Adjacency list from progrex index

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adjl_from_progindex   *Build the network from the already processed Progress Index file*

Description

adjl_from_progindex is able to use the output file from original campari software (or the output file from gen_annotation) in order to generate again the minimum spanning tree.

Usage

adjl_from_progindex(prog_index_file)

Arguments

prog_index_file

        Progress index file location. This should be of the kind "PROGIDX_000000000001.dat" (original campari) or "REPIX_000000000001.dat" (CampaRi).

Details

For details, please refer to the main documentation of the original campari software http://campari.sourceforge.net/documentation.html

Value

adjl_from_progindex will return a minimum spanning tree: degree list, connectivity matrix and weights

See Also

gen_progindex, gen_annotation

Examples

```
## Not run:
adjl <- adjl_from_progindex(fil = "PROGIDX_000000000001.dat")

## End(Not run)
```

Index