dCheckParallel

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dCheckParallel	*Function to check whether parallel computing should be used and how*

Description

dCheckParallel is used to check whether parallel computing should be used and how

Usage

```
dCheckParallel(multicores = NULL, verbose = T)
```

Arguments

multicores
: an integer to specify how many cores will be registered as the multicore parallel backend to the 'foreach' package. If NULL, it will use a half of cores available in a user's computer

verbose
: logical to indicate whether the messages will be displayed in the screen. By default, it sets to true for display

Value

TRUE for using parallel computing; FALSE otherwise

Note

Whether parallel computation with multicores is used is system-specific (now only Linux or Mac OS). Also, it will depend on whether these two packages "foreach" and "doMC" have been installed. It can be installed via: source("http://bioconductor.org/biocLite.R"); biocLite(c("foreach","doMC")).

See Also

[dRWR](), [dRWRcontact](), [dRWRpipeline](), [dDAGtermSim](), [dDAGgeneSim]()

Examples

```
dCheckParallel(multicores=2)
```

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