# Physics 305 Demo Notebook 3: Calculating the Mean Square Displacement (MSD) from a Time-Series - Brownian Motion

The mean square deviation or MSD is defined to be:  $\langle x^2 \rangle - \langle x \rangle^2$ . This is also known as the *structure function*.

Recall that for any two points  $x_i$  and  $x_j$ , we define the lag time  $\tau$  to be  $t_i - t_i$ .

Then, for Brownian motion, MSD =  $2D\tau$  (Eq. 1), where D is the diffusion coefficient. Recall also that the PDF for a given lag time  $\tau$  is a Gaussian with a standard deviation of MSD<sup>1/2</sup> =  $\sqrt{2D\tau}$  (as derived in class and illustrated in Demo Notebook #2).

Let us now also define  $\Delta$  to be the number of time steps equivalent to the lag time  $\tau$ , i.e.,  $\tau = \Delta * dt$ , where dt is the size of the time step.

We introduce the empirical definition of the MSD as a function of  $\Delta$  (Eq. 2):

$$MSD(\Delta) = \frac{1}{n-\Delta} \sum_{j=0}^{j=n-1-\Delta} (x_{j+\Delta} - x_j)^2$$

Here, we will calculate the MSD as a function of lag time (using Eq. 2) for standard Brownian motion samples and compare them with the analytical result. Recall that for standard Brownian motion, D=1/2, so we expect (from Eq. 1):

$$MSD(\tau) = 2D\tau = \tau.$$

This is simply a linear relation with a slope of 1.

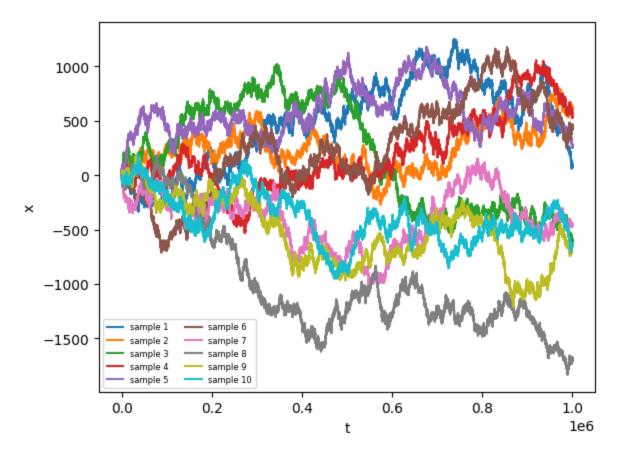
# **Step 1: Generate Brownian motion samples**

```
In [2]: # import libraries
   import numpy as np
   from scipy.stats import norm
   import matplotlib.pyplot as plt
%matplotlib inline
```

```
In [3]: # Set random seed
        np.random.seed(seed=17)
        # Set parameters
        n samp = 10 # no. of samples/realizations
        n = int(1e6) # no. of timesteps
        dt = 1. # size of time step
        sd = np.sqrt(dt) # standard deviation
        t = dt*np.arange(n) # time
        # Initialize array to hold the BM samples
        x_samp = np.zeros((n, n_samp))
        # Loop over realizations
        for i in np.arange(n samp):
          # Generate random numbers from Gaussian distribution centered at 0 and
        standard deviation sd
          rnd = norm.rvs(size = n, scale = sd)
          # Get cumulative sum of the elements of the array of random numbers
          x samp[:, i] = np.cumsum(rnd)
```

```
In [4]: # Plot BM samples
    for i in np.arange(n_samp):
        plt.plot(t, x_samp[:, i],label="sample %d" % (i+1))
        plt.xlabel("t")
        plt.ylabel("x")
        plt.legend(loc="lower left", ncol=2, fontsize=6)
```

Out[4]: <matplotlib.legend.Legend at 0x7974e8fe9f60>



# Step 2: Generate the MSD

First, let us define the range of lag times to calculate the MSD for. Recall that when we calculated the displacement probability distributions (PDFs) in Demo Notebook #2, we selected  $\tau = 10.10^2, 10^3, 10^4, 10^5, 9 \times 10^5$ .

We found that for  $\tau = 10^3$  and lower, there is a very good match between the calculated and analytical results, even for a single realization (but this does not necessarily hold for larger lag times).

Here, let us calculate the MSD for  $1 \le \tau \le 9 \times 10^5$  and indicate the regime  $\tau < 10^3$  (to remind us that small-number statistics is affecting large lag times).

Computationally, we will do this for an grid of  $\tau$  values equally-spaced in log scale. Let us also perform the calculation for all  $n_{\rm samp}=10$  samples.

```
In [5]: n tau = 100
        tau_vals = np.logspace(0, np.log10(9e5), n_tau)
        delta vals = np.round(tau vals/dt).astype(int)
        tau vals
Out[5]: array([1.00000000e+00, 1.14853402e+00, 1.31913040e+00, 1.51506614e+00,
               1.74010500e+00, 1.99856979e+00, 2.29542540e+00, 2.63637416e+00,
               3.02796541e+00, 3.47772129e+00, 3.99428121e+00, 4.58756786e+00,
               5.26897775e+00, 6.05160020e+00, 6.95046871e+00, 7.98284976e+00,
               9.16857453e+00, 1.05304198e+01, 1.20945453e+01, 1.38909968e+01,
               1.59542824e+01, 1.83240361e+01, 2.10457788e+01, 2.41717930e+01,
               2.77621265e+01, 3.18857468e+01, 3.66218650e+01, 4.20614578e+01,
               4.83090152e+01, 5.54845475e+01, 6.37258903e+01, 7.31913530e+01,
               8.40627589e+01, 9.65489385e+01, 1.10889740e+02, 1.27360639e+02,
               1.46278027e+02, 1.68005291e+02, 1.92959792e+02, 2.21620885e+02,
               2.54539126e+02, 2.92346846e+02, 3.35770298e+02, 3.85643611e+02,
               4.42924807e+02, 5.08714209e+02, 5.84275575e+02, 6.71060375e+02,
               7.70735671e+02, 8.85216138e+02, 1.01670085e+03, 1.16771551e+03,
               1.34116099e+03, 1.54036903e+03, 1.76916623e+03, 2.03194761e+03,
               2.33376095e+03, 2.68040385e+03, 3.07853501e+03, 3.53580219e+03,
               4.06098910e+03, 4.66418414e+03, 5.35697416e+03, 6.15266707e+03,
               7.06654744e+03, 8.11617014e+03, 9.32169752e+03, 1.07062867e+04,
               1.22965345e+04, 1.41229882e+04, 1.62207325e+04, 1.86300631e+04,
               2.13972612e+04, 2.45754825e+04, 2.82257777e+04, 3.24182659e+04,
               3.72334812e+04, 4.27639199e+04, 4.91158168e+04, 5.64111866e+04,
               6.47901669e+04, 7.44137108e+04, 8.54666784e+04, 9.81613878e+04,
               1.12741693e+05, 1.29487670e+05, 1.48720995e+05, 1.70811122e+05,
               1.96182384e+05, 2.25322143e+05, 2.58790146e+05, 2.97229287e+05,
               3.41377948e+05, 3.92084187e+05, 4.50322027e+05, 5.17210168e+05,
               5.94033474e+05, 6.82267654e+05, 7.83607611e+05, 9.00000000e+05])
```

Similar to what we did when calculating the PDFs, we get the displacements for all au values and samples.

```
In [6]: # initialize array to store displacements
        # note that the size n is larger than the actual number of values to be
        stored
        # values are initialized to NaN (which are not included in the MSD calcu
        lation)
        dx tau = np.empty((n, n tau, n samp))*np.nan
        for i samp in np.arange(n samp):
          for i, tau in enumerate(tau vals):
            delta = delta vals[i]
             # get truncated copy of x, ending in initial data point of the last
        pair
            x_trunc = x_samp[:-1*delta, i samp]
            # get shifted copy of x, starting from end data point of the first p
        air
            x_shift = x_samp[delta:, i_samp]
            # get displacements
            dx = x \text{ shift } - x \text{ trunc}
            # store in output array
            dx tau[:len(dx), i, i samp] = dx
             #print(i, tau, delta, len(dx))
```

```
In [7]: np.shape(dx_tau)
Out[7]: (1000000, 100, 10)
```

Next, we get the mean of the squared displacements for each value of  $\tau$ :

```
In [18]: # initialize array to store MSD values
    msd_tau = np.empty((n_tau, n_samp))*np.nan

for i_samp in np.arange(n_samp):
    for i, tau in enumerate(tau_vals):
        dx2_sum = np.nansum(dx_tau[:, i, i_samp]**2) # returns sum treating
    NaNs as zero
        denom = n-delta_vals[i]
        msd_tau[i, i_samp] = dx2_sum/denom
```

Finally, we plot the MSD as a function of  $\tau$  for the different samples and overlay the analytical result MSD  $(\tau) = \tau$ .

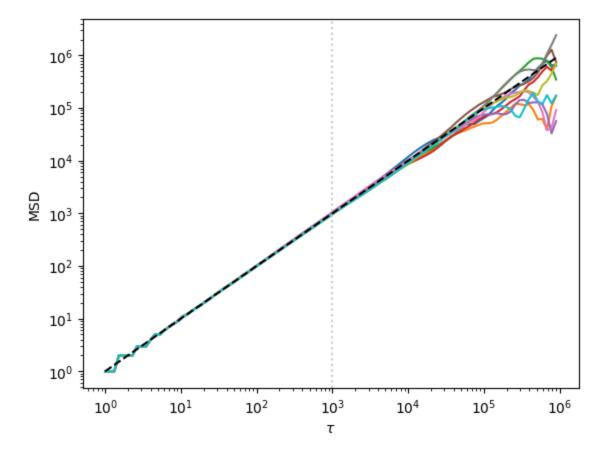
```
In [24]: # plot MSD for all realizations
for i_samp in np.arange(n_samp):
    plt.plot(tau_vals, msd_tau[:, i_samp])

# plot analytical result
    msd_theo = tau_vals
    plt.plot(tau_vals, msd_theo, 'k--')

# plot vertical line at tau=10^3
    tau_ref = 1e3
    plt.axvline(tau_ref, color='0.8', ls=':')

plt.xscale("log")
    plt.yscale("log")
    plt.xlabel(r"$\tau$")
    plt.ylabel("MSD")
```

Out[24]: Text(0, 0.5, 'MSD')



We find deviations from the analytical result for large values of  $\tau$ , as expected from the smaller number of data point pairs used in this regime.

Zooming in on  $\tau < 10^3$ , we find deviations from the analytical result for individual simulations, as expected, but a good match to the average MSD curve (over all realizations).

```
In [31]: # get the average MSD over all realizations
    msd_mean = np.mean(msd_tau, axis=1)
    np.shape(msd_mean)
Out[31]: (100,)
```

```
In [32]: # plot MSD for all realizations
    for i_samp in np.arange(n_samp):
        plt.plot(tau_vals, msd_tau[:, i_samp])

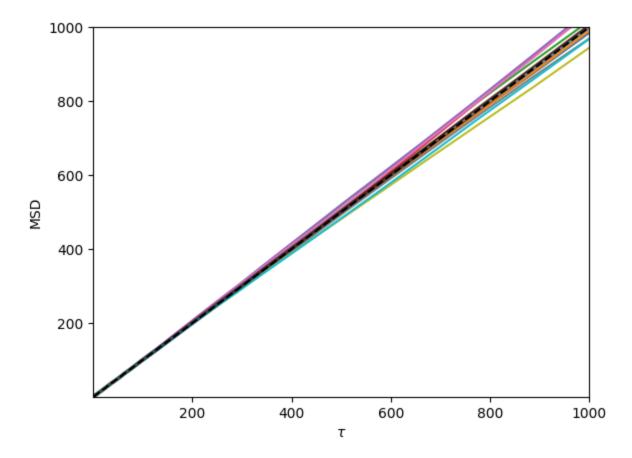
# plot average MSD over all realizations
    plt.plot(tau_vals, msd_mean, color='k', lw=3, alpha=0.5)

# plot analytical result
    msd_theo = tau_vals
    plt.plot(tau_vals, msd_theo, 'k--')

# plt.xscale("log")
    #plt.yscale("log")
    plt.xlabel(""$\tau$")
    plt.ylabel("MSD")

plt.xlim((1, 1e3))
    plt.ylim((1, 1e3))
```

## Out[32]: (1.0, 1000.0)



```
In [42]: # plot residuals from analytical result

# plot MSD for all realizations
for i_samp in np.arange(n_samp):
    plt.plot(tau_vals, msd_tau[:, i_samp]-msd_theo)

# plot average MSD over all realizations
plt.plot(tau_vals, msd_mean-msd_theo, color='k', lw=3, alpha=0.5)

# plot horizontal line at zero
plt.axhline(0., color='k', ls=':')

plt.xlabel(r"$\tau$")
plt.ylabel("MSD")

plt.xlim((1, 1e3))
plt.ylim((-55, 55))
```

## Out[42]: (-55.0, 55.0)

