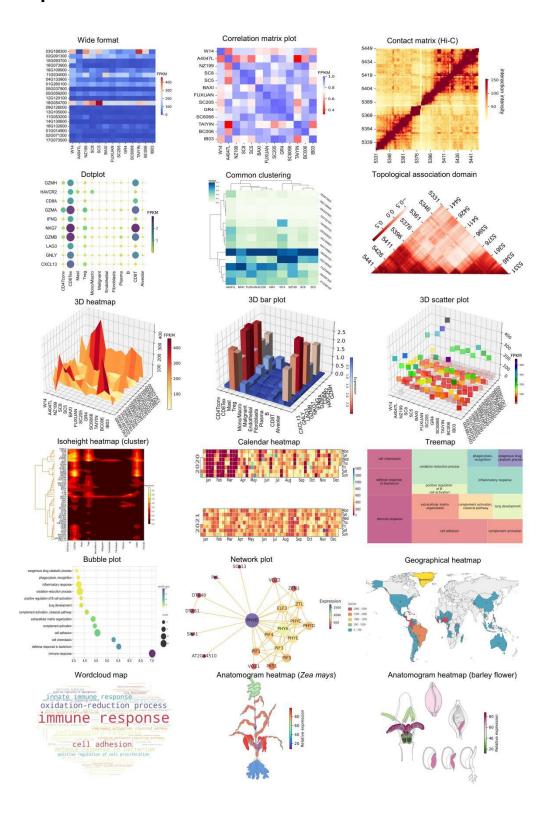
# DataColor v1.0 Software Operation Manual

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# Sample



# **Overview**

With the rapid advancement of high-throughput histology technologies, the need for multidimensional and big data visualization is increasing along with the rapid increase of big biological data. Technological high speed innovation researchers have developed various bioinformatics software, software packages and other tools to draw visualization graphs, but also require users to have a basic knowledge of various programming languages. More and more users need comprehensive, fast, easy and intuitive retrieval of the information contained in big data. Easier visualization of large histological data to solve biological problems is extremely important.

For this purpose, we designed the heat map visualization software DataColor. the tool provides comprehensive adjustment parameters and integrates a large number of general-purpose base and clustering heat maps, as well as isometric heat maps, 3D heat maps, 3D clustering heat maps, 3D scatter heat maps, tree maps, calendar heat maps, bubble maps, network maps, geographic heat maps, word cloud maps, anatomical heat maps, etc., and allows comparison on the same platform of different 25 results under different drawing tools on the same platform.

# **Download and install DataColor**

1.Github download link

https://github.com/frankgenome/DataColor/releases/tag/Software

Link to manual and test data

https://github.com/frankgenome/DataColor

2.figshare download link

https://figshare.com/account/home#/projects/169160

# **Core programs**

# Introduction to the input and output files

The input file is a csv file, which stores tabular data (numbers and text) in plain text, and is recommended to be opened using WORDPAD or Notepad, or by saving a new file and opening it in EXCEL. format files.

The output files are images, scalar images such as png and jpg, and svg vector images, and can also be produced as in pdf format.

# 1. Sample collection

# 1.1 Calendar plotter

The Calendar plotter is an intuitive way to visualise data when doing feature analysis of time-series data.

#### Select

Menu -> Sample collection -> Calendar plotter

# Input

Csv format or Excel format target series file

Date	Counts
2/24/2003	2871
5/7/2003	2765.9
7/1/2003	3884.34
8/25/2003	3746.7
10/10/2003	5205.27
10/28/2003	3479.76

Fig: Input file.

# Output

Generate image in the left canvas

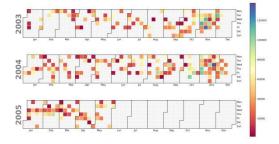


Fig: Calendar plotter output image

#### **Detailed instructions for use**

1. First of all, click the file button on the right, select the Csv format or Excel

format file that needs to be visualized.

- 2. The default parameters have been set, if you need to adjust the parameters, please see the introduction below.
  - 1) Show data displays the data values of each square on the heat map
  - 2) The colour map contains 166 types and is divided into four main categories: Sequential colormaps: continuous colormaps, Diverging colormaps: discrete colormaps at both ends, Qualitative colormaps: discrete colormaps, Miscellaneous colormaps: Other colormaps. For example, 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r colour map is a horizontal flip of the original colour bar.
  - 3) Border widths are the dividing lines for each month of the Calendar plotter, if you need to adjust the colour, just select a value greater than 0 for the width.
  - 4) 'Year lable colour' sets the colour of the month label.
  - 5) Title can set the label.
- 3. After completing all configurations, click the 'Run' button to display or update the Treemap, which will appear in the canvas on the left.
- 4. Users can right-click the "Save" export option to export heat maps, and DataColor provides 900-1000dpi images for users to use. There are four image formats: png, jpg, pdf, and svg, to meet different needs.

# 1.2 Geographical map

Geographical plotter are map-like heat maps that can be displayed in a visual way with colour values based on the values of each geographical location in the data.

#### Select

Menu -> Sample collection -> Geographical map

# Input

Csv format or Excel format target series file

Location	Values
China	95. 1
Canada	23. 2
Brazil	43.3
Russia	66. 4
United	88. 5
States	00. 0
Australia	55

Location	Values
河南	45. 23
安徽	10
黑龙江	11
吉林	3
北京	37. 56
江苏	16
陕西	11
山西	12
河北	21

Fig: Input file.

# Output

Generate image in the left canvas

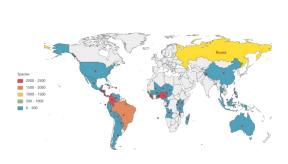


Fig: Geographical map output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. The default parameters have been set, if you need to adjust the parameters, please see the introduction below
  - 1) Type input the map type, such as "world", "china", "贵州", etc.
  - 2) Width and Height set the width and height of the map.
  - 3) Scale-min and Scale-max to customise the total range of values for the Color bar.
  - 4) Color selects whether the map has a colour bar or not.

- 5) Color bar sets its orientation selection.
- 6) Title, table, Color bar can set the label name.
- 3.After completing all the configurations, click on the 'Run' button to display or update the Map, which will pop up as a web page where the user can click on the download button to save the image.

# 1.3 Scatter plot

Datacolor adds the Scatter plot function, which refers to the distribution of data points on the plane of the Cartesian coordinate system. 23 different shapes such as point, diamond, circle, and star can be used to draw the graph. In addition to the color representing the data, the size of the point and the depth of the color are consistent, and at the same time represent the size of the value. And it can realize point synchronous enlargement and synchronous reduction.

# Select

Menu -> Sample collection -> Scatter plot

# Input

Csv format or Excel format target series file

-4	Α	В	C	D	E
1	ID	CD4Tconv	CD8Tex	Mast	Treg
2	CXCL13	0. 16739	2. 34559	0.03696	0.54081
3	GNLY	0. 17927	1. 98292	0.09965	0. 221
4	LAG3	0.07972	1. 51753	0.00396	0. 35516
5	GZMB	0. 46595	2.88772	0. 10262	0.37069
6	NKG7	0.38711	2. 98404	0.1641	0.71816
7	IFNG	0. 23486	1.37227	0.03247	0. 27498
8	GZMA	1.00936	2.82031	0.08614	0.91043
9	CD8A	0. 16879	1.63692	0.02098	0. 33116

Fig: Input file.

# Output

Generate image in the left canvas

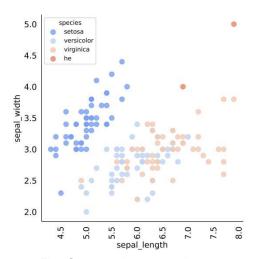


Fig: Scatter plot output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. The default parameters have been set, if you need to adjust the parameters, please see the introduction below.
  - 1) Set the X-axis and Y-axis on and off, font size, color, rotation angle and step size

- 2) Color bar sets its direction, label font size, label name, size, color, position, and font size.
- 3) The color map contains 166 types, which are mainly divided into the following four categories: Sequential colormaps: continuous color map, Diverging colormaps: color map diverging at both ends, Qualitative colormaps: discretized color map, Miscellaneous colormaps: other color maps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc., each suffix \_r color map is the original color bar flipped horizontally.
- 4) Dot shape sets 23 dot shapes
- 5) Dot size magnification can adjust the size of the point
- 6) The Transparency parameter can adjust the transparency of the point color
- 3. After completing all configurations, click the 'Run' button to display or update the Treemap, which will appear in the canvas on the left.
- 4. Users can right-click the "Save" export option to export heat maps, and DataColor provides 900-1000dpi images for users to use. There are four image formats: png, jpg, pdf, and svg, to meet different needs.

# 2 Structure and evolution

# 2.1 Hi-C contact matrix

High-throughput chromosome conformation capture is an experimental technique for studying genome-wide three-dimensional conformations and analysing chromatin fragment interactions.Hi-C aims to understand the three-dimensional conformation of chromatin in the nucleus, to obtain chromatin fragments in close spatial proximity or with interactions in the nucleus and to better study chromatin interactions within or between chromatin. interactions and genome-wide regulation of gene regulatory elements.If you are hic format files, before using this function you need to use HiCexplorer software hicConvertFormat to convert hic/ hicpro matrix files to ginteractions format, and then to txt or csv three column format files.

#### Select

Menu -> Structure and evolution -> Hi-C contact matrix

## Input

TxT format target series file



Fig: txt formatted file.

# **Output**

Generate image in the left canvas

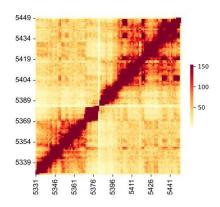


Fig: Hi-C contact matrix output image

## **Detailed instructions for use**

1. First click on the file button on the right and select txt format file you need to visualize.

Default parameters have been set, if you need to adjust the parameters, please see the following introduction

- 1) Set the X-axis and Y-axis on/off, font, font size, color, rotation angle and step length.
- 2) Scale-min and Scale-max can customize the value range of Color bar, or you can choose None, which is automatically defined by the system.
- 3) show data can display the data value of each square on the heat map, and also can adjust the display font size of the data value.
- 4) Scale lable is divided into out, into, and in, which adjusts the orientation of the label teeth of X and Y coordinate axes.
- 5) Cell square is to turn every small square generated by the heat map into a square.
- 6) Color bar sets its direction, label font size, label name, size, and color.
- 7) Color map contains 166 kinds, mainly divided into the following four categories: Sequential colormaps: continuous color maps, Diverging colormaps: color maps with dispersion at both ends, Qualitative colormaps: discrete color maps, Miscellaneous colormaps: Other colormaps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r color map is the original color bar horizontally flipped.
- 8) Line widths is the split line of the heat map, if you need to be able to adjust the color, the width will be greater than 0 value can be selected.
- 3. Once all configurations are complete, click on the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. You can export the plotter by right clicking on the 'Save' export option. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 2.2 Topologically assocaited domain

Genome interoperation mapping is essentially a symmetric matrix with equal information on both sides of the diagonal. The intensity of the interactions ranges from weak to strong, with the colour of the cells transitioning from white to red. There are small triangular regions repeated out of the bottom edge, almost all of which are red inside, indicating a high frequency of interactions between chromatin segments within these regions, such regions are called self-interaction regions, while the frequency of interactions between adjacent triangular regions is low, with red triangular regions corresponding to interactions between regions within TADs, and black regions corresponding to inter-TAD The red triangular areas correspond to the interactions within the TAD, while the black areas correspond to the interactions between TADs. The red triangular region corresponds to the interactions of the intra-TAD region, while the black region corresponds to the interactions between TADs. The triangular interactions are presented on the triangular interactions map, which corresponds to a number of small red triangles on the bottom edge, while the triangles correspond to interactions that are all white. format file, you will need to use HiCexplorer's hicConvertFormat to convert the .hic/ hicpro matrix file to ginteractions format, and then to a three-column file in txt or csv before using this function.

#### Select

Menu -> Structure and evolution -> Topologically assocaited domain Input

TxT format target series file

index	1	index2	freq
5331	5331	154	
5331	5332	154	
5331	5333	129	
5331	5334	87	
5331	5335	58	
5331	5336	69	
5331	5337	65	
5331	5338	48	

Fig: Input file.

# Output

# Generate image in the left canvas

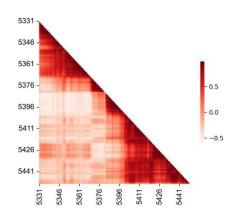


Fig: Topologically assocaited domain output image

#### **Detailed instructions for use**

1. First click on the file button on the right and select txt format file you need to visualize.

Default parameters have been set, if you need to adjust the parameters, please see the following introduction

- 9) Set the X-axis and Y-axis on/off, font, font size, color, rotation angle and step length.
- 10) Scale-min and Scale-max can customize the value range of Color bar, or you can choose None, which is automatically defined by the system.
- 11) show data can display the data value of each square on the heat map, and also can adjust the display font size of the data value.
- 12) Scale lable is divided into out, into, and in, which adjusts the orientation of the label teeth of X and Y coordinate axes.
- 13) Cell square is to turn every small square generated by the heat map into a square.
- 14) Color bar sets its direction, label font size, label name, size, and color.
- 15) Color map contains 166 kinds, mainly divided into the following four categories: Sequential colormaps: continuous color maps, Diverging colormaps: color maps with dispersion at both ends, Qualitative colormaps: discrete color maps, Miscellaneous colormaps: Other colormaps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r color map is the original color bar horizontally flipped.
- 16) Line widths is the split line of the heat map, if you need to be able to adjust the color, the width will be greater than 0 value can be selected.
- 3. Once all configurations are complete, click on the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. You can export the plotter by right clicking on the 'Save' export option. Four

image formats are available: png, jpg, pdf and svg to suit different needs.

# 2.3 Chr. density plotter

The Chromosome Density tool is a mapping tool for representing protein-protein interactions, gene regulatory networks, and metabolic pathways. This facilitates the study of functional associations between biomolecules, regulation of biological pathways, and mechanisms of disease pathogenesis.

## Select

Menu -> Structure and evolution -> Chr. density plotter

# Input

Csv format or Excel format target series file

4	A	В	C	D	7.4	Δ	В
1	Chromosome	Start	End	Count	4		70
2	1	0	100000	24	1	Chromosome	Length
3	1	100000	200000	33	2	1	30427671
4	1	200000	300000	31	3	2	19698289
5	1	300000	400000	28	1		
6	1	400000	500000	37	4	3	23459830
7	1	500000	600000	29	5	4	18585056
8	1	600000	700000	31	6	5	26975502

Fig: Input file.

# **Output**

Generate image in the left canvas

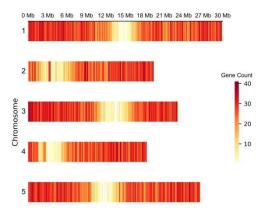


Fig: Chr. density plotter output image

# **Detailed instructions for use**

- 1. First, click the file button on the right and select the Csv or Excel file that you want to visualize
- 2. The default parameters have been set. If you need to adjust the parameters, please see the following

# Detailed usage:

- 1) Set the X-axis and Y-axis on and off, font, font size, color, rotation angle and step size
- 2) Scale-min and Scale-max can customize the value range of Color bar, or select None, which will be automatically defined by the system.

- 3) Scale lable is divided into out, into, and in, and adjusts the orientation of the label teeth of the X and Y coordinate axes.
- 4) Color bar sets its direction, label font size, label name, size and color.
- 5) The color map contains 166 types, which are mainly divided into the following four categories: Sequential colormaps: continuous color map, Diverging colormaps: color map diverging at both ends, Qualitative colormaps: discretized color map, Miscellaneous colormaps: other color maps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc., each suffix \_r color map is the original color bar flipped horizontally.
- 6) Title and Y-axis lable set the input text
- 3. After completing all configurations, click the "Run" button to display or update the heat map. The heat map will appear in the left canvas.
- 4. You can right-click Save to export the heat map. DataColor provides six data ranges from 300 dpi to 1000dpi. There are four image formats: png,.jpg, pdf,.SVG to meet different needs.

# 2.4 Density peak plotter

Density peak plot mainly shows the same X-axis data, which can be time series, genetic data, etc., corresponding to different Y-axis data, and clearly shows the relationship between different data and variables.

#### Select

Menu -> Structure and evolution -> Density peak plotter

# Input

Csv format or Excel format target series file

4	Α	В	C
1	Ks1	Ks2	Ks3
2	2.608285	1.726104	2. 158059
3	2.883082	1.730459	3.411821
4	3. 528264	1.754804	3.083901
5	3.894024	0.809481	2.645047
6	3.460463	1. 780281	2.704113

Fig: Input file.

# **Output**

Generate image in the left canvas

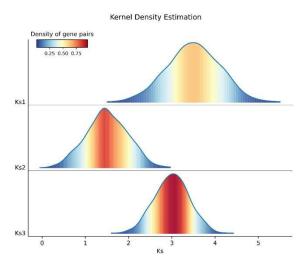


Fig: Density peak plotter output image

# **Detailed instructions for use**

- 1. First, click the file button on the right and select the Csv or Excel file that you want to visualize
- 2. The default parameters have been set. If you need to adjust the parameters, please see the following

# Detailed usage:

- 1) Set X axis and Y axis on and off, font, size, color, rotation Angle
- 2) Color bar sets its direction, label size, label name, size, color, position, and size.

- 3) color map contains 166 kinds, mainly divided into the following four categories: Sequential colormaps: continuous colormap, Diverging colormaps: divergent colormap at both ends, Qualitative colormaps: Miscellaneous colormaps: miscellaneous colormaps. For example, 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc., each suffix \_r color map is the original color bar flipped horizontally.
- 4) Vertical spacing Sets the distance between subplots
- 5) Y-axis lable sets the rotation direction and font size
- 6) X-axis lable sets the input word and size
- 3. After completing all configurations, click the "Run" button to display or update the heat map. The heat map will appear in the left canvas.
- 4. You can right-click Save to export the heat map. DataColor provides six data ranges from 300 dpi to 1000dpi. There are four image formats: png,.jpg, pdf,.SVG to meet different needs.

# 3 Expression analysis

# 3.1 2D plotter

# 3.1.1 Wide format

Wide Format is a convenient heat map where the colour of the cells varies according to their values. This method has no clustering and can be chosen if the collation of data is not required.

## **Select**

Menu -> Expression analysis -> 2D plotter -> Wide format

# Input

Csv format or Excel format target series file

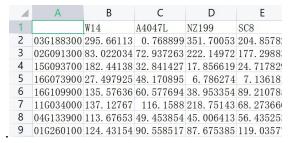


Fig: Input file.

# **Output**

Generate image in the left canvas

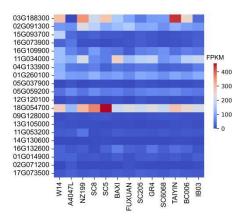


Fig: Wide Format output image

#### Detailed instructions for use

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, if you need to adjust the parameters, please see the following introduction
  - 1) Set the X-axis and Y-axis on/off, font, font size, color, rotation angle and step length.
  - 2) Scale-min and Scale-max can customize the value range of Color bar, or you can choose None, which is automatically defined by the system.
  - 3) show data can display the data value of each square on the heat map, and also can adjust the display font size of the data value.
  - 4) Scale lable is divided into out, into, and in, which adjusts the orientation of the label teeth of X and Y coordinate axes.
  - 5) Cell square is to turn every small square generated by the heat map into a square.
  - 6) Color bar sets its direction, label font size, label name, size, and color.
  - 7) Color map contains 166 kinds, mainly divided into the following four categories: Sequential colormaps: continuous color maps, Diverging colormaps: color maps with dispersion at both ends, Qualitative colormaps: discrete color maps, Miscellaneous colormaps: Other colormaps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r color map is the original color bar horizontally flipped.
  - 8) Line widths is the split line of the heat map, if you need to be able to adjust the color, the width will be greater than 0 value can be selected.
- 3. After finishing all the configurations, click the "Run" button to display or update the heat map, which will appear in the canvas on the left.

Users can export the heat map by right-clicking the "Save" export option, and DataColor provides 6 levels of images from 300-1000dpi for users to use. There are four image formats: png, .jpg, pdf, and .svg, to meet different needs.

# 3.1.2 Correlation Matrix

Correlation Matrix is used to find the pairwise correlation of all columns in a data frame and to discover which variable is correlated with another variable by visualisation.

#### Select

Menu -> Expression analysis -> 2D plotter -> Correlation matrix Input

Csv format or Excel format target series file

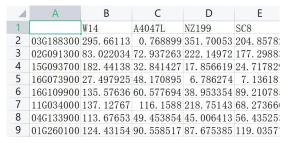


Fig:Input file.

# Output

Generate image in the left canvas

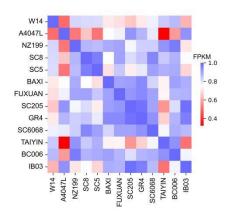


Fig: Correlation Matrix output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, if you need to adjust the parameters please see Wide Format parameters.
- 3. Once all configurations are complete, click on the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. You can export the plotter by right clicking on the 'Save' export option. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 3.1.3 Half Correlation matrix

Correlation Matrix is used to find the pairwise correlation of all columns in a data frame and to discover which variable is correlated with another variable by visualisation.

#### Select

Menu -> Expression analysis -> 2D plotter -> Half Correlation matrix Input

Csv format or Excel format target series file

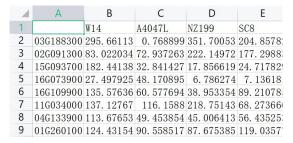


Fig: Input file.

# Output

Generate image in the left canvas

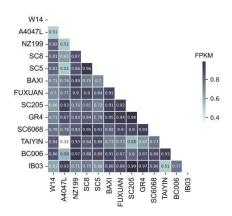


Fig: Half Correlation matrix output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, if you need to adjust the parameters please see Wide Format parameters.
- 3. Once all configurations are complete, click on the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. You can export the plotter by right clicking on the 'Save' export option. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 3.1.4 Circle heatmap

Circle plotter, as the name suggests, is a form of heat map. The advantage of Circle plotter is that it can display multiple aspects on a single graph, which is suitable for the study of multiple groups or multiple omics, and can reveal the change rules and connections of different omics.

#### Select

Menu -> Expression analysis -> 2D plotter -> Circle heatmap

# Input

Csv format or Excel format target series file

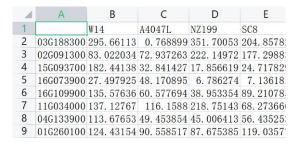


Fig:Input file.

# **Output**

Generate image in the left canvas

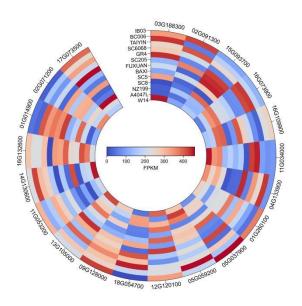


Fig: Circle heatmap output image

- 1. First, click the file button on the right and select the Csv or Excel file that you want to visualize
- 2. The default parameters have been set. If you need to adjust the parameters,

please see the following description

- 1) Set X axis and Y axis on and off, font, size, color, rotation Angle.
- 2) show data can display the data value of each square on the heat map, and can also adjust the display size of the data value.
- Color bar sets its direction, label size, label name, height and width, and color.
- 4) color map contains 166 kinds, mainly divided into the following four categories: Sequential colormaps: continuous colormap, Diverging colormaps: divergent colormap at both ends, Qualitative colormaps: Miscellaneous colormaps: miscellaneous colormaps. For example, 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc., each suffix \_r color map is the original color bar flipped horizontally.
- 5) Border widths are the dividing lines of the heat map. The color can be adjusted if necessary, and the width can be selected as a value greater than 0.
- 3. After completing all configurations, click the "Run" button to display or update the heat map. The heat map will appear in the left canvas.
- 4. You can right-click Save to export the heat map. DataColor provides six data ranges from 300 dpi to 1000dpi. There are four image formats: png,.jpg, pdf,.SVG to meet different needs.

# 3.1.5 Normalization (Column/ Row)

The design of the normalised class heat map parameters is consistent with the base class heat map, unlike the mean normalisation equation we set up in the programming of the normalised class heat map. Normalisation is a fundamental task in data mining. Different evaluation indicators often have different magnitudes and magnitude units, such a situation will affect the results of data analysis, in order to eliminate the influence of magnitudes between indicators, data normalisation is needed to address the comparability between data indicators. After the raw data has been standardised, the indicators are at the same order of magnitude and are suitable for comprehensive comparative evaluation. The mean normalisation equation gives the mean and standard deviation of the raw data to standardise the data. Column is selected for row normalisation; Row is selected for column normalisation.

#### **Select**

Menu -> Expression analysis -> 2D plotter -> Normalization by Col\Normalization by Row

#### Input

Csv format or Excel format target series file

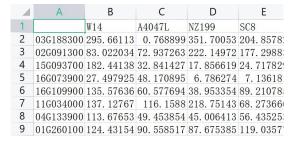


Fig: Input file.

## Output

Generate image in the left canvas

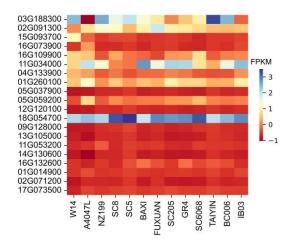


Fig: Normalization by Col\Normalization by Row output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, if you need to adjust the parameters please see Wide Format parameters.
- 3. Once all configurations are complete, click on the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. You can export the plotter by right clicking on the 'Save' export option. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 3.1.6 Common clustering

Common clustering is a heat map with tree-like clustering for users who want to cluster their data. This type of graph can be simply understood as a cluster of data values that are close to each other using a distance algorithm on top of the original base plotter. Seven clustering methods and 22 distance measures are provided. **Select** 

Menu -> Expression analysis -> 2D plotter -> Common clustering

# Input

Csv format or Excel format target series file

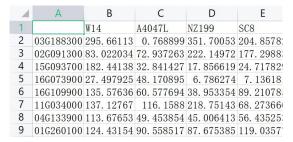


Fig: Input file.

## **Output**

Generate image in the left canvas

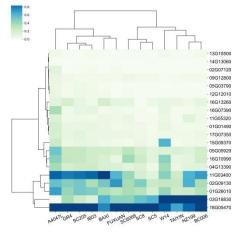


Fig: Common clustering output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, if you need to adjust the parameters please see the following introduction.
  - 1) X-axis and Y-axis set the X-axis and Y-axis on/off, font, font size, colour, rotation angle and step length.

- 2) Row cluster and Col cluster control the clustering of rows and columns on and off.
- 3) Scale-min and Scale-max allow you to define the range of values for the Color bar, or you can choose None, which is automatically defined by the system.
- 4) show data allows you to display the data values of each square on the heat map, and to adjust the font size of the data values.
- 5) Metric allows you to choose from 7 clustering methods.
- 6) Method allows you to choose from 22 distance measures.
- 7) Cell square converts each of the resulting small squares of the heat map into a square.
- 8) Color bar sets its orientation, label font size, label name, size and colour.
- 9) The colour map contains 166 types and is divided into the following four main categories: Sequential colormaps: continuous colormaps, Diverging colormaps: colormaps with dispersion at both ends, Qualitative colormaps: discrete colormaps, Miscellaneous colormaps. Other colormaps. For example, 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r colour map is a horizontal flip of the original colour bar.
- 10) Line widths are the dividing lines of the heat map, if you need to adjust the colour, select a value greater than 0 for the width.
- 11) Column comment enables column commenting, comment column name enters the name of the column to comment on, which is also the name of the comment column in the diagram.
- 12) Classify color1, Classify color2, Classify color3 select what colour.
- 13) Tree style sets the shape, width and colour of the clustering tree.
- 14) Robust opens the possibility to use quantile to calculate the colour mapping range.
- 3. Once all configurations are complete, click the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. The user can export the plotter by right clicking on the 'Save' export option, DataColor offers a range of images from 900-1000 dpi for use. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 3.1.7 Standard clustering

Standard clustering is a normalised approach to clustering plotters. The dimension is normalised for rows or columns, i.e. each row or column is subtracted from the minimum value of the data and divided by the maximum value of the data.

#### Select

Menu -> Expression analysis -> 2D plotter -> Standard clustering

# Input

Csv format or Excel format target series file

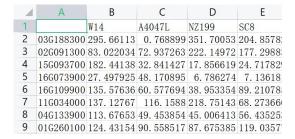


Fig: Input file.

## **Output**

Generate image in the left canvas

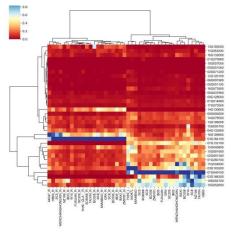


Fig: Standard clustering output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, if you need to adjust parameters please see Cluster parameters.
- 3. Once all the configuration is complete, click on the "Run" button to display or update the heat map, which will appear in the canvas on the left.

4. You can export the plotter by right clicking on the 'Save' export option. Four image formats are available: png, jpg, pdf and svg to suit different needs.				

# 3.1.8 Z-score clustering

The Z-score is a normalisation of the clustering heat map. the Z-score equation gives the mean and standard deviation of the original data to normalise the data.

#### Select

Menu -> Expression analysis -> 2D plotter -> Z-score clustering Input

Csv format or Excel format target series file

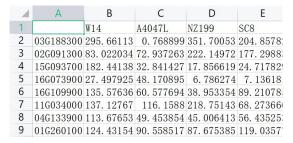


Fig:Input file.

# Output

Generate image in the left canvas

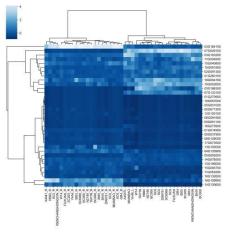


Fig: Z-score clustering output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, if you need to adjust parameters please see Cluster parameters.
- 3. Once all the configuration is complete, click on the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. You can export the plotter by right clicking on the 'Save' export option. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 3.2 3D plotter

# 3.2.1 3D non-cluster

The 3D plotter displays three dimensions of data at the same time, making the results of complex data analysis much clearer and easier to obtain directly from the graph.

#### Select

Menu -> Expression analysis -> 3D plotter -> 3D non-cluster

# Input

Csv format or Excel format target series file

4	Α	В	C	D	E
1	ID	CD4Tconv	CD8Tex	Mast	Treg
2	CXCL13	0. 16739	2. 34559	0.03696	0.54081
3	GNLY	0.17927	1. 98292	0.09965	0. 221
4	LAG3	0.07972	1.51753	0.00396	0.35516
5	GZMB	0.46595	2.88772	0.10262	0.37069
6	NKG7	0.38711	2. 98404	0.1641	0.71816
7	IFNG	0. 23486	1. 37227	0.03247	0.27498
8	GZMA	1.00936	2.82031	0.08614	0.91043
9	CD8A	0. 16879	1.63692	0.02098	0. 33116

Fig: Input file.

# Output

Generate image in the left canvas

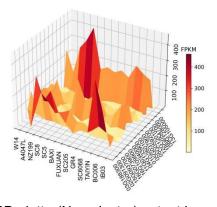


Fig: 3D plotter(Non-cluster) output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. The default parameters have been set, if you want to adjust the parameters please see the following introduction.
  - 1) X-axis, Y-axis, Z-axis set the X-axis, Y-axis, Z-axis on and off, font, font size, colour, rotation angle
  - 2) Color bar sets the direction, label size, label name, size and colour.
  - 3) The colour map contains 166 types, mainly divided into the following

four categories: Sequential colormaps: continuous colormaps, Diverging colormaps: discrete colormaps at both ends, Qualitative colormaps: discrete colormaps, Miscellaneous colormaps. Other colormaps. For example, 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r colour map is a horizontal flip of the original colour bar.

- 4) Number of colour bar scales set the number of scales
- 5) Bar lable and Bar lable size can set the label and label size of the Color bar.
- 6) Title, X-lable, Y-lable, Z-lable can set the label.
- 3. Once all configurations are completed, click the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. Users can export the plotter by right clicking on the 'Save' export option. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 3.2.2 3D cluster by Row

The 3D cluster by Row performs column clustering on top of the 3D plotter.

Seven clustering methods and 22 distance metrics are provided.

#### Select

Menu -> Expression analysis -> 3D plotter -> 3D Cluster by Row

# Input

Csv format or Excel format target series file

4	А	В	С	D	E
1	ID	CD4Tconv	CD8Tex	Mast	Treg
2	CXCL13	0. 16739	2. 34559	0.03696	0.54081
3	GNLY	0.17927	1. 98292	0.09965	0. 221
4	LAG3	0.07972	1.51753	0.00396	0.35516
5	GZMB	0.46595	2.88772	0.10262	0.37069
6	NKG7	0.38711	2. 98404	0.1641	0.71816
7	IFNG	0. 23486	1. 37227	0.03247	0. 27498
8	GZMA	1.00936	2.82031	0.08614	0. 91043
9	CD8A	0.16879	1.63692	0.02098	0.33116

Fig: Input file.

# **Output**

Generate image in the left canvas

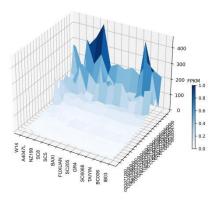


Fig: 3D cluster by Row output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. The default parameters have been set, if you want to adjust the parameters please see the following introduction.
  - 1) X-axis, Y-axis, Z-axis set the X-axis, Y-axis, Z-axis on and off, font, font size, colour, rotation angle.
  - 2) Metric can choose 7 types of clustering methods.
  - 3) Method allows you to choose 22 distance metrics.
  - 4) Color bar sets its direction, label font, label name, size and colour.

- 5) The colour map contains 166 types, mainly divided into the following four categories: Sequential colormaps: continuous colour maps, Diverging colormaps: colour maps with dispersion at both ends, Qualitative colormaps: discrete colour maps, Miscellaneous colormaps. Other colormaps. For example, 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r colour map is a horizontal flip of the original colour bar.
- 6) Number of colour bar scales set the number of scales
- 7) Bar lable and Bar lable size can set the label and label size of the Color bar
- 8) Title, X-lable, Y-lable, Z-lable can set the label
- 3. Once all configurations are completed, click the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. The user can export the plotter by right clicking on the 'Save' export option, DataColor offers 900-1000dpi images for the user to use. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 3.2.3 3D cluster by Col

The 3D plotter (Cluster by Col) performs row clustering on top of the 3D plotter. Seven clustering methods and 22 distance metrics are provided.

### Select

Menu -> Expression analysis -> 3D plotter -> Cluster by Col

## Input

Csv format or Excel format target series file

4	А	В	C	D	E
1	ID	CD4Tconv	CD8Tex	Mast	Treg
2	CXCL13	0. 16739	2. 34559	0.03696	0.54081
3	GNLY	0.17927	1. 98292	0.09965	0. 221
4	LAG3	0.07972	1.51753	0.00396	0.35516
5	GZMB	0.46595	2.88772	0.10262	0.37069
6	NKG7	0.38711	2. 98404	0.1641	0.71816
7	IFNG	0. 23486	1. 37227	0.03247	0. 27498
8	GZMA	1.00936	2.82031	0.08614	0.91043
9	CD8A	0.16879	1. 63692	0. 02098	0. 33116

Fig: Input file.

## **Output**

Generate image in the left canvas

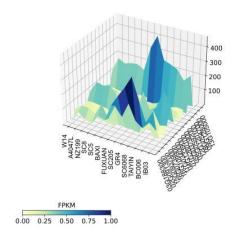


Fig: 3D cluster by Col output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, to adjust parameters see 3D cluster by Col parameters.
- 3. Once all configurations are completed, click the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. Users can export the plotter by right clicking on the 'Save' export option.

Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 3.2.4 3D scatter plotter

The 3D Scatter plotter is very similar to the 3D plotter in that all data is represented by a square scatter in 3D coordinates and the colour is represented by the corresponding data in 2D. The most characteristic feature is that the 3D plotter when viewed from an overhead perspective is extremely similar to the 2D plotter presentation, ensuring the accuracy of the plotter. The third data can be represented both by colour differences and by the height in 3D space.

### Select

Menu -> Expression analysis -> 3D plotter -> 3D scatter plotter Input

Csv format or Excel format target series file

4	Α	В	C	D	E
1	ID	CD4Tconv	CD8Tex	Mast	Treg
2	CXCL13	0. 16739	2. 34559	0.03696	0.54081
3	GNLY	0.17927	1. 98292	0.09965	0. 221
4	LAG3	0.07972	1. 51753	0.00396	0.35516
5	GZMB	0. 46595	2.88772	0.10262	0.37069
6	NKG7	0.38711	2. 98404	0. 1641	0.71816
7	IFNG	0. 23486	1. 37227	0.03247	0. 27498
8	GZMA	1.00936	2.82031	0.08614	0.91043
9	CD8A	0. 16879	1.63692	0. 02098	0. 33116

Fig: Input file.

## Output

Generate image in the left canvas

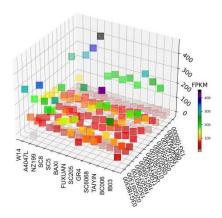


Fig: 3D scatter plotter output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, to adjust parameters please see 3D plotter parameters.
- 3. Once all configurations are completed, click the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. Users can export the plotter by right clicking on the 'Save' export option. Four image formats are available: png, jpg, pdf and svg to suit different needs.

## 3.2.5 3D Bar plotter

The 3D Bar plotter, much like the 3D Scatter plotter, represents all data as columns in 3D coordinates, with the colours being represented by their 2D counterparts, the data being represented both by colour differences and by the height of the columns.

#### Select

Menu -> Expression analysis -> 3D plotter -> 3D Bar plotter]

### Input

Csv format or Excel format target series file

	Α	В	C	D	E
1	ID	CD4Tconv	CD8Tex	Mast	Treg
2	CXCL13	0. 16739	2. 34559	0.03696	0.54081
3	GNLY	0.17927	1. 98292	0.09965	0. 221
4	LAG3	0.07972	1. 51753	0.00396	0.35516
5	GZMB	0.46595	2.88772	0.10262	0.37069
6	NKG7	0.38711	2. 98404	0. 1641	0.71816
7	IFNG	0. 23486	1. 37227	0.03247	0. 27498
8	GZMA	1.00936	2.82031	0.08614	0. 91043
9	CD8A	0. 16879	1. 63692	0.02098	0. 33116

Fig: Input file.

## **Output**

Generate image in the left canvas

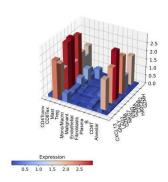


Fig: 3D Bar plotter output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. Default parameters have been set, to adjust parameters please see 3D plotter parameters.
- 3. Once all configurations are completed, click the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. Users can export the plotter by right clicking on the 'Save' export option. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 3.3 Isoheight plotter

## 3.3.1 Isoheight plotter (non-cluster)

Isoheight plotter (Non-cluster) is a plotter model that we have created by combining contour maps with tree clustering. Isoheight plotters use a two-dimensional format to represent three-dimensional data. This type of plotter is better suited to large amounts of biological data, and the differences in colour allow us to see the differences between data at a macro level.

#### Select

Menu -> Expression analysis -> Isoheight plotter -> Isoheight plotter (Non-cluster)

### Input

Csv format or Excel format target series file

-4	А	В	C	D	E
1	ID	CD4Tconv	CD8Tex	Mast	Treg
2	CXCL13	0. 16739	2. 34559	0.03696	0.54081
3	GNLY	0.17927	1. 98292	0.09965	0. 221
4	LAG3	0.07972	1. 51753	0.00396	0.35516
5	GZMB	0. 46595	2.88772	0. 10262	0.37069
6	NKG7	0. 38711	2. 98404	0. 1641	0.71816
7	IFNG	0. 23486	1. 37227	0.03247	0. 27498
8	GZMA	1.00936	2.82031	0.08614	0.91043
9	CD8A	0. 16879	1.63692	0.02098	0. 33116

Fig: Input file.

### **Output**

Generate image in the left canvas

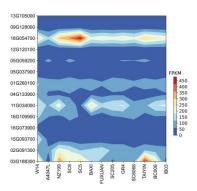


Fig: Isoheight plotter (Non-cluster) output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. The default parameters have been set, if you need to adjust the parameters, please see the introduction below
  - 1) X-axis and Y-axis set the X-axis and Y-axis on and off, font, font size,

- color, rotation angle and step size.
- 2) Row cluster and Col cluster control the opening and closing of the clustering dendrogram of rows and columns.
- 3) Scale-min and Scale-max can customize the total value range of the Color bar, and Scale interval sets the value range of each color block on the Color bar, and can change the fineness of the color to adjust the drawing effect.
- 4) Show data can display the data value of each grid on the heat map, and can also adjust the display font size of the data value.
- 5) Color bar sets its direction, label font size, label name, size and color.
- 6) There are 166 color maps, which are mainly divided into the following four categories: Sequential colormaps: continuous color maps, Diverging colormaps: color maps that diverge at both ends, Qualitative colormaps: discrete color maps, Miscellaneous colormaps: other color maps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc., each suffix \_r color map is the original color bar flipped horizontally.
- 7) Number of color bar scales Set the number of scales.
- 8) Bar lable and Bar lable size can set the label and label font size of Color bar.
- 3. After completing all configurations, click the "Run" button to display or update the plotter, and the plotter will appear in the left canvas.
- 4. Users can right-click the "Save" export option to export heat maps, and DataColor provides 900-1000dpi images for users to use. There are four image formats: png, jpg, pdf, and svg, to meet different needs.

# 3.3.2 Isoheight plotter (Cluster)

Isoheight plotter is a heat map mode that we combine contour map and dendritic clustering. The contour heat map uses two-dimensional form to reflect three-dimensional data. This type of heat map is more suitable for a large amount of biological data, and the high expression value is surrounded by clustering. Through the color difference, we can see the difference between the data macroscopically. The clustering effect is reflected by a variety of tree colors, which is also one of the highlights of the contour heat map.

#### Select

Menu -> Expression analysis -> Isoheight plotter -> Isoheight plotter (Cluster)
Input

Csv format or Excel format target series file

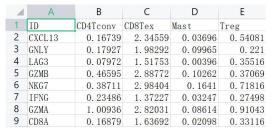


Fig: Input file.

## Output

Generate image in the left canvas

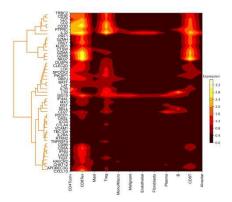


Fig: Isoheight plotter (Cluster) output image

### **Detailed instructions for use**

1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.

- 2. The default parameters have been set, if you need to adjust the parameters, please see the introduction below
  - 1) X-axis and Y-axis set the X-axis and Y-axis on and off, font, font size, color, rotation angle and step size.
  - 2) Row cluster and Col cluster control the opening and closing of the clustering dendrogram of rows and columns.
  - 3) Scale-min and Scale-max can customize the total value range of the Color bar, and Scale interval sets the value range of each color block on the Color bar, and can change the fineness of the color to adjust the drawing effect.
  - 4) Show data can display the data value of each grid on the heat map, and can also adjust the display font size of the data value.
  - 5) Metric can choose 7 clustering methods.
  - 6) Method can choose 22 distance measures.
  - 7) Color bar sets its direction, label font size, label name, size and color.
  - 8) There are 166 color maps, which are mainly divided into the following four categories: Sequential colormaps: continuous color maps, Diverging colormaps: color maps that diverge at both ends, Qualitative colormaps: discrete color maps, Miscellaneous colormaps: other color maps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc., each suffix \_r color map is the original color bar flipped horizontally.
  - 9) Number of color bar scales Set the number of scales.
  - 10) Bar lable and Bar lable size can set the label and label font size of Color bar.
- 3. After completing all configurations, click the "Run" button to display or update the plotter, and the plotter will appear in the left canvas.
- 4. Users can right-click the "Save" export option to export heat maps, and DataColor provides 900-1000dpi images for users to use. There are four image formats: png, jpg, pdf, and svg, to meet different needs.

## 4 Functional omics

## 4.1 Anatomogram plotter

Displaying tissue information effectively in multicellular organisms can be a time-consuming and laborious process. With different colours representing the expression of tissues in anatomical maps, it is easy to spot differences between tissues or tissues and immediately provide the biological context for these observations and grasp the visualisation results more quickly.

#### Select

Menu+-> Functional omics -> -> Anatomogram plotter

### Input

Csv format or Excel format target series file

Organs	Value
root	3
flower	2
leaf	2
fruit	3

Fig: Input file.

### Output

Generate image in the left canvas

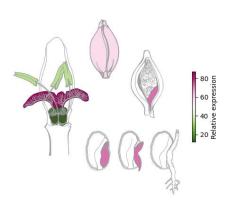


Fig: Anatomogram plotter output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. The default parameters have been set, if you need to adjust the parameters, please see the introduction below
  - 1) Type selects the desired base map for the deconvolution map
  - 2) The colour map contains 166 types, mainly divided into the following four categories: Sequential colormaps: continuous colour maps,

Diverging colormaps: colour maps with dispersion at both ends, Qualitative colormaps: discrete colour maps, Miscellaneous colormaps. Other colormaps. For example, 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r colour map is a horizontal flip of the original colour bar.

- 3) Color bar sets its orientation selection.
- 4) Title, table, Color bar can set the label name.
- 3. Once all configurations are complete, click the 'Run' button to display or update the Anatomogram plotter, which will appear in the canvas on the left.
- 4. The user can export the heat map by right clicking on the 'Save' export option, DataColor offers 900-1000 dpi images for the user to use. Four image formats are available: png, jpg, pdf and svg to suit different needs.

# 4.2 Bubble plotter

Bubble plotter is a multivariate chart and a variant of scatter plot. Bubble plot is a combination of scatter plot and percentage area plot. bubble plot can analyze the correlation between data by the position, area and color size of bubbles. GO and KEGG enrichment data are common scenarios for Bubble plot, where the color of the bubble indicates the p-value (or q-value, etc.) and the size indicates the number of genes.

### Select

Menu -> Functional omics -> Bubble plotter

### Input

Csv format or Excel format target series file

4	A	В	C
1	Description	PValue	Count
2	immune response	9.61E-08	20
3	defense response to bacterium	3.02E-06	11
4	cell chemotaxis	5.14E-06	8
5	cell adhesion	2.73E-05	17
6	complement activation	3.56E-05	8

Fig: Input file.

### Output

Generate image in the left canvas

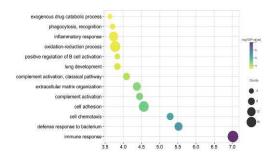


Fig: Bubble plotter output image

- 1. First of all, click the file button on the right, select the Excel format or Csv format file that needs to be visualized
- 2. Default parameters have been set, if you want to adjust the parameters, please see the following introduction

- 1) Set the X-axis and Y-axis on and off, font, font size, color, rotation angle and step length
- 2) Color bar sets its direction, label size, label name, size, color, position, font size.
- 3) Bar lable2 is used as the label of size bar, and Bar lable2 size parameter is added.' Distance between entries in the legend' parameter can set the size bar internal label distance
- 4) color map contains 166 kinds, mainly divided into the following four categories: Sequential colormaps: continuous color maps, Diverging colormaps: color maps with dispersion at both ends, Qualitative colormaps: discrete color maps, Miscellaneous colormaps: Other colormaps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r color map is a horizontal flip of the original color bar.
- 5) Dot shape sets 15 kinds of dot shapes.
- 6) Dot size magnification can adjust the size of the dot.
- 7) Transparency parameter can adjust the transparency of the dot color.
- 3. After finishing all the configurations, click "Run" button to display or update Bubble plot, Bubble plot will appear in the left canvas.
- 4. Users can export the heat map by right-clicking the "Save" export option, and DataColor provides 6 levels of images from 300-1000 dpi for users to use. There are four image formats: png, .jpg, pdf, and .svg, to meet different needs.

# 4.3 Dot plotter

Datacolor adds Dotplot function, which can use point, diamond, circle, star and other 23 different shapes to draw heat map, increasing the richness of heat map graphic display, except the color represents the data, where the size of the dot and the shade of the color are the same, while representing the size of the value. And you can achieve synchronous zooming in and out of points.

### **Select**

Menu -> Functional omics -> Dotplot

### Input

Csv format or Excel format target series file

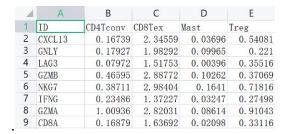


Fig: Input file.

## **Output**

Generate image in the left canvas

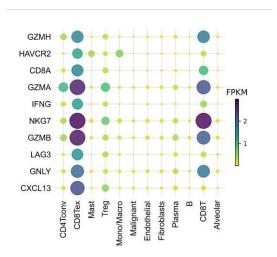


Fig: Dotplot output image

### **Detailed instructions for use**

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized
- 2. Default parameters have been set, if you need to adjust the parameters, please see below.

#### Detailed use method

- 1) Set the X-axis and Y-axis on and off, font, font size, color, rotation angle and step length.
- 2) Color bar sets its direction, label size, label name, size, color, position, font size.
- 3) The color map contains 166 types, mainly divided into the following four categories: Sequential colormaps: continuous color maps, Diverging colormaps: discrete color maps at both ends, Qualitative colormaps: discrete color maps, Miscellaneous colormaps: Other colormaps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r color map is a horizontal flip of the original color bar.
- 4) Dot shape sets 23 kinds of dot shapes.
- 5) Dot size magnification can adjust the size of the dot.
- 6) Transparency parameter can adjust the transparency of the dot color.
- 3. After finishing all the configurations, click the "Run" button to display or update the heat map, which will appear in the canvas on the left.
- 4. Users can export the heat map by right-clicking the "Save" export option, and DataColor provides 6 levels of images from 300-1000dpi for users to use. There are four image formats: png, .jpg, pdf, and .svg, to meet different needs.

# 4.4 Network plotter

Network plot function is a graphical model that is composed of two factors: connecting lines and nodes. different nodes that are linked, connected by 1 or more lines, and plotting the node size according to the number of correlations. dataColor adds six different ways of node layout, including random, graphical and algorithmic layout, to give users a better custom node experience. It is often used in differential co-expression network analysis to show the correlation of different genes.

#### Select

Menu -> Functional omics -> Network plot

### Input

Csv format or Excel format target series file

Date	Counts
2/24/2003	2871
5/7/2003	2765.9
7/1/2003	3884.34
8/25/2003	3746.7
10/10/2003	5205.27
10/28/2003	3479.76

Fig: Input file.

## **Output**

Generate image in the left canvas

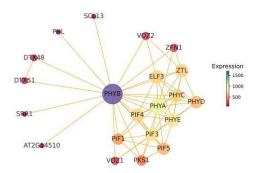


Fig:Network plot output image

- 1. First of all, click the file button on the right, select the Excel format or Csv format file that needs to be visualized
- 2. Have set the default parameters, if you need to adjust the parameters, please see the following introduction
  - 1) Set the font size and color of the node.

- 2) Color bar sets its direction, label font size, label name, size, color, position, font size.
- 3) Color map contains 166 kinds, mainly divided into the following four categories: Sequential colormaps: continuous color maps, Diverging colormaps: color maps with dispersion at both ends, Qualitative colormaps: discrete color maps, Miscellaneous colormaps: Other colormaps. Such as 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r color map is a horizontal flip of the original color bar.
- 4) Nodes size magnification can adjust the size of the points.
- 5) Transparency parameter can adjust the transparency of the point color.
- 6) Nodes arrangement parameter adds six different node layout methods, including random, graphical and algorithmic layout, etc.
- 3. After finishing all the configurations, click "Run" button to display or update the Network plot, which will appear in the canvas on the left.
- 4. Users can export the heat map by right-clicking the "Save" export option, and DataColor provides 6 levels of images from 300-1000 dpi for users to use. There are four image formats: png, .jpg, pdf, and .svg, to meet different needs.

# 4.5 Treemap

Treemap diagrams, also known as rectangular tree mapping, rectangular tree mapping and rectangular tree mapping, refer to a method of displaying tree data using nested rectangles. This method of presentation allows different categories to be presented in different coloured blocks and allows the size of the blocks to be compared to the size of each category. The larger the range of the blocks, the larger and more numerous the category.

### Select

Menu -> Functional omics -> Treemap

### Input

Csv format or Excel format target series file

Description	PValue
dendritic cell chemotaxis	0.0133971
receptor-mediated endocytosis	0.013738177
proteolysis	0.016467098
negative regulation of growth	0.01661592
positive regulation of fibroblast	0.018995878
negative regulation of protein ki	0.02032845
positive regulation of cardiac mu:	0.021990955
platelet degranulation	0.023126906
response to mechanical stimulus	0.023974023

Fig: Input file.

## Output

Generate image in the left canvas

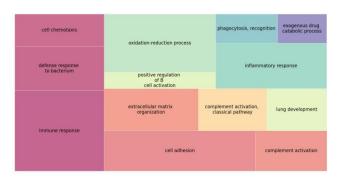


Fig: Treemap output image

- 1.First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. The default parameters have been set, if you need to adjust the parameters, please see the introduction below
  - 1) Show lable displays the data values of each square on the Treemap

- and allows you to adjust the display size of the lable.
- 2) The colour map contains 166 types and is divided into four main categories: Sequential colormaps: continuous colormaps, Diverging colormaps: discrete colormaps at both ends, Qualitative colormaps: discrete colormaps, Miscellaneous colormaps. Other colormaps. For example, 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix r colour map is a horizontal flip of the original colour bar.
- 3) Border widths are the dividing lines for each cell of the Treemap, if you need to adjust the colour, select a value greater than 0 for the width.
- 4) Transparency sets the transparency of the colour bar, ranging from 0 to 1.
- 5) Title sets the label and font size.
- 3. After completing all configurations, click the 'Run' button to display or update the Treemap, which will appear in the canvas on the left.
- 4. Users can right-click the "Save" export option to export heat maps, and DataColor provides 900-1000dpi images for users to use. There are four image formats: png, jpg, pdf, and svg, to meet different needs.

# 4.6 Wordcloud plotter

A Wordcloud, also known as a word cloud, is a visual representation of text data, consisting of a cloud-like coloured graph of words, used to display large amounts of text data. The importance of each word is displayed in font size or colour.

The Wordcloud is mainly used for frequency analysis of keyword occurrences in text content and is suitable for visualisation of text content mining. Words that appear more frequently in the word cloud are presented in a larger format and words that appear less frequently are presented in a smaller format. The word cloud is essentially a point graph, the result of drawing text with a specific style at the corresponding coordinate points.

#### Select

Menu -> Functional omics -> Wordcloud plotter

### Input

Csv format or Excel format target series file

Description	Count
immune response	20
defense response to bacterium	11
cell chemotaxis	8
cell adhesion	17
complement activation	8
extracellular matrix organization	11
complement activation, classical pathway	8
lung development	7
positive regulation of B cell activation	5
oxidation-reduction process	18

Fig:Input file.

### Output

Generate image in the left canvas



Fig: Wordcloud plotter output image

- 1. First of all, click the file button on the right, select the Csv format or Excel format file that needs to be visualized.
- 2. The default parameters have been set, if you need to adjust the parameters, please see the introduction below
  - 1) Width and Height set the width and height of the output image.
  - 2) Background colour parameter selects the colour of the background
  - 3) Word distance sets the distance between words.
  - 4) Maximum word count and maximum font size allows you to customise the total range of font size values.
  - 5) Colour map contains 166 types, mainly divided into the following four categories: Sequential colormaps: continuous colormaps, Diverging colormaps: discrete colormaps, Qualitative colormaps: discrete colormaps, Miscellaneous colormaps. Other colormaps. For example, 'Accent', 'Accent\_r', 'Blues', 'Blues\_r', 'BrBG', 'BrBG\_r', etc. Each suffix \_r colour map is a horizontal flip of the original colour bar.
  - 6) 'The color is the same as the background image color' asks if the output image font colour is the same as the background image.
- 3. Once all configurations are complete, click the 'Run' button to display or update the Wordcloud Map, which will appear in the canvas on the left.
- 4. The user can export the heat map by right clicking on the 'Save' export option, DataColor offers 900-1000 dpi images for the user to use. Four image formats are available: png, jpg, pdf and svg to suit different needs.