

Package ‘slurm’

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Title Running and Parsing Slurm Commands

Description User-friendly functions which parse output of command line programs used to query Slurm.
Morris A. Jette and Tim Wickberg (2023)
<[doi:10.1007/978-3-031-43943-8_1](https://doi.org/10.1007/978-3-031-43943-8_1)> describe Slurm in detail.

Imports data.table (>= 1.11.6), nc (>= 2019.11.22)

SystemRequirements Slurm command line programs (sacct, squeue)

Suggests testthat, ggplot2, R.utils

NeedsCompilation no

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Contents

slurm-package	2
int.pattern	2
na.as.zero	2
sacct	3
sacct-2019-11-21	3
sacct-job13936577	3
sacct-multitype	4
sacct.pattern.list	4
sacct_fields	4
sacct_fread	5
sacct_lines	5
sacct_tasks	6
sjob	7
sjob_dt	8

sq.jobs	8
task.pattern	9

Index **10**

slurm-package *Running and Parsing SLURM Commands*

Description

User-friendly wrappers for SLURM commands.

Author(s)

Toby Dylan Hocking

int.pattern *int pattern*

Description

Match one or more digits and convert to integer.

Usage

"int.pattern"

na.as.zero *na as zero*

Description

Convert empty string to zero and use as.integer otherwise.

Usage

na.as.zero(int.or.empty)

Arguments

int.or.empty int.or.empty

Value

Integer vector.

Author(s)

Toby Hocking <toby.hocking@r-project.org> [aut, cre]

sacct	<i>sacct</i>
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Description

Run `sacct_lines` then `sacct_tasks`.

Usage

`sacct(...)`

Arguments

... ...

Value

Same as result of `sacct_tasks`.

Author(s)

Toby Hocking <toby.hocking@r-project.org> [aut, cre]

sacct-2019-11-21	<i>Output of sacct for thousands of jobs</i>
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Description

The data come from running `sacct` on a SLURM system.

Format

This "csv" file results from `sacct -P` which uses the vertical bar | as the separator. There are five columns and six rows (including header).

sacct-job13936577	<i>Output of sacct for one job with several tasks</i>
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Description

The data come from running `sacct` on a SLURM system.

Format

This "csv" file results from `sacct -P` which uses the vertical bar | as the separator. There are five columns and 709 rows (including header).

sacct-multitype	<i>Output of sacct for three job with several tasks</i>
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Description

The data come from running sacct on a SLURM system.

Format

This "csv" file results from sacct -P which uses the vertical bar | as the separator. There are five columns and six rows (including header).

sacct.pattern.list	<i>sacct pattern list</i>
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Description

Named list of patterns for parsing [sacct](#) fields.

Usage

```
"sacct.pattern.list"
```

sacct_fields	<i>sacct fields</i>
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Description

Get current fields from [sacct](#).

Usage

```
sacct_fields()
```

Value

character vector.

Author(s)

Toby Hocking <toby.hocking@r-project.org> [aut, cre]

sacct_fread	<i>sacct_fread</i>
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Description

Run fread on the output of [sacct](#).

Usage

```
sacct_fread(...)
```

Arguments

... ...

Value

Data table with the same number of rows as the output of the [sacct](#) command, and additional columns that result from parsing the columns with [sacct.pattern.list](#).

Author(s)

Toby Hocking <toby.hocking@r-project.org> [aut, cre]

Examples

```
library(slurm)
sacct_fread(text="JobID|ExitCode|State|MaxRSS|Elapsed
18473217_1|0:0|RUNNING||00:03:47
18473217_1.extern|0:0|RUNNING||00:03:47")
```

sacct_lines	<i>sacct_lines</i>
-------------	--------------------

Description

Run [sacct](#) with args and parse output as a data.table

Usage

```
sacct_lines(args, format.fields = c("JobID",
  "ExitCode", "State",
  "MaxRSS", "Elapsed"),
  delimiter = "\t")
```

Arguments

args	character string passed to <code>sacct</code> command line, e.g. <code>-j123</code> for selecting job ID 123.
format.fields	character vector of field names to pass to <code>sacct -format</code> . Use <code>sacct_fields</code> to get all fields.
delimiter	passed as <code>-delimiter</code> .

Value

Same as `sacct_fread`.

Author(s)

Toby Hocking <toby.hocking@r-project.org> [aut, cre]

sacct_tasks	<i>sacct tasks</i>
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Description

Summarize output from `sacct_fread`.

Usage

```
sacct_tasks(match.dt)
```

Arguments

match.dt	match.dt
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Value

data.table with one row per job/task.

Author(s)

Toby Hocking <toby.hocking@r-project.org> [aut, cre]

Examples

```
library(slurm)
sacct.csv.gz <- system.file(
  "data", "sacct-job13936577.csv.gz", package="slurm", mustWork=TRUE)
if(requireNamespace("R.utils")){
  sacct.dt <- sacct_fread(sacct.csv.gz)
  task.dt <- sacct_tasks(sacct.dt)
  print(task.dt[State_batch != "COMPLETED"])
  if(require(ggplot2)){
```

```
ggplot()+  
  geom_point(aes(  
    hours, megabytes, fill=State_batch),  
    shape=21,  
    data=task.dt)+  
  scale_fill_manual(values=c(  
    COMPLETED=NA,  
    FAILED="red"))+  
  scale_x_log10()+  
  scale_y_log10()  
  }  
}
```

sjob

sjob

Description

Run [sacct](#) and summarize State/ExitCode values for given job IDS

Usage

```
sjob(job.id = sq.jobs(),  
     tasks.width = 11)
```

Arguments

job.id	job.id
tasks.width	tasks.width

Value

Data table from [sjob_dt](#).

Author(s)

Toby Hocking <toby.hocking@r-project.org> [aut, cre]

sjob_dt *sjob dt*

Description

Get summary dt from [sacct](#) dt.

Usage

```
sjob_dt(time.dt, tasks.width = 11)
```

Arguments

time.dt	time.dt
tasks.width	tasks.width

Value

data.table summarizing State/ExitCode distribution over jobs

Author(s)

Toby Hocking <toby.hocking@r-project.org> [aut, cre]

Examples

```
library(slurm)
sacct.csv.gz <- system.file(
  "data", "sacct-job13936577.csv.gz", package="slurm", mustWork=TRUE)
if(requireNamespace("R.utils")){
  task.dt <- sacct_fread(sacct.csv.gz)
  print(summary.dt <- sjob_dt(task.dt))
}
```

sq.jobs *sq jobs*

Description

get currently running jobs

Usage

```
sq.jobs(args = "-u $USER")
```


task.pattern

9

Arguments

args args

Value

character scalar: job1,job2,etc

Author(s)

Toby Hocking <toby.hocking@r-project.org> [aut, cre]

<code>task.pattern</code>	<i>task pattern</i>
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Description

Pattern for either one task or a range.

Usage

"task.pattern"

Index

* datasets

- sacct-2019-11-21, 3
- sacct-job13936577, 3
- sacct-multitype, 4

* package

- slurm-package, 2

int.pattern, 2

na.as.zero, 2

sacct, 3, 4–8

sacct-2019-11-21, 3

sacct-job13936577, 3

sacct-multitype, 4

sacct.pattern.list, 4, 5

sacct_fields, 4, 6

sacct_fread, 5, 6

sacct_lines, 3, 5

sacct_tasks, 3, 6

sjob, 7

sjob_dt, 7, 8

slurm (slurm-package), 2

slurm-package, 2

sq.jobs, 8

task.pattern, 9