

Downscale alignments

The **Downscale alignments** task can be invoked on data node containing bam/sam files, e.g. *Aligned reads* data node. The only parameter to specify is the percentage of the alignments to keep in the results, the range of the parameter should be between 0 to 100 (Figure 1). All the samples in the input data node will use the same parameter. The output data node contains bam files with a subset of the alignments.

Downscale alignments to percentage

Specify the percentage of output to report for each individual sample.

80

Figure 2. Default filter alignments settings

Additional Assistance

If you need additional assistance, please visit [our support page](#) to submit a help ticket or find phone numbers for regional support.

Your Rating:

Results:

7 rates