a plotly | Graphing Libraries (https://plotly.com/)(/graphing-libraries/)



Python (/python) > (/python/) > Manhattan Plot

¿utm_campaign=studio_cloud_launch&utm_content=sidebar)

• Suggest an edit to this page(https://github.com/plotly/plotly.py/edit/doc-prod/doc/python/bio-manhattanplot.md)

Manhattan Plot in Python

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Plotly Studio: Transform any dataset into an interactive data application in minutes with Al. Sign up for early access now. (https://plotly.com/studio/?utm_medium=graphing_libraries&utm_campaign=studio_early_access&utm_content=sidebar)

Manhattan Plot

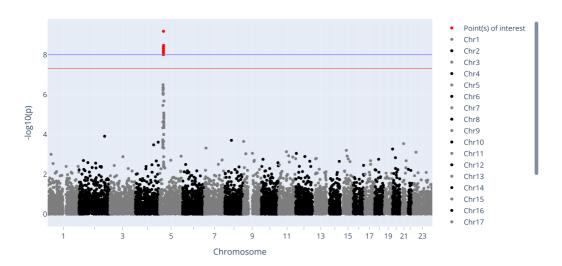
ManhattanPlot allows you to visualize genome-wide association studies (GWAS) efficiently. Using WebGL under the hood, you can interactively explore overviews of massive datasets comprising hundreds of thousands of points at once, or take a closer look at a small subset of your data. Hover data and click data are accessible from within the Dash app.

```
import pandas as pd
import dash_bio

df = pd.read_csv('https://raw.githubusercontent.com/plotly/dash-bio-docs-files/master/manhattan_data.csv')

dash_bio.ManhattanPlot(
    dataframe=df,
)
```

Manhattan Plot



Highlighted points color, and colors of the suggestive line and the genome-wide line

Change the color of the points that are considered significant.



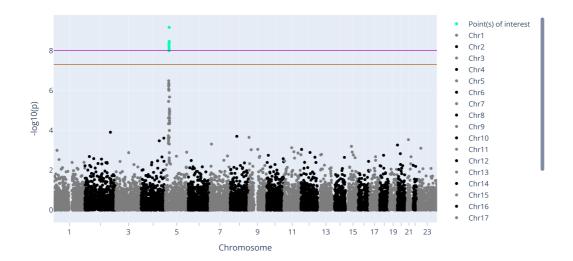
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```
import pandas as pd
import dash_bio

df = pd.read_csv('https://raw.githubusercontent.com/plotly/dash-bio-docs-files/master/manhattan_data.csv')

dash_bio.ManhattanPlot(
    dataframe=df,
    highlight_color='#00FFAA',
    suggestiveline_color='#AAAOAA',
    genomewideline_color='#AA5500'
)
```

Manhattan Plot





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ManhattanPlot with Dash

```
from dash import Dash, dcc, html, Input, Output
import dash_bio as dashbio
                                                                                                                                     DOWNLOAD
import pandas as pd
app = Dash(__name__)
df = pd.read_csv('https://git.io/manhattan_data.csv')
app.layout = html.Div([
    'Threshold value',
    dcc.Slider(
       id='slider',
       min=1,
       max=10,
       marks={
          i: {'label': str(i)} for i in range(10)
        value=6
    ),
    html.Br(),
    html.Div(
        dcc.Graph(
            id='graph',
            figure=dashbio.ManhattanPlot(
Threshold value
      Manhattan Plot
                                                                                                       Point(s) of interest
                                                                                                       Chr1
                                                                                                       Chr2
                                                                                                       Chr3
                                                                                                       Chr4
                                                                                                       Chr5
     -log10(p)
                                                                                                       Chr6
                                                                                                       Chr7
                                                                                                       Chr8
                                                                                                       Chr9
                                                                                                       Chr10
                                                                                                       Chr11
                                                                                                       Chr12
                                                                                                       Chr13
                                               Chromosome
```



What About Dash?

<u>Dash (https://dash.plot.ly/)</u> is an open-source framework for building analytical applications, with no Javascript required, and it is tightly integrated with the Plotly graphing library.

Learn about how to install Dash at https://dash.plot.ly/installation (https://dash.plot.ly/installation).

Everywhere in this page that you see fig.show(), you can display the same figure in a Dash application by passing it to the figure argument of the <u>Graph component</u> (https://dash.plot.ly/dash-core-components/graph) from the built-in dash_core_components package like this:

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```
import plotly.graph_objects as go # or plotly.express as px
fig = go.Figure() # or any Plotly Express function e.g. px.bar(...)
# fig.add_trace( ... )
# fig.update_layout( ... )

from dash import Dash, dcc, html

app = Dash()
app.layout = html.Div([
    dcc.Graph(figure=fig)
])

app.run(debug=True, use_reloader=False) # Turn off reloader if inside Jupyter
```





(https://dash.plotly.com/tutorial?utm_medium=graphing_libraries&utm_content=python_footer)

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