments**

configuration	EPANET configuration, representing an EPANET input file, as returned by <a href="#">readEpanetInputFile</a>
---------------	---

**Value**

named vector of numeric representing output file size in bytes, kB (rounded) and MB (rounded), respectively

---

plotCalibration	<i>Plot Calibration</i>
-----------------	-------------------------

---

**Description**

Plot Calibration

**Usage**

```
plotCalibration(newRes)
```

**Arguments**

newRes	expects data.frame object "newRes" as input parameter (is automatically produced by calibrateModel()). Required columns: <i>Qerror</i> , <i>calibrationRunNumber</i> , <i>pch</i> , <i>colors</i> , <i>pumpNames</i>
--------	--

---

plotCurves	<i>Plot Curves</i>
------------	--------------------

---

**Description**

Plot Curves

**Usage**

```
plotCurves(curves, curveNames = unique(curves$ID), ...)
```

**Arguments**

curves	data frame containing pump curves, as returned by <a href="#">readEpanetInputFile</a> in list element <i>CURVES</i>
curveNames	names of curves to be plotted. Default: all names in column <i>ID</i> of <i>curves</i>
...	additional arguments passed to <i>xyplot</i>

---

plotModel

*Plot Model*


---

### Description

plot EPANET model network. Allows plotting of changed pipe ids, which are defined as vector in parameter "changedPipeIDs"

### Usage

```
plotModel(
  inpdat,
  pch = 16,
  cex = 0.2,
  xlab = "",
  ylab = "",
  main = defaultMain(inpdat, 1),
  zoomToPumps = FALSE,
  changedPipeIDs = NULL,
  ...
)
```

### Arguments

inpdat	EPANET input file
pch	Either an integer specifying a symbol or a single character to be used as the default in plotting points. See <a href="#">points</a> for possible values and their interpretation (default: 16)
cex	A numerical value giving the amount by which plotting text and symbols should be magnified relative to the default. This starts as 1 when a device is opened, and is reset when the layout is changed, e.g. by setting mfrow. (default: 0.2)
xlab	a title for the x axis (default: "")
ylab	a title for the y axis (default: "")
main	an overall title for the plot (default: <code>kwb.epanet:::defaultMain(inpdat, 1)</code> )
zoomToPumps	should plot zoomed to pumps (default: FALSE)
changedPipeIDs	optional, vector with changed pipe IDs (default: NULL)
...	additional arguments passed to <a href="#">plot</a>



---

plotOptimisationResults *Plot Optimisation Results*

---

**Description**

Plot Optimisation Results

**Usage**

```
plotOptimisationResults(
  totalEnergy,
  name = "",
  pumpsToReplace = "",
  userConstraints,
  currentOperation,
  ...
)
```

**Arguments**

totalEnergy	list element <i>energyTotal</i> as retrieved by runOptimisationStrategy
name	name of optimisation scenario (default: "")
pumpsToReplace	optional vector with pump_ids to be replaced (default: "")
userConstraints	list of userConstraints with elements <i>namesOfWellsWithQualityProblems</i> , <i>waterDemand</i>
currentOperation	list of currentOperation with elements <i>SpecificEnergy</i> , (specific energy demand: kwh/m3) and <i>Label</i> ("Specific energy demand of current operation")
...	additional arguments passed to <a href="#">scatter2D</a>

---

plotPumpPerformance *Plot Pump Performance*

---

**Description**

Plot time series of discharge, head and pump performance

**Usage**

```
plotPumpPerformance(xCols, yCols, pumpPerformanceTimeSeries)
```

**Arguments**

xCols	vector of columns contained in data.frame retrieved by getPumpPerformance() to be used as for x axis plotting, e.g. c("step", "Q")
yCols	vector of columns contained in data.frame retrieved by getPumpPerformance() to be used as for y axis plotting, e.g. c("Eff", "specEn", "En")
pumpPerformanceTimeSeries	pump performance time series as retrieved by getPumpPerformance()

**See Also**

[getPumpPerformance](#)

---

readEpanetInputFile    *Read EPANET Input File*

---

**Description**

Read EPANET Input File

**Usage**

```
readEpanetInputFile(inpfile, dbg = FALSE)
```

**Arguments**

inpfile	full path to EPANET input file
dbg	if TRUE, debug messages are shown. Default: FALSE

**Value**

list with elements representing the different sections of the EPANET input file. The names of the list elements correspond to the names of the sections that were found in the input file. Each list element is a data frame containing the content of the corresponding section.

**See Also**

[availableSections](#), [readEpanetOutputFile](#)

---

readEpanetOutputFile    *Read EPANET Output File*

---

### Description

Read EPANET Output File

### Usage

```
readEpanetOutputFile(  
  outfile,  
  read.prolog = TRUE,  
  read.energyUse = TRUE,  
  read.dynamicResults = TRUE,  
  read.epilog = TRUE  
)
```

### Arguments

outfile	full path to EPANET output file
read.prolog	if TRUE, the "Prolog" section is read from the output file and contained in the output list
read.energyUse	if TRUE, the "Energy Use" section is read from the output file and contained in the output list
read.dynamicResults	if TRUE, the "Extended Period" section is read from the output file and contained in the output list
read.epilog	if TRUE, the "Epilog" section is read from the output file and contained in the output list

### Value

list with elements *prolog* (if read.prolog = TRUE), *energyUse* (if read.energyUse = TRUE), *dynamicResults* (if read.dynamicResults = TRUE) and *epilog* (if read.epilog = TRUE), containing the different parts of the output file, as described in the documentation of the EPANET Toolkit.

### See Also

[readEpanetInputFile](#)

---

```
readResultsFromReportFile
```

*Read Results From Report File*

---

### Description

reads an EPANET report file and returns a list with node and link data

### Usage

```
readResultsFromReportFile(reportFile, warn = TRUE)
```

### Arguments

reportFile	full path to report file, generated by EPANET
warn	if TRUE, a warning is given if no Node or Link results were found in the report file.

### Value

list with elements *nodeData* containing a data frame with node results and *linkData* containing a data frame with link results

---

```
replaceCurves
```

*Replace Curves In CURVES Data*

---

### Description

Replace curves in CURVES data

### Usage

```
replaceCurves(curves, newCurves)
```

### Arguments

curves	data frame representing curves, with columns <i>ID</i> , <i>X_Value</i> , <i>Y_Value</i> , as returned by <a href="#">readEpanetInputFile</a> in list element <i>CURVES</i>
newCurves	new curves data frame representing curves, with columns <i>ID</i> , <i>X_Value</i> , <i>Y_Value</i> ,

---

replaceCurveSectionInInputFile  
*Replace Curve Section In Input File*

---

**Description**

Replace Curve Section In Input File

**Usage**

```
replaceCurveSectionInInputFile(inpfile, newCurves)
```

**Arguments**

inpfile	full path to EPANET input file
newCurves	modified CURVES section. Data frame as returned by <a href="#">readEpanetInputFile</a> in list element <i>curves</i>

**Value**

vector of character(s) representing the rows of the input file

---

replaceOneCurve      *Replace One Curve In CURVES Data*

---

**Description**

Replace One Curve In CURVES Data

**Usage**

```
replaceOneCurve(curves, curveName, x, y)
```

**Arguments**

curves	data frame representing curves, with columns <i>ID</i> , <i>X_Value</i> , <i>Y_Value</i> , as returned by <a href="#">readEpanetInputFile</a> in list element <i>CURVES</i>
curveName	name of the curve to be replaced
x	vector of x values of the curve
y	vector of y values of the curve

**Value**

data frame in which the lines corresponding to the curve named *curveName* are replaced with x and y values given in *x* and *y*, respectively

---

reportEnergyUse	<i>Report Energy Use</i>
-----------------	--------------------------

---

**Description**

Report Energy Use

**Usage**

```
reportEnergyUse(outdat)
```

**Arguments**

outdat	output data read from EPANET out-file
--------	---------------------------------------

---

reportVariable	<i>Report Variable Definition</i>
----------------	-----------------------------------

---

**Description**

Report variable definition for argument *variables* of [setReportOptions](#)

**Usage**

```
reportVariable(
  name = "Elevation",
  yes = TRUE,
  below = NA,
  above = NA,
  precision = 2
)
```

**Arguments**

name	name of variable. Node variables that can be reported on include: "Elevation", "Demand", "Head", "Pressure", "Quality". Link variables include: "Length", "Diameter", "Flow", "Velocity", "Headloss", "LinkQuality", "LinkStatus", "Setting" (Roughness for pipes, speed for pumps,, pressure/flow setting for valves), "Reaction" (reaction rate), "FFactor" (friction factor)
yes	shall the variable be importet on?
below	if set to a non-NA value (default: NA) only values below the given value will be reported on
above	if set to a non-NA value (default: NA) only values above the given value will be reported on
precision	the variable will be reported on with the given precision (number of decimal places). Default: 2

---

runEpanet

*Run EPANET With Given Input File*


---

### Description

Run EPANET With Given Input File

### Usage

```
runEpanet(
  inpfile,
  returnOutput = FALSE,
  epanet.dir = getEpanetInstallationPath(),
  intern = FALSE,
  write.output = TRUE,
  ...,
  dbg = FALSE
)
```

### Arguments

inpfile	full path to EPANET input file
returnOutput	if TRUE, the output is read from the generated output file and returned
epanet.dir	path to EPANET installation directory. Default: getEpanetInstallationPath()
intern	a logical, indicates whether to make the output of the command an R object.
write.output	if TRUE, EPANET will write a binary output file, else not
...	further arguments passed to <a href="#">readEpanetOutputFile</a> , such as: <i>read.prolog</i> , <i>read.energyUse</i> , <i>read.dynamicResults</i> , <i>read.epilog</i> , see there.
dbg	if TRUE, debug messages are shown. Default: FALSE

---

runEpanetConfiguration

*Run EPANET INP configuration*


---

### Description

Run EPANET INP configuration

**Usage**

```
runEpanetConfiguration(
  inpdat,
  name = "tmpEpanet",
  returnOutput = write.output,
  write.output = TRUE,
  ...,
  dbg = FALSE
)
```

**Arguments**

inpdat	input data as retrieved by <a href="#">readEpanetInputFile</a>
name	name of input file to be generated in tempdir()
returnOutput	if TRUE, the output is read from the output file (if generated, see <i>write.output</i> ) and returned. Default: value of <i>write.output</i>
write.output	if TRUE, EPANET will write a binary output file, else not
...	further arguments passed to <a href="#">runEpanet</a> and finally to <a href="#">readEpanetOutputFile</a> , such as: <i>read.prolog</i> , <i>read.energyUse</i> , <i>read.dynamicResults</i> , <i>read.epilog</i> , see there.
dbg	if TRUE, debug messages are shown. Default: FALSE

---

runEpanetGUI

*Run Epanet GUI*


---

**Description**

Run Epanet GUI

**Usage**

```
runEpanetGUI(inpfile = "", epanet.dir = getEpanetInstallationPath())
```

**Arguments**

inpfile	path to EPANET input file
epanet.dir	EPANET directory (default: <code>getEpanetInstallationPath()</code> )



---

`runEpanetOnCommandLine`*Run Epanet On Command Line*

---

**Description**

Run Epanet On Command Line

**Usage**

```
runEpanetOnCommandLine(  
  inpfile,  
  epanet.exe,  
  intern = FALSE,  
  write.output = TRUE,  
  dbg = FALSE  
)
```

**Arguments**

<code>inpfile</code>	path to EPANET input file
<code>epanet.exe</code>	path to EPANET executable
<code>intern</code>	a logical, indicates whether to make the output of the command an R object. (default: FALSE)
<code>write.output</code>	if TRUE, EPANET will write a binary output file, else not
<code>dbg</code>	if TRUE, debug messages are shown. Default: FALSE

---

`runOptimisationStrategy`*Run Optimisation Strategy*

---

**Description**

Run Optimisation Strategy

**Usage**

```
runOptimisationStrategy(  
  configuration,  
  newCurvesData,  
  optimisationStrategy,  
  operationSchemes = wellFieldOperationSchemes(getNamesOfPumps(configuration)),  
  showLivePlot = FALSE  
)
```

**Arguments**

- configuration EPANET configuration, representing an EPANET input file, as returned by [readEpanetInputFile](#)
- newCurvesData list with elements *Pump* (data frame with columns "ID", "X\_VALUE" and "Y\_VALUE") of pump curves, *GlobalPumpEfficiency* (data frame with columns "ID", "X\_VALUE" and "Y\_VALUE") and sublist *PumpNamePrefix* (with elements *PumpCurves* = "TDH" and *GlobalPumpEfficiency* = "Eff" )
- optimisationStrategy list with elements *name* (name of optimisation strategy), *shortName* (short name of optimisation strategy) and *pumpsToReplace* (vector with pump-ids to be replaced, if none: "")
- operationSchemes possible wellfield operation schemes. Default: wellFieldOperationSchemes(getNamesOfPumps(configuration))
- showLivePlot (default: FALSE)

**Value**

- list with elements *energyTotal* and *energyPerPump*

---

setEpanetInstallationPath  
*Set Epanet Installation Path*

---

**Description**

Set Epanet Installation Path

**Usage**

```
setEpanetInstallationPath(epanet.dir)
```

**Arguments**

- epanet.dir full path to MS Access database or ODBC database name

---

setReportOptions	<i>Set REPORT Option in EPANET configuration</i>
------------------	--

---

### Description

Set REPORT Option in EPANET configuration

### Usage

```
setReportOptions(
    configuration,
    pagesize = 0,
    file = "",
    status = "NO",
    summary = "YES",
    messages = "YES",
    energy = "NO",
    nodes = "NONE",
    links = "NONE",
    variables = defaultReportVariables()
)
```

### Arguments

configuration	EPANET configuration, representing an EPANET input file, as returned by <a href="#">readEpanetInputFile</a>
pagesize	pagesize sets the number of lines written per page of the output report. The default is 0, meaning that no line limit per page is in effect.
file	file supplies the name of a file to which the output report will be written. If the file name contains spaces then it must be surrounded by double quotes. If not supplied then the Report file, as specified in the second parameter of the ENopen (or ENepanet) function will be used.
status	status determines whether hydraulic status messages are written to the Report file. If YES is selected the messages will identify those network components that change status during each time step of the simulation. If FULL is selected, then convergence information will also be included from each trial of each hydraulic analysis. This level of detail is only useful for de-bugging networks that become hydraulically unbalanced. The default is NO.
summary	summary determines whether a summary table of number of network components and key analysis options is generated. The default is YES.
messages	messages determines whether error and warning messages generated during a hydraulic/water quality analysis are written to the Report file. The default is YES.
energy	energy determines if a table reporting average energy usage and cost for each pump is provided. The default is NO.

nodes	nodes identifies which nodes will be reported on. You can either list individual node ID labels or use the keywords NONE or ALL. Additional NODES lines can be used to continue the list. The default is NONE.
links	links identifies which links will be reported on. You can either list individual link ID labels or use the keywords NONE or ALL. Additional LINKS lines can be used to continue the list. The default is NONE.
variables	list of report variables as defined by <a href="#">reportVariable</a> , e.g. list(reportVariable(name = "VELOCITY", above = 3.0, precision = 4), reportVariable(name = "F-FACTOR", precision = 4)). Default: <a href="#">defaultReportVariables</a>

---

setTimeParameter      *Set EPANET' Time-Related Simulation Parameters*

---

### Description

Set EPANET' time-related simulation parameters. See EPANET reference for the meaning of the parameters

### Usage

```
setTimeParameter(
  configuration,
  duration = "",
  hydraulic.timestep = "",
  quality.timestep = "",
  rule.timestep = "",
  pattern.timestep = "",
  pattern.start = "",
  report.timestep = "",
  report.start = "",
  start.clocktime = "",
  statistic = ""
)
```

### Arguments

configuration	EPANET configuration, representing an EPANET input file, as returned by <a href="#">readEpanetInputFile</a>
duration	(default: "")
hydraulic.timestep	(default: "")
quality.timestep	(default: "")
rule.timestep	(default: "")
pattern.timestep	(default: "")

```

pattern.start (default: "")
report.timestep
              (default: "")
report.start (default: "")
start.clocktime
             (default: "")
statistic    (default: "")

```

**Value**

return configuration with modified [TIMES] parameterisation

---

setWellFieldOperation *Set Well Field Operation*

---

**Description**

Set Well Field Operation

**Usage**

```
setWellFieldOperation(config, operationScheme)
```

**Arguments**

config            EPANET configuration, representing an EPANET input file, as returned by [readEpanetInputFile](#)

operationScheme    one possible wellfield operation scheme (e.g. wellFieldOperationSchemes(getNamesOfPumps(configuration)))

**Value**

config with modified well field operation rules

---

showProperties            *Show Properties*

---

**Description**

Show node and link properties available in EPANET output

**Usage**

```
showProperties(outdata)
```

**Arguments**

outdata            list structure with EPANET results as retrieved by [readEpanetOutputFile](#)

---

wellFieldOperationSchemes  
*Well Field Operation Schemes*

---

### Description

Well Field Operation Schemes

### Usage

```
wellFieldOperationSchemes(  
    pumpNames,  
    niceLabels = list(searchPattern = "pmp", searchReplacement = "p")  
)
```

### Arguments

pumpNames	Vector of pumpNames (used in EPANET input file section "PUMPS")
niceLabels	Optionally nice labeling. List with elements <i>searchPattern</i> and <i>searchReplacement</i> . Default: list(searchPattern = "pmp", searchReplacement = "p")

---

writeCurves                      *Write Curves*

---

### Description

Write Curves

### Usage

```
writeCurves(  
    epanetConfig,  
    pumpCurves = NULL,  
    efficiencyCurves = NULL,  
    drawdownCurves = NULL,  
    deleteOldCurves = TRUE,  
    dbg = TRUE  
)
```

**Arguments**

epanetConfig EPANET configuration, as retrieved by readEpanetInputFile  
 pumpCurves list(data = data.frame(ID, X\_Value, Y\_Value), label = "")  
 efficiencyCurves list(data = data.frame(ID, X\_Value, Y\_Value), label = "")  
 drawdownCurves list(data = data.frame(ID, X\_Value, Y\_Value), label = "")  
 deleteOldCurves should all curves in 'epanetConfig' be deleted before writing?  
 dbg show debug messages? (default: TRUE)

---

writeDrawdownCurves *Write Drawdown Curves*

---

**Description**

Write Drawdown Curves

**Usage**

```

writeDrawdownCurves(
  epanetConfig,
  DD,
  curveNamePrefix = "dd",
  deleteOldCurves = FALSE,
  dbg = TRUE
)

```

**Arguments**

epanetConfig EPANET configuration, as retrieved by readEpanetInputFile  
 DD data.frame(ID, X\_Value, Y\_Value)  
 curveNamePrefix prefix to be used in the curve name (default: "dd")  
 deleteOldCurves if TRUE all curves in epanetConfig\$CURVES will be deleted before adding new ones (default: FALSE)  
 dbg show debug messages? (default: TRUE)

**Value**

Modified EPANET configuration

---

writeEfficiencyCurves *Write Efficiency Curves*

---

### Description

Write Efficiency Curves

### Usage

```
writeEfficiencyCurves(
  epanetConfig,
  Eff,
  curveNamePrefix = "Eff",
  deleteOldCurves = FALSE,
  dbg = TRUE
)
```

### Arguments

epanetConfig	EPANET configuration, as retrieved by readEpanetInputFile
Eff	data.frame(ID, X_Value, Y_Value)
curveNamePrefix	prefix to be used in the curve name (default: "Eff")
deleteOldCurves	if TRUE all curves in epanetConfig\$CURVES will be deleted before adding new ones (default: FALSE)
dbg	show debug messages? (default: TRUE)

### Value

Modified EPANET configuration

---

writeEpanetInputFile *Write Epanet Input File*

---

### Description

Write Epanet Input File

### Usage

```
writeEpanetInputFile(inpdat, infile, dbg = FALSE)
```



**Arguments**

inpdatt	input data to be saved in EPANET's input file format
inpfile	full path to input file to be created
dbg	if TRUE, debug messages are shown. Default: FALSE

---

```
writeInputFileWithNewCurveSection
```

*Write Input File With New Curve Section*

---

**Description**

Write Input File With New Curve Section

**Usage**

```
writeInputFileWithNewCurveSection(
  inpfile,
  newCurves,
  inpfile.new = sub("(\\.[^\\.]*)$", "_new\\1", inpfile)
)
```

**Arguments**

inpfile	full path to EPANET input file
newCurves	modified CURVES section. Data frame as returned by <a href="#">readEpanetInputFile</a> in list element <i>curves</i>
inpfile.new	full path to modified EPANET input file. (default: <code>&lt;inpfile-without-extension_new&gt;.inp</code> )

**Value**

full path to created input file

---

```
writePumpCurves
```

*Write Pump Curves*

---

**Description**

Write Pump Curves

**Usage**

```
writePumpCurves(  
  epanetConfig,  
  TDH,  
  curveNamePrefix = "TDH",  
  deleteOldCurves = FALSE,  
  dbg = TRUE  
)
```

**Arguments**

<code>epanetConfig</code>	EPANET configuration, as retrieved by <code>readEpanetInputFile</code>
<code>TDH</code>	<code>data.frame(ID, X_Value, Y_Value)</code>
<code>curveNamePrefix</code>	(default: "TDH")
<code>deleteOldCurves</code>	if TRUE all curves in <code>epanetConfig\$CURVES</code> will be deleted before adding new ones (default: FALSE)
<code>dbg</code>	show debug messages? (default: TRUE)

**Value**

Modified EPANET configuration

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