

Reference Documentation for ODGI

odgi (1)

Erik Garrison

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NAME

odgi - dynamic succinct variation graph tool

SYNOPSIS

odgi build -g graph.gfa -o graph.og

odgi stats -i graph.og -S

DESCRIPTION

Odgi is a set of tools that manipulate variation graphs. TODO Explain data structure. TODO Link to vg paper. TODO Link to handle graph paper. TODO Explain purpose. TODO Give summary of the capabilities of the set of tools.

COMMANDS

Each command has its own man page which can be viewed using e.g. **man odgi_build.1**. Below we have a brief summary of syntax and sub-command description.

odgi build [-g, --gfa=FILE] [-o, --out=FILE] [OPTION]...

The odgi build(1) command constructs a succinct variation graph from a GFA. Currently, only GFA1 is supported. For details of the format please see <https://github.com/GFA-spec/GFA-spec/blob/master/GFA1.md>.

odgi stats [-i, --idx=FILE] [OPTION]...

The odgi stats(1) command produces statistics of a variation graph. Among other metrics, it can calculate the #nodes, #edges, #paths and the total nucleotide length of the graph. Various histogram summary options complement the tool. If [-B, --bed-multicov=BED] is set, the metrics will be produced for the intervals specified in the BED.

LIMITATIONS

BUGS

Refer to the **odgi** issue tracker at <https://github.com/vgteam/odgi/issues>.

AUTHORS

Erik Garrison from the University of California Santa Cruz wrote the whole odgi tool. Despite small code contributions, Simon Heumos from the Quantitative Biology Center Tübingen wrote odgi pathindex, odgi panpos, odgi server, and the documentation.

RESOURCES

Project web site: <https://github.com/vgteam/odgi>

Git source repository on GitHub: <https://github.com/vgteam/odgi>

GitHub organization: <https://github.com/vgteam>

Discussion list / forum: <https://github.com/vgteam/odgi/issues>

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odgi build(1)

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NAME

odgi_build - construct a dynamic succinct variation graph

SYNOPSIS

odgi build [-g, --gfa=*FILE*] [-o, --out=*FILE*] [*OPTION*]...

DESCRIPTION

The odgi build(1) command constructs a succinct variation graph from a GFA. Currently, only GFA1 is supported. For details of the format please see <https://github.com/GFA-spec/GFA-spec/blob/master/GFA1.md>.

OPTIONS

Graph Files IO

-g, --gfa=*FILE*

GFA1 file containing the nodes, edges and paths to build a dynamic succinct variation graph from.

-o, --out=*FILE*

Write the dynamic succinct variation graph to this file. A file ending with *.og* is recommended.

Graph Sorting

-s, --sort

Apply a general topological sort to the graph and order the node ids accordingly. A bidirected adaptation of Kahn's topological sort (1962) is used, which can handle components with no heads or tails. Here, both heads and tails are taken into account.

Processing Information

-p, --progress

Print progress updates to stdout.

-d, --debug

Verbosely print graph information to stderr. This includes the maximum node_id, the minimum node_id, the handle to node_id mapping, the deleted nodes and the path metadata.

--trace

Include backtrace information when reporting errors.

-v, --verbose

Verbosely print processing information to stderr, including debug-level log messages.

-w, --warnings

Turn on script warnings (applies to executed code).

-t, --timings

Print timings report to stderr (time to read, parse, and convert).

Program Information

-h, --help

Print a help message for **odgi build**.

EXIT STATUS

0

Success.

1

Failure (syntax or usage error; parameter error; file processing failure; unexpected error).

BUGS

Refer to the **odgi** issue tracker at <https://github.com/vgteam/odgi/issues>.

AUTHORS

odgi build was written by Erik Garrison.

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odgi stats(1)

Erik Garrison

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NAME

odgi_stats - metrics describing variation graphs

SYNOPSIS

odgi stats [-i, --idx=*FILE*] [*OPTION*]...

DESCRIPTION

The odgi stats(1) command produces statistics of a variation graph. Among other metrics, it can calculate the #nodes, #edges, #paths and the total nucleotide length of the graph. Various histogram summary options complement the tool. If [-B, --bed-multicov=*BED*] is set, the metrics will be produced for the intervals specified in the BED.

OPTIONS

Graph Files IO

-i, --idx=FILE

File containing the succinct variation graph to create statistics from. The file name usually ends with *.og*.

Summary Options

-S, --summarize

Summarize the graph properties and dimensions. Print to stdout the *#nucleotides*, *#nodes*, *#edges* and *#paths* of the graph.

-b, --base-content

Describe the base content of the graph. Print to stdout the *#A*, *#C*, *#G* and *#T* of the graph.

-C, --coverage

Provide a histogram of path coverage over bases in the graph. Print three tab-delimited columns to stdout: **type**, **cov**, **N**. **type** is one of *full* or *uniq* and determines if the histogram corresponds to the full graph or only to a unique paths graph. **cov** implies the *#paths*. **N** implies the *#nucleotides*.

-V, --set-coverage

Provide a histogram of coverage over unique set of paths. Print two tab-delimited columns to stdout: **cov**, **sets**. **cov** implies *#nucleotides*. **sets** lists the unique set of paths in a comma separated list. Sets with a **cov** of one and no paths in **sets** are listed, too.

-M, --multi-coverage

Provide a histogram of coverage over unique multisets of paths. Print two tab-delimited columns to stdout: **cov**, **sets**. **cov** implies *#nucleotides*. **sets** lists the unique multisets of paths in a comma separated list. Multisets with a **cov** of one and no paths in **sets** are listed, too.

BED Interval

-B, --bed-multicov=BED

For each BED entry, provide a table of path coverage over unique multisets of paths in the graph. Each unique multiset of paths overlapping a given BED interval is described in terms of its length relative to the total interval, the number of path traversals and unique paths involved in these traversals.

Threading

-t, --threads=N

Number of threads to use.

Program Information

-h, --help

Print a help message for **odgi stats**.

EXIT STATUS

0

Success.

1

Failure (syntax or usage error; parameter error; file processing failure; unexpected error).

BUGS

Refer to the **odgi** issue tracker at <https://github.com/vgteam/odgi/issues>.

AUTHORS

odgi stats was written by Erik Garrison.

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odgi sort(1)

Erik Garrison

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NAME

odgi_sort - sort a variation graph

SYNOPSIS

odgi sort [-i, --idx=*FILE*] [*OPTION*]...

DESCRIPTION

The odgi sort(1) command produces sorts a succinct a variation graph.

OPTIONS

Graph Files IO

-i, --idx=*FILE*

File containing the succinct variation graph to sort. The file name usually ends with *.og*.

-o, --out=*FILE*

Write the sorted dynamic succinct variation graph to this file. A file ending with *.og* is recommended.

-s, --sort-order=*FILE*

File containing the sort order. Each line contains one node identifier.

Topological Sorts

Threading

-t, --threads=*N*

Number of threads to use for parallel sorting in SGD. Only specify this argument in combination

with **-S, --linear-sgd**. No multi-threading support for any other sorting algorithm.

Processing Information

-P, --progress

Print sort progress to stdout.

Program Information

-h, --help

Print a help message for **odgi sort**.

EXIT STATUS

0

Success.

1

Failure (syntax or usage error; parameter error; file processing failure; unexpected error).

BUGS

Refer to the **odgi** issue tracker at <https://github.com/vgteam/odgi/issues>.

AUTHORS

odgi sort was written by Erik Garrison.

RESOURCES

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