

cov-matrix-processing

July 1, 2022

```
[1]: import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline

[2]: cov_nercome = np.loadtxt("../output/Patchy_V6C_BOSS_DR12_NGC_z1_cov_nercome_500.
    ↪matrix")
cov_sample = np.loadtxt("../output/Patchy_V6C_BOSS_DR12_NGC_z1_cov_sample_500.
    ↪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/
    ↪C_2048_BOSS_DR12_NGC_z1_V6C_1_1_1_1_1_10_200_200_prerecon.matrix")

[3]: print(cov_nercome)
print(cov_sample)
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[[ 2.08632987e+08  2.31300315e+07 -9.53980998e+05 ... -2.57146234e+04
   3.22483402e+04  6.82058944e+04]
 [ 2.31300315e+07  4.92548896e+07  4.72947080e+06 ...  9.52509150e+02
   7.48281007e+03 -2.97310946e+04]
 [-9.53980998e+05  4.72947080e+06  1.70951338e+07 ...  1.55172277e+04
   1.31942194e+04  1.19134763e+04]
 ...
 [-2.57146234e+04  9.52509150e+02  1.55172277e+04 ...  6.48180100e+03
   1.14824933e+03  4.36267351e+02]
 [ 3.22483402e+04  7.48281007e+03  1.31942194e+04 ...  1.14824933e+03
   6.09538365e+03  1.19427703e+03]
 [ 6.82058944e+04 -2.97310946e+04  1.19134763e+04 ...  4.36267351e+02
   1.19427703e+03  5.90490079e+03]]
[[ 2.03256210e+08  2.26312165e+07 -8.10168108e+05 ... -2.56605953e+04
   3.12592410e+04  6.58264930e+04]
 [ 2.26312165e+07  4.70756413e+07  4.54771244e+06 ...  3.46957861e+02
   7.10159306e+03 -2.86043497e+04]
 [-8.10168108e+05  4.54771244e+06  1.61021470e+07 ...  1.47426460e+04
   1.28943477e+04  1.12660773e+04]
 ...
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[-2.56605953e+04  3.46957861e+02  1.47426460e+04 ...  3.93458356e+03
 9.11664424e+02  3.91148844e+02]
[ 3.12592410e+04  7.10159306e+03  1.28943477e+04 ...  9.11664424e+02
 3.65193234e+03  9.30709554e+02]
[ 6.58264930e+04 -2.86043497e+04  1.12660773e+04 ...  3.91148844e+02
 9.30709554e+02  3.51887142e+03]]

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[4]: indices = np.concatenate((np.arange(40), np.arange(40)+80, np.arange(40)+160))
cov_real = (cov_real_preproc[indices, :])[:, indices]

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[5]: print(cov_real)

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[[ 2.39531896e+08  1.91051563e+07  1.44038639e+05 ...  9.68083967e+03
 -1.21684291e+04  2.21567909e+04]
 [ 1.91051563e+07  5.15256710e+07  4.78376836e+06 ...  9.13608077e+03
 9.26311893e+03 -1.41639754e+04]
 [ 1.44038639e+05  4.78376836e+06  1.67772970e+07 ... -5.49127765e+02
 -2.28200815e+03  6.76773412e+03]
 ...
 [ 9.68083967e+03  9.13608077e+03 -5.49127765e+02 ...  4.22979902e+03
 1.28931823e+03  3.78440184e+02]
 [-1.21684291e+04  9.26311893e+03 -2.28200815e+03 ...  1.28931823e+03
 3.88691287e+03  1.00320951e+03]
 [ 2.21567909e+04 -1.41639754e+04  6.76773412e+03 ...  3.78440184e+02
 1.00320951e+03  3.62984716e+03]]

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[6]: evals_nercome = np.linalg.eigvalsh(cov_nercome)
evals_sample = np.linalg.eigvalsh(cov_sample)
evals_real = np.linalg.eigvalsh(cov_real)

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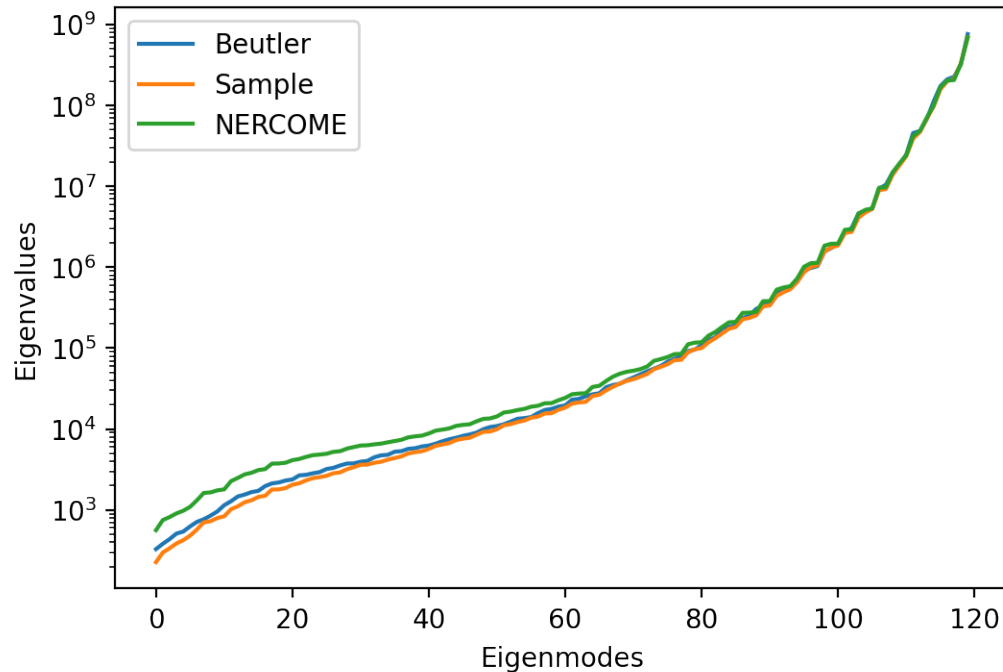
[7]: plt.figure(dpi=200)
plt.plot(evals_real, label="Beutler")
plt.plot(evals_sample, label="Sample")
plt.plot(evals_nercome, label="NERCOME")
plt.yscale("log")
plt.legend()
plt.xlabel("Eigenmodes")
plt.ylabel("Eigenvalues")

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[7]: Text(0, 0.5, 'Eigenvalues')

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[8]: nercome_rel_errors = (cov_nercome-cov_real)/np.abs(cov_real)
print(nercome_rel_errors)
```

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[[-1.28997057e-01  2.10669579e-01 -7.62309091e+00 ... -3.65623895e+00
  3.65016461e+00  2.07832910e+00]
 [ 2.10669579e-01 -4.40708743e-02 -1.13503743e-02 ... -8.95742039e-01
 -1.92193243e-01 -1.09906427e+00]
 [-7.62309091e+00 -1.13503743e-02  1.89444609e-02 ...  2.92579550e+01
  6.78184590e+00  7.60334559e-01]
 ...
 [-3.65623895e+00 -8.95742039e-01  2.92579550e+01 ...  5.32413470e-01
 -1.09413559e-01  1.52803982e-01]
 [ 3.65016461e+00 -1.92193243e-01  6.78184590e+00 ... -1.09413559e-01
  5.68181193e-01  1.90456246e-01]
 [ 2.07832910e+00 -1.09906427e+00  7.60334559e-01 ...  1.52803982e-01
  1.90456246e-01  6.26762926e-01]]
```

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[9]: 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:
            nercome_neg_count += 1
```

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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]}")

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NERCOME overestimated 8066 elements
NERCOME underestimated 6334 elements
Maximum relative error is 23462.389040678867
Maximum relative error index is (80, 94)
NERCOME: 312399.69904777897, real: -13.315481811674502

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[10]: sample_rel_errors = (cov_sample-cov_real)/np.abs(cov_real)
print(sample_rel_errors)

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[[-0.15144407  0.18456066 -6.62465819 ... -3.65065802  3.56888056
  1.97093985]
 [ 0.18456066 -0.08636529 -0.04934518 ... -0.96202334 -0.23334752
 -1.01951422]
 [-6.62465819 -0.04934518 -0.04024188 ... 27.84738773  6.65043895
  0.66467492]
 ...
 [-3.65065802 -0.96202334 27.84738773 ... -0.06979421 -0.29290969
  0.03358169]
 [ 3.56888056 -0.23334752  6.65043895 ... -0.29290969 -0.06045428
 -0.07226802]
 [ 1.97093985 -1.01951422  0.66467492 ...  0.03358169 -0.07226802
 -0.03057312]]

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[11]: 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:
            sample_neg_count += 1

print(f"Sample overestimated {sample_pos_count} elements")
print(f"Sample underestimated {sample_neg_count} elements")

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index_max_sample = np.unravel_index(np.abs(sample_rel_errors).argmax(),
    ↪sample_rel_errors.shape)

print(f"Maximum relative error is {sample_rel_errors[index_max_sample]}")
print(f"Maximum relative error index is {index_max_sample}")
print(f"Sample: {cov_sample[index_max_sample]}, real:
    ↪{cov_real[index_max_sample]}")

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Sample overestimated 7434 elements
Sample underestimated 6966 elements
Maximum relative error is 24707.78037823806
Maximum relative error index is (80, 94)
Sample: 328982.6847514654, real: -13.315481811674502

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[12]: 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}")

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MSE NERCOME: 1.6255763973991568e+16
MSE sample: 1.664069018003966e+16

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[ ]:

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