

```
In [71]: import pandas as pd

sim = pd.read_json("/Users/massoudmaher/data/tpois_ran_walk.json")
```

```
In [72]: def cast_n_sort(df_dict):
    df = pd.DataFrame(df_dict)
    df.index = df.index.astype("int")
    return df.sort_index()

sim["cn_data"] = [cast_n_sort(x) for x in sim["cn_data"]]
sim["plinkage"] = [cast_n_sort(x) for x in sim["plinkage"]]
sim["clustering"] = [cast_n_sort(x) for x in sim["clustering"]]
#
#display(sim.columns)
print(sim.shape)

(100, 17)
```

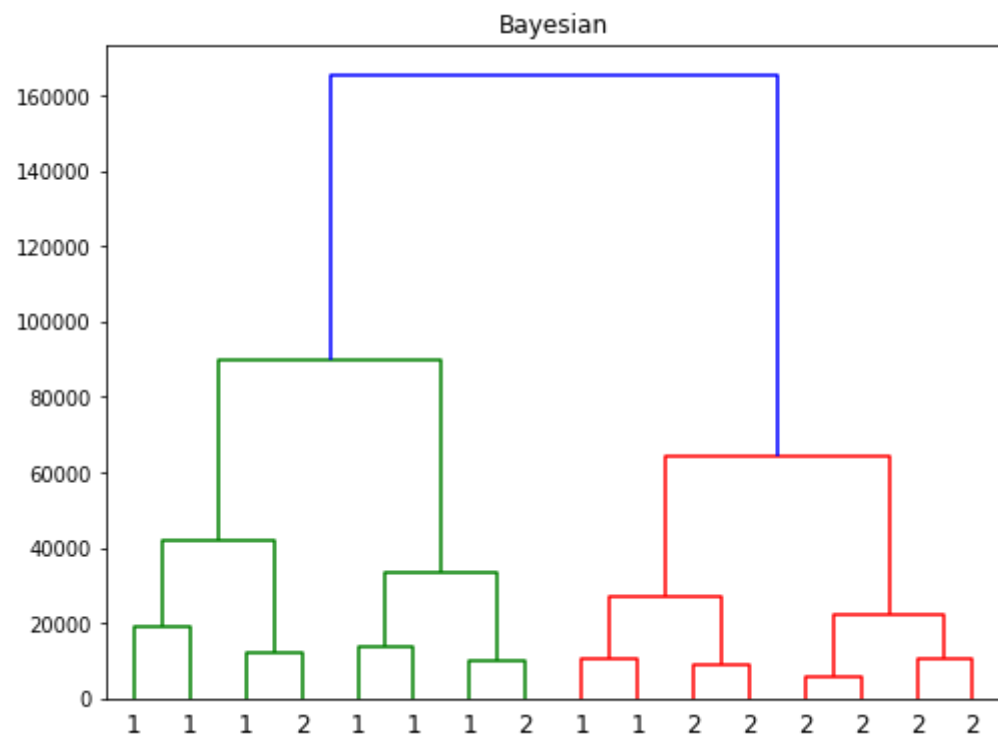
## Example clustering

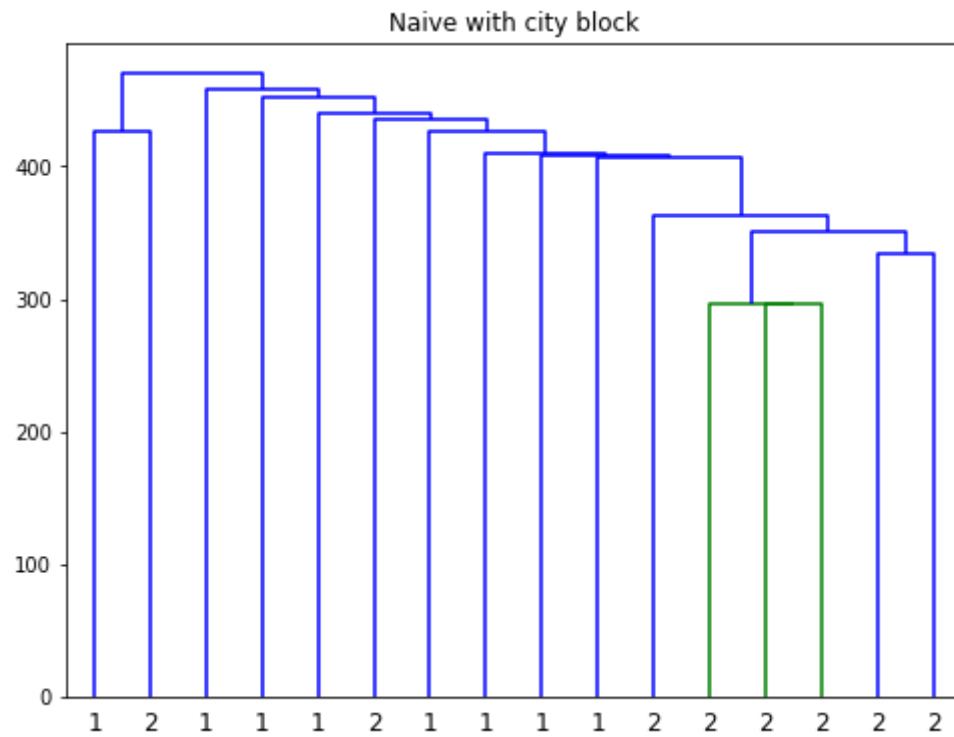
```
In [73]: # Bayesian hierarchical
from scipy.cluster.hierarchy import dendrogram
from scgenome.cncluster import bayesian_cluster
from scgenome.simulation import get_plot_data
import matplotlib.pyplot as plt
import numpy as np

eg_ind = 50
cn_data = sim.loc[eg_ind, "cn_data"].copy()
n_states = sim.loc[eg_ind, "max_cn"]
alpha = sim.loc[eg_ind, "alpha"]
bhc_linkage = sim.loc[eg_ind, "plot_data"]
bhc_plinkage = sim.loc[eg_ind, "plinkage"]
naive_linkage = sim.loc[eg_ind, "naive_linkage"]
cluster_ids = list(pd.Series(sim.loc[eg_ind, "cell_id"]).str[2])

fig = plt.figure(figsize=(8, 6))
f = dendrogram(bhc_linkage, labels=cluster_ids)
txt = fig.axes[0].set_title("Bayesian")

fig = plt.figure(figsize=(8, 6))
f = dendrogram(naive_linkage, labels=cluster_ids)
txt = fig.axes[0].set_title("Naive with city block")
```





```
In [74]: all_sim = pd.concat(list(sim["plinkage"]))
all_sim.head()
```

Out[74]:

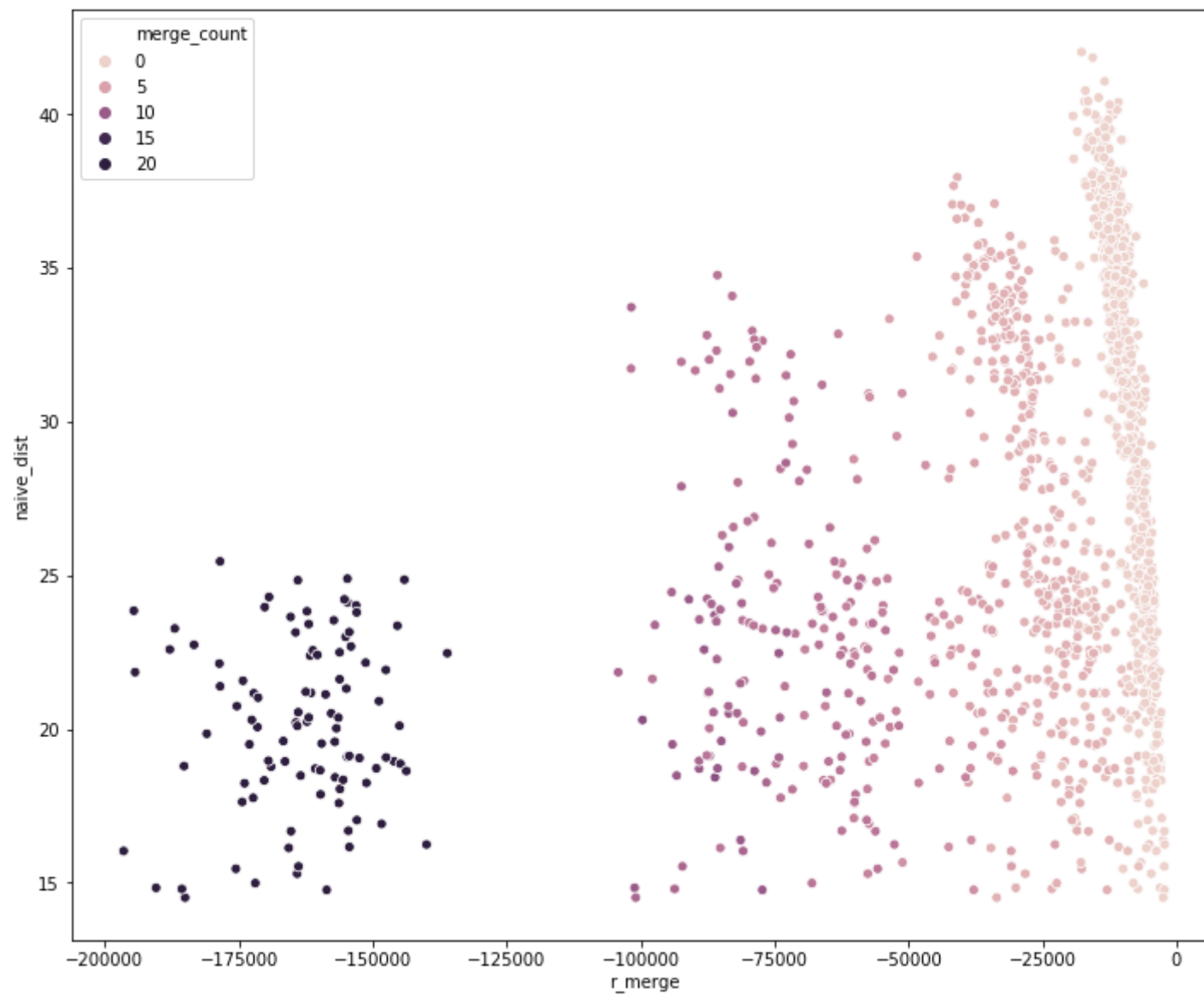
	i	j	r_merge	naive_dist	log_like	i_count	j_count	merge_count	dist
0	9	11	-5594.562331	28.193182	-5597.167501	1	1	2	5594.562331
1	8	12	-6778.561168	24.120467	-6781.166338	1	1	2	6778.561168
2	10	14	-7547.105025	27.789437	-7549.710195	1	1	2	7547.105025
3	7	13	-9479.377614	34.107227	-9481.982784	1	1	2	9479.377614
4	1	15	-9797.192799	35.982038	-9799.797970	1	1	2	9797.192799

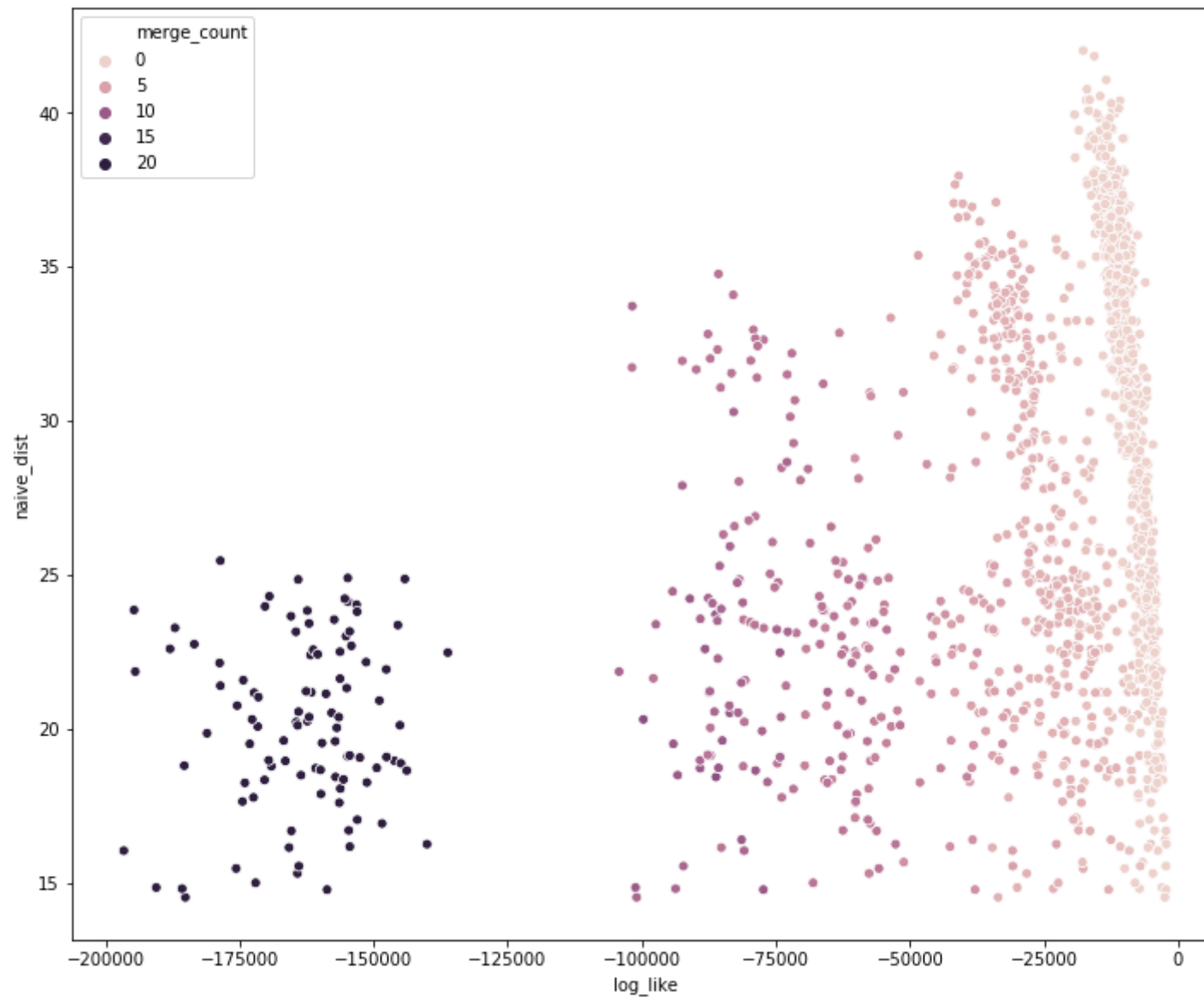
```
In [75]: import seaborn as sns
import matplotlib.pyplot as plt

fig = plt.figure(figsize=(12,10))
ax = sns.scatterplot(data=all_sim, x="r_merge", y="naive_dist", hue="merge_count")

fig = plt.figure(figsize=(12,10))
sns.scatterplot(data=all_sim, x="log_like", y="naive_dist", hue="merge_count")
```

Out[75]: <matplotlib.axes.\_subplots.AxesSubplot at 0x14dec7f10>





```
In [76]: from scipy.stats import pearsonr
print(pearsonr(all_sim["r_merge"], all_sim["naive_dist"])[0]**2 )
print(pearsonr(all_sim["log_like"], all_sim["naive_dist"])[0]**2)
```

```
0.13313831584341068
0.13316749727381114
```

For ease, going to only use 500 bin

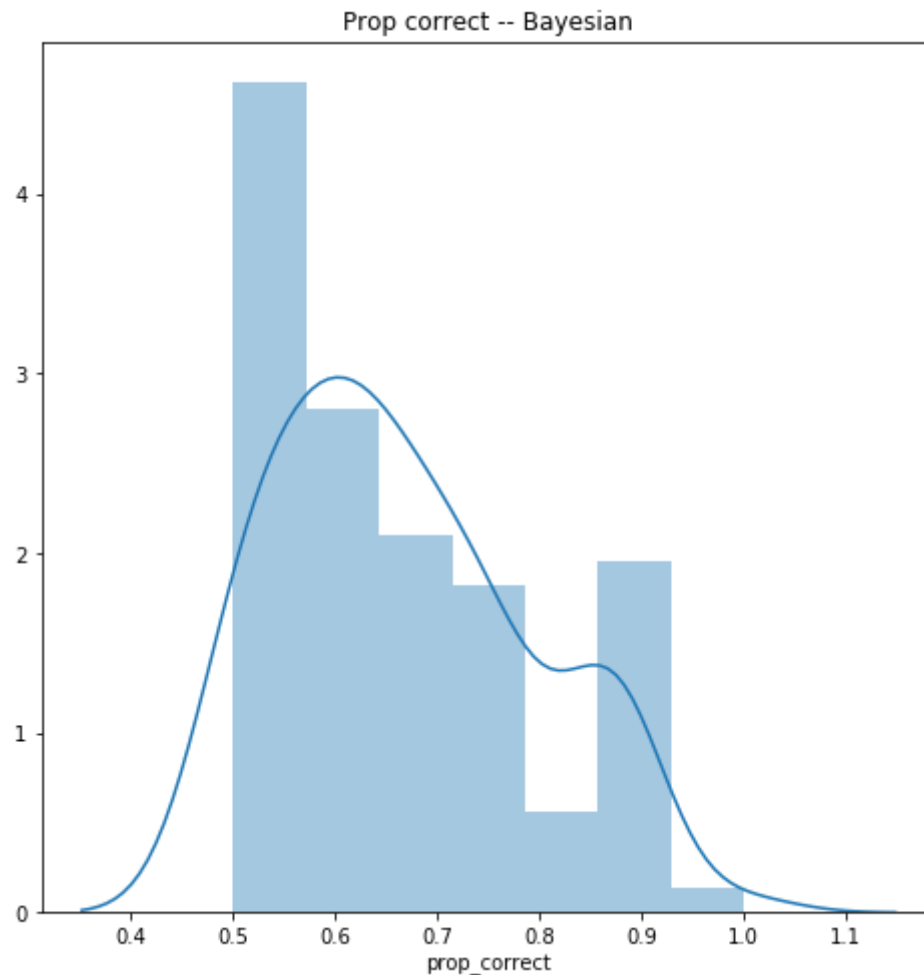


```
In [78]: import seaborn as sns
fig = plt.figure(figsize=(8,8))
sns.distplot(sim["prop_correct"])
plt.title("Prop correct -- Bayesian")
```

/Users/massoudmaher/Documents/Code/scgenome/venv/lib/python3.7/site-packages/scipy/stats/stats.py:1713: FutureWarning: Using a non-tuple sequence for multidimensional indexing is deprecated; use `arr[tuple(seq)]` instead of `arr[seq]`. In the future this will be interpreted as an array index, `arr[np.array(seq)]`, which will result either in an error or a different result.

```
return np.add.reduce(sorted[indexer] * weights, axis=axis) / sumval
```

```
Out[78]: Text(0.5,1,'Prop correct -- Bayesian')
```



In [79]:

```
Out[79]: 0      0.8750
          1      0.6250
          10     0.6250
          11     0.6250
          12     0.5000
          13     0.6250
          14     0.6250
          15     0.5000
          16     0.5625
          17     0.7500
          18     0.8750
          19     0.6250
          2      0.5625
          20     0.5625
          21     0.6250
          22     0.7500
          23     0.6250
          24     0.6875
          25     0.6875
          26     0.6250
          27     0.8750
          28     0.6250
          29     0.5000
          3      0.6875
          30     0.5625
          31     0.7500
          32     0.5000
          33     0.8750
          34     0.5625
          35     1.0000
          ...
          72     0.8750
          73     0.5000
          74     0.7500
          75     0.6250
          76     0.8750
          77     0.5000
          78     0.6250
          79     0.6875
          8      0.8750
          80     0.5625
          81     0.6250
          82     0.5625
```

```
83    0.5000
84    0.8125
85    0.8750
86    0.7500
87    0.5000
88    0.6875
89    0.7500
9    0.5000
90    0.7500
91    0.5625
92    0.6250
93    0.6250
94    0.6250
95    0.5000
96    0.8750
97    0.7500
98    0.5625
99    0.8750
Name: prop_correct, Length: 100, dtype: float64
```

In [ ]: