

# Package ‘CARNIVAL’

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**Type** Package

**Title** A CAusal Reasoning tool for Network Identification (from gene expression data) using Integer VALue programming.

**Version** 0.9.1

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**Description** An upgraded causal reasoning tool from Melas et al. in R with updated assignments of TFs' weights from PROGENy scores. Cplex parameters can be freely adjusted and multiple solutions from cplex can be obtained and aggregated

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**Imports** doParallel, igraph, tidyr, dplyr, readr, tidyverse, viper

**Suggests** knitr, rmarkdown,

**VignetteBuilder** knitr

**RoxygenNote** 6.1.1

**BiocViews**

**NeedsCompilation** no

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AddPerturbationNode	AddPerturbationNode
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## Description

Introduces a perturbation node connecting periphery nodes without a target in the prior knowledge network.

## Usage

```
AddPerturbationNode(network)
```

## Arguments

network	The original prior knowledge network
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## Value

A list of updated network with perturbation nodes and re-assigned list of input to include only the interactions from the perturbation node.

---

all\_constraints\_wLoop all\_constraints\_wLoop

---

**Description**

all\_constraints\_wLoop

**Usage**

```
all_constraints_wLoop(c0 = c0, c1 = c1, c2 = c2, c3 = c3,
  c4 = c4, c5 = c5, c6 = c6, c7 = c7, c8 = c8, c9 = c9)
```

**Arguments**

c0	Constraints number 0
c1	Constraints number 1
c2	Constraints number 2
c3	Constraints number 3
c4	Constraints number 4
c5	Constraints number 5
c6	Constraints number 6
c7	Constraints number 7
c8	Constraints number 8
c9	Constraints number 9

**Value**

The list of all LP constraints with removed failed-to-write constraints that returns NaN

---

assignPROGENyScores assignPROGENyScores

---

**Description**

assignPROGENyScores

**Usage**

```
assignPROGENyScores(progeny = progeny, progenyMembers = progenyMembers,
  id = "uniprot")
```

**Arguments**

progeny	Contains the progeny scores as obtained from runPROGENy.
progenyMembers	Contains the list of members for each PROGENy pathway.
id	Contains the members identifiers (default: uniprot).

**Value**

This function is used to account for the PROGENy scores in the objective function.

---

```
buildDataMatrix    buildDataMatrix
```

---

**Description**

```
buildDataMatrix
```

**Usage**

```
buildDataMatrix(data = data, pknList = pknList, inputs = inputs)
```

**Arguments**

data	Contains the measured data.
pknList	Contains the background network which serves as a prior knowledge and which we train.
inputs	Contains the list of targets as inputs.

**Value**

This function returns the data matrix containing the data for running CARNIVAL and a set of identifiers for Targets, Measured and Un-measured nodes.

---

```
create_variables_all    create_variables_all
```

---

**Description**

```
create_variables_all
```

**Usage**

```
create_variables_all(pknList = pknList, dataMatrix = dataMatrix)
```

**Arguments**

pknList	Contains the background network which serves as a prior knowledge and which we train.
dataMatrix	Contains the matrix which stores the data for running CARNIVAL and a set of identifiers for Targets, Measured and Un-measured nodes.

**Value**

This function returns the identifiers of all the variables used in the ILP formulation together with an explanation about the meaning of each of them. Also it returns a list of useful identifiers.

---

exportResult	exportResult
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---

**Description**

Extract and export the optimisation results from the cplex solution file (XML) as files and variables for further plotting functions

**Usage**

```
exportResult(cplexSolutionFileName = cplexSolutionFileName,
             variables = variables, conditionIDX = conditionIDX,
             pknList = pknList, dir_name = dir_name, inputs = inputs,
             measurements = measurements, Export_all = Export_all,
             writeIndividualResults = F)
```

**Arguments**

cplexSolutionFileName	Path to the cplex solution file (XML)
variables	The list of mapping indices of LP variables
conditionIDX	The number of experimental condition to be processed
pknList	The provided prior knowledge network
dir_name	The name of directory to store results
inputs	The list of known or potential target of perturbation
measurements	The discretised observations (here transcription factor activities) of values [-1,0,1]
Export_all	An option to define whether all detailed mapped LP variables will be written as individual files
writeIndividualResults	An option to define whether the results of individual solutions will be written; if FALSE, only the global combined solution will be written

**Value**

Output files of ILP solutions and a list of networks and node activities to be written into figures

---

files2res	files2res
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---

**Description**

Read in a list of exported/written network and node activities files into a variable for plotting

**Usage**

```
files2res(counterlist)
```

**Arguments**

counterlist Indices of model solutions to be read-in

**Value**

A list of variables for plotting functions

---

generate\_measfile generate\_measfile

---

**Description**

This function generates the measurement file for each condition which is required as one of the input file

**Usage**

```
generate_measfile(measurements, topnumber = NULL,
write2folder = "./measurements")
```

**Arguments**

measurements A dataframe describing the measurements (e.g. inferred TF activities). Columns/colnames should correspond to perturbations, row/rownames to measurements.

topnumber If given, only the top number of measurements will be written out.

write2folder Path to outputfolder

**Value**

A n\*m dataframe describing the inferred TF activities, where n is the number of Tfs and m the number of conditions.

---

genesymbol2uniprot genesymbol2uniprot

---

**Description**

This function converts the gene symbol rownames of a dataframe to uniprot rownames. In case one gene symbol maps to two uniprot IDs, the row will be duplicated.

**Usage**

```
genesymbol2uniprot(df, map, geneID = 1, uniprotID = 2)
```

**Arguments**

df A vector of genes.

map A dataframe with a uniprot ID column and a genesymbol column.

geneID Column index of the gene symbol column in map.

uniprotID Column index of the uniprot column in map.

**Value**

A vector of genes or uniprot IDs.

---

```
load_CARNIVAL_examples
      load_CARNIVAL_examples
```

---

**Description**

load\_CARNIVAL\_examples

**Usage**

```
load_CARNIVAL_examples(CARNIVAL_example)
```

**Arguments**

CARNIVAL\_example

The number of CARNIVAL example to be loaded; 1 = Toy model, 2 = EGF-SBVimprover, 3 = APAP-TGGs

**Value**

The input variables for CARNIVAL including network, measurement, input target +/- pathways scores

---

```
runPROGENy      runPROGENy This function is designed to compute progeny pathway scores and assess their significance using a gene sampling based permutation strategy, for a series of experimental samples/contrasts.
```

---

**Description**

runPROGENy This function is designed to compute progeny pathway scores and assess their significance using a gene sampling based permutation strategy, for a series of experimental samples/contrasts.

**Usage**

```
runPROGENy(df, weight_matrix, k = 10000, z_scores = T,
            get_nulldist = F)
```

**Arguments**

df	A data.frame of n*m+1 dimension, where n is the number of omic features to be considered and m is the number of samples/contrasts. The first column should be the identifiers of the omic features. These identifiers must be coherent with the identifiers of the weight matrix.
weight_matrix	A progeny coefficient matrix. the first column should be the identifiers of the omic features, and should be coherent with the identifiers provided in df.
k	The number of permutations to be performed to generate the null-distribution used to estimate significance of progeny scores. Default value is 10000.
z_scores	If true, provides z-scores. If false, provides significance scores (1-pval).
get_nulldist	If get_nulldist is true, then the function will return the null model dataframe that was used.

**Value**

This function returns a list of two elements. The first element is a dataframe of p\*m+1 dimensions, where p is the number of progeny pathways, and m is the number of samples/contrasts. Each cell represent the significance of a progeny pathway score for one sample/contrast. The significance ranges between -1 and 1. The significance is equal to  $x^{*2}-1$ , x being the quantile of the progeny pathway score with respect to the null distribution. Thus, this significance can be interpreted as the equivalent of 1-p.value (two sided test over an empirical distribution) with the sign indicating the direction of the regulation. The second element is the null distribution list (a null distribution is generated for each sample/contrast).

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run_dorothea	run_dorothea
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---

**Description**

This function infers TF activities based on the DoRothEA regulon and the viper function. The regulon can be filtered by confidence level (A-E).

**Usage**

```
run_dorothea(df, regulon, confidence_level = c("A", "B", "C"),
  write2file = NULL)
```

**Arguments**

df	A n*m dataframe describing the contrast t-statistics, where n is the number of genes and m the number of conditions.
regulon	DoRothEA regulon list
confidence_level	A vector describing which confidence levels of the DoRothEA regulon to use
write2file	Path to outputfile for all TF_activities if desired

**Value**

A n\*m dataframe describing the inferred TF activities, where n is the number of Tfs and m the number of conditions.



---

Uniprot2GeneSymbol	Uniprot2GeneSymbol
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---

**Description**

Conversion of Uniprot ID (e.g. from Omnipath) to official gene symbol in the plotting step

**Usage**

```
Uniprot2GeneSymbol(res)
```

**Arguments**

res	A list of variables in Uniprot ID (network and node activities)
-----	---

**Value**

A list of mapped variables from Uniprot to official gene symbol for a better readability

---

WriteDOTfig	WriteDOTfig
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---

**Description**

This function takes results from ILP optimisation and write out a figure in DOT format

**Usage**

```
WriteDOTfig(res, idxModel = 0, dir_name, inputs, measurements,
            UP2GS = F)
```

**Arguments**

res	A list containing optimised network structures and node activities from ILP optimisation
idxModel	(optional) The index or indices of optimised model to be plotted (Default idxModel=0; plot the combined and average network)
dir_name	The directory name that stores CARNIVAL results (as a subfolder of the main "results" folder)
inputs	A named vector of inputs and their node states
measurements	A named vector of measurements and their state values
UP2GS	(T/F) A parameter defining whether to convert the input format UniprotID to Gene Symbol

**Value**

DOT figures stored in the directory "results/dir\_name" with the prefix "ActivityNetwork\_model"

---

 writeLPFile

 writeLPFile
 

---

### Description

Write a list of linear programming (LP) constraints into a file which will be read by interactive cplex solver to perform network optimisation.

### Usage

```
writeLPFile(data = data, pknList = pknList, inputs = inputs,
  alphaWeight = 1, betaWeight = 0.2, scores = scores, mipGAP = 0.1,
  poolrelGAP = 0.01, limitPop = 100, poolCap = 100,
  poolIntensity = 0, poolReplace = 2, timelimit = 1800,
  measWeights = NULL, repIndex, condition = "")
```

### Arguments

data	The discretised observations (here transcription factor activities) of values [-1,0,1]
pknList	The prior knowledge network
inputs	The list of known or potential target of perturbation
alphaWeight	The weight for mismatch penalty between discretised observations and predicted model states
betaWeight	The weight for network size (node) penalty
scores	The continuous pathway scores from PROGENy
mipGAP	The minimal integer programming percentage gap to be accepted as a solution
poolrelGAP	The allowed relative percentage gap between the best solution and the equivalent solutions in the solution pool
limitPop	The number of allowed solutions to be populated
poolCap	The number of solutions to be kept in the pool of solution
poolIntensity	The intensity of the search in solution space
poolReplace	The replacement strategy of the solutions in the solution pool
timelimit	The allowed amount of time (in seconds) for the optimisation
measWeights	The continuous weight of observations (here transcription factor activities) - to replace the default alphaWeight if assigned; Note: take only positive values!
repIndex	The indexing of optimisation - useful in case more than one experiment is performed
condition	The free variable which could be assigned for additional study e.g. to vary the effect of betaWeight in a loop

### Value

An integer programming file containing the description of ILP optimisation problem and a cplex command file to communicate with the interactive version of cplex solver

---

write_binaries	write_binaries
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---

**Description**

write\_binaries

**Usage**

write\_binaries(variables = variables)

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
-----------	---

**Value**

This code writes the list of binary variables (xp, xm, up & um).

---

write_boundaries	write_boundaries
------------------	------------------

---

**Description**

write\_boundaries

**Usage**

write\_boundaries(variables = variables, oF = oF)

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
oF	Is the objective function of the formulation.

**Value**

This code writes the boundaries of each variable.

---

```
write_constraints_1_all
      write_constraints_1_all
```

---

**Description**

```
write_constraints_1_all
```

**Usage**

```
write_constraints_1_all(variables = variables)
```

**Arguments**

`variables`        Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.

**Value**

This code writes the list of constraints (1) of the ILP problem for all the conditions.

---

```
write_constraints_2_all
      write_constraints_2_all
```

---

**Description**

```
write_constraints_2_all
```

**Usage**

```
write_constraints_2_all(variables = variables)
```

**Arguments**

`variables`        Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.

**Value**

This code writes the list of constraints (2) of the ILP problem for all the conditions.

---

```
write_constraints_3_all
write_constraints_3_all
```

---

**Description**

```
write_constraints_3_all
```

**Usage**

```
write_constraints_3_all(variables = variables)
```

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
-----------	---

**Value**

This code writes the list of constraints (3) of the ILP problem for all the conditions.

---

```
write_constraints_4_all
write_constraints_4_all
```

---

**Description**

```
write_constraints_4_all
```

**Usage**

```
write_constraints_4_all(variables = variables)
```

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
-----------	---

**Value**

This code writes the list of constraints (4) of the ILP problem for all the conditions.

---

```
write_constraints_5_all
      write_constraints_5_all
```

---

**Description**

```
write_constraints_5_all
```

**Usage**

```
write_constraints_5_all(variables = variables)
```

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
-----------	---

**Value**

This code writes the list of constraints (5) of the ILP problem for all the conditions.

---

```
write_constraints_6  write_constraints_6
```

---

**Description**

```
write_constraints_6
```

**Usage**

```
write_constraints_6(variables = variables, dataMatrix = dataMatrix,
  inputs = inputs)
```

**Arguments**

variables	The list of mapping indices of LP variables. variables Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
dataMatrix	Contains the matrix which stores the data for running CARNIVAL and a set of identifiers for Targets, Measured and Un-measured nodes.
inputs	Contains the list of targets as inputs.

**Value**

This code writes the list of constraints (6) of the ILP problem for all the conditions.

---

write\_constraints\_7    write\_constraints\_7

---

**Description**

write\_constraints\_7

**Usage**

```
write_constraints_7(variables = variables, dataMatrix = dataMatrix,  
                  inputs = inputs)
```

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
dataMatrix	Contains the matrix which stores the data for running CARNIVAL and a set of identifiers for Targets, Measured and Un-measured nodes.
inputs	Contains the list of targets as inputs.

**Value**

This code writes the list of constraints (7) of the ILP problem for all the conditions.

---

write\_constraints\_8    write\_constraints\_8

---

**Description**

write\_constraints\_8

**Usage**

```
write_constraints_8(variables = variables, inputs = inputs,  
                  pknList = pknList)
```

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
inputs	Contains the list of targets as inputs.
pknList	Contains the background network which serves as a prior knowledge and which we train.

**Value**

This code writes the list of constraints (8) of the ILP problem for all the conditions.

---

```
write_constraints_objFunction_all
      write_constraints_objFunction_all
```

---

**Description**

write\_constraints\_objFunction\_all

**Usage**

```
write_constraints_objFunction_all(variables = variables,
      dataMatrix = dataMatrix)
```

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
dataMatrix	Contains the matrix which stores the data for running CARNIVAL and a set of identifiers for Targets, Measured and Un-measured nodes.

**Value**

This function returns the list of constraints associated with the 'Absolute Difference' variables and which measure the mis-fit between inferred and measured data.

---

```
write_generals      write_generals
```

---

**Description**

write\_generals

**Usage**

```
write_generals(variables = variables, oF = oF)
```

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
oF	Is the objective function of the formulation.

**Value**

This code writes all the variables.



---

```
write_loop_constraints
write_loop_constraints
```

---

**Description**

```
write_loop_constraints
```

**Usage**

```
write_loop_constraints(variables = variables, pknList = pknList,
inputs = inputs)
```

**Arguments**

variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
pknList	Contains the background network which serves as a prior knowledge and which we train.
inputs	Contains the list of targets as inputs.

**Value**

This function writes the constraints preventing self-activation of nodes in the network due to positive feedback loops.

---

```
write_objective_function
write_objective_function
```

---

**Description**

```
write_objective_function
```

**Usage**

```
write_objective_function(dataMatrix = dataMatrix,
variables = variables, alphaWeight = alphaWeight,
betaWeight = betaWeight, scores = scores,
measWeights = measWeights, conditionIDX = conditionIDX)
```

**Arguments**

dataMatrix	Contains the matrix which stores the information for each node in the PKN, i.e. activity of the nodes which are measured, at each condition.
variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
alphaWeight	The weighting factor of the measurement.

betaWeight	The weighting factor of the network size.
scores	The provided PROGENy scores.
measWeights	A weighting factor for the measurements.
conditionIDX	The index of the current condition being considered.

**Value**

This code writes the objective function of the ILP problem for one specific condition.

---

```
write_objective_function_all
        write_objective_function_all
```

---

**Description**

`write_objective_function_all`

**Usage**

```
write_objective_function_all(dataMatrix = dataMatrix,
    variables = variables, alphaWeight = alphaWeight,
    betaWeight = betaWeight, scores = scores,
    measWeights = measWeights, conditionIDX = conditionIDX)
```

**Arguments**

dataMatrix	Contains the matrix which stores the information for each node in the PKN, i.e. activity of the nodes which are measured, at each condition.
variables	Contains the list of variables as used to formulate the ILP problem, explanations for each variable and a list of useful indices.
alphaWeight	The weighting factor of the measurement.
betaWeight	The weighting factor of the network size.
scores	The provided PROGENy scores.
measWeights	A weighting factor for the measurements.
conditionIDX	The index of the current condition being considered.

**Value**

This code writes the objective function of the ILP problem for all the conditions.

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