Chapter 8 - Comparison of taxa and period of building origin

November 10, 2020

```
[1]: import pandas as pd
      import geopandas as gpd
      import seaborn as sns
      import matplotlib.pyplot as plt
      import husl
      from legendgram import legendgram
      import mapclassify
      from matplotlib_scalebar.scalebar import ScaleBar
      from matplotlib.colors import ListedColormap
      from shapely.geometry import Point
      from tqdm import tqdm
 [2]: clusters = pd.read csv('/Users/martin/Dropbox/Academia/Data/Geo/Prague/
       →Clustering/complete data/200218_clusters_complete_n20.csv', index_col=0)
 [3]: years = pd.read_csv('/Users/martin/Dropbox/Academia/Data/Geo/Prague/Validation/
       →origin_year.csv', index_col=0)
 [4]: years['year'] = pd.Categorical(years['ROK_PUVODNI'],
                                     categories=[1840, 1880, 1920, 1950, 1970, 1990, L
       \rightarrow2012],
                                     ordered=True)
 [5]: | joined = clusters.merge(years[['uID', 'year']], on='uID', how='left')
 [7]: joined.head(4)
        uID cluster year
 [7]:
           0
                   1 1990
                   11 1840
      1
           1
      2
           2
                   13 1970
           3
                   18 1950
[32]: buildings = gpd.read_file('/Users/martin/Dropbox/Academia/Data/Geo/Prague/
       →Clustering/geometry.gpkg', layer='buildings')
```

```
[33]: buildings = buildings.merge(joined, on='uID', how='left')
[46]: buildings.year.unique()
[46]: [1990, 1840, 1970, 1950, NaN, 1920, 2012, 1880]
      Categories (7, int64): [1840 < 1880 < 1920 < 1950 < 1970 < 1990 < 2012]
 []:
 [7]: import numpy as np
      def show_values_on_bars(axs):
          def _show_on_single_plot(ax):
              for p in ax.patches:
                  _x = p.get_x() + p.get_width() / 2
                  _y = p.get_y() + p.get_height() + 0.02
                  value = '{:.2f}'.format(p.get_height())
                  ax.text(_x, _y, value, ha="center")
          if isinstance(axs, np.ndarray):
              for idx, ax in np.ndenumerate(axs):
                  _show_on_single_plot(ax)
          else:
              _show_on_single_plot(axs)
      pal = [husl.husl to hex(*color) for color in colors]
 []: # historical core
      data = joined.loc[joined['cluster'].isin([11])]['year'].
       →value_counts(sort=False, normalize=True)
      sns.set(context="paper", style="ticks", rc={'patch.force_edgecolor': False})
      fig, ax = plt.subplots(figsize=(10, 5))
      sns.barplot(ax=ax, x=data.index, y=data, order=data.index, palette=pal)
      sns.despine(offset=10)
      plt.ylabel('frequency')
      plt.xlabel('historical period')
      plt.ylim(0, 1)
      show_values_on_bars(ax)
 []: fig, ax = plt.subplots(2, 2, figsize=(10, 7))
      data = joined.loc[joined['cluster'].isin([11])]['year'].
       →value_counts(sort=False, normalize=True)
      sns.barplot(ax=ax[0, 0], x=data.index, y=data, order=data.index, palette=pal)
      sns.despine(offset=10)
      ax[0,0].set_ylabel('frequency')
```

```
ax[0,0].set_ylim(0, 1)
     show_values_on_bars(ax[0, 0])
     data = joined.loc[joined['cluster'].isin([5])]['year'].value_counts(sort=False,_
     →normalize=True)
     sns.barplot(ax=ax[0, 1], x=data.index, y=data, order=data.index, palette=pal)
     sns.despine(offset=10)
     ax[0,1].set_ylabel('frequency')
     ax[0,1].set_xlabel('historical period')
     ax[0,1].set_title('cluster 5')
     ax[0,1].set_ylim(0, 1)
     show_values_on_bars(ax[0, 1])
     data = joined.loc[joined['cluster'].isin([12])]['year'].
     →value_counts(sort=False, normalize=True)
     sns.barplot(ax=ax[1, 0], x=data.index, y=data, order=data.index, palette=pal)
     sns.despine(offset=10)
     ax[1,0].set_ylabel('frequency')
     ax[1,0].set_xlabel('historical period')
     ax[1,0].set_title('cluster 12')
     ax[1,0].set_ylim(0, 1)
     show_values_on_bars(ax[1, 0])
     data = joined.loc[joined['cluster'].isin([13])]['year'].
     →value counts(sort=False, normalize=True)
     sns.barplot(ax=ax[1, 1], x=data.index, y=data, order=data.index, palette=pal)
     sns.despine(offset=10)
     ax[1,1].set_ylabel('frequency')
     ax[1,1].set_xlabel('historical period')
     ax[1,1].set_title('cluster 13')
     ax[1,1].set_ylim(0, 1)
     show_values_on_bars(ax[1, 1])
     plt.tight_layout()
     plt.savefig('figures/PRG_cluster_origin_subplot.pdf')
[]: fig, ax = plt.subplots(1, 2, figsize=(10, 3.5))
     data = joined.loc[joined['cluster'].isin([11, 15, 5])]['year'].
     →value_counts(sort=False, normalize=True)
     sns.barplot(ax=ax[0], x=data.index, y=data, order=data.index, palette=pal)
     sns.despine(offset=10)
     ax[0].set_ylabel('frequency')
     ax[0].set_xlabel('historical period')
     ax[0].set_title('compact core')
```

ax[0,0].set_xlabel('historical period')

ax[0,0].set_title('cluster 11')

```
ax[0].set_ylim(0, 1)
      show_values_on_bars(ax[0])
      data = joined.loc[joined['cluster'].isin([1, 19])]['year'].
      →value_counts(sort=False, normalize=True)
      sns.barplot(ax=ax[1], x=data.index, y=data, order=data.index, palette=pal)
      sns.despine(offset=10)
      ax[1].set ylabel('frequency')
      ax[1].set_xlabel('historical period')
      ax[1].set_title('large scale industry')
      ax[1].set_ylim(0, 1)
      show_values_on_bars(ax[1])
      plt.tight_layout()
      plt.savefig('figures/PRG_branch_origin_subplot.pdf')
[43]: # save all clusters
      for cl in range(20):
          data = joined.loc[joined['cluster'].isin([cl])]['year'].
       →value_counts(sort=False, normalize=True)
          sns.set(context="paper", style="ticks", rc={'patch.force_edgecolor': False})
          fig, ax = plt.subplots(figsize=(10, 5))
          sns.barplot(ax=ax, x=data.index, y=data, order=data.index, palette=pal)
          sns.despine(offset=10)
          plt.ylabel('frequency')
          plt.xlabel('historical period')
          plt.ylim(0, 1)
          show_values_on_bars(ax)
```

plt.savefig('figures/PRG_cluster_' + str(cl) + '_origin.' + ext,__

for ext in ['pdf', 'png']:

→bbox_inches='tight')
plt.close()

```
plt.savefig('figures/PRG_compact_' + str(cl) + '_origin.' + ext,⊔

⇔bbox_inches='tight')
```

years and their clusters

```
[82]: colors = [(257, 71, 27), (98, 93, 78), (14, 79, 58), (26, 0, 50), (75, 90, 85), [
      \hookrightarrow (347, 72, 60), (246, 79, 60)]
      cols = []
      for col in colors:
          pal = sns.light_palette(col, input="husl", n_colors=5)
          for rgb in pal[1:]:
              cols.append(rgb)
      symbology = \{0: cols[10],
                    1: cols[14],
                    2: cols[20],
                    3: cols[9],
                    4: cols[22],
                    5: cols[1],
                    6: cols[18],
                    7: cols[23],
                    8: cols[11],
                    9: cols[21],
                    10: cols[0],
                    11: cols[3],
                    12: cols[7],
                    13: cols[22],
                    14: cols[6],
                    15: cols[2],
                    16: cols[19],
                    17: cols[23],
```

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18: cols[17],
19: cols[15]}
```

These have to be interpeted, but some taxa show remarkable link between year of construction and urban pattern. YAY!

```
[8]: import scipy.stats as ss
import numpy as np

def cramers_v(x, y):
    confusion_matrix = pd.crosstab(x,y)
    chi2 = ss.chi2_contingency(confusion_matrix)[0]
    n = confusion_matrix.sum().sum()
    phi2 = chi2/n
    r,k = confusion_matrix.shape
    phi2corr = max(0, phi2-((k-1)*(r-1))/(n-1))
    rcorr = r-((r-1)**2)/(n-1)
    kcorr = k-((k-1)**2)/(n-1)
    return np.sqrt(phi2corr/min((kcorr-1),(rcorr-1)))
```

```
[9]: cramers_v(joined.cluster, joined.year)
```

[9]: 0.35805157150989764

The resulting value of 0.5 indicates moderate relationship between clustering and historical origin.

Chi-square test of independence of variables in a contingency table

```
[10]: confusion_matrix = pd.crosstab(joined.cluster, joined.year)
chi, p, dof, exp = ss.chi2_contingency(confusion_matrix)
```

```
[11]: p
```

```
[11]: 0.0
[12]: dof
[12]: 114
[13]: chi
[13]: 106700.50861461915
[15]: joined.cluster.size
[15]: 140315
    p-value is < 0.001</pre>
```

there is a significant dependency between variables