# metagene: a package to produce metagene plots

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Package: metagene (http://bioconductor.org/packages/metagene)

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# 1 Introduction

This package produces metagene-like plots to compare the behavior of DNA-interacting proteins at selected groups of features. A typical analysis can be done in viscinity of transcription start sites (TSS) of genes or at any regions of interest (such as enhancers). Multiple combinations of group of features and/or group of bam files can be compared in a single analysis. Bootstraping analysis is used to compare the groups and locate regions with statistically different enrichment profiles. In order to increase the sensitivity of the analysis, alignment data is used instead of peaks produced with peak callers (i.e.: MACS2 or PICS). The metagene package uses bootstrap to obtain a better estimation of the mean enrichment and the confidence interval for every group of samples.

This vignette will introduce all the main features of the metagene package.

# 2 Loading the metagene package

library(metagene)

# 3 Inputs

## 3.1 Alignment files (BAM files)

There is no hard limit in the number of BAM files that can be included in an analysis (but with too many BAM files, memory may become an issue). BAM files must be indexed. For instance, if you use a file names file.bam, a file named file.bam.bai or file.bai must be present in the same directory.

The path (relative or absolute) to the BAM files must be in a vector:

```
bam_files <- get_demo_bam_files()
bam_files</pre>
```

- ## [1] "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1\_re
  p1.bam"
- ## [2] "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1\_re
  p2.bam"
- ## [3] "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2\_re
  p1.bam"
- ## [4] "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2\_re
  p2.bam"
- ## [5] "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/ctrl.bam"

For this demo, we have 2 samples (each with 2 replicates). It is also possible to use a named vector to add your own names to each BAM files:

```
named_bam_files <- bam_files
names(named_bam_files) <- letters[seq_along(bam_files)]
named_bam_files</pre>
```

```
##
## "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1_rep1.b
am"
##
 b
## "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1_rep2.b
##
## "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2_rep1.b
am"
##
  d
## "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2_rep2.b
am"
##
          "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/ctrl.b
##
am"
```

Using named BAM files can simplify the use of the metagene helper functions and the creation of the design.

## 3.2 Genomic regions

#### 3.2.1 BED files

To compare custom regions of interest, it is possible to use a list of one or more BED files.

```
regions <- get_demo_regions()
regions

## [1] "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/list1.be
d"
## [2] "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/list2.be
d"</pre>
```

The name of the files (without the extension) will be used to name each groups.

metagene also support the narrowPeak (https://genome.ucsc.edu/FAQ/FAQformat.html#format12) and the broadPeak (https://genome.ucsc.edu/FAQ/FAQformat.html#format13).

#### 3.2.2 GRanges or GRangesList objects - Regions

As an alternative to a list of BED files, **Granges** or **GrangesList** objects can be used.

#### 3.2.3 Available datasets

Some common datasets are already available with the metagene package:

- promoters\_hg19
- promoters\_hg18
- promoters\_mm10
- promoters\_mm9

data(promoters\_hg19)
promoters\_hg19

##	GRanges of	object wi	th 23056 rar	nges and $1$ m $_{ m c}$	etadata c	olumn:
##	9	seqnames		ranges :	strand	gene_id
##		<rle></rle>		<iranges></iranges>	<rle>  </rle>	<character></character>
##	1	chr19	[ 58873215,	58875214]	-	1
##	10	chr8	[ 18247755,	18249754]	+	10
##	100	chr20	[ 43279377,	43281376]	-	100
##	1000	chr18	[ 25756446,	25758445]	-	1000
##	10000	chr1	[244005887,	244007886]	-	10000
##						
##	9991	chr9	[115094945,	115096944]	-	9991
##	9992	chr21	[ 35735323,	35737322]	+	9992
##	9993	chr22	[ 19108968,	19110967]	-	9993
##	9994	chr6	[ 90538619,	90540618]	+	9994
##	9997	chr22	[ 50963906,	50965905]	-	9997
##		=				
##	seqinfo	o: 93 seq	juences (1 ci	ircular) fro	n hg19 ge	nome

For more details about each datasets, please refer to their documentation (i.e.: ?promoters\_hq19).

## 3.3 Design groups

A design group contains a set of BAM files that, when put together, represent a logical analysis. Furthermore, a design group contains the relationship between every BAM files present. Samples (with or without replicates) and controls can be assigned to a same design group. There can be as many groups as necessary. A BAM file can be assigned to more than one group.

To represent the relationship between every BAM files, design groups must have the following columns:

- The list of paths to every BAM files related to an analysis.
- One column per group of files (replicates and/or controls).

There is two possible way to create design groups, by reading a file or by directly creating a design object in R.

#### 3.3.1 Design groups from a file

Design groups can be loaded into the metagene package by using a text file. As the relationship between BAM files as to be specified, the following columns are mandatory:

- First column: The list of paths (absolute or relative) to every BAM files for all the design groups, the BAM filenames or the BAM names (if a named BAM. file was used).
- Following columns: One column per design group (replicates and/or controls). The column can take only three values:
  - 0: ignore file
  - 1: input
  - 2: control

The file must also contain a header. It is recommanded to use <code>samples</code> for the name of the first column, but the value is not checked. The other columns in the design file will be used for naming design groups, and must be unique.

Samples	align1	align2
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1_rep1.bam	1	0
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1_rep2.bam	1	0
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2_rep1.bam	0	1
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2_rep2.bam	0	1
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/ctrl.bam	2	2

#### 3.3.2 Design groups from R

It is not obligatory to use a design file, you can create the design data.frame using your prefered method (as long as the restrictions on the values mentioned previously are respected).

For instance, the previous design data.frame could have been create directly in R:

Samples	align1	align2
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1_rep1.bam	1	0
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1_rep2.bam	1	0
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2_rep1.bam	0	1
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2_rep2.bam	0	1
/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/ctrl.bam	2	2

# 4 Analysis steps

A typical metagene analysis will consist steps:

- Extraction the read count of every BAM files in selected regions.
- Conversion in coverage.
- Noise removal
- Normalization of the coverage values.
- Table production.
- Data frame production.
- Generation of the metagene plot.

### 4.1 Minimal analysis

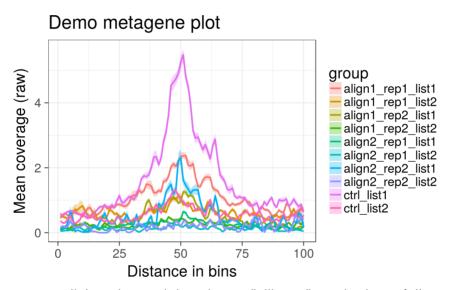
A minimal metagene analysis can be performed in 2 steps:

```
1. Initialization (the new function).
```

2. plot

```
regions <- get_demo_regions()
bam_files <- get_demo_bam_files()
# Initialization
mg <- metagene$new(regions = regions, bam_files = bam_files)
# Plotting
mg$plot(title = "Demo metagene plot")
## produce data table : ChIP-Seq
## produce data frame : ChIP-Seq</pre>
```

## Plot : ChIP-Seq



As you can see, it is not mandatory to explicitly call each step of the metagene analysis. For instance, in the previous example, the plot function call the other steps automatically with default values (the next section will describe the steps in more details).

In this specific case, the plot is messy since by default *metagene* (http://bioconductor.org/packages/metagene) will produce a curve for each possible combinations of BAM file and regions. Since we have 5 BAM files and 2 regions, this gives us 10 curves.

If we want more control on how every step of the analysis are performed, we have to call each functions directly.

# 4.2 Complete analysis

In order to fully control every step of a metagene analysis, it is important to understand how a complete analysis is performed. If we are satisfied with the default values, it is not mandatory to explicitly call every step (as was shown in the previous section).

#### 4.2.1 Initialization

During this step, the coverages for every regions specified are extracted from every BAM files. More specifically, a new **GRanges** is created by combining all the regions specified with the regions param of the **new** function.

```
regions <- get_demo_regions()
bam_files <- get_demo_bam_files()
mg <- metagene$new(regions = regions, bam_files = bam_files)</pre>
```

#### 4.2.2 Producing the table

To produce the table, coverages (produced from Genomics regions (.BED), Alignment Files (.BAM) and Design Sheet) was treated for noise removal and normalized. Furthermore, to reduce the computation time during the following steps, the positions are also binned. Regions, designs, bins, associated values and orientation of strands are pulled into a data.table called 'table' and accessible thanks to the getter get\_table.

We can control the size of the bins with the bin\_count argument. By default, a bin\_count of 100 will be used during this step.

```
mg$produce_table()
## produce data table : ChIP-Seq
```

We can also use the design we produced earlier to remove background signal and combine replicates:

```
mg$produce_table(design = design)
## produce data table : ChIP-Seq
```

#### 4.2.3 Producing the data.frame

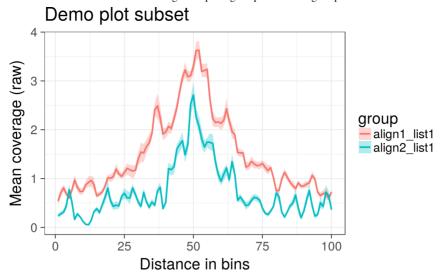
The metagene plot are produced using the ggplot2 package, which require a data.frame as input. During this step, the values of the ribbon are calculated. Metagene uses "bootstrap" to obtain a better estimation of the mean of enrichment for every positions in each groups.

```
mg$produce_data_frame()
## produce data frame : ChIP-Seq
```

#### 4.2.4 Plotting

During this step, metagene will use the data.frame to plot the calculated values using ggplot2. We show a subset of the regions by using the region\_names and design\_names parameter. The region\_names correspond to the names of the regions used during the initialization. The design\_name will vary depending if a design was added. If no design was added, this param correspond to the BAM name or BAM filenames. Otherwise, we have to use the names of the columns from the design.

```
mg$plot(region_names = "list1", title = "Demo plot subset")
## Plot : ChIP-Seq
```



# 5 Manipulating the metagene objects

#### 5.1 Getters

Multiple getters functions are available to access the data that is stored in a metagene object.

#### 5.1.1 get\_table

To get the data.table containing regions, designs, bins, values at bins and orientation of strands.

```
mg <- get_demo_metagene()
mg$produce_table()

## produce data table : ChIP-Seq
mg$get_table()</pre>
```

```
##
         region
                      design bin value strand
      1: list1 align1_rep1
                              1 0.00
##
      2: list1 align1_rep1
                              2
                                  0.75
##
       3: list1 align1_rep1
                             3 1.55
          list1 align1_rep1
                               4 1.15
##
      5:
          list1 align1_rep1
                                  1.25
##
## 49996:
          list2
                        ctrl
                              96
                                  0.00
  49997:
          list2
                              97
                                  0.00
                        ctrl
          list2
  49998:
                                  0.00
                        ctrl
                              98
## 49999:
          list2
                        ctrl
                              99
                                  0.00
## 50000:
                        ctrl 100 0.00
          list2
```

#### 5.1.2 get\_matrices

To get the data.table as matrices (the former data structure)

```
mg <- get_demo_metagene()
mg$produce_table()</pre>
```

## produce data table : ChIP-Seq

m <- mg\$get\_matrices()
# m\$list1\$ctrl\$input to access to region 'list1' and 'ctrl' design</pre>

#### 5.1.3 get\_data\_frame

get\_data\_frame = function(region\_names = NULL, design\_names = NULL) To get the data.frame containing regions and design

mg <- get\_demo\_metagene()
mg\$produce\_table()

## produce data table : ChIP-Seq
mg\$produce\_data\_frame()

## produce data frame : ChIP-Seq</pre>

mg\$get\_data\_frame()

##		region	design	hin	value	strand	qinf	qsup
	1	_	align1_rep1		0.449	3 CT ATIG		0.48680500
	2		align1_rep1		0.552	*		0.60416625
##	3		align1_rep1		0.609	*		0.64951750
##	4		align1_rep1		0.534	*		0.57067125
##	5		align1_rep1		0.522	*		0.55359500
##	6		align1_rep1		0.589	*		0.64152625
##	7		align1_rep1		0.704	*		0.77301625
##	8		align1_rep1	8	0.591	*	0.54773750	0.65927375
##	9		align1_rep1	9	0.562	*	0.50557250	0.63969625
##	10		align1_rep1	10	0.488	*	0.44454125	0.52765750
##	11	list1	align1_rep1	11	0.614	*	0.56367625	0.67129375
##	12	list1	align1_rep1	12	0.745	*	0.65208875	0.83588000
##	13	list1	align1_rep1	13	0.772	*	0.66964625	0.89126500
##	14	list1	align1_rep1	14	0.659	*	0.55825250	0.76395375
##	15	list1	align1_rep1	15	0.481	*	0.42948750	0.55225000
##	16	list1	align1_rep1	16	0.558	*	0.50034000	0.60948125
##	17	list1	align1_rep1	17	0.614	*	0.57199125	0.66838625
##	18	list1	align1_rep1	18	0.708	*	0.66254500	0.75875250
##	19	list1	align1_rep1	19	0.847	*	0.79283875	0.90719750
##			align1_rep1		0.745	*		0.78670500
##	21		align1_rep1	21	0.731	*	0.68396500	0.76956375
##			align1_rep1		0.795	*		0.87101875
##			align1_rep1		0.772	*		0.81412750
##			align1_rep1		0.866	*		0.92642750
##			align1_rep1		0.872	*		0.91655000
##			align1_rep1		0.841	*		0.91615625
	27		align1_rep1		0.808	*		0.90423875
	28		align1_rep1		0.835	*		0.94235500
	29		align1_rep1		0.915	*		0.98224625
	30		align1_rep1		1.123	*		1.24534625
	31		align1_rep1		1.161	*		1.37410875
##	32		align1_rep1		1.122	*		1.24831625
##	33 34		align1_rep1		1.301 1.235	*		1.42765250 1.33553125
##	35		<pre>align1_rep1 align1_rep1</pre>		1.285	*		1.33333123
##			align1_rep1		1.497	*		1.58780125
	37		align1_rep1		1.480	*		1.59286375
	38		align1_rep1		1.433	*		1.57895875
##	39		align1_rep1		1.234	*		1.38964250
##			align1_rep1		1.227	*		1.29442250
##			align1_rep1		1.302	*		1.37638125
##			align1_rep1		1.267	*		1.35204750
##			align1_rep1		1.584	*		1.69500375
##			align1_rep1		1.602	*		1.67481000
##			align1_rep1		1.727	*		1.83965000
##	46		align1_rep1		2.020	*		2.10974375
##	47	list1	align1_rep1	47	2.120	*	2.03103500	2.21784250
##	48		align1_rep1	48	2.044	*	1.95802250	2.14372000
##	49	list1	align1_rep1	49	2.196	*	2.11362375	2.28838500
##	50	list1	align1_rep1	50	2.340	*	2.22450125	2.44324250
##	51	list1	align1_rep1	51	2.367	*	2.27733500	2.48304750
##	52	list1	align1_rep1	52	2.374	*	2.20454000	2.50833000
##	53		align1_rep1		2.086	*		2.22979875
##			align1_rep1		2.053	*		2.16354875
##			align1_rep1		2.126	*		2.23719750
	56		align1_rep1		1.710	*		1.79946500
##	57		align1_rep1		1.429	*		1.53952625
	58		align1_rep1		1.425	*		1.52454875
##	59		align1_rep1		1.618	*		1.68257125
##	60		align1_rep1		1.707	*		1.82527500
##	61		align1_rep1		1.716	*		1.80780625
##	62	I1st1	align1_rep1	62	1.699	*	1.59463500	1.83498250

```
##
  63
         list1 align1_rep1
                             63 1.383
                                            * 1.29198875 1.48455500
##
   64
         list1 align1_rep1
                             64 1.316
                                              1.22987500 1.44782625
##
   65
         list1 align1_rep1
                             65 1.293
                                              1.23328000 1.35369125
##
   66
         list1 align1_rep1
                             66 1.136
                                            * 1.05875250 1.19035250
##
   67
         list1 align1_rep1
                             67 1.095
                                              1.05683625 1.14662125
##
   68
         list1 align1_rep1
                             68 1.028
                                              0.96354750 1.11742875
                                              0.84407125 0.95505000
##
   69
         list1 align1_rep1
                             69 0.895
         list1 align1_rep1
##
   70
                             70 0.825
                                              0.78094250 0.88992875
##
   71
         list1 align1_rep1
                             71 0.898
                                              0.82789125 0.96091000
##
   72
         list1 align1_rep1
                             72 0.856
                                              0.77976875 0.90984875
##
   73
         list1 align1_rep1
                             73 0.879
                                              0.81262250 0.94256500
##
   74
         list1 align1_rep1
                             74 0.899
                                              0.83854750 0.96353875
##
   75
         list1 align1_rep1
                             75 0.958
                                              0.89171375 1.01254875
##
   76
                                              0.97860250 1.12890375
         list1 align1_rep1
                             76 1.048
  77
##
         list1 align1_rep1
                             77 0.963
                                              0.90519125 1.05092625
##
  78
         list1 align1_rep1
                             78 0.941
                                              0.87531125 1.02200750
##
   79
         list1 align1_rep1
                             79 0.823
                                              0.79011125 0.88422375
##
   80
         list1 align1_rep1
                             80 0.564
                                              0.52652625 0.60308375
##
   81
         list1 align1_rep1
                             81 0.486
                                              0.42711000 0.54408125
##
   82
                                              0.67725000 0.79184125
         list1 align1_rep1
                             82 0.730
   83
                                              0.76671875 0.86981875
##
         list1 align1_rep1
                             83 0.820
##
   84
         list1 align1_rep1
                             84 0.711
                                              0.68028250 0.75114125
##
   85
         list1 align1_rep1
                             85 0.702
                                              0.64181250 0.75697000
##
   86
         list1 align1_rep1
                             86 0.664
                                              0.61660750 0.73234125
##
   87
         list1 align1_rep1
                             87 0.508
                                              0.45423000 0.56466250
##
   88
         list1 align1_rep1
                             88 0.538
                                              0.49992500 0.58529375
   89
         list1 align1_rep1
                             89 0.582
                                              0.54993000 0.61298000
##
   90
         list1 align1_rep1
                                              0.66581000 0.73482250
##
                             90 0.701
   91
                                              0.57640375 0.67905250
##
         list1 align1_rep1
                             91 0.628
##
   92
         list1 align1_rep1
                             92 0.649
                                              0.61858625 0.69771875
##
   93
         list1 align1_rep1
                             93 0.900
                                              0.82738375 0.94997000
##
   94
         list1 align1_rep1
                             94 0.830
                                              0.76371875 0.90351000
##
   95
         list1 align1_rep1
                             95 0.590
                                              0.53027000 0.63925375
         list1 align1_rep1
##
   96
                             96 0.557
                                              0.52818375 0.61661500
   97
                                              0.52451000 0.62274750
##
         list1 align1_rep1
                             97 0.575
##
   98
         list1 align1_rep1
                             98 0.525
                                              0.46769125 0.57051000
  99
                             99 0.404
                                              0.36301375 0.45440875
##
         list1 align1_rep1
##
   100
         list1 align1_rep1 100 0.537
                                              0.49653875 0.58144625
##
   101
         list1 align1_rep2
                              1 0.092
                                              0.06266000 0.11710250
##
  102
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```

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##
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                             90 0.148
```

```
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                                            * 0.22314875 0.28102500
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                                            * 0.13780000 0.18268250
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                                            * 0.40082000 0.45725375
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```

```
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##
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                             10 0.020
                                              0.01100000 0.02600000
##
  311
         list1 align2_rep2
                             11 0.015
                                              0.01098625 0.01811625
  312
                                              0.01379000 0.02778625
##
         list1 align2_rep2
                             12 0.021
##
  313
         list1 align2_rep2
                             13 0.029
                                              0.01738875 0.03822625
   314
##
         list1 align2_rep2
                             14 0.153
                                              0.11906000 0.18900250
##
   315
         list1 align2_rep2
                             15 0.220
                                              0.18722500 0.26220000
##
  316
         list1 align2_rep2
                             16 0.177
                                              0.14986750 0.20835625
##
  317
         list1 align2_rep2
                             17 0.342
                                              0.28971500 0.42179000
                                            * 0.34072000 0.52633500
## 318
         list1 align2_rep2
                             18 0.435
```

```
##
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         list1 align2_rep2
                             19 0.603
                                            * 0.47731500 0.76962250
##
   320
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                             20 0.400
                                              0.31052750 0.50549000
##
   321
         list1 align2_rep2
                             21 0.277
                                              0.21283375 0.34110000
##
  322
         list1 align2_rep2
                             22 0.287
                                              0.21272000 0.37319000
##
  323
         list1 align2_rep2
                             23 0.277
                                              0.21984500 0.33166125
  324
##
         list1 align2_rep2
                             24 0.480
                                              0.41511500 0.54483500
   325
                                              0.38086500 0.56655375
##
         list1 align2_rep2
                             25 0.477
   326
##
         list1 align2_rep2
                             26 0.364
                                              0.30227375 0.46605250
##
   327
         list1 align2_rep2
                             27 0.373
                                              0.27401750 0.47422750
##
  328
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                             28 0.558
                                              0.42475000 0.68322750
##
   329
                                              0.33243875 0.48158625
         list1 align2_rep2
                             29 0.397
##
   330
         list1 align2_rep2
                             30 0.266
                                              0.17946750 0.36129750
##
   331
                             31 0.259
                                              0.21608000 0.31114625
         list1 align2_rep2
##
   332
                                              0.45559500 0.65817500
         list1 align2_rep2
                             32 0.555
##
  333
                                              0.20627000 0.29554500
         list1 align2_rep2
                             33 0.247
##
  334
         list1 align2_rep2
                                              0.13799625 0.20304500
                             34 0.163
##
   335
         list1 align2_rep2
                             35 0.275
                                              0.22476500 0.33957500
##
   336
         list1 align2_rep2
                             36 0.155
                                              0.12709500 0.17698750
   337
##
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                             37 0.412
                                              0.36738875 0.46506500
##
   338
                             38 0.509
                                              0.43763000 0.58550750
         list1 align2_rep2
   339
                                              0.09921500 0.15548375
##
         list1 align2_rep2
                             39 0.128
   340
                                              0.22883750 0.29698750
##
         list1 align2_rep2
                             40 0.261
##
   341
         list1 align2_rep2
                             41 0.901
                                              0.80188250 1.00612875
                                              0.73175625 0.90068750
##
   342
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                             42 0.811
##
   343
         list1 align2_rep2
                             43 0.915
                                              0.80296875 1.10424625
##
  344
         list1 align2_rep2
                             44 1.259
                                              1.09269125 1.34008750
##
  345
         list1 align2_rep2
                             45 1.376
                                              1.25218625 1.54948750
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         list1 align2_rep2
                                              1.36039500 1.65173375
##
                             46 1.493
##
   347
                                              1.02717000 1.34210250
         list1 align2_rep2
                             47 1.185
##
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         list1 align2_rep2
                             48 1.324
                                              1.21736625 1.46350875
##
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                             49 2.221
                                              1.96501000 2.45733750
##
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         list1 align2_rep2
                             50 2.301
                                              2.02763125 2.49898875
##
   351
         list1 align2_rep2
                             51 1.610
                                              1.44991000 1.76580625
         list1 align2_rep2
##
   352
                             52 1.464
                                              1.32279875 1.65947875
  353
##
         list1 align2_rep2
                             53 1.342
                                              1.20659250 1.53793375
##
  354
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                                              1.10241750 1.30627375
                             54 1.217
  355
                                              1.22505625 1.42766625
##
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                             55 1.312
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##
         list1 align2_rep2
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                                              1.17259500 1.42706500
##
   357
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                                              1.15079875 1.43373000
##
   358
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                             58 0.944
##
  359
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                             59 0.709
                                              0.63740750 0.85016875
##
   360
                             60 0.642
                                              0.51576375 0.73992750
         list1 align2_rep2
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                                              0.69331875 0.90469625
##
         list1 align2_rep2
                             61 0.807
   362
                                              0.72300125 0.94425875
##
         list1 align2_rep2
                             62 0.846
   363
         list1 align2_rep2
##
                             63 0.693
                                              0.61848375 0.75595375
##
   364
         list1 align2_rep2
                             64 1.135
                                              0.94392500 1.28190750
##
   365
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                                              0.72752875 0.93070875
##
  366
                                              0.23601125 0.35371375
         list1 align2_rep2
                             66 0.297
##
   367
         list1 align2_rep2
                             67 0.370
                                              0.32851250 0.43227750
##
   368
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                                              0.27023000 0.41040125
                             68 0.341
##
   369
                                              0.21366750 0.39522000
         list1 align2_rep2
                             69 0.306
##
   370
         list1 align2_rep2
                             70 0.371
                                              0.26843250 0.46816375
##
   371
                                              0.34668250 0.46461375
         list1 align2_rep2
                             71 0.412
##
   372
         list1 align2_rep2
                             72 0.378
                                              0.29207875 0.45677750
##
   373
         list1 align2_rep2
                             73 0.346
                                              0.29554750 0.39990875
##
  374
         list1 align2_rep2
                             74 0.292
                                              0.21467500 0.34851875
##
  375
         list1 align2_rep2
                             75 0.140
                                              0.10736000 0.16992000
  376
                                              0.04562625 0.09582375
##
         list1 align2_rep2
                             76 0.069
                             77 0.105
##
  377
         list1 align2_rep2
                                              0.06585000 0.14606250
   378
##
         list1 align2_rep2
                             78 0.137
                                              0.09075875 0.19299250
##
   379
         list1 align2_rep2
                             79 0.137
                                              0.10954125 0.16047625
##
   380
         list1 align2_rep2
                             80 0.472
                                              0.38380750 0.57019375
##
  381
         list1 align2_rep2
                             81 0.456
                                              0.37481375 0.52461375
  382
                                            * 0.11811000 0.19064375
##
         list1 align2_rep2
                             82 0.157
```

```
##
   383
         list1 align2_rep2
                              83 0.046
                                             * 0.03214750 0.06504750
##
   384
         list1 align2_rep2
                              84 0.151
                                               0.11694875 0.19586750
##
   385
         list1 align2_rep2
                              85 0.193
                                               0.15564000 0.22789500
##
   386
         list1 align2_rep2
                              86 0.220
                                               0.16840125 0.27937500
##
   387
         list1 align2_rep2
                              87 0.499
                                               0.38941125 0.58585250
   388
                              88 0.337
                                               0.25910000 0.42441625
##
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                              89 0.140
                                               0.11383750 0.18119750
##
         list1 align2_rep2
   390
                                               0.16346625 0.23735375
##
         list1 align2_rep2
                              90 0.203
##
   391
         list1 align2_rep2
                              91 0.378
                                               0.30216125 0.46222625
   392
##
         list1 align2_rep2
                              92 0.556
                                               0.46215625 0.65201875
##
   393
                                               0.23894625 0.34445375
         list1 align2_rep2
                              93 0.284
##
   394
         list1 align2_rep2
                              94 0.081
                                               0.05326875 0.10098000
##
   395
                              95 0.138
                                               0.09431250 0.18834000
         list1 align2_rep2
   396
##
                                               0.27149625 0.46225125
         list1 align2_rep2
                              96 0.361
   397
                                               0.23026875 0.35186875
##
         list1 align2_rep2
                              97 0.298
##
   398
         list1 align2_rep2
                              98 0.606
                                               0.46890000 0.75939000
##
   399
         list1 align2_rep2
                              99 0.536
                                               0.42349000 0.62201500
##
   400
         list1 align2_rep2 100 0.174
                                               0.12288250 0.20352000
                                               0.54950875 0.63605750
##
   401
         list1
                       ctrl
                               1 0.584
##
   402
         list1
                               2 0.515
                                               0.47355625 0.55711125
                       ctrl
##
   403
         list1
                               3 0.592
                                               0.54083875 0.65412500
                       ctrl
   404
                                               0.60926375 0.70741250
##
         list1
                       ctrl
                               4 0.654
##
   405
         list1
                       ctrl
                               5 0.651
                                               0.59533750 0.73836875
##
   406
                                               0.61857875 0.70402000
         list1
                               6 0.660
                       ctrl
##
   407
         list1
                       ctrl
                               7 0.614
                                               0.57836500 0.64459000
##
   408
         list1
                       ctrl
                               8 0.574
                                               0.53395250 0.60348625
##
   409
         list1
                               9 0.567
                                               0.53409000 0.60137750
                       ctrl
                              10 0.568
   410
         list1
                                               0.53877500 0.61865250
##
                       ctrl
                              11 0.644
   411
                       ctrl
                                               0.60202250 0.69862125
##
         list1
##
   412
         list1
                       ctrl
                              12 0.504
                                               0.46330375 0.54284250
##
   413
         list1
                       ctrl
                              13 0.487
                                               0.45792875 0.52105000
##
   414
         list1
                       ctrl
                              14 0.530
                                               0.49300000 0.56547000
##
   415
         list1
                       ctrl
                              15 0.620
                                               0.57278750 0.67980375
##
   416
         list1
                       ctrl
                              16 0.708
                                               0.65791125 0.76573625
##
   417
                              17 0.798
                                               0.74673625 0.84015625
         list1
                       ctrl
##
   418
                              18 0.796
                                               0.75204625 0.84173500
         list1
                       ctrl
   419
                              19 0.855
                                               0.80476750 0.91989750
##
         list1
                       ctrl
                                               0.74509500 0.85757375
##
   420
         list1
                       ctrl
                              20 0.790
##
   421
         list1
                       ctrl
                              21 0.792
                                               0.69310625 0.83648625
##
   422
                              22 0.854
                                               0.79229000 0.93744000
         list1
                       ctrl
##
   423
         list1
                       ctrl
                              23 0.786
                                               0.72707500 0.84102750
##
   424
         list1
                              24 0.984
                                               0.91007125 1.06155000
                       ctrl
   425
                              25 1.143
                                               1.07071250 1.25113375
##
         list1
                       ctrl
   426
                                               0.83945250 0.99961750
##
         list1
                       ctrl
                              26 0.919
##
   427
         list1
                       ctrl
                              27 0.940
                                               0.80788125 1.04076250
##
   428
         list1
                       ctrl
                              28 1.023
                                               0.94403875 1.12652750
##
   429
         list1
                       ctrl
                              29 1.112
                                               1.02974875 1.19297875
                                               1.02371875 1.14231000
##
  430
         list1
                              30 1.076
                       ctrl
##
   431
         list1
                              31 1.020
                                               0.96138500 1.08849625
                       ctrl
##
   432
         list1
                              32 1.217
                                               1.14070125 1.28220500
                       ctrl
##
   433
                                               1.30627250 1.45078375
         list1
                       ctrl
                              33 1.373
##
   434
         list1
                       ctrl
                              34 1.663
                                               1.58499500 1.72300250
   435
##
         list1
                              35 1.750
                                               1.63692750 1.82463500
                       ctrl
##
   436
         list1
                       ctrl
                              36
                                 1.727
                                               1.62515125 1.83417750
##
   437
         list1
                       ctrl
                              37
                                 1.786
                                               1.67645500 1.86019875
##
   438
         list1
                       ctrl
                              38 2.013
                                               1.90335750 2.17886875
##
   439
                              39 2.331
                                               2.15666875 2.43813125
         list1
                       ctrl
   440
                       ctrl
                              40 2.336
                                               2.19115875 2.48718125
##
         list1
                                               1.99723875 2.23116250
##
   441
         list1
                       ctrl
                              41 2.092
                                               2.18400000 2.37780000
##
   442
         list1
                       ctrl
                              42 2.298
##
   443
         list1
                              43 2.821
                                               2.65019500 3.02451750
                       ctrl
##
   444
         list1
                       ctrl
                              44 3.097
                                               2.97059750 3.25868250
## 445
         list1
                       ctrl
                              45 3.621
                                               3.46419625 3.77571625
                                             * 4.32578875 4.80135625
## 446
         list1
                       ctrl
                              46 4.533
```

```
##
  447
         list1
                        ctrl
                              47 4.631
                                              * 4.44690750 4.89002250
##
   448
         list1
                        ctrl
                              48 4.739
                                               4.56665125 4.90769000
   449
##
         list1
                        ctrl
                              49 4.906
                                               4.73844375 5.15870250
##
   450
         list1
                              50 5.254
                                              * 5.01375125 5.41991250
                        ctrl
##
   451
         list1
                        ctrl
                              51 5.477
                                               5.25071500 5.69075875
                                               4.58632375 4.98943000
   452
                              52 4.813
##
         list1
                        ctrl
   453
         list1
                              53 4.509
                                               4.32015875 4.75271125
##
                        ctrl
   454
                                               3.75091500 4.11633750
##
         list1
                        ctrl
                              54 3.935
##
   455
         list1
                        ctrl
                              55 3.140
                                               2.94684750 3.28498375
##
   456
         list1
                        ctrl
                              56 2.912
                                               2.77331375 3.03176875
##
   457
                                               2.79071750 3.06654375
         list1
                        ctrl
                              57 2.945
##
   458
         list1
                        ctrl
                              58 2.748
                                               2.61760000 2.85646750
##
   459
                        ctrl
                              59 2.555
                                               2.43193250 2.68259500
         list1
##
   460
                                               2.55511000 2.84429625
         list1
                        ctrl
                              60 2.708
   461
                                              * 2.33305125 2.63798500
##
         list1
                        ctrl
                              61 2.500
##
   462
         list1
                              62 2.279
                                               2.11350750 2.43116375
                        ctrl
##
   463
         list1
                        ctrl
                              63 2.672
                                               2.50559250 2.86405625
##
   464
         list1
                        ctrl
                              64 2.878
                                               2.68513375 3.03311500
                                               2.12615125 2.39585500
##
   465
         list1
                        ctrl
                              65 2.269
##
   466
         list1
                              66 1.775
                                               1.63098750 1.92174250
                        ctrl
   467
                                               1.57132875 1.79744750
##
         list1
                        ctrl
                              67 1.684
                                               1.47735000 1.65645375
##
   468
         list1
                        ctrl
                              68
                                 1.558
##
   469
         list1
                        ctrl
                              69 1.326
                                               1.25149000 1.42175000
##
   470
                        ctrl
                              70 1.319
                                               1.20242875 1.44584000
         list1
##
   471
         list1
                        ctrl
                              71 1.376
                                               1.29725375 1.45540500
##
   472
         list1
                        ctrl
                              72 1.121
                                              * 1.05367125 1.19552500
##
   473
         list1
                              73 1.067
                                               1.00035125 1.14457125
                        ctrl
                              74 1.111
   474
         list1
                                               1.03866500 1.18546250
##
                        ctrl
##
   475
                              75 1.069
                                               0.99934625 1.13350875
         list1
                        ctrl
##
   476
         list1
                        ctrl
                              76 1.086
                                               1.02163250 1.16895750
##
   477
         list1
                        ctrl
                              77 0.938
                                               0.88032750 1.01783375
                                               0.76477500 0.90000500
##
   478
         list1
                        ctrl
                              78 0.836
##
   479
         list1
                        ctrl
                              79 0.831
                                               0.77222375 0.87638500
##
   480
         list1
                        ctrl
                              80 0.868
                                               0.78972250 0.95342125
##
   481
                              81 0.918
                                               0.85825375 1.02384125
         list1
                        ctrl
##
   482
                              82 0.690
                                               0.63176875 0.71988750
         list1
                        ctrl
   483
                        ctrl
                                               0.71686375 0.80852000
##
         list1
                              83 0.764
                                               0.89545625 1.03117125
##
   484
         list1
                        ctrl
                              84 0.957
                                               0.75758875 0.91073250
##
   485
         list1
                        ctrl
                              85 0.839
##
   486
                                               0.91234250 1.07130750
         list1
                        ctrl
                              86 0.984
##
   487
         list1
                        ctrl
                              87 1.081
                                               0.97427125 1.16472500
##
   488
         list1
                              88 0.868
                                               0.81232750 0.94926875
                        ctrl
   489
                                               0.71216625 0.84667625
##
         list1
                        ctrl
                              89 0.779
   490
                                               0.95296375 1.08006250
##
         list1
                        ctrl
                              90 1.013
##
   491
                        ctrl
                              91 0.984
                                               0.92696125 1.03916875
         list1
##
   492
         list1
                        ctrl
                              92 0.995
                                               0.93194125 1.07852750
##
   493
         list1
                              93 1.035
                                               0.95807375 1.09797250
                        ctrl
##
   494
         list1
                                               0.74440000 0.87826625
                        ctrl
                              94 0.810
##
   495
         list1
                              95 0.902
                                               0.83384750 0.96924500
                        ctrl
                              96 1.034
##
   496
                                               0.92961250 1.15518750
         list1
                        ctrl
##
   497
                              97 0.903
                                               0.79692500 0.96869000
         list1
                        ctrl
##
   498
         list1
                        ctrl
                              98 0.808
                                               0.74956500 0.87561500
   499
                                               0.81756625 0.95778375
##
         list1
                              99 0.876
                        ctrl
##
   500
         list1
                        ctrl
                             100 0.655
                                               0.60142125 0.71916125
   501
##
         list2 align1_rep1
                               1 0.525
                                               0.46844000 0.56933625
##
   502
         list2 align1_rep1
                               2 0.495
                                               0.43355875 0.57123000
##
   503
         list2 align1_rep1
                               3 0.469
                                               0.42296875 0.52139000
##
   504
                                               0.40471875 0.50872125
         list2 align1_rep1
                               4 0.442
   505
##
         list2 align1_rep1
                               5 0.482
                                               0.40589250 0.56621625
   506
                                               0.47225375 0.80433750
##
         list2 align1_rep1
                               6 0.627
                               7 0.719
                                               0.53801250 0.89457625
##
   507
         list2 align1_rep1
##
   508
         list2 align1_rep1
                               8 0.704
                                               0.54020375 0.91339250
##
   509
         list2 align1_rep1
                               9 0.777
                                               0.67160250 0.91258375
                              10 0.825
                                              * 0.67686250 0.98583500
##
   510
         list2 align1_rep1
```

```
##
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         list2 align1_rep1
                             11 0.668
                                            * 0.52540875 0.77734875
##
   512
         list2 align1_rep1
                             12 0.564
                                              0.48744375 0.63908000
##
   513
         list2 align1_rep1
                             13 0.497
                                              0.40874250 0.60845000
##
  514
         list2 align1_rep1
                             14 0.580
                                              0.48856125 0.66320375
##
  515
         list2 align1_rep1
                             15 0.652
                                              0.53380000 0.75968375
                                              0.34659500 0.51914250
   516
##
         list2 align1_rep1
                             16 0.419
##
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```

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   723
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##
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##
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##
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##
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                             38 0.105
##
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##
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         list2 align2_rep1
##
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  766
##
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```

```
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                                            * 0.02539375 0.04061000
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##
##
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##
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##
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##
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##
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                                              0.00000000 0.00000000
##
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##
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##
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                             25 0.060
##
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##
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##
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##
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                                            * 0.02512750 0.05320000
## 830
                             30 0.038
         list2 align2_rep2
```

```
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                                            * 0.06024500 0.10053750
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##
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##
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##
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                                              0.05200000 0.10310000
                                            * 0.01676000 0.03000000
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```

```
##
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                                              * 0.06907000 0.10497000
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##
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##
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                        ctrl
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##
   903
         list2
                        ctrl
                               3 0.311
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   904
##
         list2
                        ctrl
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   905
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         list2
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                        ctrl
##
   906
         list2
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##
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         list2
                        ctrl
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##
   908
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         list2
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                               8 0.340
   909
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##
         list2
                        ctrl
                               9 0.403
##
   910
         list2
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                        ctrl
##
   911
         list2
                        ctrl
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##
   912
         list2
                        ctrl
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   913
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##
         list2
                        ctrl
                              13 0.461
##
   914
         list2
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                        ctrl
##
   915
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                        ctrl
   916
                                               0.35306125 0.42893000
##
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                              16 0.388
##
   917
         list2
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##
   918
                        ctrl
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##
   919
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##
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##
   921
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                        ctrl
   922
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##
                        ctrl
##
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##
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##
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##
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##
   927
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##
   928
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## 918		
## 920	## 918	ctrl_list2
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               ctrl_list2
## 1000
               ctrl_list2
```

#### 5.1.4 get\_params

The various parameters used during the initialization of the metagene object, the production of the table and the production of the plot are saved and can be accessed with the get\_params function:

```
mg <- get_demo_metagene()
mg$get_params()</pre>
```

```
## $padding_size
## [1] 0
##
## $verbose
## [1] FALSE
##
## $bam_files
##
                                                               align1_r
ep1
## "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1_rep1.b
am"
##
                                                              align1_r
ep2
## "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align1_rep2.b
am"
##
                                                              align2_r
ep1
## "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2_rep1.b
am"
##
                                                              align2_r
ep2
## "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/align2_rep2.b
am"
##
                                                                      c
trl
          "/tmp/Rtmp5YXNbI/Rinst4a2e231599bd/metagene/extdata/ctrl.b
##
am"
##
## $force_seglevels
## [1] FALSE
##
## $flip_regions
## [1] FALSE
##
## $assay
## [1] "chipseq"
## $df_needs_update
## [1] TRUE
##
## $df_arguments
## [1] ""
##
## $table_needs_update
## [1] TRUE
```

#### 5.1.5 get\_design

To get the design that was used to produce the last version of the table, you can use the <code>get\_design</code> function:

```
mg$produce_table(design = get_demo_design())

## produce data table : ChIP-Seq

## Alternatively, it is also possible to add a design without produc
ing the
## table:
mg$add_design(get_demo_design())
mg$get_design()
```

#### 5.1.6 get\_bam\_count

To get the number of aligned read in a BAM file, you can use the get\_bam\_count function:

```
mg$get_bam_count(bam_files[1])
## [1] 4635
```

#### 5.1.7 get\_regions

To get all the regions, you can use the get\_regions function:

```
mg$get_regions()
```

```
## GRangesList object of length 2:
## GRanges object with 50 ranges and 0 metadata columns:
##
          segnames
                                   ranges strand
##
             <Rle>
                                <IRanges> <Rle>
##
      [1]
              chr1
                     [16103663, 16105662]
##
      [2]
                     [23921318, 23923317]
              chr1
                    [34848977, 34850976]
##
      [3]
              chr1
##
      [4]
              chr1
                    [36368182, 36370181]
                     [36690488, 36692487]
##
      [5]
              chr1
##
      . . .
              . . .
              chr1 [172081530, 172083529]
##
     [46]
##
     [47]
              chr1 [172081796, 172083795]
##
     [48]
              chr1 [172147016, 172149015]
##
     [49]
              chr1 [172205805, 172207804]
##
     [50]
              chr1 [172260642, 172262641]
##
## ...
## <1 more element>
## seqinfo: 1 sequence from an unspecified genome; no seqlengths
```

It is also possible to extract a subset of the regions with the get\_regions
function:

```
mg$get_regions(region_names = c(regions[1]))
```

```
## GRangesList object of length 1:
## $1ist1
## GRanges object with 50 ranges and 0 metadata columns:
##
          segnames
                                    ranges strand
##
             <Rle>
                                 <IRanges> <Rle>
##
      [1]
              chr1
                    [16103663, 16105662]
##
      [2]
              chr1
                    [23921318, 23923317]
##
      [3]
              chr1
                     [34848977, 34850976]
##
      Γ41
              chr1
                     [36368182, 36370181]
                     [36690488, 36692487]
      [5]
##
              chr1
##
      . . .
              . . .
                                              . . .
##
              chr1 [172081530, 172083529]
     [46]
##
     [47]
              chr1 [172081796, 172083795]
##
     [48]
              chr1 [172147016, 172149015]
              chr1 [172205805, 172207804]
##
     [49]
              chr1 [172260642, 172262641]
##
     Γ50]
##
## ----
## seginfo: 1 sequence from an unspecified genome; no seglengths
```

#### 5.1.8 get\_raw\_coverages

To get the coverages produced during the initialization of the metagene object, you can use the get\_raw\_coverages function. Please note that to save space, metagene will only extract the coverages in the regions provided.

```
coverages[[1]]
## RleList of length 22
## $chr10
## integer-Rle of length 130694993 with 1 run
     Lengths: 130694993
##
##
     Values :
##
## $chr11
## integer-Rle of length 122082543 with 1 run
     Lengths: 122082543
##
##
     values :
##
## $chr12
## integer-Rle of length 120129022 with 1 run
     Lengths: 120129022
##
##
     values :
##
## $chr13
## integer-Rle of length 120421639 with 1 run
##
     Lengths: 120421639
     Values:
##
##
## $chr14
## integer-Rle of length 124902244 with 1 run
##
     Lengths: 124902244
     values:
##
##
## ...
## <17 more elements>
```

coverages <- mg\$get\_raw\_coverages()</pre>

length(coverages)

## [1] 5

It is also possible to extract a subset of all the coverages by providing the filenames:

coverages <- mg\$get\_raw\_coverages(filenames = bam\_files[1:2])
length(coverages)</pre>

## [1] 2

#### 5.1.9 get\_normalized\_coverages

The get\_normalized\_coverages function works exactly like the get\_raw\_coverages function except that it returns the coverages in read per million aligned (RPM).

# 5.2 Chaining functions

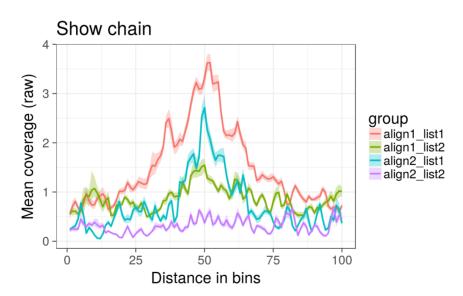
Every function of metagene (except for the getters) invisibly return a pointer to itself. This means that the functions can be chained:

```
rg <- get_demo_regions()
bam <- get_demo_bam_files()
d <- get_demo_design()
title <- "Show chain"
mg <- metagene$new(rg, bam)$produce_table(design = d)$plot(title = t
itle)</pre>
```

## produce data table : ChIP-Seq

## produce data frame : ChIP-Seq

## Plot : ChIP-Seq



## 5.3 Copying a metagene object

To copy a metagene object, you have to use the clone function:

mg\_copy <- mg\$clone()</pre>

# 6 Managing large datasets

While metagene try to reduce it's memory usage, it's possible to run into memory limits when working with multiple large datasets (especially when there is a lot of regions with a large width).

One way to avoid this is to analyse each dataset seperately and then merge just before producing the metagene plot:

```
mg1 <- metagene$new(bam_files = bam_files, regions = regions[1])
mg1$produce_data_frame()
## produce data table : ChIP-Seq
## produce data frame : ChIP-Seq
mg2 <- metagene$new(bam_files = bam_files, regions = regions[2])
mg2$produce_data_frame()
## produce data table : ChIP-Seq
## produce data frame : ChIP-Seq</pre>
```

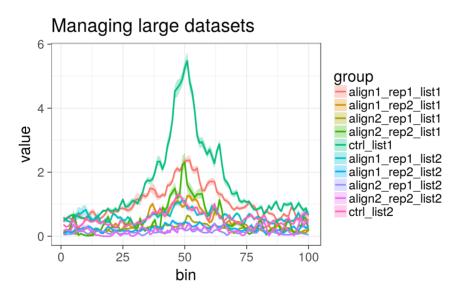
Then you can extract the data.frame s and combine them with rbind:

```
df1 <- mg1$get_data_frame()
df2 <- mg2$get_data_frame()
df <- rbind(df1, df2)</pre>
```

Finally, you can use the plot\_metagene function to produce the metagene plot:

```
p <- plot_metagene(df)
## Plot : ChIP-Seq</pre>
```

p + ggplot2::ggtitle("Managing large datasets")



# 7 Comparing profiles with permutations

It is possible to compare two metagene profiles using the permutation\_test function provided with the metagene package. Please note that the permutation tests functionality is still in development and is expected to change in future releases.

The first step is to decide which profiles we want to compare and extract the corresponding tables:

```
tab <- mg$get_table()
tab0 <- tab[which(tab$region == "list1"),]
tab1 <- tab0[which(tab0$design == "align1"),]
tab2 <- tab0[which(tab0$design == "align2"),]</pre>
```

Then we defined to function to use to compare the two profiles. For this, a companion package of metagene named similaRpeak (http://bioconductor.org/packages/similaRpeak) provides multiple metrics.

For this example, we will prepare a function to calculate the RATIO\_NORMALIZED\_INTERSECT between two profiles:

```
library(similaRpeak)
perm_fun <- function(profile1, profile2) {
    sim <- similarity(profile1, profile2)
    sim[["metrics"]][["RATIO_NORMALIZED_INTERSECT"]]
}</pre>
```

We then compare our two profiles using this metric:

```
## [1] 0.7387951
```

To check if this value is significant, we can permute the two tables that were used to produce the profile and calculate their RATIO\_NORMALIZED\_INTERSECT:

Finally, we check how often the calculated value is greater than the results of the permutations:

```
## [1] 0.021
```