Student 4 CMA ES

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1 Covariance Matrix Adaptive ES

Note this is a simplified version of the original algorithm. Make sure to check the original paper before applying this optimizer to a real-world problem!

```
[]: %matplotlib notebook
  from math import sin, cos, sqrt, pi
  from matplotlib import cm
  import numpy as np
  import matplotlib.pyplot as plt
  from matplotlib.patches import Ellipse
  import matplotlib.transforms as transforms

[]: def loss_function(x, a=10):
    dummy = a * len(x)
    for ii in range(len(x)):
        dummy += x[ii] ** 2 - a * cos(2 * pi * x[ii])
    return dummy
```

```
def plot_loss(ax):
    x = np.linspace(-5, 5, 200)
    y = np.linspace(-5, 5, 200)
    X, Y = np.meshgrid(x, y)

Z = np.zeros_like(X)
    for ii in range(X.shape[0]):
        for jj in range(X.shape[1]):
            Z[ii][jj] = loss_function([X[ii][jj], Y[ii][jj]])
    img = ax.contour(X, Y, Z, levels=30, cmap=cm.coolwarm)
    plt.colorbar(img, ax=ax)

    return ax
```

```
[]: def error_ellipse(ax, xc, yc, cov, sigma=3, **kwargs):
    """
    https://github.com/megbedell/plot_tools/blob/master/error_ellipse.py
    Plot an error ellipse contour over your data.
```

```
Inputs:
   ax : matplotlib Axes() object
   xc : x-coordinate of ellipse center
  yc : x-coordinate of ellipse center
  cov : covariance matrix
  sigma : # sigma to plot (default 1)
   additional kwargs passed to matplotlib.patches.Ellipse()
  w, v = np.linalg.eigh(cov) # assumes symmetric matrix
  order = w.argsort()[::-1]
  w, v = w[order], v[:, order]
  theta = np.degrees(np.arctan2(*v[:, 0][::-1])) # * unpacks argument_
\rightarrow instead of [0]
  ellipse = Ellipse(
       xy=(xc, yc),
       width=2.0 * sigma * np.sqrt(w[0]),
       height=2.0 * sigma * np.sqrt(w[1]),
       angle=theta,
       **kwargs
  ellipse.set facecolor("none")
  ax.add_artist(ellipse)
  return ax
```

```
[]: def covar_helper(x1, mu_x1, x2, mu_x2):
    if len(x1) != len(x2):
        raise Exception

tmp = 0
    for ii in range(len(x1)):
        tmp += (x1[ii] - mu_x1) * (x2[ii] - mu_x2)

return tmp / len(x1)
```

```
[]: # Simplified CMA ES
    n_evolutions = 20
    n_population = 100
    best_perc = 0.25
    start = [3.5, 3.5]
    init_population = [start] * n_population
    sig_0 = 0.65
    population = np.random.normal(init_population, sig_0)

centers = [start]

# Inits for plot
```

```
fig = plt.figure()
ax = fig.gca()
axes = plt.gca()
axes.set_xlim([-5, 5])
axes.set_ylim([-5, 5])
plt.ion()
fig.show()
fig.canvas.draw()
plot_loss(ax)
for episode in range(n_evolutions):
   if episode > 0:
        # confidence_ellipse(cov, mean_old, ax)
        ax = error_ellipse(
            ax, mean_old[0], mean_old[1], cov, ec="green", zorder=9999
    # Population Update Strategy
    # Todo: store the loss of each particle in a list named scores
    # Todo: Find the best best_perc percent particles with respect to their_
→loss.
            Store these particles in a numpy array named best_population
    # Todo: Find the mean of the best population and save it as mean
   centers.append(list(mean))
   if episode == 0:
       mean_old = mean
   sig_xx = covar_helper(
       best_population[:, 0], mean_old[0], best_population[:, 0], mean_old[0]
   )
    sig_xy = covar_helper(
       best_population[:, 0], mean_old[0], best_population[:, 1], mean_old[1]
    sig_yy = covar_helper(
       best_population[:, 1], mean_old[1], best_population[:, 1], mean_old[1]
```

```
# Todo: build a covariance matrix with the entries above and draw a new_
population around the mean calculated above.

mean_old = mean

# Plot population
fig.canvas.draw()

centers = np.array(centers)

plt.plot(centers[:,0], centers[:,1], ".-g", label="Centers")
plt.plot(centers[0,0], centers[0,1], "or", label="Start")
plt.plot(centers[-1,0], centers[-1,1], "xr", label="End")
plt.legend()
```

[]: