cov-matrix-processing

July 19, 2022

[1]: import numpy as np

```
import matplotlib.pyplot as plt
     from matplotlib.offsetbox import AnchoredText
     %matplotlib inline
[2]: n = 250
     cov_nercome = np.loadtxt(f"../output/data/
      →Patchy_V6C_BOSS_DR12_NGC_z1_cov_nercome_{n}_avg10000.matrix")
     cov_sample = np.loadtxt(f"../output/data/
      →Patchy_V6C_BOSS_DR12_NGC_z1_cov_sample_{n}.matrix")
     # 'Real' covariance matrix before processing.
     # The term 'real' here indicates that this is the matrix that we use to compare
     # our estimates to.
     cov_real_preproc = np.loadtxt("../data/
      G_2048_BOSS_DR12_NGC_z1_V6C_1_1_1_1_1_10_200_200_prerecon.matrix")
                                                Traceback (most recent call last)
     <ipython-input-2-37c9f68b0221> in <module>
            1 n = 250
      ---> 2 cov_nercome = np.loadtxt(f"../output/data/
       →Patchy_V6C_BOSS_DR12_NGC_z1_cov_nercome_{n}_avg10000.matrix")
            3 cov_sample = np.loadtxt(f"../output/data/
       →Patchy_V6C_BOSS_DR12_NGC_z1_cov_sample_{n}.matrix")
           5 # 'Real' covariance matrix before processing.
      ~/opt/anaconda3/lib/python3.8/site-packages/numpy/lib/npyio.py in loadtxt(fname
       dtype, comments, delimiter, converters, skiprows, usecols, unpack, ndmin,
       ⇔encoding, max_rows)
          979
                          fname = os fspath(fname)
          980
                     if _is_string_like(fname):
                          fh = np.lib._datasource.open(fname, 'rt', encoding=encoding)
      --> 981
                          fencoding = getattr(fh, 'encoding', 'latin1')
         982
          983
                          fh = iter(fh)
```

```
~/opt/anaconda3/lib/python3.8/site-packages/numpy/lib/_datasource.py in_u
       ⇔open(path, mode, destpath, encoding, newline)
          267
          268
                  ds = DataSource(destpath)
      --> 269
                  return ds.open(path, mode, encoding=encoding, newline=newline)
          270
         271
      ~/opt/anaconda3/lib/python3.8/site-packages/numpy/lib/_datasource.py in_
       →open(self, path, mode, encoding, newline)
          621
                                                    encoding=encoding, newline=newlin
          622
                      else:
      --> 623
                         raise IOError("%s not found." % path)
          624
          625
     OSError: ../output/data/Patchy_V6C_BOSS_DR12_NGC_z1_cov_nercome_250_avg10000.
       →matrix not found.
[]: print(cov_nercome)
     print(cov_sample)
[]: indices = np.concatenate((np.arange(40), np.arange(40)+80, np.arange(40)+160))
     cov_real = (cov_real_preproc[indices, :])[:, indices]
[]: print(cov_real)
[]: MSE_NERCOME = np.trace((cov_nercome-cov_real)@(cov_nercome-cov_real).T)
     MSE sample = np.trace((cov sample-cov real)@(cov sample-cov real).T)
     print(f"MSE NERCOME: {MSE_NERCOME}")
     print(f"MSE sample: {MSE_sample}")
[]: evals nercome = np.linalg.eigvalsh(cov nercome)
     evals_sample = np.linalg.eigvalsh(cov_sample)
     evals real = np.linalg.eigvalsh(cov real)
[]: plt.figure(dpi=200)
     plt.plot(evals_real, label="Beutler")
     plt.plot(evals_sample, label="Sample")
     plt.plot(evals_nercome, label="NERCOME")
     plt.yscale("log")
     plt.legend()
     plt.xlabel("Eigenmodes")
     plt.ylabel("Eigenvalues")
     parameters = AnchoredText(
         fr"$n = {n}$" "\n"
```

```
fr"MSE NERCOME $= {np.format_float_scientific(MSE_NERCOME, precision=3)}$"u
      \hookrightarrow"\n"
         fr"MSE Sample $={np.format_float_scientific(MSE_sample, precision=3)}$",
         frameon=False, loc="lower right", pad=0.5)
     plt.setp(parameters.patch, facecolor='white', alpha=0.5)
     plt.gca().add artist(parameters)
[ ]: nercome_rel_errors = (cov_nercome-cov_real)/np.abs(cov_real)
     print(nercome_rel_errors)
[]: nercome_pos_count = 0
     nercome_neg_count = 0
     for row in nercome_rel_errors:
         for error in row:
             if error >= 0:
                 nercome_pos_count += 1
             elif error < 0:</pre>
                 nercome_neg_count += 1
     print(f"NERCOME overestimated {nercome_pos_count} elements")
     print(f"NERCOME underestimated {nercome neg count} elements")
     index max nercome = np.unravel index(np.abs(nercome rel errors).argmax(),
      →nercome_rel_errors.shape)
     print(f"Maximum relative error is {nercome_rel_errors[index_max_nercome]}")
     print(f"Maximum relative error index is {index_max_nercome}")
     print(f"NERCOME: {cov_nercome[index_max_nercome]}, real:__
      →{cov_real[index_max_nercome]}")
[]: sample_rel_errors = (cov_sample-cov_real)/np.abs(cov_real)
     print(sample_rel_errors)
[]: sample_pos_count = 0
     sample_neg_count = 0
     for row in sample rel errors:
         for error in row:
             if error >= 0:
                 sample_pos_count += 1
             elif error < 0:</pre>
                 sample_neg_count += 1
     print(f"Sample overestimated {sample_pos_count} elements")
     print(f"Sample underestimated {sample_neg_count} elements")
     index max sample = np.unravel index(np.abs(sample rel_errors).argmax(),__
      ⇔sample_rel_errors.shape)
```

[]: