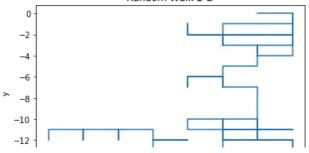
Random walk and self-avoiding random walk

2-D Random Walk - the usual way

A 2-D normal Random Walk is propagated in a 2-D(x-y) plane. It consists of motion in 4 directions i.e.all of Up-Down-Left-Right. Espescially the walker is allowed to come back to points already visited.

```
In [1]:
         import numpy as np
         import matplotlib.pyplot as plt
         import random
         from numpy import linalg as LA
         from scipy.optimize import curve fit
In [2]:
         class ClassicalRandomWalk:
             # Constructor for setting up the grid
             def __init__(self, size_grid, start_pos):
                 # pos for storing the coordinates of the position of the walker in a size gridx2 matrix
                 self. pos = np.zeros((size grid+1,2))
                 # Set initial start position
                 self.\_pos[0][0] = start\_pos[0]
                 self._pos[0][1] = start_pos[1]
                 # We assume four directions of movement
                 self.__direction = ["NORTH", "SOUTH", "EAST", "WEST"]
             def x positions(self):
                 return self. pos[:, 0]
             def y_positions(self):
                 return self.__pos[:, 1]
             def simulate walk(self, steps):
                 for i in range(1,steps+1):
                     # Randomly choosing the direction of movement
                     step = random.choice(self.__direction)
                     # Updating the direction with respect to the direction of motion choosen
                     if step == "EAST":
                         self.\_pos[i] = [self.\_pos[i-1][0] + 1, self.\_pos[i-1][1]]
                     elif step == "WEST":
                         self.__pos[i] = [self.__pos[i-1][0] - 1, self.__pos[i-1][1]]
                     elif step == "NORTH":
                         self._pos[i] = [self._pos[i-1][0], self._pos[i-1][1] + 1]
                         self.\_pos[i] = [self.\_pos[i-1][0], self.\_pos[i-1][1] - 1]
                 return self.__pos[steps][:]
             def plot trajectory(self):
                 plt.title("Random Walk 2-D")
                 plt.xlabel('x'
                 plt.ylabel('y')
                 plt.plot(self.x_positions(), self.y_positions())
                 plt.show()
In [3]:
         # the number of steps we want to simulate, so increase the value of n increses the complexity of the shown of
         steps = 100
         # Run the simulation and plot the results.
         classical random walk = ClassicalRandomWalk(steps, [0,0])
         classical_random_walk.simulate_walk(steps)
         classical_random_walk.plot_trajectory()
                            Random Walk 2-D
```

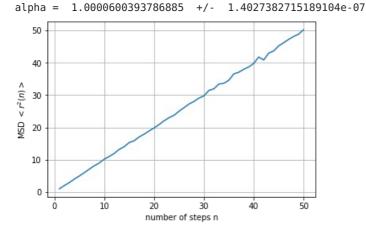


```
-14 - -6 -5 -4 -3 -2 -1 0 1
```

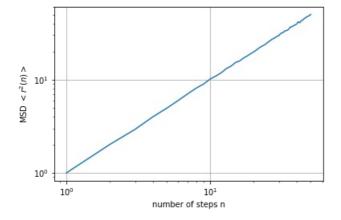
```
In [4]:
         # Calculation of the mean squared displacement (MSD) depending on the number of steps n
         # number of timesteps
         num\_timesteps = 50
         # Number of simulations being done for the given timestep
         num trajectories = 10000
         msd = np.zeros(num_timesteps)
         for timestep in range(1,num timesteps+1):
             # Array for caching the L2-norm squared of the
             # end point of each trajectory being simulated
             msd_single_ensemble = np.zeros(num_trajectories)
             # Calculate the ensemble mean for the actual timestep
             for i in range(num_trajectories):
                 classical random walk = ClassicalRandomWalk(timestep, [0,0])
                 end_pos_n = classical_random_walk.simulate_walk(timestep)
                 msd_single_ensemble[i] = LA.norm(end_pos_n, 2)**2
             msd[timestep-1] = 1. / num trajectories * np.sum(msd single ensemble)
```

```
In [5]:
# Plot the dependency between number of timesteps and MSD
plt.plot(np.arange(1,num_timesteps+1), msd)
plt.xlabel('number of steps n')
plt.ylabel('MSD $<r^2(n)>$')
plt.grid()

popt, pcov = curve_fit(lambda x,a:x**a, np.arange(1,num_timesteps+1), msd)
print("alpha = " , float(popt), " +/- ", float(pcov))
```



```
In [6]:
    # Plot the above dependency in a log-log-scale
    plt.plot(np.arange(1,num_timesteps+1), msd)
    plt.xlabel('number of steps n')
    plt.ylabel('MSD $<r^2(n)>$')
    plt.xscale('log')
    plt.yscale('log')
    plt.grid()
```



2-D Random walk - the self-avoiding random walk (SAW)

In the so-called self-avoiding random walk (SAW), the walker is not allowed to do so and hence does not cross its own path.

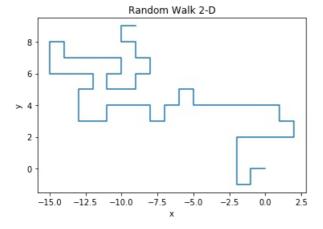
```
In [7]:
         class SelfAvoidingRandomWalk:
              # Constructor for setting up the grid
              def init (self, size grid, start pos):
                  # size of the grid on which the random walk is performed
                  self. size grid = size grid + 1;
                  # pos for storing the coordinates of the position of the walker in a size gridx2 matrix
                  self.__pos = np.zeros((self.__size_grid,2))
                  # Set initial start position
                  self._pos[0][0] = start_pos[0]
                  self.__pos[0][1] = start_pos[1]
                  # We assume four directions of movement
                  self. direction = ["NORTH", "SOUTH", "EAST", "WEST"]
                  # For the self avoiding Random Walk we introduce a grid for checking if node already has been visite
                  self. visited pos = np.zeros((size grid+1, size grid+1))
                  # Mark initial start position node as visited
                  self.__visited_pos[int(self.__pos[0][0])][int(self.__pos[0][1])] = 1
              def x positions(self):
                  return self. pos[:, 0]
              def y positions(self):
                  return self.__pos[:, 1]
              def simulate walk(self, steps):
                  for i in range(1, steps+1):
                      check validity = self.simulation step(i)
                      if (check_validity == False):
                           # Reset all arrays
                           self.__pos = np.zeros((self.__size grid,2))
                           self. visited pos = np.zeros((self. size grid, self. size grid))
                           # Restart computation
                           self.simulate walk(steps)
                  return self.__pos[steps][:]
              def simulation step(self, step num):
                  #Flag in order to check if node has already been visited
                  step positive = False
                  # Array for identify dead streets and restart computation
                  dead end = np.zeros(4);
                  while(step positive == False):
                      # Randomly choosing the direction of movement
                      step = random.choice(self. direction)
                       # Updating the direction with respect to the direction of motion choosen
                      if step == "EAST":
                           if(self.__visited_pos[int(self.__pos[step_num-1][0] + 1)][int(self.__pos[step_num-1][1])] ==
    self.__pos[step_num] = [self.__pos[step_num-1][0] + 1, self.__pos[step_num-1][1]]
                               self.__visited_pos[int(self.__pos[step_num-1][0] + 1)][int(self.__pos[step_num-1][1])] =
                               step positive = True
                           else:
                               dead end[0] = 1
                      elif step == "WEST":
                           if(self.__visited_pos[int(self.__pos[step_num-1][0] - 1)][int(self.__pos[step_num-1][1])] ==
    self.__pos[step_num] = [self.__pos[step_num-1][0] - 1, self.__pos[step_num-1][1]]
                               self.__visited_pos[int(self.__pos[step_num-1][0] - 1)][int(self.__pos[step_num-1][1])] =
                               step positive = True
                           else:
                               dead_end[1] = 1
                      elif step == "NORTH":
                           if(self.__visited_pos[int(self.__pos[step_num-1][0])][int(self.__pos[step_num-1][1] + 1)] ==
```

 $self._pos[step_num] = [self._pos[step_num-1][0], self._pos[step_num-1][1] + 1]$

```
self. visited pos[int(self. pos[step num-1][0])][int(self. pos[step num-1][1] + 1)];
                  step positive = True
             else:
                  dead end[2] = 1
         else:
             if(self.__visited_pos[int(self.__pos[step_num-1][0])][int(self.__pos[step_num-1][1] - 1)] ==
    self.__pos[step_num] = [self.__pos[step_num-1][0], self.__pos[step_num-1][1] - 1]
                  self.__visited_pos[int(self.__pos[step_num-1][0])][int(self.__pos[step_num-1][1] - 1)] =
                  step_positive = True
              else:
                  dead end[3] = 1
         # Check whether we land in a dead end
         if((dead_end == 1).all() == True):
              return False
    # Simulation step succeed
    return True
def plot trajectory(self):
    plt.title("Random Walk 2-D")
    plt.xlabel('x')
    plt.ylabel('y')
    plt.plot(self.x positions(), self.y positions())
    plt.show()
```

```
# the number of steps we want to simulate, so increase the value of n increases the complexity of the shown
steps = 60;

# Run the simulation and plot the results.
self_avoiding_random_walk = SelfAvoidingRandomWalk(steps, [0,0])
self_avoiding_random_walk.simulate_walk(steps)
self_avoiding_random_walk.plot_trajectory()
```



```
In [9]:
         \# Calculation of the mean squared displacement (MSD) depending on the number of steps n
         # number of timesteps
         num timesteps = 10
         # Number of simulations being done for the given timestep
         num trajectories = 1500
         msd = np.zeros(num timesteps)
         for timestep in range(1,num_timesteps+1):
             # Array for caching the L2-norm squared of the
             # end point of each trajectory being simulated
             msd_single_ensemble = np.zeros(num_trajectories)
             # Calculate the ensemble mean for the actual timestep
             for i in range(num_trajectories):
                 self avoiding random walk = SelfAvoidingRandomWalk(timestep, [0,0])
                 end_pos_n = self_avoiding_random_walk.simulate_walk(timestep)
                 msd single ensemble[i] = LA.norm(end pos n, 2)**2
             msd[timestep-1] = 1. / num_trajectories * np.sum(msd_single_ensemble)
```

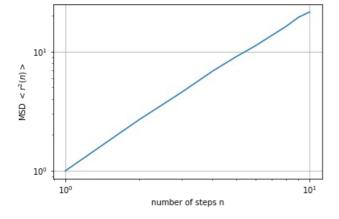
```
In [10]: # Plot the dependency between number of timesteps and MSD
plt.plot(np.arange(1,num_timesteps+1), msd)
plt.xlabel('number of steps n')
```

```
plt.ylabel('MSD $<r^2(n)>$')
plt.grid()

popt, pcov = curve_fit(lambda x,a:x**a, np.arange(1,num_timesteps+1), msd)
print("alpha = " , float(popt), " +/- ", float(pcov))
```

```
alpha = 1.3423752826087423 +/- 1.1830030562571881e-05
```

```
# Plot the above dependency in a log-log-scale
plt.plot(np.arange(1,num_timesteps+1), msd)
plt.xlabel('number of steps n')
plt.ylabel('MSD $<r^2(n)>$')
plt.xscale('log')
plt.yscale('log')
plt.grid()
```



Summary

As we see, the exponent for the classsical random walk is $\alpha = 1$, which corresponds to a normal diffusion process.

For the self avoiding random walk the exponent is $\alpha = 1.34$, which corresponds to a superdiffusion process. So probably, a self avoiding random walk is a good way to model a superdiffusion process.

The higher exponent for the self avoiding random walk is also quite obvious, since a particle must diffuse away faster.