Package 'CNORfuzzy'

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| Title Addon to CellNOptR: Fuzzy Logic |
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| Date 2013-8-28 |
| Author M. Morris, T. Cokelaer |
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| Description This package is an extension to CellNOptR. It contains additional functionality needed to simulate and train a prior knowledge network to experimental data using constrained fuzzy logic (cFL, rather than Boolean logic as is the case in CellNOptR). Additionally, this package will contain functions to use for the compilation of multiple optimization results (either Boolean or cFL). |
| License GPL-2 |
| Depends R (>= 2.15.0), CellNOptR (>= 1.4.0), nloptr (>= 0.8.5) |
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Description

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This package does optimisation of constrained Fuzzy logic networks of signalling pathways based on a previous knowledge network and a set of data collected upon perturbation of some of the nodes in the network.

Details

Package: CNOR
Type: Package
Version: 1.4.0
Date: 2013-08-28
License: GPL-2
LazyLoad: yes

Depends: R (>= 2.15.0), CellNOptR (>= 1.3.29), nloptr (>= 0.8.5)

Author(s)

M.K. Morris

Maintainer: T. Cokelaer < cokelaer@ebi.ac.uk>

References

- 1. J. Saez-Rodriguez, L. G. Alexopoulos, J. Epperlein, R. Samaga, D. A. Lauffenburger, S. Klamt and P. K. Sorger. Discrete logic modeling as a means to link protein signaling networks with functional analysis of mammalian signal transduction, Molecular Systems Biology, 5:331, 2009.
- 2. Morris MK, Saez-Rodriguez J, Clarke DC, Sorger PK, Lauffenburger DA (2011). Training Signaling Pathway Maps to Biochemical Data with Constrained Fuzzy Logic: Quantitative Analysis of Liver Cell Responses to Inflammatory Stimuli. PLoS Comput Biol. 7(3): e1001099.

See Also

CellNOptR package.

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Examples

```
# Get data from CellNOptR package
data(CNOlistToy,package="CellNOptR")
data(ToyModel,package="CellNOptR")

# Use the default parameters and set Data and Model
paramsList=defaultParametersFuzzy()
paramsList$data<-CNOlistToy
paramsList$model<-ToyModel

## Not run:
    # Run the simulator
    Res = CNORwrapFuzzy(data=CNOlistToy, model=ToyModel, paramsList=paramsList)

## End(Not run)</pre>
```

CNORwrapFuzzy

CNORfuzzy analysis wrapper

Description

This function is a wrapper around the whole CNOR Fuzzy analysis. It performs the following steps:

- 1. Plot the CNOlist
- 2. Checks data to model compatibility
- 3. Pre-processing steps
- 4. Prepare for simulation (see prep4simFuzzy)
- 5. Optimisation using Fuzzy transfer function (see gaDiscreteT1)
- 6. Refinement and reduction steps (see getRefinedModel and reduceFuzzy).

Usage

```
CNORwrapFuzzy(data, model, paramsList=NULL, verbose=TRUE)
```

Arguments

data a CNOlist structure (as created by makeCNOlist) that contains the data that you

will use (see readMIDAS and readSIF from CellNOptR).

model that you want to optimise

paramsList Use defaultParametersFuzzy function to create a template. Entries are 3-types:

(i) GA algorithm parameters for the optimisation, (ii) Fuzzy parameters for the transfer functions and (iii) internal optimisation parameters within the refine-

ment step.

See defaultParametersFuzzy function for details on the parameters.

verbose

4 compileMultiRes

Details

If you do not provide a parameter list argument, defaultParametersFuzzy is called internally to populate the paramsList argument.

Value

This function returns an object containing the results that can be used by other functions such as compileRes.

Author(s)

M.K. Morris

References

CNORWrap function from CellNOptR package

See Also

defaultParametersFuzzy, compileMultiRes

Examples

```
# Load some data
data(CNOlistToy, package="CellNOptR")
data(ToyModel, package="CellNOptR")
# Get some default parameters to play with, limiting the duration of the GA
# algorithm and optimisation step
paramsList = defaultParametersFuzzy()
paramsList$maxTime = 20
paramsList$optimisation$maxtime = 10
results = CNORwrapFuzzy(CNOlistToy, ToyModel, paramsList)
```

 ${\tt compile} {\tt MultiRes}$

Compiles results from multiple runs and produces graph for choosing Post Refinement Threshold

Description

This function takes a list of objects returned by CNORwrapfuzzy (run using identical parameters, models, and data) and packages them together so they can be compared with plotMeanFuzzyFit and writeFuzzyNetwork. Because almost all training of cFL models are underdetermined problems, analyzing multiple runs together is essential.

Usage

```
compileMultiRes(allRes, tag=NULL, show=TRUE)
```

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Arguments

allRes list of objects returned by the CNORwrapFuzzy function.

tag If provided, save the results in 3 files. Each file starts with the string "file-

name" that is provided. (<tag>_allRes.RData, <tag>_allFinalMSEs.RData and

<tag>_allFinalNumParams.RData)

show plot the MSE and mean number of parameters versus threshold. Can be switch

off if show=FALSE

Author(s)

M.K. Morris, T. Cokelaer

Examples

```
data(ToyModel, package="CellNOptR")
data(CNOlistToy,package="CellNOptR")
paramsList = defaultParametersFuzzy(CNOlistToy, ToyModel)
N = 10
allRes = list()
## Not run:
    for (i in 1:N){
        Res = CNORwrapFuzzy(CNOlistToy, ToyModel, paramsList)
        allRes[[i]] = Res
    }

summary = compileMultiRes(allRes)
summary$allFinalMSEs
summary$allFinalNumParams

# You can save the resuls in files using the tag argument
compileMultiRes(allRes, "output")

## End(Not run)
```

computeScoreFuzzy

Compute Score of a model compared to the data for a given intString.

Description

compute and return score of a model (cut using a bitstring).

Usage

```
computeScoreFuzzy(CNOlist, model,simList=NULL, indexList=NULL, paramsList,
  intString=NULL, sizeFac=0.0001,NAFac=1)
```

Arguments

cNolist a CNolist on which the score is based (based on valueSignals[[2]], i.e. data at

t1)

model a model list

simList a list that contains additional fields for the simulator, as created by prep4sim

applied to the model above

indexList a list of indexes of species stimulated/inhibited/signals, as produced by indexfinder

applied on the model and CNOlist above

paramsList list of parameters. See defaultParametersFuzzy

intString a bitstring of the same size as the number of reactions in the model above sizeFac the scaling factor for the size term in the objective function, default to 0.0001 NAFac the scaling factor for the NA term in the objective function, default to 1

Value

score See gaBinaryT1 for details

Author(s)

T. Cokelaer

defaultParametersFuzzy

Create a list of default parameters

Description

Parameters are required at different levels in the Fuzzy optimisation. This function provides a list with all parameters that are necessary.

Usage

defaultParametersFuzzy(data=NA, model=NA, nTF=7)

Arguments

data the CNOlist that contains the data that you will use

model that you want to optimise

nTF number of discrete values to be used for each transfer function parameter.

Details

The list contains 3 types of parameters:

- Fuzzy parameters (e.g, Type1Funs, Type2Funs, RedThresh, DoRefinement)
- GA parameters similar to those used in CellNOptR package (see gaDiscreteT1 or defaultParametersFuzzy
- optimisation parameters related to the refinement step.

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- algorithm='NLOPT_LN_SBPLX'
- 2. xtol_abs=0.001
- 3. maxEval=1000
- 4. maxTime=5*60
- In addition, you can set Model and Data (CNOlist).

Value

params a list of default parameters.

Author(s)

T. Cokelaer

Examples

```
data(ToyModel,package="CellNOptR")
data(CNOlistToy,package="CellNOptR")
params = defaultParametersFuzzy(CNOlistToy, ToyModel)
```

gaDiscreteT1

Genetic algorithm used to optimise a cFL model

Description

This function is the genetic algorithm to be used to optimise a cFL model by fitting to data containing one time point.

Usage

```
gaDiscreteT1(CN0list, model, paramsList, initBstring=NULL, sizeFac=0.0001,
    NAFac=1, popSize=50, pMutation=0.5, maxTime=60, maxGens=500,
    stallGenMax=100, selPress=1.2, elitism=5, relTol=0.1,
    verbose=TRUE,maxSizeHashTable = 1000)
```

Arguments

CNOlist a CNOlist on which the score is based (based on valueSignals[[2]], i.e. data at t1)

model a Model list

paramsList CellNOptR software parameters (this functions uses transfer functions to choose

from)

initBstring an initial bitsring to be tested, should be of the same size as the number of

reactions in the model above.

sizeFac the scaling factor for the size term in the objective function, default to 0.0001 NAFac the scaling factor for the NA term in the objective function, default to 1.

popSize the population size for the genetic algorithm, default set to 50

pMutation the mutation probability for the genetic algorithm, default set to 0.5

maxTime the maximum optimisation time in seconds, default set to 60

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maxGens the maximum number of generations in the genetic algorithm, default set to 500. stallGenMax the maximum number of stall generations in the genetic algorithm, default to

100.

selPress the selective pressure in the genetic algorithm, default set to 1.2.

elitism the number of best individuals that are propagated to the next generation in the

genetic algorithm, default set to 5.

relTol the relative tolerance for the best bitstring reported by the genetic algorithm,

i.e.how different from the best solution can solutions be to be reported as well,

default set to 0.1.

verbose logical (default to TRUE) do you want the statistics of each generation to be

printed on the screen?

maxSizeHashTable

a hash table is use to store bitstring and related score. This allows the GA to be very efficient is the case of small models. The size of the hash table is 5000 by

default, which may be too large for large models

Details

The GA procedure is implemented based on the gaBinaryT1 in CellNOptR (see those man pages for a basic description). Necessary extensions to optimize a string of numbers rather than zero and one have been made. Additionally, since the scoring function is defined inside the function, it has also been altered for this function.

The parameters are similar to those used in CellNOptR and the returned list contains similar results as well.

Value

This function returns a list with elements:

bString The best bitstring

results A matrix with columns "Generation", "Best_score", "Best_bitString", "Stall_Generation",

"Avg_Score_Gen", "Best_score_Gen", "Best_bit_Gen", "Iter_time".

stringsTol The bitstrings whose scores are within the tolerance

stringsTolScores

The scores of the above-mentioned strings

Author(s)

M. Morris based on gaBinaryT1 function by C. Terfve (CellNOptR package)

See Also

```
prep4simFuzzy, simFuzzyT1
```

Examples

```
data(CNOlistToy,package="CellNOptR")
data(ToyModel,package="CellNOptR")

#pre-process model
model <- preprocessing(CNOlistToy, ToyModel, verbose=FALSE)</pre>
```

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```
#set parameters
paramsList <- defaultParametersFuzzy(CNOlistToy, ToyModel)

# the GA algorithm
ToyT1opt<-gaDiscreteT1(
   CNOlist=CNOlistToy,
   model=model,
   paramsList=paramsList,
   maxTime=3,
   verbose=FALSE)</pre>
```

getRefinedModel

Refinement of Parameters of cFL model

Description

Performs refinement of cFL model parameters

Usage

```
getRefinedModel(res,CNOlist,cutModel, cutSimList, refParams)
```

Arguments

res Optimum returned by gaDiscreteT1

cNolist a CNolist on which the score is based (based on valueSignals[[2]], i.e. data at

T1)

cutModel Model (with unnesssary edges cut in reduceFuzzy or interpretDiscreteGA

cutSimList Fields for simulation based on cutModel (again, cut with reduceFuzzy or inter-

pretDiscreteGA)

refParams parameter list object as returned by defaultParametersFuzzy.

Details

After the discreteGA chooses transfer functions from a discrete set of transfer functions and removing interactions inconsistent with the data, this function 'refines' the parameters by using optim to go to the local minimum of error to data

Value

refModel A refined model

finalSet Set of final fuzzy parameter

MSE The MSE value

Author(s)

M.K. Morris

See Also

```
gaDiscreteT1, reduceFuzzy, interpretDiscreteGA
```

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Description

This function takes the integer string output by the discrete genetic algorithm for training a cFL model and generates the corresponding model based on the Fuzzy parameters.

Usage

interpretDiscreteGA(model, paramsList, intString, bitString=NULL)

Arguments

model PKN trained (same model input as to gaDiscrete).

paramsList List containing parameters (see defaultParametersFuzzy). Only the fuzzy pa-

rameters are used.

intString Integer string resulting from gaDiscrete (in bString field of gaDiscreteT1 out-

put).

bitString (optional) if you want to cut additional interactions from the model. Used in

reduceFuzzy function.

Details

After the discreteGA chooses transfer functions from a discrete set of transfer functions and removing interactions inconsistent with the data, this function interprets the output and returns an actual model using these transfer functions as well as a model from which logical redundancy was cut.

Value

model The selected initial model based on the provided bitstring.

simList The corresponding data related to Model field bitString The bitstring corresponding to the Model field

cutModel Same as Model but redundant reactions are also removed.

cutSimList The corresponding data related to cutModel field cutBitString The corresponding bitstring related to cutModel field.

Author(s)

M.K. Morris

See Also

gaDiscreteT1

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Examples

```
data(ToyModel, package="CellNOptR")
data(CNOlistToy,package="CellNOptR")
paramsList = defaultParametersFuzzy()
## Not run:

# preprocessing (see CNORwrapFuzzy or gaDiscreteT1)
    T1opt = gaDiscreteT1(....) # see CNORwrapFuzzy or gaDiscreteT1 for details
    interpretDiscreteGA(ToyModel, paramsList, T1opt$bString)

## End(Not run)
```

plotMeanFuzzyFit

Simulates models returned from multiple cFL runs and plots mean fit to data

Description

Uses post refinement threshold (selection threshold) to choose reduced refined model resulting from each run. Simulates model and plots result and fit to data

Usage

```
plotMeanFuzzyFit(postRefThresh, allFinalMSEs, allRes, plotPDF=FALSE, tag=NULL,
show=TRUE, plotParams=list(cex=0.8, cmap_scale=1))
```

Arguments

postRefThresh Post refinement threshold (selection threshold) chosen from plot produced by

compileMultiRes

.

allFinalMSEs matrix containing MSEs produced by compileMultiRes allRes list containing results of several CNORwrapFuzzy runs

plotPDF TRUE or FALSE depending on if a PDF file should be saved

tag String to include in filename of PDF plot

show If the plot should be displayed

plotParams a list of option related to the PDF and plotting outputs. (1) cex is the font size of

the header. (2) cmap_scale below 1 allows to put more emphasizes on low errors (default 1 means all colors have the same weight). See plotOptimResultsPan

from CellNOptR for other fields.

Value

This function does not have any output, it just plots and saves results if applicable.

Author(s)

M.K. Morris

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Examples

```
data(ToyModel, package="CellNOptR")
data(CNOlistToy,package="CellNOptR")
paramsList = defaultParametersFuzzy(CNOlistToy, ToyModel)
N = 10
allRes = list()

## Not run:
for (i in 1:N){
   Res = CNORwrapFuzzy(CNOlistToy, ToyModel, paramsList)
   allRes[[i]] = Res
}

summary = compileMultiRes(allRes)
plotMeanFuzzyFit(0.1, summary$allFinalMSEs, allRes)

## End(Not run)
```

prep4simFuzzy

Prepare a model for simulation

Description

Adds to the model some fields that are used by the simulation engine and calls prep4sim function from CellNOptR package

Usage

```
prep4simFuzzy(model, paramsList, verbose=TRUE)
```

Arguments

model A model, as output by readSIF, normally pre-processed but that is not a require-

ment of this function.

paramsList A parameters list that must contain at least the CNOlist parameter in the *Data*

field (param = list(data=cnolist)) and possibly optional arguments related to the

fuzzy logic (see defaultParametersFuzzy)

verbose a verbose option set to TRUE by default.

Details

This adds fields that are necessary for the simulation engine for both Boolean and constrained Fuzzy logic simulation.

Value

In addition to the fields returned by the prep4sim function of CellNOptR, this function appends the following fields:

finalCube stores, for each reac(row) the location of its inputs (col)

ixNeg stores, for each reac(row) and each input (col) whether it is a negative input

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ignoreCube logical matrix of the same size as the 2 above, that tells whether the particular

cell is filled or not

maxIx row vector that stores, for each reac, the location of its output modelname stores the name of the model from which these fields were derived

Author(s)

C. Terfve, modified by M.K. Morris and T. Cokelaer

See Also

```
simFuzzyT1
```

Examples

```
data(ToyModel, package="CellNOptR")
data(CNOlistToy, package="CellNOptR")
params <- defaultParametersFuzzy()
params$data = CNOlistToy
fields4sim <- prep4simFuzzy(ToyModel, params)</pre>
```

reduceFuzzy

Remove unnecessary interactions from cFL model

Description

Determine if interactions in cFL model are necessary to fit the data

Usage

```
reduceFuzzy(firstCutOff, CNOlist, model, res, params )
```

Arguments

firstCutOff Threshold for removing or replacing and edge. If the score doesn't get any worse

than this, its removed or replaced

cNolist a CNolist on which the score is based (based on valueSignals[[2]], i.e. data

at t1)

model Model input into gaDiscreteT1 (uncut)

res Optimum returned by gaDiscrete

params parameters (as generated from defaultParametersFuzzy (only Fuzzy parameters

required)

Details

After gaDiscrete removes interactions that are inconsistent with the data, this function asks if they are necessary to fit the data. For all interactions, it asks if they can be removed without making the score worse by a value of the firstCutOff. For AND interactions, it asks if they can be replaced with an OR gate.

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Value

redModel
redSimList
bitString
intString
MSE

Author(s)

M.K. Morris

See Also

gaDiscreteT1

simFuzzyT1

Simulation of a cFL model

Description

This is the simulator, inspired from CNOfuzzySimEngv23 in the Matlab CellNOpt, to be used on one time point simulations

Usage

```
simFuzzyT1(CNOlist, model, simList)
```

Arguments

CNOlist a CNOlist

model a Model that only contains the reactions to be evaluated

simList as created by prep4simFuzzy, that has also already been cut to contain

only the reactions to be evaluated

Details

I'm not sure if this applies to the one I wrote, which was based on I think an older version by Camille maybe? Differences from the BoolSimEngMKM simulator include: the valueInhibitors has not been previously flipped; the function outputs the values across all conditions for all species in the model, instead of only for the signal species. This is because then the output of this function can be used as initial values for the version of the simulator that works on time point 2 (not implemented in this version).

If you would like to compute the output of a model that contains some of the gates in the model but not all, we suggest that you use the function SimulateT1 and specify in the bStringT1 argument which gates you want to be included. Indeed, SimulateT1 is a wrapper around simulatorT1 that takes care of cutting the model for you before simulating it.

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Value

This function outputs a single matrix of format similar to valueSignals in the CNOlist but that contains an output for each species in the model. This matrix is the simulated equivalent of valueSignals at time 1, if you consider only the columns given by indexSignals.

Author(s)

M.K. Morris based on function by C. Terfve from CellNOptR package.

See Also

SimulateT1 from CellNOptR

Examples

```
#This computes the output of the full model, which is normally not done on a stand alone basis, but if you have a
    data(CNOlistToy,package="CellNOptR")
    data(ToyModel,package="CellNOptR")

paramsList <- defaultParametersFuzzy(data=CNOlistToy, model=ToyModel)

indicesToy<-indexFinder(CNOlistToy,ToyModel,verbose=TRUE)
ToyFields4Sim<-prep4simFuzzy(ToyModel,paramsList)

Sim<-simFuzzyT1(
    CNOlist=CNOlistToy,
    model=ToyModel,
    simList=ToyFields4Sim)</pre>
```

writeFuzzyNetwork

Despict the network results of training a cFL model to data in multiple runs.

Description

Fuzzy network results output.

Usage

```
writeFuzzyNetwork(postRefThresh, allFinalMSEs, allRes, tag=NULL,verbose=FALSE)
```

Arguments

| postRefThresh | Post refinement threshold | (selection threshold) | chosen from | plot produced by |
|---------------|---------------------------|-----------------------|-------------|------------------|
| | | | | |

compileMultiRes

allFinalMSEs matrix containing MSEs produced by compileMultiRes allRes list containing all results produced by compileMultiRes

tag String to include in filename of pdf plot verbose If extra warnings should be displayed

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Details

The weights of the edges are computed as the mean across models using post refinement threshold (selection threshold) to choose reduced refined model resulting from each run.

As with writeNetwork, this function maps back the edges weights from the optimised (expanded and compressed) model to the original model. The mapping back only works if the path has length 2 at most (i.e. you have node1-comp1-comp2-node2, where comp refer to nodes that have been compressed).

Value

This function does not have any output, it just writes a SIF file, an edge attribute file, and a node attribute file

Note

The mapback of this function is still an open question, even in the Matlab version. Future developments will include more robust versions of the mapping back algorithm, probably as a separate mapback function.

Author(s)

M.K. Morris based on code by C. Terfve

See Also

writeNetwork

Examples

```
data(ToyModel, package="CellNOptR")
    data(CNOlistToy,package="CellNOptR")
    paramsList = defaultParametersFuzzy(CNOlistToy, ToyModel)
    N = 10
    allRes = list()

## Not run:
    for (i in 1:N){
        Res = CNORwrapFuzzy(CNOlistToy, ToyModel, paramsList)
        allRes[[i]] = Res
    }

    summary = compileMultiRes(allRes)
    summary$allFinalMSEs
    summary$allFinalNumParams

    writeFuzzyNetwork(postRefThresh, summary$allFinalMSEs, allRes)

## End(Not run)
```

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