main

August 31, 2021

1 Heatmap of sharing

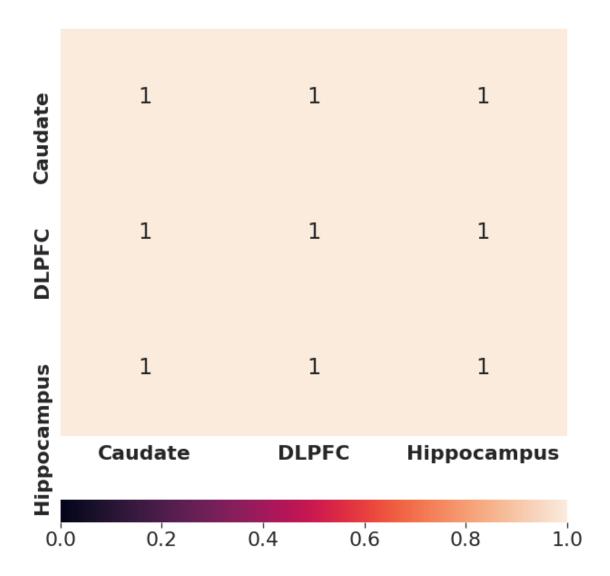
```
[1]: import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
from rpy2.robjects import r, pandas2ri
```

```
[2]: pandas2ri.activate()
```

1.1 Sign match only

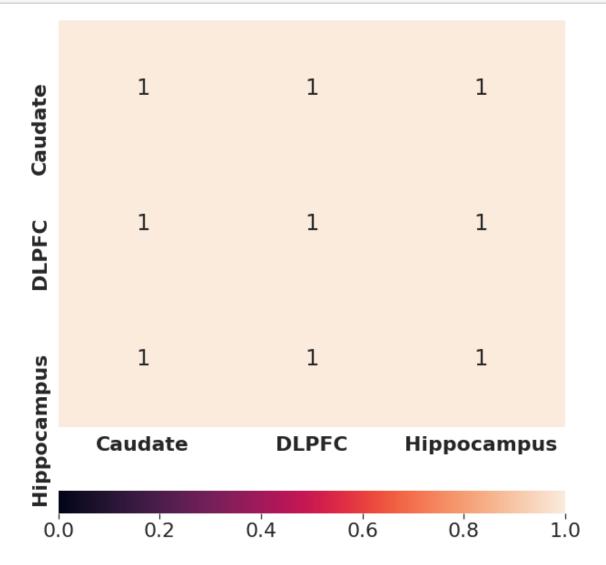
1.1.1 Genes

```
[3]: Caudate DLPFC Hippocampus
Caudate 1.0 1.0 1.0
DLPFC 1.0 1.0 1.0
Hippocampus 1.0 1.0 1.0
```



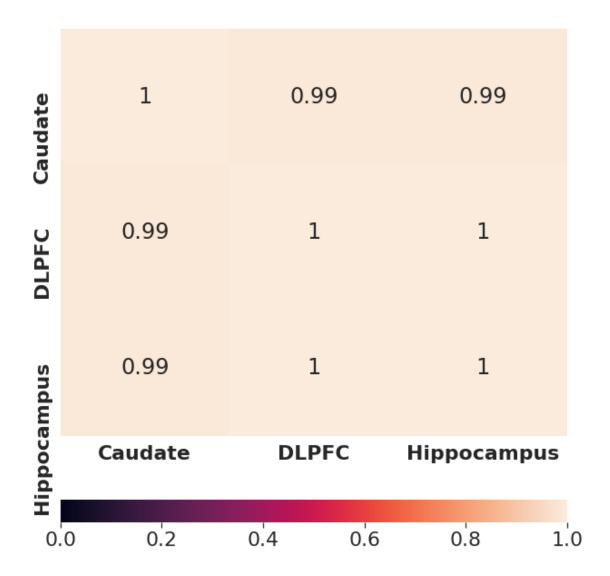
1.1.2 Transcripts

```
[5]:
                   Caudate
                                DLPFC
                                       Hippocampus
     Caudate
                  1.000000
                             0.998077
                                                1.0
     DLPFC
                  0.998077
                             1.000000
                                                1.0
     Hippocampus
                  1.000000
                             1.000000
                                                1.0
```



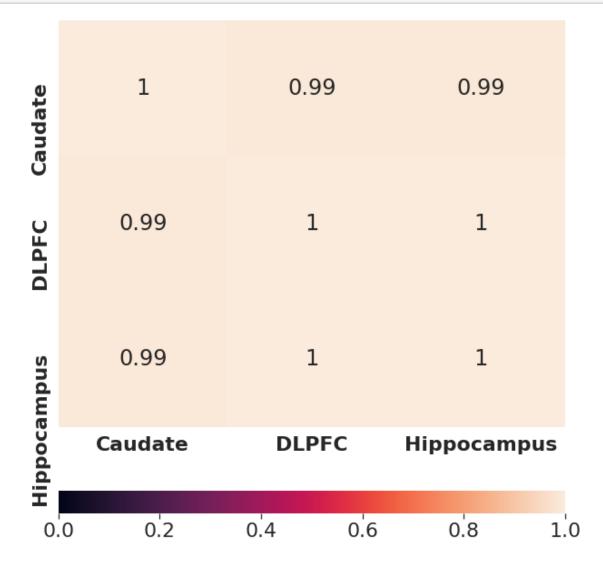
1.1.3 Exons

```
[7]: mat = r('''load("../../ m/exons/mashr meta results.RData"); mashr::
     df = pd.DataFrame(np.array(mat), index=["Caudate", "DLPFC", "Hippocampus"],
                      columns=["Caudate", "DLPFC", "Hippocampus"])
    df
[7]:
                  Caudate
                             DLPFC Hippocampus
    Caudate
                 1.000000 0.989175
                                       0.990476
    DLPFC
                 0.989175 1.000000
                                       1.000000
    Hippocampus 0.990476 1.000000
                                       1.000000
[8]: sns.set(font_scale=2)
    grid_kws = {"height_ratios": (.9, .05), "hspace": .3}
    f, (ax, cbar_ax) = plt.subplots(2, gridspec_kw=grid_kws, figsize=(10,10))
    chart = sns.heatmap(df, ax=ax, vmin=0, vmax=1,
                        annot=True, cbar_ax=cbar_ax,
                        cbar_kws={"orientation": "horizontal"})
    chart.set_yticklabels(chart.get_yticklabels(), fontweight="bold")
    chart.set_xticklabels(chart.get_xticklabels(), fontweight="bold")
    sns_plot = chart.get_figure()
    sns_plot.savefig("eQTL_sharing_heatmap_signOnly_exon.pdf")
    sns_plot.savefig("eQTL_sharing_heatmap_signOnly_exon.png")
```



1.1.4 Junctions

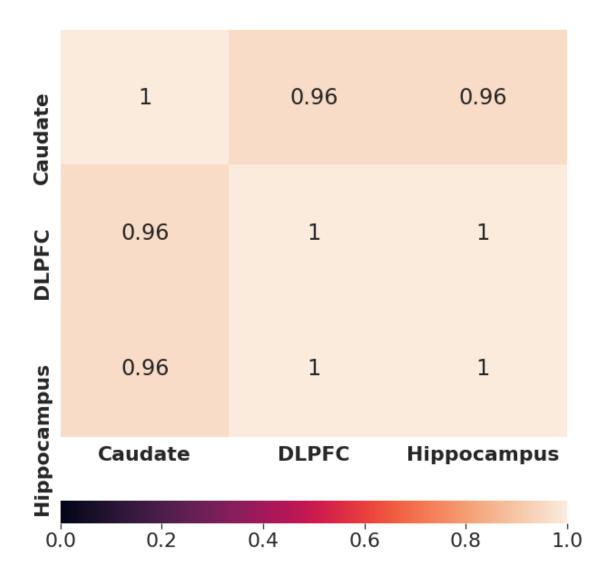
```
[9]: Caudate DLPFC Hippocampus
Caudate 1.000000 0.991266 0.991228
DLPFC 0.991266 1.000000 1.000000
Hippocampus 0.991228 1.000000 1.000000
```



1.2 Factor 0.5

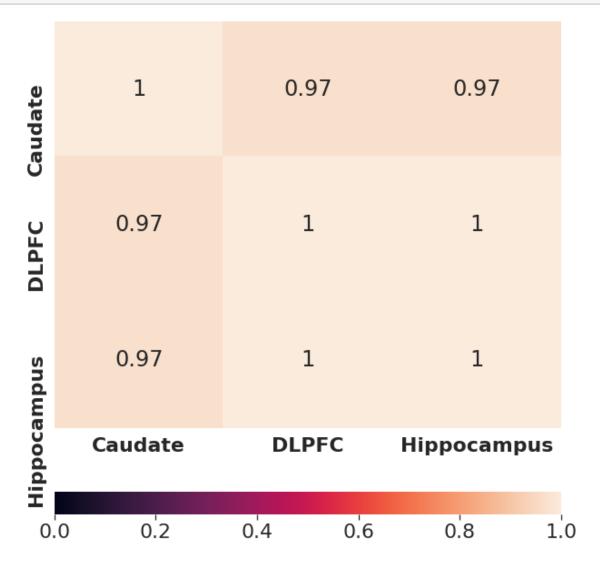
1.2.1 Genes

```
[11]: mat = r('''load("../../_m/genes/mashr_meta_results.RData"); mashr::
     df = pd.DataFrame(np.array(mat), index=["Caudate", "DLPFC", "Hippocampus"],
                       columns=["Caudate", "DLPFC", "Hippocampus"])
     df
[11]:
                   Caudate
                               DLPFC Hippocampus
     Caudate
                  1.000000 0.960784
                                        0.960784
     DLPFC
                  0.960784 1.000000
                                        1.000000
     Hippocampus 0.960784 1.000000
                                        1.000000
[12]: sns.set(font_scale=2)
     grid_kws = {"height_ratios": (.9, .05), "hspace": .3}
     f, (ax, cbar_ax) = plt.subplots(2, gridspec_kw=grid_kws, figsize=(10,10))
     chart = sns.heatmap(df, ax=ax, vmin=0, vmax=1,
                         annot=True, cbar_ax=cbar_ax,
                         cbar kws={"orientation": "horizontal"})
     chart.set_yticklabels(chart.get_yticklabels(), fontweight="bold")
     chart.set_xticklabels(chart.get_xticklabels(), fontweight="bold")
     sns plot = chart.get figure()
     sns_plot.savefig("eQTL_sharing_heatmap_gene.pdf")
     sns_plot.savefig("eQTL_sharing_heatmap_gene.png")
```



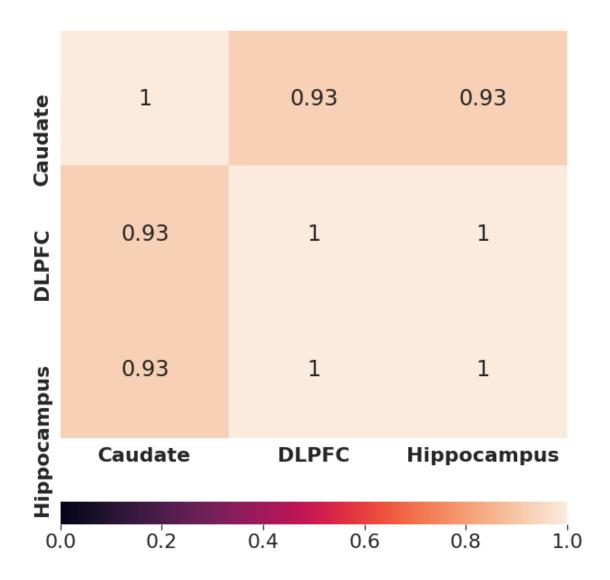
1.2.2 Transcripts

```
[13]: Caudate DLPFC Hippocampus
Caudate 1.000000 0.971154 0.969231
DLPFC 0.971154 1.000000 1.000000
Hippocampus 0.969231 1.000000 1.000000
```



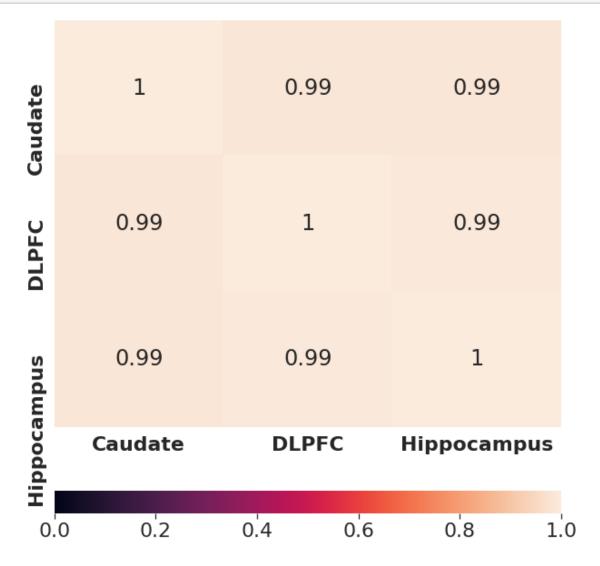
1.2.3 Exons

```
[15]: mat = r('''load("../../ m/exons/mashr meta results.RData"); mashr::
      df = pd.DataFrame(np.array(mat), index=["Caudate", "DLPFC", "Hippocampus"],
                       columns=["Caudate", "DLPFC", "Hippocampus"])
     df
[15]:
                   Caudate
                               DLPFC Hippocampus
     Caudate
                  1.000000 0.929635
                                        0.927891
     DLPFC
                  0.929635 1.000000
                                        0.998192
     Hippocampus 0.927891 0.998192
                                        1.000000
[16]: sns.set(font_scale=2)
     grid_kws = {"height_ratios": (.9, .05), "hspace": .3}
     f, (ax, cbar_ax) = plt.subplots(2, gridspec_kw=grid_kws, figsize=(10,10))
     chart = sns.heatmap(df, ax=ax, vmin=0, vmax=1,
                         annot=True, cbar_ax=cbar_ax,
                         cbar_kws={"orientation": "horizontal"})
     chart.set_yticklabels(chart.get_yticklabels(), fontweight="bold")
     chart.set_xticklabels(chart.get_xticklabels(), fontweight="bold")
     sns_plot = chart.get_figure()
     sns_plot.savefig("eQTL_sharing_heatmap_exon.pdf")
     sns_plot.savefig("eQTL_sharing_heatmap_exon.png")
```



1.2.4 Junctions

```
[17]: Caudate DLPFC Hippocampus
Caudate 1.000000 0.9869 0.986842
DLPFC 0.986900 1.0000 0.995000
Hippocampus 0.986842 0.9950 1.000000
```



[]: