

### Chapter 31-13: Parabolic PDEs Using the Monte Carlo Method.

Monte Carlo Methods are a class of computational solutions that can be applied to various problems which rely on repeated random sampling to provide generally approximate solutions. Monte Carlo is a stochastic approach, in which a series of simulations (trials), representing the analyzed problem, with randomly selected input values, are performed. Among these trials, a specified number of properly defined successes is achieved. The ratio between the number of success trials to the number of all trials, scaled by dimensional quantity (e.g., area or function value) allows for the estimation of the unknown solution, providing the number of trials is large enough.

At the address: <https://srome.github.io/On-Solving-Partial-Differential-Equations-with-Brownian-Motion-in-Python/> (<https://srome.github.io/On-Solving-Partial-Differential-Equations-with-Brownian-Motion-in-Python/>) there is an interesting blog about doing pde solving with Brownian motion approximation. However, BM is not exactly Monte Carlo, is it? Actually, the two stochastic approaches are sometimes covered in the same place, for example: [http://www.columbia.edu/~mh2078/MonteCarlo/MCS\\_Generate\\_RVars.pdf](http://www.columbia.edu/~mh2078/MonteCarlo/MCS_Generate_RVars.pdf) ([http://www.columbia.edu/~mh2078/MonteCarlo/MCS\\_Generate\\_RVars.pdf](http://www.columbia.edu/~mh2078/MonteCarlo/MCS_Generate_RVars.pdf)). So it will be assumed here that using BM is essentially the same as using MC. And besides, it took 7.5 hrs for a 13700K processor to spit out the graphics for it, so they are not about to be wasted.

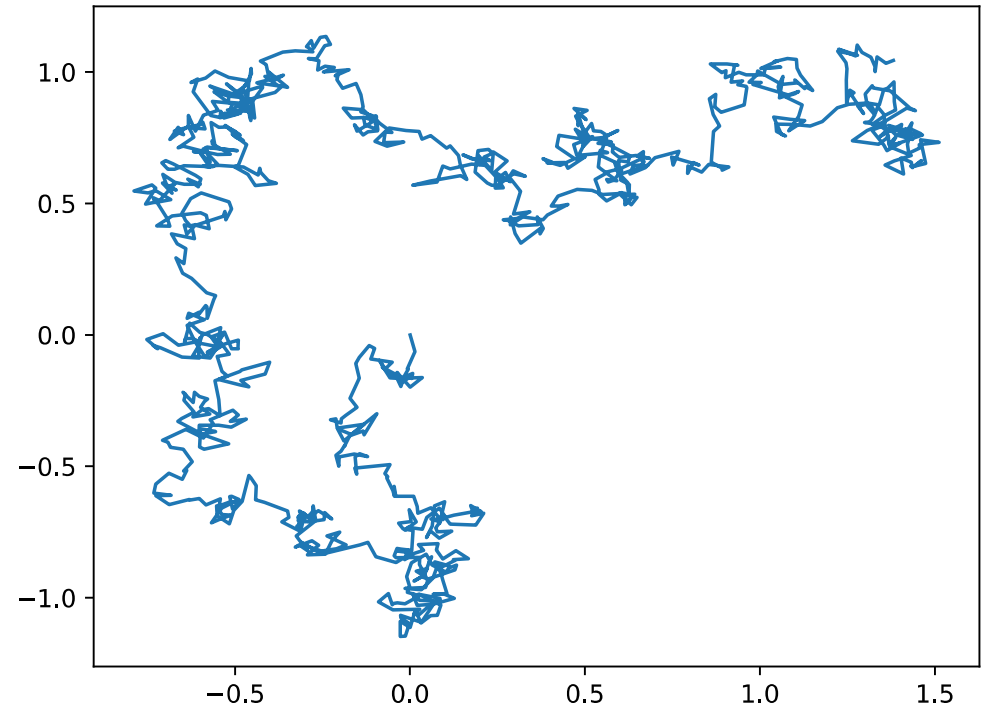
Perhaps an opportunity is here to reaffirm that the Laplace equation is in the parabolic category.

In [7]:

```
1  %%matplotlib inline
2  import numpy as np
3  import matplotlib.pyplot as plt
4
5  t=1
6  n = 1000
7  delta_t = np.sqrt(t/n)
8
9  # Create each dW step
10 dw1 = delta_t * np.random.normal(0,1,size=n)
11 dw2 = delta_t * np.random.normal(0,1,size=n)
12 W = np.zeros((n+1,2))
13
14 # Add W_{j-1} + dW_{j-1}
15 W[1:,0] = np.cumsum(dw1)
16 W[1:,1] = np.cumsum(dw2)
17
18 plt.plot(W[:,0],W[:,1], '-')
```

Out[7]:

```
[<matplotlib.lines.Line2D at 0x2a2a88dea90>]
```



Above: a random Brownian walk, which has the same appearance as a Monte Carlo generated walk.

In [8]:

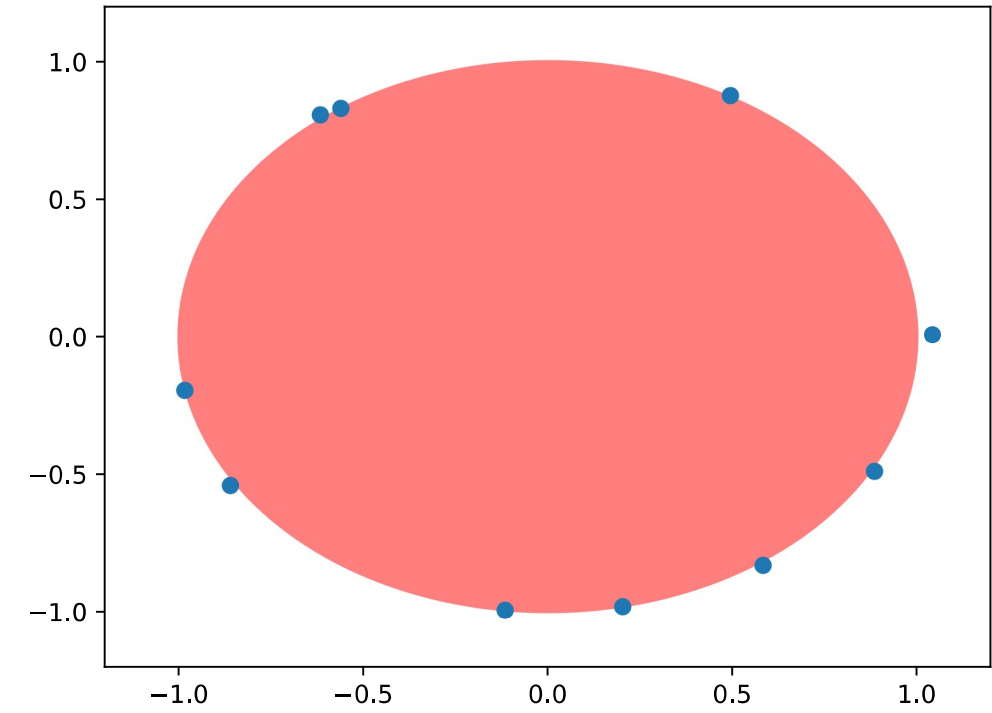
```
1  def check_if_exit(v):
2      # Takes a vector v=(x,y)
3      # Checks if v has intersected with the boundary of D
4      if np.linalg.norm(v,2) >=1:
5          return True
6      return False
7
8  def simulate_exit_time(v):
9      # Simulates exit time starting at v=(x,y), returns exit position
10     delta_t = np.sqrt(.001)
11     exit = False
12     x = v.copy()
13     while not exit:
```

```

14         x = x + delta_t * np.random.normal(0,1,size=2)
15         exit = check_if_exit(x)
16         return x
17
18 v=np.array((0,0)) # The origin
19 exit_times = np.array([simulate_exit_time(v) for k in range(0,10) ])
20
21
22 circle1=plt.Circle((0,0),1,color='r', alpha=.5)
23 plt.gcf().gca().add_artist(circle1)
24 plt.axis([-1.2, 1.2, -1.2, 1.2])
25 plt.scatter(exit_times[:,0],exit_times[:,1])
26
27

```

Out[8]: <matplotlib.collections.PathCollection at 0x2a2a8a943a0>



Above: simulated exit points on an ellipse, fairly similar to the epsilon criterion for deciding when a random step reaches the domain boundary in the WOS (walk on spheres) version of popular pde solvers.

In [ ]:

```

In [4]: 1      np.random.seed(8) #Side Infinity
2
3      def check_if_exit(v):
4          # Takes a vector v=(x,y)
5          # Checks if v has intersected with the boundary of D
6          if np.linalg.norm(v,2) >=1:
7              return True
8          return False
9
10     def simulate_exit_time(v):
11         # Simulates exit time starting at v=(x,y), returns exit position
12         delta_t = np.sqrt(.001)
13         exit = False
14
15         # Copy because simulation modifies in place
16         if hasattr(v,'copy'): # For NumPy arrays
17             x = v.copy()
18         else:
19             x = np.array(v) # We input a non-NumPy array
20         while not exit:
21             x += delta_t * np.random.normal(0,1,size=2) # += modifies in place
22             exit = check_if_exit(x)
23         return x
24
25     v=np.array((.5,.5)) # The origin
26     u = lambda x : np.linalg.norm(x,2)*np.cos(np.arctan2(x[1],x[0]))
27     f = lambda x : np.cos(np.arctan2(x[1],x[0]))
28
29     def get_exp_f_exit(starting_point, n_trials):
30         return np.mean([f(simulate_exit_time(starting_point)) for k in range(0,n_trials)])
31
32     exp_f_exit = get_exp_f_exit(v,2000) # Expected value of f(Exit(x,d))
33     print('The value u(v) = %s\nThe value of Exp(f(Exit))=%s' %(u(v), exp_f_exit))
34
35

```

The value u(v) = 0.5000000000000001  
The value of Exp(f(Exit))=0.4975601145395467

```

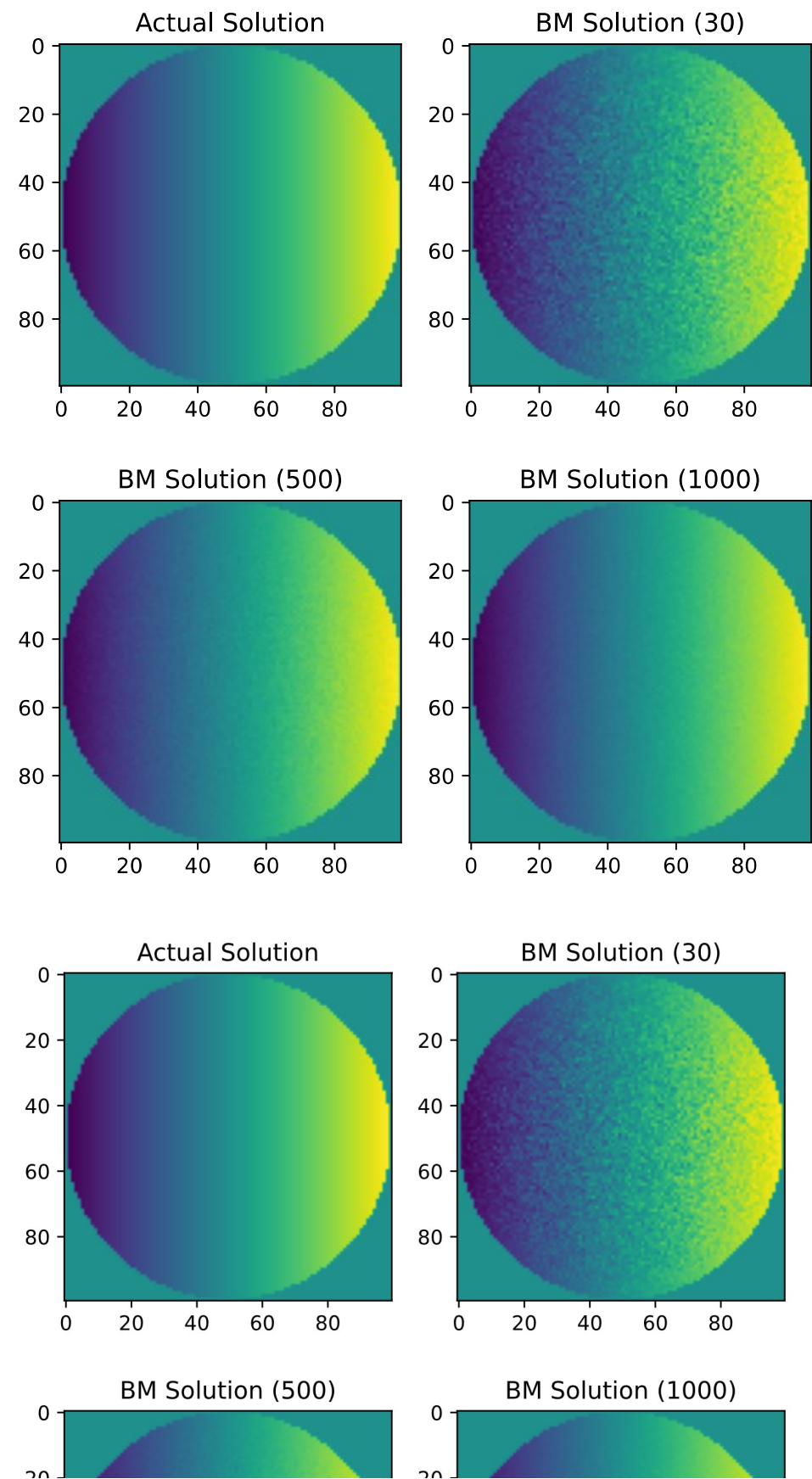
In [5]: 1      lin = np.linspace(-1, 1, 100)
2      x, y = np.meshgrid(lin, lin)
3      print(x.shape)
4      u_vec = np.zeros(x.shape)
5      bm_vec_30 = np.zeros(x.shape)
6      bm_vec_500 = np.zeros(x.shape)
7      bm_vec_1000 = np.zeros(x.shape)
8

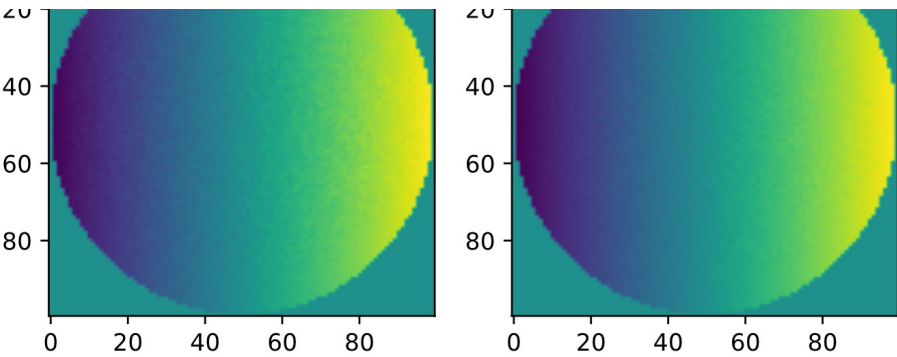
```

```
9      # Convert u to a solution in x,y coordinates
10     u_x = lambda x,y : np.linalg.norm(np.array([x,y]),2)*np.cos(np.arctan2(y,x))
11
12     # Calculate actual and approximate solution for (x,y) in D
13     for k in range(0,x.shape[0]):
14         for j in range(0,x.shape[1]):
15             x_t = x[k,j]
16             y_t = y[k,j]
17
18             # If the point is outside the circle, the solution is undefined
19             if np.sqrt((x_t)**2 + (y_t)**2) > 1:
20                 continue
21
22             # Calculate function value at this point for each image
23             u_vec[k,j] = u_x(x_t,y_t)
24             bm_vec_30[k,j] = get_exp_f_exit((x_t,y_t),30)
25             bm_vec_500[k,j] = get_exp_f_exit((x_t,y_t),500)
26             bm_vec_1000[k,j] = get_exp_f_exit((x_t,y_t),1000)
27
28     fig = plt.figure()
29     ax = fig.add_subplot(121)
30     plt.imshow(u_vec)
31     plt.title('Actual Solution')
32
33     ax = fig.add_subplot(122)
34     plt.title('BM Solution (30)')
35     plt.imshow(bm_vec_30)
36
37     fig = plt.figure()
38
39     ax = fig.add_subplot(121)
40     plt.title('BM Solution (500)')
41     plt.imshow(bm_vec_500)
42
43     ax = fig.add_subplot(122)
44     plt.title('BM Solution (1000)')
45     plt.imshow(bm_vec_1000)
46
```

(100, 100)

Out[5]: <matplotlib.image.AxesImage at 0x2a2a852cc10>





Above: Brownian motion solutions, rather similar to the walk on spheres output for a parabolic pde performed with Monte Carlo.

The long processing time practically guarantees that Github will never show this result, so it is shown here in static form.

In [ ]: