## 04 Spatial autocorrelation

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Computational notebook 04 for Morphological tessellation as a way of partitioning space: Improving consistency in urban morphology at the plot scale.

Fleischmann, M., Feliciotti, A., Romice, O. and Porta, S. (2020) 'Morphological tessellation as a way of partitioning space: Improving consistency in urban morphology at the plot scale', Computers, Environment and Urban Systems, 80, p. 101441. doi: 10.1016/j.compenvurbsys.2019.101441.

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Note: notebook has been cleaned and released retroactively. It is likely that different versions of packages were initially used, but we made sure that the results remained unaltered.

## Data

The source of the data used within the research is the Amtliche Vermessung dataset accessible from the Zurich municipal GIS open data portal (https://maps.zh.ch). From it can be extracted the cadastral layer (Liegenschaften\_Liegenschaft\_Area) and the layer of buildings (all features named Gebäude). All data are licensed under CC-BY 4.0.

Source data: Vektor-Übersichtsplan des Kantons Zürich, 13.03.2018, Amt für Raumentwicklung Geoinformation / GIS-Produkte, Kanton Zürich, https://opendata.swiss/de/dataset/vektor-ubersichtsplan1

Data structure:

import esda
import libpysal

```
data/
single_uids.csv - IDs of buildings being alone on a single plot (QGIS generated)

cadastre/
blg_cadvals.shp - Cadastral values spatially joined to buildings to allow 1:1 comparise

tessellation/
{k}_tessellation.shp - tessellation layers

[3]: import matplotlib.pyplot as plt
import geopandas as gpd
```

```
import multiprocessing
     import pandas as pd
[4]: esda.__version__, gpd.__version__, libpysal.__version__, pd.__version__
[4]: ('2.2.1', '0.7.0', '4.2.2', '1.0.3')
[]: blg = gpd.read_file('data/cadastre/blg_cadvals.shp')
     characters = ['area', 'lal', 'circom', 'shapeix', 'rectan', 'fractal',
                   'orient', 'freq', 'car', 'gini_area', 'gini_car', 'Reach']
     singleuids = pd.read_csv('data/single_uids.csv')
     singles = singleuids['2'].to list()
     def worker(k):
         print(k)
         file = gpd.read_file('data/tessellation/{k}_tessellation.shp'.format(k=k))
         file_s = file.loc[file['uID'].isin(singles)].copy()
         file_m = file.loc[~file['uID'].isin(singles)].copy()
         for ch in characters:
             print(ch)
             try:
                 local_moran = esda.Moran_Local(file[[ch]], weights)
                 file['m_{{}}'.format(ch)] = local_moran.q
                 file['p_{}'.format(ch)] = local_moran.p_sim
                 single moran = esda.Moran Local(file s[[ch]], weights s)
                 file.loc[file['uID'].isin(singles), 'ms_{}'.format(ch)] =_u
      →single_moran.q
                 file.loc[file['uID'].isin(singles), 'ps_{}'.format(ch)] =__
     →single_moran.p_sim
                 multi_moran = esda.Moran_Local(file_m[[ch]], weights_m)
                 file.loc[~file['uID'].isin(singles), 'mm {}'.format(ch)] = [
      →multi_moran.q
                 file.loc[~file['uID'].isin(singles), 'pm_{{}}'.format(ch)] =__
      →multi_moran.p_sim
             # plot local autocorrelation(local moran, file, ch)
             # plt.savefig('files/moran/{k}_{ch}.png'.format(k=k, ch=ch))
             # plt.gcf().clear()
             except Exception:
                 print('missing, skipped')
         file.to_file('data/tessellation/{k}_tessellation.shp'.format(k=k))
         print('saved')
     weights = libpysal.weights.DistanceBand.from_dataframe(blg, 200)
     single = blg.loc[blg['uID_left'].isin(singles)].copy()
```

from splot.esda import plot\_local\_autocorrelation

```
weights_s = libpysal.weights.DistanceBand.from_dataframe(single, 200)
multi = blg.loc[~blg['uID_left'].isin(singles)].copy()
weights_m = libpysal.weights.DistanceBand.from_dataframe(multi, 200)
for ch in characters:
    print(ch)
    local_moran = esda.Moran_Local(blg[[ch]], weights)
    blg['m_{}'.format(ch)] = local_moran.q
    blg['p_{}'.format(ch)] = local_moran.p_sim
    single_moran = esda.Moran_Local(single[[ch]], weights_s)
    blg.loc[blg['uID_left'].isin(singles), 'ms_{{}}'.format(ch)] = single_moran.q
    blg.loc[blg['uID_left'].isin(singles), 'ps_{}'.format(ch)] = single_moran.
\rightarrow p_sim
    multi_moran = esda.Moran_Local(multi[[ch]], weights_m)
    blg.loc[~blg['uID_left'].isin(singles), 'mm_{}'.format(ch)] = multi_moran.q
    blg.loc[~blg['uID_left'].isin(singles), 'pm_{{}}'.format(ch)] = multi_moran.
\hookrightarrow p_sim
blg.to_file('data/cadastre/blg_cadvals.shp')
```

[]: