GNG documentation

June 4, 2014

GNG

Constructor of GrowingNeuralGas object

Description

Construct GNG object

Usage

GNG

Arguments

beta coefficient. Decrease the error variables of all node nodes by this fraction. For-

getting rate. Default 0.99

alpha Alpha coefficient. Decrease the error variables of the nodes neighboring to the

newly inserted node by this fraction. Default 0.5

uniformgrid.optimization

TRUE/FALSE. You cannot use utility option with this, so please pass FALSE

here then.

lazyheap.optimization

TRUE/FALSE. You cannot use utility option with this, so please pass FALSE

here then.

max.nodes Maximum number of nodes (after reaching this size it will continue running, but

won't add new nodes)

eps_n Default 0.05 eps_v Default 0.0006

dataset.type Dataset type. Possibilities gng.dataset.bagging, gng.dataset.bagging.prob, gng.dataset.sequential

experimental_utility_option

EXPERIMENTAL Utility option gng.experimental.utility.option.off/gng.experimental.utility.option.basi

experimental_utility_k

EXPERIMENTAL Utility option constant

Growing-Neural-Gas

```
load_model_filename
```

Set to path to file from which load serialized model

```
experimental_vertex_extra_data
```

if TRUE each example should have additional coordinate, that will be voted in graph. Each node (that you can get using node function) will have extra_data field that will be equal to mean of samples around given node. If used with probability dataset example layout is <vertex position> <vertex_extra_data> <sampling_probability>, for example c(0.3, 0.6, 0.7, 100, 0.7)

Format

2

NULL

Examples

```
# Default GNG instance, without optimitzations and vertex dimensionality 3
gng <- GNG(dataset_type=gng.dataset.bagging.prob, max_nodes=max_nodes, dim=3)
# Construct GNG loaded from file with uniform grid
gng <- GNG(dataset_type=gng.dataset.bagging.prob, max_nodes=max_nodes, dim=3,
uniformgrid_optimization=TRUE, lazyheap_optimization=FALSE,
uniformgrid_boundingbox_sides=c(3,3,3), uniformgrid_boundingbox_origin=c(-0.5,-0.5,-0.5),
load_model_filename="sphere_simple.bin")</pre>
```

Growing-Neural-Gas

Efficient C++ implementation of online clustering algorithm Growing Neural Gas.

Description

This package contains fast C++ implementation of online clustering algorithm Growing Neural Gas. It produces topological graph, that you can easily convert to igraph, or you can dump your model to optimized binary file and load it later on.

Details

Package: Growing-Neural-Gas

Type: Package Version: 0.6

Date: 2014-06-03 License: MIT License

For overview of usage please see exemplary code in demo/ folder.

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References

~~ Literature or other references for background information ~~

Examples

```
gng <- GNG(dataset_type=gng.dataset.bagging.prob, max_nodes=max_nodes, dim=3)
insert_examples(gng, preset=gng.preset.sphere, N=10000, prob=0.8)
run(gng)
insert_examples(gng, preset=gng.preset.box, N=10000, prob=0.8)
plot(gng, mode=gng.plot.2d.errors, vertex.color=gng.plot.color.cluster, layout=gng.plot.layout.igraph.v2d)</pre>
```

centroids.gng

centroids

Description

Using infomap.communities finds communities and for each community pick node with biggest betweenness score

Usage

```
centroids(gng)
```

Format

NULL

Details

Get centroids

```
# Print position of the first centroid
print(node(gng, centroids(gng)[1])$pos)
```

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convert_igraph.gng

convert_igraph

Description

Converts to igraph (O(n) method, writing intermediately to disk)

Usage

```
convert_igraph(gng)
```

Format

NULL

Examples

```
convert_igraph(gng)
```

dump_model.gng

dump_model

Description

Writes graph to a disk space efficient binary format. It can be used in GNG constructor to reconstruct graph from file.

Usage

```
dump_model(gng)
```

Arguments

filename

Dump destination

Format

NULL

Details

Dump model to binary

```
dump_model(gng, graph.bin)
```

error_statistics.gng 5

```
error_statistics.gng error_statistics
```

Description

Gets vector with errors for every second of execution

Usage

```
error_statistics(gng)
```

Format

NULL

Examples

```
error_statistics(gng)
```

insert_examples.gng insert_examples

Description

Insert (inefficiently) examples to algorithm dataset. For efficient insertion use gng\$inser_examples, and for setting memory pointer use gng\$set_memory_move_examples

Usage

```
insert_examples(gng, M)
```

Arguments

examples Matrix with examples of dimensionality N rows x C columns, where C columns

= dim (passed as parameter to GNG object) + 1 or 0 (1 if vertex_extra_data is

TRUE) + 1 or 0 (1 if dataset_type=gng.dataset.bagging.prob).

preset Use only if you are adding exemplary dataset. Possibilities: gng.preset.sphere,

gng.preset.box. You can specify preset params: N=1000, center=c(0.5,0.5,0.5),

prob=-1.

Format

NULL

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Note

Complicated memory layout of the matrix is due to need for memory efficiency. In the future versions you can expect wrapper simplifying addition and also streaming from disk file

Examples

#Add preset examples with probability of being sampled (this assumed GNG was created with gng.dataset.bagging.prob insert_examples(gng, preset=gng.preset.sphere)

```
#Insert efficiently examples
M <- matrix(0, ncol=3, nrow=10)
M[1,] = c(4,5,6)
gng$insert_examples(M)

#Set memory of the algorithm to point in memory. Note: you cannot remove this matrix while in execution or you will experience memory error gng$set_memory_move_examples(M)</pre>
```

mean_error.gng

mean_error

Description

Gets mean error of the graph (note: blocks the execution, O(n))

Usage

```
mean_error(gng)
```

Format

NULL

```
mean_error(gng)
```

node.gng 7

node.gng

node

Description

Retrieves node from resulting graph

Usage

```
node(gng, 10)
```

Arguments

gng_id

Id of the node to retrieve. This is the id returned by functions like predict, or centroids

Format

NULL

Details

Get GNG node

Examples

```
print(node(gng, 10)$pos)
```

pause.gng

pause

Description

Pause algorithm

Usage

pause(gng)

Format

NULL

Examples

pause(gng)

8 plot.gng

Description

Plot resulting graph using igraph plotting, or using rgl 3d engine.

Usage

plot(gng)

Arguments

mode gng.plot.rgl3d (3d plot), gng.plot.2d (igraph plot) or gng.plot.2d.errors (igraph

plot with mean error log plot)

layout to be used when plotting. Possible values: gng.plot.layour.igraph.v2d

(first two dimensions), gng.plot.layout.igraph.auto (auto layout from igraph) gng.plot.layout.igraph.fruchterman.fast (fast fruchterman reingold layout),or any

function accepting igraph graph and returning layout

vertex.color how to color vertexes. Possible values: gng.plot.color.cluster(vertex color is set

to fastgreedy.community clustering), gng.plot.color.extra(rounds to integer extra

dim if present), gng.plot.color.none(every node is white),

Format

NULL

Details

Plot GNG

Note

If you want to "power-use" plotting and plot for instance a subgraph, you might be interested in exporting igraph with convert_igraph function and plotting it using/reusing function from this package: .visualizeIGraph2d

```
# Plots igraph using first 2 coordinates and colors according to clusters
plot(gng, mode=gng.plot.2d.errors, layout=gng.plot.layout.v2d, vertex.color=gng.plot.color.cluster)
# Plot rgl (make sure you have installed rgl library)
plot(gng, mode=gng.plot.rgl, layout=gng.plot.layout.v2d, vertex.color=gng.plot.color.cluster)
# For more possibilities see gng.plot.* constants
```

run.gng

run.gng

run

Description

Run algorithm (in parallel)

Usage

run(gng)

Format

NULL

Examples

run(gng)

terminate.gng

pause

Description

Terminate algorithm

Usage

terminate(gng)

Format

NULL

Examples

terminate(gng)

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