Figure_5

February 14, 2024

```
[1]: import numpy as np
              import pandas as pd
              import seaborn as sns
              import matplotlib.pyplot as plt
              from matplotlib.lines import Line2D
              from matplotlib.patches import Patch
              from pycirclize import Circos, sector
              from pycirclize.parser import Matrix
              from itertools import permutations
[2]: table_5_A = pd.read_excel("Table_5.xlsx", sheet_name="Fig5 panel a")
              table_5_B = pd.read_excel("Table_5.xlsx", sheet_name="Fig 5 panel b")
              table 5 C = pd.read excel("Table 5.xlsx", sheet name="Fig5 panel c")
[3]: GCFs = []
              pGCFs= []
              for i in range(len(table_5_C)):
                          GCFs.append("GCFs")
                          pGCFs.append("pGCFs")
              table_5_C["GCFs"] = GCFs
              table_5_C["pGCFs"] = pGCFs
              table_5_C_1 = table_5_C[["REDgroup", "Number of actual GCFs", "GCFs"]].
                  orename(columns={"GCFs": "groups", "Number of actual GCFs": "Number of under of und
                  GCFs"})
              table_5_C_2 = table_5_C[["REDgroup", "Number of potential GCFs", "pGCFs"]].
                  orename(columns={"pGCFs" : "groups", "Number of potential GCFs" : "Number of
                 GCFs"})
              new_table_5_C = pd.concat((table_5_C_1, table_5_C_2))
```

```
[5]: | # my_liste = [ (('Actinobacteriota', 7603.0, 24012.0), ('Actinobacteriota', u
      →40421.0, 24012.0)),
           (('Streptomyces', 126.0, 6034.0), ('Streptomyces', 11942.0, 6034.0)),
           (('Streptomyces', 32.0, 126.0), ('SB', 311.0, 217.0)),
     #
           (('Actinobacteriota', 309.0, 7603.0), ('Streptomyces', 19236.0, 11942.0)),
           (('Actinobacteriota', 93.0, 309.0), ('SB', 527.0, 311.0)),
     #
           (('Actinobacteriota', 36.0, 93.0), ('SC', 156.0, 99.0)),
     #
           (('Actinobacteriota', 0.0, 36.0), ('SD', 92.0, 56.0)),
     #
           (('Streptomyces', 17.0, 32.0), ('SC', 99.0, 84.0)),
           (('Streptomyces', 0.0, 17.0), ('SD', 56.0, 39.0)),
     #
           (('SD', 0.0, 16.0), ('SD', 32.0, 16.0)),
     #
           (('SC', 0.0, 1.0), ('SD', 33.0, 32.0)),
     #
           (('SC', 1.0, 40.0), ('SC', 79.0, 40.0)),
     #
           (('SB', 0.0, 6.0), ('SD', 39.0, 33.0)),
     #
           (('SB', 6.0, 11.0), ('SC', 84.0, 79.0)),
           (('SB', 11.0, 114.0), ('SB', 217.0, 114.0))
     # ]
[6]: # bosliste = []
     # sorted list = ["Streptomyces", "SB", "SC", "SD", "Actinobacteriota"]
     # for permutation in permutations(sorted_list):
           matrix = Matrix.parse_fromto_table(table_5_B)
           ax = Circos.initialize_from_matrix(matrix,
     #
                                           order=permutation,
     #
                                           space=3,
     #
                                           r_lim=(99,100),
     #
                                           cmap="tab10",
     #
                                           link \ kws=dict(direction=0, r1=97, r2=97)
     #
           bosliste.append(ax)
[7]: # table 5 B = table 5 B.rename({"source taxon":"from", "target taxon":
      → "to", "Number of shared GCFs": "value"}, axis="columns")
[8]: | # my liste = [ (('Actinobacteriota', 7603.0, 24012.0), ('Actinobacteriota', u
      →40421.0, 24012.0)),
     #
           (('Streptomyces', 126.0, 6034.0), ('Streptomyces', 11942.0, 6034.0)),
           (('Streptomyces', 32.0, 126.0), ('SB', 311.0, 217.0)),
           (('Actinobacteriota', 309.0, 7603.0), ('Streptomyces', 19236.0, 11942.0)),
           (('Actinobacteriota', 93.0, 309.0), ('SB', 527.0, 311.0)),
           (('Actinobacteriota', 36.0, 93.0), ('SC', 156.0, 99.0)),
     #
           (('Actinobacteriota', 0.0, 36.0), ('SD', 92.0, 56.0)),
     #
           (('Streptomyces', 17.0, 32.0), ('SC', 99.0, 84.0)),
     #
           (('Streptomyces', 0.0, 17.0), ('SD', 56.0, 39.0)),
```

```
# (('SD', 0.0, 16.0), ('SD', 32.0, 16.0)),

# (('SC', 0.0, 1.0), ('SD', 33.0, 32.0)),

# (('SC', 1.0, 40.0), ('SC', 79.0, 40.0)),

# (('SB', 0.0, 6.0), ('SD', 39.0, 33.0)),

# (('SB', 6.0, 11.0), ('SC', 84.0, 79.0)),

# (('SB', 11.0, 114.0), ('SB', 217.0, 114.0))

# ]
```

```
[9]: # bosliste = []
     # sorted_list = ['SD', 'SC', 'SB', 'Streptomyces', 'Actinobacteriota']
     # for permutation in permutations(sorted_list):
           matrix = Matrix.parse_fromto_table(table_5_B)
     #
           mylst = np.random.shuffle(matrix.all_names)
     #
           matrix = matrix.sort(mylst)
     #
           ax = Circos.initialize from matrix(matrix,
     #
                                            start = -150, end = 210,
                                            space=3,
                                            r_lim=(99,100),
     #
     #
                                            cmap="tab10",
     #
                                            link kws=dict(direction=0, r1=97, r2=97)
     #
           bosliste.append(ax)
```

```
[10]: # for i, all_ax in enumerate(bosliste):
# all_ax.savefig(f"fig{i}.png",dpi=21)
```

```
[11]: # matrix_df = pd.DataFrame(matrix_data, index=row_names, columns=col_names)

# circos = Circos.initialize_from_matrix(
# second_Figure.transpose(),
# space=5,
# cmap="RdPu",
# label_kws=dict(size=12),
# )

# circos.plotfig();
```

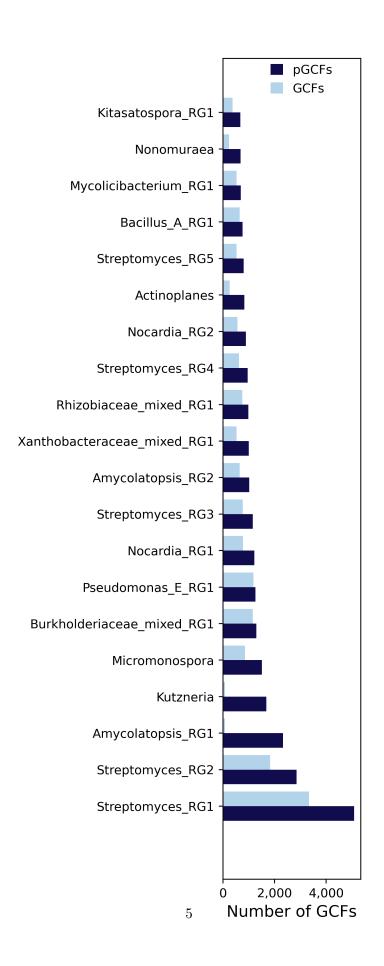
FIGURE 5 C i,ii

```
[12]: f, ax = plt.subplots(figsize=(2, 12), dpi=400)

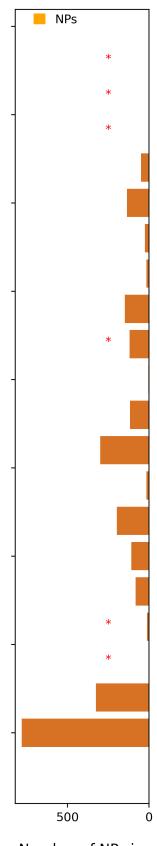
new_table_5_C['groups'] = pd.Categorical(new_table_5_C['groups'],__
categories=['GCFs', 'pGCFs'])
```

```
ax = sns.barplot(x="Number of GCFs",
                 y="REDgroup",
                 data = new_table_5_C,
                 hue="groups",
                 palette=["#070159","#a9d4f5"],
                 orient='h',
                 hue_order=["pGCFs","GCFs"])
ax.invert_yaxis()
ax.xaxis.set_ticks([0, 2000, 4000])
ax.set_ylim([-2, 20.48])
ax.set_ylabel(None)
ax.set_xlabel("Number of GCFs", fontsize=12.9)
ax.set_xticklabels(["0", "2,000", "4,000"])
sns.move_legend(ax, "lower center",bbox_to_anchor=(.57, 0.942), ncol=1,_

→title=None, frameon=False)
ax.set_xticks([0,2000,4000]);
f.savefig('output_figure_5_C_i_Topic_10.png', dpi=400)
```



```
[13]: | legend_elements = [Patch(facecolor='orange', edgecolor='orange', label='NPs')]
      f, ax = plt.subplots(figsize=(2, 12), dpi=400)
      ax = sns.barplot(x="Number of NPs in NPASS", y="REDgroup", data = table_5_C,__
       ⇔orient='h',color="#f56d05")
      ax.legend(handles=legend_elements, loc="lower center", bbox_to_anchor=(.3, 0.
       →966), ncol=1, title=None, frameon=False)
      plt.text(250,1.9, "*", ha='center', va='bottom', color="red")
      plt.text(250,2.9, "*", ha='center', va='bottom', color="red")
      plt.text(250,10.9, "*", ha='center', va='bottom', color="red")
      plt.text(250,16.9, "*", ha='center', va='bottom', color="red")
      plt.text(250,17.9, "*", ha='center', va='bottom', color="red")
      plt.text(250,18.9, "*", ha='center', va='bottom', color="red")
      ax.invert_xaxis()
      ax.invert_yaxis()
      ax.set(yticklabels=[])
      ax.set(ylabel=None)
      yticks = [i*20/8 \text{ for } i \text{ in } range(0,9)]
      ax.set_yticks(yticks)
      ax.set_xticks([0,500])
      ax.text(800, -3.9, "Number of NPs in\nNPASS database", fontsize=12, __
       →multialignment="left")
      ax.set_ylim([-2, 20.48])
      ax.set_xlabel(None)
      ax.set_ylabel(None)
      plt.show()
      f.savefig('output_figure_5_C_ii_Topic_10.png', dpi=400)
```



Number of NPs in NPASS database

$\textbf{0.0.1} \quad \textbf{END OF FIGURE 5} \ A, B, C_i, C_{ii}$