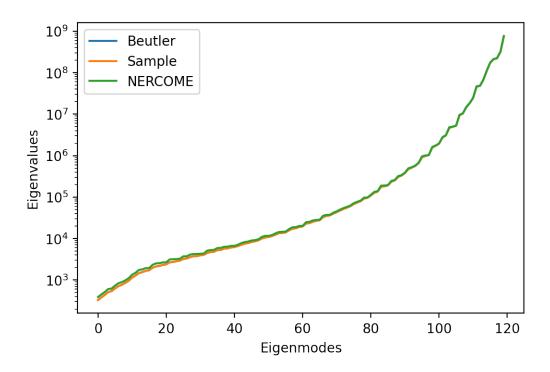
cov-matrix-processing

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[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_2048.matrix")
    cov_sample = np.loadtxt("../output/Patchy_V6C_BOSS_DR12_NGC_z1_cov_sample_2048.
     ⇔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)
   -1.24516461e+04 2.24203892e+04]
    [ 1.86024630e+07 5.16992164e+07 4.74738440e+06 ... 9.42837843e+03
      9.52712278e+03 -1.41014722e+04]
    [ 9.93829745e+04 4.74738440e+06 1.69697364e+07 ... -4.73772115e+02
     -2.34027554e+03 6.80012721e+03]
    1.36829796e+03 4.09849815e+02]
    [-1.24516461e+04 9.52712278e+03 -2.34027554e+03 ... 1.36829796e+03
      4.35627101e+03 1.08072377e+03]
    [ 2.24203892e+04 -1.41014722e+04 6.80012721e+03 ... 4.09849815e+02
      1.08072377e+03 4.07136743e+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]
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[ 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]]
[4]: indices = np.concatenate((np.arange(40), np.arange(40)+80, np.arange(40)+160))
    cov real = (cov real preproc[indices, :])[:, indices]
[5]: print(cov_real)
    -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]
    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]]
[6]: evals_nercome = np.linalg.eigvalsh(cov_nercome)
    evals_sample = np.linalg.eigvalsh(cov_sample)
    evals_real = np.linalg.eigvalsh(cov_real)
[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")
[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)
    [[-0.00922118 -0.02631192 -0.31002559 ... -0.04869081 -0.02327474
       0.01189695]
     0.00441283]
     [-0.31002559 -0.00760571 0.01147023 ... 0.1372279 -0.02553338
       0.0047864 ]
     [-0.04869081 \quad 0.03199377 \quad 0.1372279 \quad ... \quad 0.1122456
                                                         0.06125698
       0.08299761]
     [-0.02327474 0.02850054 -0.02553338 ... 0.06125698
       0.07726627]
      \hbox{ [ 0.01189695 \ 0.00441283 \ 0.0047864 \ \dots \ 0.08299761 \ 0.07726627 } 
       0.12163605]]
[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:</pre>
                 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]}")
     NERCOME overestimated 8846 elements
     NERCOME underestimated 5554 elements
     Maximum relative error is -962.7092816081185
     Maximum relative error index is (54, 25)
     NERCOME: -35.440281134271565, real: -0.03677486749440995
[10]: sample_rel_errors = (cov_sample-cov_real)/np.abs(cov_real)
     print(sample rel errors)
     0.00000000e+00 -3.28384992e-16]
      [ 3.89977475e-16 -1.44599390e-16 1.94683878e-16 ... 1.99099532e-16
       5.89107001e-16 -2.56847298e-16]
      [7.47606155e-15 1.94683878e-16 2.22043533e-16 ... 4.14063338e-16
      -2.39130093e-15 -1.34386884e-16]
      [-5.63687490e-16 -1.99099532e-16 8.28126676e-16 ... -2.15020784e-16
       1.76351866e-16 -4.50613502e-16]
     -1.16994480e-16 1.13323126e-16]
      [-1.64192496e-16 2.56847298e-16 -1.34386884e-16 ... -1.50204501e-16
       1.13323126e-16 0.00000000e+00]]
[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:</pre>
                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(),_u
       ⇒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]}")
     Sample overestimated 8087 elements
     Sample underestimated 6313 elements
     Maximum relative error is 7.58384684111064e-12
     Maximum relative error index is (25, 54)
     Sample: -0.03677486749413106, real: -0.03677486749440995
[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}")
     MSE NERCOME: 52247720917254.195
     MSE sample: 2.2537432416564147e-14
 []:
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