# FittingMethods

May 1, 2023

## 1 Fitting Methods

Here we will explore the various fitting methods in AutoProf. You have already encountered some of the methods, but here we will take a more systematic approach and discuss their strengths/weaknesses. Each method will be applied to the same problem with the same initial conditions so you can see how they operate.

```
[1]: import torch
import numpy as np
import matplotlib.pyplot as plt
from matplotlib.patches import Ellipse
from scipy.stats import gaussian_kde as kde
from scipy.stats import norm

%matplotlib inline
import autoprof as ap
```

```
[2]: # Setup a fitting problem. You can ignore this cell to start, it just makes
      \rightarrowsome test data to fit
     def true_params():
         # just some random parameters to use for fitting. Feel free to play around
      →with these to see what happens!
         sky_param = np.array([1.5])
         sersic_params = np.array([
             [ 68.44035491, 65.58516735, 0.54945988, 127.19794926*np.pi/180,
                                                                                     2.
      \rightarrow 14513004,
                   22.05219055, 2.45583024],
             [ 54.00353786, 41.54430634,
                                             0.40203928, 82.03862521*np.pi/180,
                                                                                     2.
      <del>-</del>88613347,
                  12.095631,
                                   2.76711163],
             [ 43.13601431, 58.3422508, 0.71894728, 167.07973506*np.pi/180,
      964371,
                5.3767236.
                                   2.41520244],
         ])
         return sersic_params, sky_param
     def init_params():
```

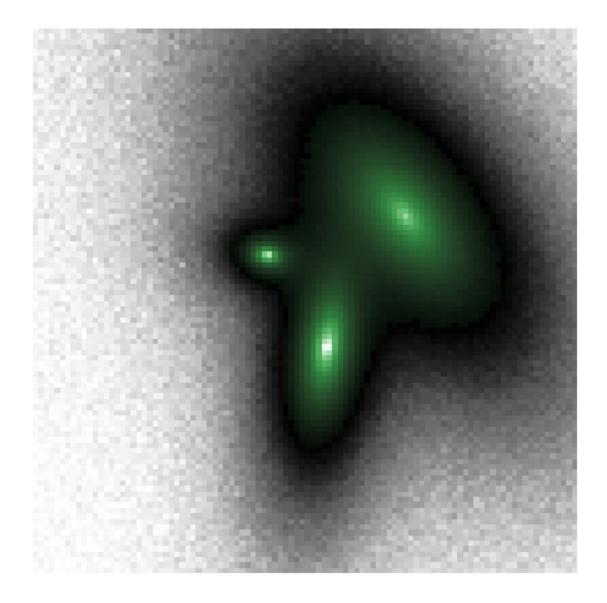
```
sky_param = np.array([1.4])
    sersic_params = np.array([
        [ 67., 66., 0.6, 130.*np.pi/180, 1.5, 25., 2.],
        [55., 40., 0.5, 80.*np.pi/180, 2., 10., 3.],
        [ 41., 60., 0.8, 170.*np.pi/180, 3., 4.,
                                                            2.],
   ])
   return sersic_params, sky_param
def initialize_model(target, use_true_params = True):
    # Pick parameters to start the model with
    if use_true_params:
       sersic_params, sky_param = true_params()
   else:
       sersic_params, sky_param = init_params()
    # List of models, starting with the sky
   model_list = [ap.models.AutoProf_Model(
       name = "sky",
       model_type = "flat sky model",
       target = target,
       parameters = {"sky": sky_param[0]},
   )]
    # Add models to the list
   for i, params in enumerate(sersic_params):
       model_list.append([
           ap.models.AutoProf_Model(
               name = f"sersic {i}",
               model_type = "sersic galaxy model",
               target = target,
               parameters = {
                   "center": [params[0],params[1]],
                   "q": params[2],
                   "PA": params[3],
                   "n": params[4],
                   "Re": params[5],
                   "Ie": params[6],
               #psf_mode = "full", # uncomment to try everything with PSF_
→blurring (takes longer)
           )
       ])
   MODEL = ap.models.Group_Model(
       name = "group",
```

```
model_list = model_list,
       target = target,
    # Make sure every model is ready to go
   MODEL.initialize()
   return MODEL
def generate_target():
   N = 100
   pixelscale = 1.
   rng = np.random.default_rng(42)
   # PSF has sigma of 2x pixelscale
   PSF = ap.utils.initialize.gaussian_psf(2, 21, pixelscale)
   PSF /= np.sum(PSF)
   target = ap.image.Target_Image(
       data = np.zeros((N,N)),
       pixelscale = pixelscale,
       psf = PSF,
   )
   MODEL = initialize_model(target, True)
    # Sample the model with the true values to make a mock image
   img = MODEL().data.detach().cpu().numpy()
    # Add poisson noise
   target.data = torch.Tensor(img + rng.normal(scale = np.sqrt(img)/2))
   target.variance = torch.Tensor(img/4)
   fig, ax = plt.subplots(figsize = (8,8))
   ap.plots.target_image(fig, ax, target)
   ax.axis("off")
   plt.show()
   return target
def corner_plot(chain, labels=None, bins=None, true_values=None,
→plot_density=True, plot_contours=True, figsize=(10, 10)):
   ndim = chain.shape[1]
   fig, axes = plt.subplots(ndim, ndim, figsize=figsize)
   plt.subplots_adjust(wspace=0., hspace=0.)
   if bins is None:
```

```
bins = int(np.sqrt(chain.shape[0]))
   for i in range(ndim):
       for j in range(ndim):
           ax = axes[i, j]
           i_range = (np.min(chain[:, i]), np.max(chain[:, i]))
           j_range = (np.min(chain[:, j]), np.max(chain[:, j]))
           if i == j:
               # Plot the histogram of parameter i
               #ax.hist(chain[:, i], bins=bins, histtype="step", range =_
\rightarrow i_range, density=True, color="k", lw=1)
               if plot_density:
                   # Plot the kernel density estimate
                   kde_x = np.linspace(i_range[0], i_range[1], 100)
                   kde y = kde(chain[:, i])(kde x)
                   ax.plot(kde_x, kde_y, color="green", lw=1)
               if true_values is not None:
                   ax.axvline(true values[i], color='red', linestyle='-', lw=1)
               ax.set_xlim(i_range)
           elif i > j:
               # Plot the 2D histogram of parameters i and j
               #ax.hist2d(chain[:, j], chain[:, i], bins=bins, cmap="Greys")
               if plot_contours:
                   # Plot the kernel density estimate contours
                   kde_ij = kde([chain[:, j], chain[:, i]])
                   x, y = np.mgrid[j_range[0]:j_range[1]:100j, i_range[0]:
→i_range[1]:100j]
                   positions = np.vstack([x.ravel(), y.ravel()])
                   kde_pos = np.reshape(kde_ij(positions).T, x.shape)
                   ax.contour(x, y, kde_pos, colors="green", linewidths=1,__
→levels=3)
               if true values is not None:
                   ax.axvline(true_values[j], color='red', linestyle='-', lw=1)
                   ax.axhline(true_values[i], color='red', linestyle='-', lw=1)
               ax.set_xlim(j_range)
               ax.set_ylim(i_range)
           else:
               ax.axis("off")
           if j == 0 and labels is not None:
```

```
ax.set_ylabel(labels[i])
            ax.yaxis.set_major_locator(plt.NullLocator())
            if i == ndim - 1 and labels is not None:
                ax.set_xlabel(labels[j])
            ax.xaxis.set_major_locator(plt.NullLocator())
    plt.show()
def corner_plot_covariance(cov_matrix, mean, labels=None, figsize=(10, 10),_
→true_values = None, ellipse_colors='g'):
    num_params = cov_matrix.shape[0]
    fig, axes = plt.subplots(num_params, num_params, figsize=figsize)
    plt.subplots_adjust(wspace=0., hspace=0.)
    for i in range(num_params):
        for j in range(num_params):
            ax = axes[i, j]
            if i == j:
                x = np.linspace(mean[i] - 3 * np.sqrt(cov_matrix[i, i]),
→mean[i] + 3 * np.sqrt(cov_matrix[i, i]), 100)
                y = norm.pdf(x, mean[i], np.sqrt(cov_matrix[i, i]))
                ax.plot(x, y, color='g')
                ax.set_xlim(mean[i] - 3 * np.sqrt(cov_matrix[i, i]), mean[i] +
 →3 * np.sqrt(cov_matrix[i, i]))
                if true values is not None:
                    ax.axvline(true_values[i], color='red', linestyle='-', lw=1)
            elif j < i:
                cov = cov_matrix[np.ix_([j, i], [j, i])]
                lambda_, v = np.linalg.eig(cov)
                lambda_ = np.sqrt(lambda_)
                angle = np.rad2deg(np.arctan2(v[1, 0], v[0, 0]))
                for k in [1, 2]:
                    ellipse = Ellipse(xy=(mean[j], mean[i]),
                                      width=lambda[0] * k * 2,
                                      height=lambda_[1] * k * 2,
                                      angle=angle,
                                      edgecolor=ellipse_colors,
                                      facecolor='none')
                    ax.add_artist(ellipse)
                # Set axis limits
                margin = 3
                ax.set_xlim(mean[j] - margin * np.sqrt(cov_matrix[j, j]),__
 →mean[j] + margin * np.sqrt(cov_matrix[j, j]))
```

```
ax.set_ylim(mean[i] - margin * np.sqrt(cov_matrix[i, i]),__
→mean[i] + margin * np.sqrt(cov_matrix[i, i]))
                if true_values is not None:
                    ax.axvline(true_values[j], color='red', linestyle='-', lw=1)
                    ax.axhline(true_values[i], color='red', linestyle='-', lw=1)
            if j > i:
                ax.axis('off')
            if i < num_params - 1:</pre>
                ax.set_xticklabels([])
            else:
                if labels is not None:
                    ax.set_xlabel(labels[j])
            ax.yaxis.set_major_locator(plt.NullLocator())
            if j > 0:
                ax.set_yticklabels([])
            else:
                if labels is not None:
                    ax.set_ylabel(labels[i])
            ax.xaxis.set_major_locator(plt.NullLocator())
    plt.show()
target = generate_target()
```



#### 1.1 Levenberg-Marquardt

This fitter is identitied as ap.fit.LM and it employs a variant of the second order Newton's method to converge very quickly to the local minimum. This is the generally accepted best algorithm for most use cases in  $\chi^2$  minimization. If you don't know what to pick, start with this minimizer. The LM optimizer bridges the gap between first-order gradient descent and second order Newton's method. When far from the minimum, Newton's method is unstable and can give wildly wrong results, so LM takes gradient descent steps. However, near the minimum it switches to the Newton's method which has "quadratic convergence" this means that it takes only a few iterations to converge to several decimal places. This can be represented as:

$$(H + LI)h = g$$

Where H is the Hessian matrix of second derivatives, L is the damping parameter, I is the identity

matrix, h is the step we will take in parameter space, and g is the gradient. We solve this linear system for h to get the next update step. The "L" scale parameter goes from L >> 1 which represents gradient descent to L << 1 which is Newton's Method. When L >> 1 the hessian is effectively zero and we get h = g/L which is just gradient descent with 1/L as the learning rate. In AutoProf the damping parameter is treated somewhat differently, but the concept is the same.

LM can handle a lot of scenarios and converge to the minimum. Keep in mind, however, that it is seeking a local minimum, so it is best to start off the algorithm as close as possible to the best fit parameters. AutoProf can automatically initialize, as discussed in other notebooks, but even that needs help sometimes (often in the form of a segmentation map).

The main drawback of LM is its memory consumption which goes as  $\mathcal{O}(PN)$  where P is the number of pixels and N is the number of parameters.

```
[3]: MODEL = initialize_model(target, False)
    fig, axarr = plt.subplots(1,4, figsize = (24,5))
    plt.subplots_adjust(wspace= 0.1)
    ap.plots.model_image(fig, axarr[0], MODEL)
    axarr[0].set_title("Model before optimization")
    ap.plots.residual_image(fig, axarr[1], MODEL, normalize_residuals = True)
    axarr[1].set_title("Residuals before optimization")

res_lm = ap.fit.LM(MODEL, verbose = 1).fit()

ap.plots.model_image(fig, axarr[2], MODEL)
    axarr[2].set_title("Model after optimization")
    ap.plots.residual_image(fig, axarr[3], MODEL, normalize_residuals = True)
    axarr[3].set_title("Residuals after optimization")
    plt.show()
```

```
L: 1.0
-----init-----
LM loss: 283.81094319158024
L: 1.0
-----iter----
LM loss: 849.0109828400438
reject
L: 11.0
-----iter----
LM loss: 192.57226752058315
accept
L: 1.2222222222223
-----iter-----
LM loss: 69.76483619240399
accept
L: 0.1358024691358025
-----iter----
LM loss: 24.256005308222395
accept
```

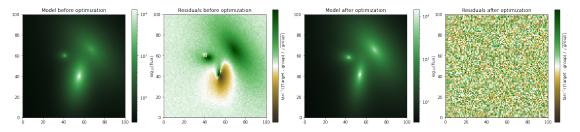
L: 0.015089163237311388 -----iter-----LM loss: 30.071938222875325 reject L: 0.16598079561042528 -----iter-----LM loss: 27.261758221743143 reject L: 1.825788751714678 -----iter----LM loss: 18.69506679958759 accept L: 0.20286541685718645 ----iter----LM loss: 28.386073296858303 reject L: 2.231519585429051 -----iter-----LM loss: 15.242646605322426 accept L: 0.2479466206032279 ----iter----LM loss: 22.387939636837555 reject L: 2.727412826635507 -----iter----LM loss: 12.461517579583184 accept L: 0.3030458696261674 ----iter----LM loss: 12.126933297954073 reject L: 3.3335045658878415 -----iter-----LM loss: 10.564772629729806 accept L: 0.3703893962097602 -----iter----LM loss: 6.123883804201112 accept L: 0.04115437735664002 ----iter----LM loss: 3.3033211897345005 accept

L: 0.004572708595182225

accept

LM loss: 1.9106270717708704

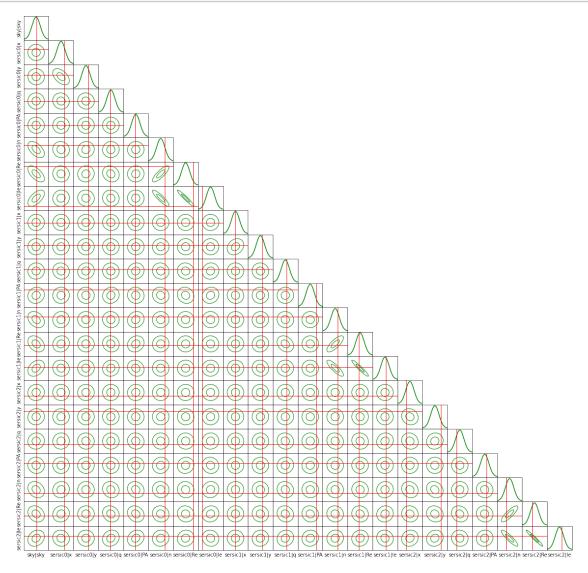
```
L: 0.000508078732798025
----iter----
LM loss: 1.4800231508207649
accept
L: 5.6453192533113885e-05
-----iter----
LM loss: 1.012878922723731
accept
L: 6.272576948123765e-06
----iter----
LM loss: 1.0127841836614357
accept
L: 6.969529942359739e-07
-----iter-----
LM loss: 1.0127841824837123
accept
```



Now that LM has found the  $\chi^2$  minimum, we can do a really neat trick. Since LM needs the hessian matrix, we have access to the hessian matrix at the minimum. This is in fact equal to the negative Fisher information matrix. If we take the matrix inverse of this matrix then we get the covariance matrix for a multivariate gaussian approximation of the  $\chi^2$  surface near the minimum. With the covariance matrix we can create a corner plot just like we would with an MCMC. We will see later that the MCMC methods (at least the ones which converge) produce very similar results!

```
param_names = list(MODEL.parameter_order())
i = 0
while i < len(param_names):
    param_names[i] = param_names[i].replace(" ", "")
    if "center" in param_names[i]:
        center_name = param_names.pop(i)
        param_names.insert(i, center_name.replace("center", "y"))
        param_names.insert(i, center_name.replace("center", "x"))
    i += 1
ser, sky = true_params()
corner_plot_covariance(
    res_lm.covariance_matrix.detach().cpu().numpy(),
    MODEL.get_parameter_vector().detach().cpu().numpy(),
    labels = param_names,</pre>
```

```
figsize = (20,20),
  true_values = np.concatenate((sky,ser.ravel()))
)
```



### 1.2 Iterative Fit (models)

An iterative fitter is identified as ap.fit.Iter, this method is generally employed for large models where it is not feasible to hold all the relevant data in memory at once. The iterative fitter will cycle through the models in a Group\_Model object and fit them one at a time to the image, using the residuals from the previous cycle. This can be a very robust way to deal with some fits, especially if the overlap between models is not too strong. It is however more dependent on good initialization than other methods like the Levenberg-Marquardt. Also, it is possible for the Iter method to get stuck in a local minimum under certain circumstances.

Note that while the Iterative fitter needs a <code>Group\_Model</code> object to iterate over, it is not necessarily true that the sub models are <code>Component\_Model</code> objects, they could be <code>Group\_Model</code> objects as well. In this way it is possible to cycle through and fit "clusters" of objects that are nearby, so long as it doesn't consume too much memory.

By only fitting one model at a time it is possible to get caught in a local minimum. For this reason it can be good to mix-and-match the iterative optimizers so they can help each other get unstuck.

```
[5]: MODEL = initialize_model(target, False)
    fig, axarr = plt.subplots(1,4, figsize = (24,5))
    plt.subplots_adjust(wspace= 0.1)
    ap.plots.model_image(fig, axarr[0], MODEL)
    axarr[0].set_title("Model before optimization")
    ap.plots.residual_image(fig, axarr[1], MODEL, normalize_residuals = True)
    axarr[1].set_title("Residuals before optimization")

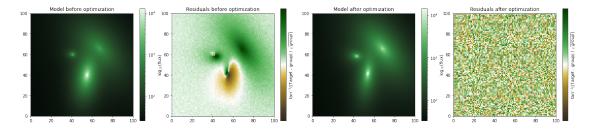
res_iter = ap.fit.Iter(MODEL, verbose = 1).fit()

ap.plots.model_image(fig, axarr[2], MODEL)
    axarr[2].set_title("Model after optimization")
    ap.plots.residual_image(fig, axarr[3], MODEL, normalize_residuals = True)
    axarr[3].set_title("Residuals after optimization")
    plt.show()
```

```
----iter----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 7.205865375699771
-----iter----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 2.4705932063631284
-----iter-----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.5805326550442294
-----iter-----
sky
sersic 0
```

```
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.2512768478124128
-----iter-----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.1127864669960037
-----iter----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.0530447882827145
-----iter-----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.0282650623186584
-----iter----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.0184151513545818
-----iter----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.0147107760651968
-----iter-----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.0134398452263913
-----iter-----
sky
```

```
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.0129958243247057
-----iter----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.012848036312808
-----iter----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.0128020490769345
-----iter----
sky
sersic 0
sersic 1
sersic 2
Update Chi^2 with new parameters
Loss: 1.012788813135204
```



#### 1.3 Iterative Fit (parameters)

This is an iterative fitter identified as ap.fit.Iter\_LM and is generally employed for large models where it is not feasible to hold all the relevant data in memory at once. This iterative fitter will cycle through chunks of parameters and fit them one at a time to the image. This can be a very robust way to deal with some fits, especially if the overlap between models is not too strong. This is very similar to the other iterative fitter, however it is necessary for certain fitting circumstances when the problem can't be broken down into individual component models. This occurs, for example, when the models have many shared (constrained) parameters and there is no obvious way to break down sub-groups of models (an example of this is discussed in the AutoProf paper).

Note that this is iterating over the parameters, not the models. This allows it to handle parameter covariances even for very large models (if they happen to land in the same chunk). However, for this to work it must evaluate the whole model at each iteration making it somewhat slower than the regular Iter fitter, though it can make up for it by fitting larger chunks at a time which makes the whole optimization faster.

By only fitting a subset of parameters at a time it is possible to get caught in a local minimum. For this reason it can be good to mix-and-match the iterative optimizers so they can help each other get unstuck.

```
[6]: MODEL = initialize_model(target, False)
    fig, axarr = plt.subplots(1,4, figsize = (24,5))
    plt.subplots_adjust(wspace= 0.1)
    ap.plots.model_image(fig, axarr[0], MODEL)
    axarr[0].set_title("Model before optimization")
    ap.plots.residual_image(fig, axarr[1], MODEL, normalize_residuals = True)
    axarr[1].set_title("Residuals before optimization")

res_iterlm = ap.fit.Iter_LM(MODEL, chunks = 11, verbose = 1).fit()

ap.plots.model_image(fig, axarr[2], MODEL)
    axarr[2].set_title("Model after optimization")
    ap.plots.residual_image(fig, axarr[3], MODEL, normalize_residuals = True)
    axarr[3].set_title("Residuals after optimization")
    plt.show()
```

```
['140099848431264:0' '140100290926240:0' '140099852362176:0'
 '140103645984896:0' '140099848431360:0' '140099848432320:0'
 '140099848430880:0' '140099848430112:0' '140099848430976:0'
 '140099848399696:0' '140099848398544:0']
chunk loss: 35.836873026504335
['140099848432224:0' '140099848431264:1' '140100290925376:0'
 '140103645984704:0' '140099848431360:1' '140099848433040:0'
 '140099848430352:0' '140099848430352:1' '140099848431552:0'
 '140099848399648:0' '140099848400800:0']
chunk loss: 14.562455505279104
Loss: 14.562455505279104
-----iter-----
['140099848431264:0' '140100290925376:0' '140100290926240:0'
 '140103645984704:0' '140103645984896:0' '140099848431360:0'
 '140099848432320:0' '140099848433040:0' '140099848430352:1'
 '140099848399648:0' '140099848400800:0']
chunk loss: 13.152171158257392
['140099848432224:0' '140099848431264:1' '140099852362176:0'
 '140099848431360:1' '140099848430880:0' '140099848430112:0'
 '140099848430976:0' '140099848430352:0' '140099848431552:0'
 '140099848399696:0' '140099848398544:0']
```

-----iter-----

```
chunk loss: 3.510773458706077
Loss: 3.510773458706077
-----iter-----
['140099848432224:0' '140099848431264:1' '140100290925376:0'
  '140099848431360:0' '140099848430880:0' '140099848433040:0'
  '140099848430352:0' '140099848430352:1' '140099848399648:0'
  '140099848399696:0' '140099848398544:0']
chunk loss: 3.353452269676198
['140099848431264:0' '140100290926240:0' '140099852362176:0'
  '140103645984704:0' '140103645984896:0' '140099848431360:1'
  '140099848432320:0' '140099848430112:0' '140099848430976:0'
  '140099848431552:0' '140099848400800:0']
chunk loss: 3.0961597807131325
Loss: 3.0961597807131325
-----iter-----
['140099848432224:0' '140099848431264:0' '140099848431264:1'
  '140100290925376:0' '140103645984896:0' '140099848431360:1'
  '140099848430112:0' '140099848433040:0' '140099848430352:0'
  '140099848399648:0' '140099848398544:0']
chunk loss: 2.8631077057761045
\[ \begin{aligned} \begin{alig
  '140099848431360:0' '140099848432320:0' '140099848430880:0'
  '140099848430976:0' '140099848430352:1' '140099848431552:0'
  '140099848399696:0' '140099848400800:0']
chunk loss: 2.7339689038865473
Loss: 2.7339689038865473
-----iter-----
['140099848432224:0' '140100290926240:0' '140099852362176:0'
  '140099848432320:0' '140099848430976:0' '140099848433040:0'
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Loss: 1.0131967739454029
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```

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Loss: 1.012888742507812
```

#### 1.4 Gradient Descent

A gradient descent fitter is identified as ap.fit.Grad and uses standard first order derivative methods as provided by PyTorch. These gradient descent methods include Adam, SGD, and LBFGS to name a few. The first order gradient is faster to evaluate and uses less memory, however it is considerably slower to converge than Levenberg-Marquardt. The gradient descent method with a small learning rate will reliably converge towards a local minimum, it will just do so slowly.

In the example below we let it run for 1000 steps and even still it has not converged. In general you should not use gradient descent to optimize a model. However, in a challenging fitting scenario the small step size of gradient descent can actually be an advantage as it will not take any unedpectedly large steps which could mix up some models, or hop over the  $\chi^2$  minimum into impossible parameter space. Just make sure to finish with LM after using Grad so that it fully converges to a reliable minimum.

```
[7]: MODEL = initialize_model(target, False)
    fig, axarr = plt.subplots(1,4, figsize = (24,5))
    plt.subplots_adjust(wspace= 0.1)
    ap.plots.model_image(fig, axarr[0], MODEL)
    axarr[0].set_title("Model before optimization")
    ap.plots.residual_image(fig, axarr[1], MODEL, normalize_residuals = True)
    axarr[1].set_title("Residuals before optimization")

res_grad = ap.fit.Grad(MODEL, verbose = 1, max_iter = 1000, optim_kwargs = \( \text{"1r": 5e-3} \)).fit()

ap.plots.model_image(fig, axarr[2], MODEL)
    axarr[2].set_title("Model after optimization")
    ap.plots.residual_image(fig, axarr[3], MODEL, normalize_residuals = True)
    axarr[3].set_title("Residuals after optimization")
    plt.show()
```

loss: 283.81094319158024 loss: 275.0393610548054 loss: 268.85931406428676 loss: 263.2779117938314 loss: 257.877334915675 loss: 252.5360252575934 loss: 247.22217833403124 loss: 241.9701048873146 loss: 236.72684346247385 loss: 231.53330760658764 loss: 226.40130852071508 loss: 221.33173382541167 loss: 216.3513607420002 loss: 211.46963084714196 loss: 206.67612250453496 loss: 201.96134579688248 loss: 197.32261129787932 loss: 192.78048552272938 loss: 188.33554270003418 loss: 183.9870655839911 loss: 179.73410690859117 loss: 175.57551803358464 loss: 171.51038920453948 loss: 167.53759711396364 loss: 163.65617554818527 loss: 159.8652348940428 loss: 156.163961996483 loss: 152.61719579475093 loss: 149.11110570683581 loss: 145.69366428214747 loss: 142.36503784991376 loss: 139.12461441907936 loss: 135.96429158199467 loss: 132.89874932029176 loss: 129.91871581018518 loss: 127.02642003435994 loss: 124.21990899280156 loss: 121.49877852316133 loss: 118.86847043127084 loss: 116.31611907280967 loss: 113.80266466016582 loss: 111.41655078144984 loss: 109.15484193103487 loss: 106.9293482488707 loss: 104.7819205352155 loss: 102.71124240337855 loss: 100.71290152999458 loss: 98.78799440108169 loss: 96.93523238749158 loss: 95.15292247895407 loss: 93.43757297433952 loss: 91.78849248086725 loss: 90.20322758570488 loss: 88.67957914006332 loss: 87.03614800064638 loss: 85.81043348397667 loss: 84.46036613864449 loss: 83.16223606790172 loss: 81.91533305248507 loss: 80.71778309850568 loss: 79.56593553098415 loss: 78.46037031794732

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- loss: 1.7384730381921822
- loss: 1.7358052977196055
- loss: 1.7369231258616409
- loss: 1.7342582601148766
- loss: 1.7353789167650537
- loss: 1.7327167612332752
- loss: 1.733840036183025
- loss: 1.731180470772833
- loss: 1.73230618252155
- loss: 1.7296491212581493
- loss: 1.7307771096393028
- loss: 1.7281224925246033
- loss: 1.7292526135757527
- loss: 1.7266004001895074
- loss: 1.727732522524705
- loss: 1.7250826870233873
- loss: 1.7262166892099464
- loss: 1.7235692164412524
- loss: 1.724704985051259
- loss: 1.7220598675563816
- loss: 1.7231972956783366
- loss: 1.7205545313992965

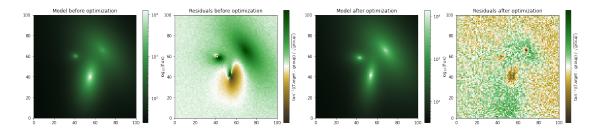
loss: 1.7216935174724703 loss: 1.7190622732431213 loss: 1.7201935562851502 loss: 1.7175646624194174 loss: 1.7187044568905698 loss: 1.7172537691872514 loss: 1.7158414595482758 loss: 1.7132151958541104 loss: 1.7143779956781056 loss: 1.7129215134761113 loss: 1.712828139310406 loss: 1.7102084276471685 loss: 1.7113750583792615 loss: 1.7087570695147463 loss: 1.709926152964093 loss: 1.7084721594406176 loss: 1.708391358222552 loss: 1.7057795883714904 loss: 1.706952711053431 loss: 1.7055014501032688 loss: 1.704141242789741 loss: 1.702792105854768 loss: 1.7026421723465028 loss: 1.7013094903996464 loss: 1.7011550839412715 loss: 1.699826383464564 loss: 1.6991091841826054 loss: 1.698359058152422 loss: 1.6976444709685174 loss: 1.6968970276451594 loss: 1.6961743285039201 loss: 1.6954404360267157 loss: 1.69472076348199 loss: 1.693988324380067 loss: 1.6932705905893128 loss: 1.6925400980075174 loss: 1.6918242199225295 loss: 1.6910955886988797 loss: 1.690381502617122 loss: 1.6896546691494152 loss: 1.688942325062543 loss: 1.6882172404046103 loss: 1.6875065981897708 loss: 1.6867832236551565 loss: 1.6860742505639392 loss: 1.6853525549467616 loss: 1.6846559680368351

loss: 1.6839250703250686

loss: 1.6832300941509544 loss: 1.6825116739757269 loss: 1.6818075593318353 loss: 1.6810907545626455 loss: 1.6803882230276483 loss: 1.679673018828669 loss: 1.6789720583862997 loss: 1.6782584431347618 loss: 1.6775590440671047 loss: 1.676847008615624 loss: 1.6761491633033039 loss: 1.6754387007420704 loss: 1.6747424035773575 loss: 1.6740335091171112 loss: 1.673338756503739 loss: 1.672631427476623 loss: 1.6719382179290352 loss: 1.6712324539127033 loss: 1.6705407882722154 loss: 1.6698365913519302 loss: 1.6691464731446175 loss: 1.668443848339766 loss: 1.6677552843125494 loss: 1.6670542402092052 loss: 1.6663672410970014 loss: 1.66566779074851 loss: 1.6649823723484412 loss: 1.6642845345336477 loss: 1.663600719194365 loss: 1.662904520161548 loss: 1.6622223388409463 loss: 1.6615278147189367 loss: 1.6614001342352485 loss: 1.6601504645557705 loss: 1.660023579200089 loss: 1.6587761403412662 loss: 1.6586500627143093 loss: 1.6574048245379838 loss: 1.656728399072158 loss: 1.6560398482961343 loss: 1.6559155714966647 loss: 1.6546746447686145 loss: 1.654551171793466 loss: 1.6533123973145778 loss: 1.6531897351546685 loss: 1.6519530934192956 loss: 1.651831245744091

loss: 1.6505967203055683

loss: 1.649927449451189 loss: 1.6492461891395451 loss: 1.649126173837288 loss: 1.6478958643406396 loss: 1.647776651548971 loss: 1.646548436830631 loss: 1.6464300305014075 loss: 1.6452038969227052 loss: 1.6450862984647026 loss: 1.6438622338493942 loss: 1.643745443933312 loss: 1.6425234375135938 loss: 1.6424074568626885 loss: 1.6411874996041054 loss: 1.6410723298647851 loss: 1.639854415252756 loss: 1.639196259936101 loss: 1.6385262372702691 loss: 1.6384128603093893 loss: 1.637199012656819 loss: 1.6370864275714128 loss: 1.6358746082643436 loss: 1.6357628208676782 loss: 1.6345530248955886 loss: 1.63444204311657



## 1.5 No U-Turn Sampler (NUTS)

Unlike the above methods, ap.fit.NUTS does not stricktly seek a minimum  $\chi^2$ , instead it is an MCMC method which seeks to explore the likelihood space and provide a full posterior in the form of random samples. The NUTS method in AutoProf is actually just a wrapper for the Pyro implementation (link here). Most of the functionality can be accessed this way, though for very advanced applications it may be necessary to manually interface with Pyro (this is not very challenging as AutoProf is fully differentiable).

The first iteration of NUTS is always very slow since it compiles the forward method on the fly, after that each sample is drawn much faster. The warmup iterations take longer as the method is exploring the space and determining the ideal step size and mass matrix for fast integration

with minimal numerical error (we only do 20 warmup steps here, if something goes wrong just try rerunning). Once the algorithm begins sampling it is able to move quickly (for an MCMC) throught the parameter space. For many models, the NUTS sampler is able to collect nearly completely uncorrelated samples, meaning that even 100 is enough to get a good estimate of the posterior.

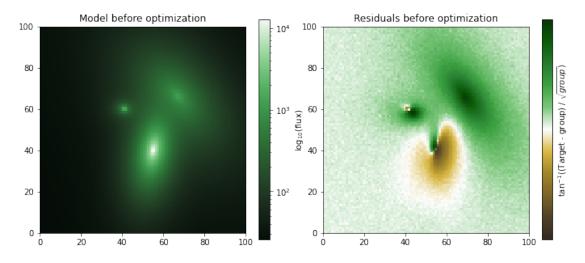
NUTS is far faster than other MCMC implementations such as a standard Metropolis Hastings MCMC. However, it is still a lot slower than the other optimizers (LM) since it is doing more than seeking a single high likelihood point, it is fully exploring the likelihood space. In simple cases, the automatic covariance matrix from LM is likely good enough, but if one really needs access to the full posterior of a complex model then NUTS is the best way to get it.

For an excellent introduction to the Hamiltonian Monte-Carlo and a high level explanation of NUTS see this review: **Betancourt 2018** 

```
[8]: MODEL = initialize_model(target, False)
    fig, axarr = plt.subplots(1,2, figsize = (12,5))
    plt.subplots_adjust(wspace= 0.1)
    ap.plots.model_image(fig, axarr[0], MODEL)
    axarr[0].set_title("Model before optimization")
    ap.plots.residual_image(fig, axarr[1], MODEL, normalize_residuals = True)
    axarr[1].set_title("Residuals before optimization")
    plt.show()

# Use LM to start the sampler at a high likelihood location, no burn-in needed!
# In general, NUTS is quite fast to do burn-in so this is often not needed
    res1 = ap.fit.LM(MODEL).fit()

# Run the NUTS sampler
    res_nuts = ap.fit.NUTS(MODEL, warmup = 20, max_iter = 100).fit()
```

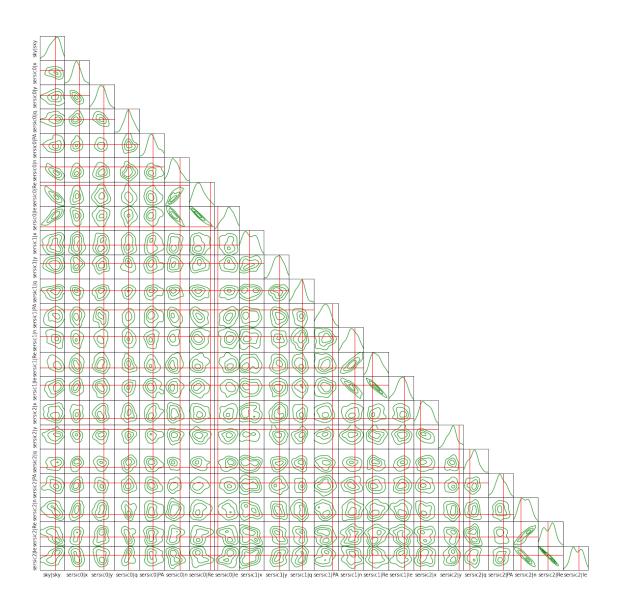


Sample: 100% | 120/120 [06:03, 3.03s/it, step size=2.45e-02, acc.

```
prob=0.653]
```

Note that there is no "after optimization" image above, because optimization was not done, it was full likelihood exploration. We can now create a corner plot with 2D projections of the 22 dimensional space that NUTS was exploring. The resulting corner plot is about what you would expect to get with 100 samples drawn from the multivariate gaussian found by LM above. If you run it again with more samples then the results will get even smoother.

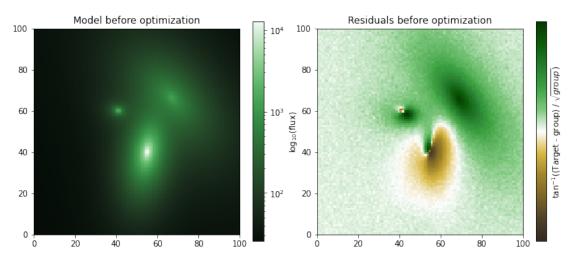
```
[9]: # corner plot of the posterior
     # observe that it is very similar to the corner plot from the LM optimization
     ⇒since this case can be roughly
     \# approximated as a multivariate gaussian centered on the maximum likelihood \sqcup
      \hookrightarrow point
     param_names = list(MODEL.parameter_order())
     i = 0
     while i < len(param_names):</pre>
         param_names[i] = param_names[i].replace(" ", "")
         if "center" in param_names[i]:
             center_name = param_names.pop(i)
             param names.insert(i, center name.replace("center", "y"))
             param_names.insert(i, center_name.replace("center", "x"))
         i += 1
     ser, sky = true_params()
     corner_plot(
         res_nuts.chain.detach().cpu().numpy(),
         labels = param_names,
         figsize = (20,20),
         true_values = np.concatenate((sky,ser.ravel()))
     )
```



## 1.6 Hamiltonian Monte-Carlo (HMC)

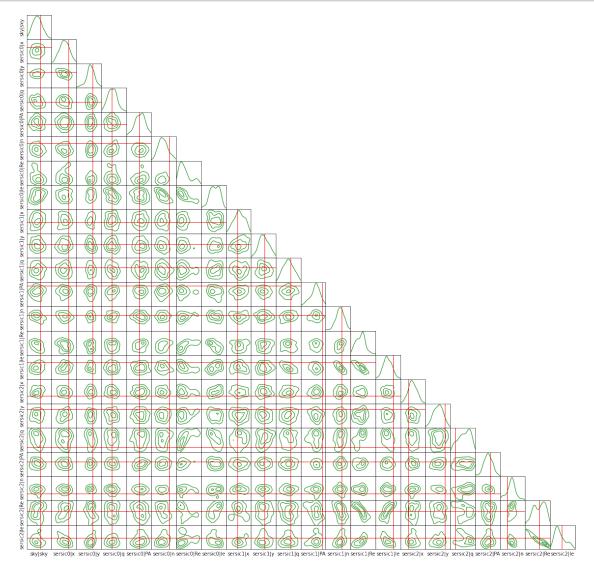
The ap.fit.HMC is a simpler variant of the NUTS sampler. HMC takes a fixed number of steps at a fixed step size following Hamiltonian dynamics. This is in contrast to NUTS which attempts to optimally choose these parameters. HMC may be suitable in some cases where NUTS is unable to find ideal parameters. Also in some cases where you already know the pretty good step parameters HMC may run faster. If you don't want to fiddle around with parameters then stick with NUTS, HMC results will still have autocorrelation which will depend on the problem and choice of step parameters.

```
[10]: MODEL = initialize_model(target, False)
fig, axarr = plt.subplots(1,2, figsize = (12,5))
plt.subplots_adjust(wspace= 0.1)
ap.plots.model_image(fig, axarr[0], MODEL)
```



Sample: 100% | 220/220 [02:26, 1.50it/s, step size=1.00e-02, acc. prob=0.939]

```
res_hmc.chain.detach().cpu().numpy(),
labels = param_names,
figsize = (20,20),
true_values = np.concatenate((sky,ser.ravel()))
)
```



## 1.7 Metropolis Hastings

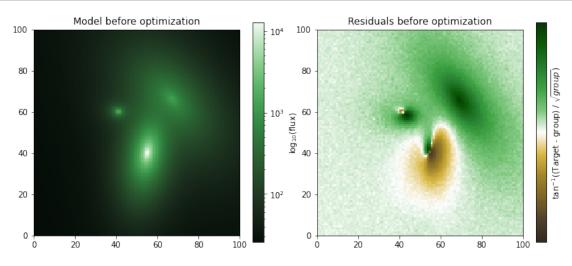
This is the classic MCMC algorithm using the Metropolis Hastngs accept step identified with ap.fit.MHMCMC. One can set the gaussian random step scale and then explore the posterior. While this technically always works, in practice it can take exceedingly long to actually converge to the posterior. This is because the step size must be set very small to have a reasonable likelihood of accepting each step, so it never moves very far in parameter space. With each subsequent sample being very close to the previous sample it can take a long time for it to wander away from its starting

point. In the example below it would take an extremely long time for the chain to converge. Instead of waiting that long, we demonstrate the functionality with 5000 steps, but suggest using NUTS for any real world problem. Still, if there is something NUTS can't handle (a function that isn't differentiable) then MHMCMC can save the day (even if it takes all day to do it).

```
[12]: MODEL = initialize_model(target, False)
    fig, axarr = plt.subplots(1,2, figsize = (12,5))
    plt.subplots_adjust(wspace= 0.1)
    ap.plots.model_image(fig, axarr[0], MODEL)
    axarr[0].set_title("Model before optimization")
    ap.plots.residual_image(fig, axarr[1], MODEL, normalize_residuals = True)
    axarr[1].set_title("Residuals before optimization")
    plt.show()

# Use LM to start the sampler at a high likelihood location, no burn-in needed!
    res1 = ap.fit.LM(MODEL).fit()

# Run the HMC sampler
    res_mh = ap.fit.MHMCMC(MODEL, verbose = 1, max_iter = 5000, epsilon = 1e-4, in the content of the content
```



100% | 5000/5000 [02:41<00:00, 31.04it/s]

Acceptance: 0.765999972820282

```
[13]: # corner plot of the posterior

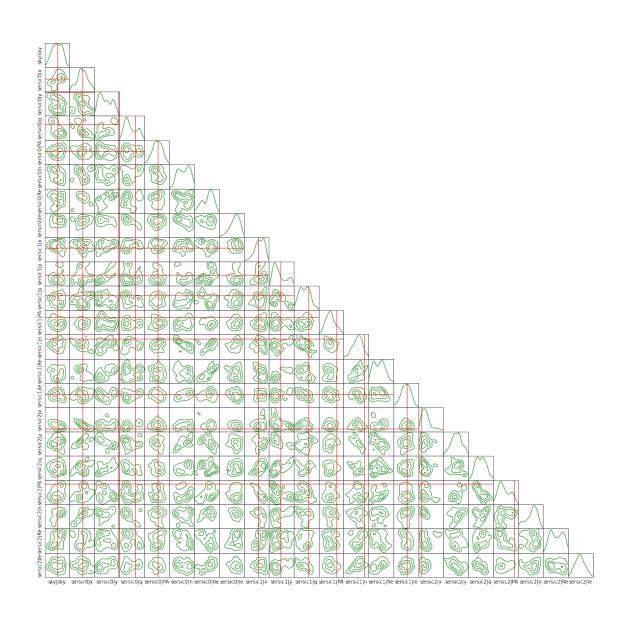
# note that, even 5000 samples is not enough to overcome the autocorrelation so⊔

→ the posterior has not converged.

# In fact it is not even close to convergence as can be seen by the multi-modal

→ blobs in the posterior since this
```

```
# problem is unimodal (except the modes where models are swapped). It is almost \Box
→never worthwhile to use this
# sampler except as a sanity check on very simple models.
param_names = list(MODEL.parameter_order())
i = 0
while i < len(param_names):</pre>
    param_names[i] = param_names[i].replace(" ", "")
    if "center" in param_names[i]:
        center_name = param_names.pop(i)
        param_names.insert(i, center_name.replace("center", "y"))
        param_names.insert(i, center_name.replace("center", "x"))
    i += 1
ser, sky = true_params()
corner_plot(
    res_mh.chain[::10], # thin by a factor 10 so the plot works in reasonable_
\rightarrow time
   labels = param_names,
    figsize = (20,20),
   true_values = np.concatenate((sky,ser.ravel()))
```



| []: |  |
|-----|--|
| []: |  |
| []: |  |