Problem 1

a.)

Expected Value/Mean:

$$\mu = \int_{-\infty}^{\infty} x f(x) dx$$
$$= \int_{0}^{a} \frac{x}{a} dx$$
$$= \frac{1}{2a} \left[x^{2} \right]_{0}^{a}$$
$$\mu = \frac{a}{2}$$

Variance:

$$\sigma^{2} = E[(X - \mu)^{2}]$$

$$= \int_{-\infty}^{\infty} (x - \mu)^{2} f(x) dx$$

$$= \int_{0}^{a} \left(x - \frac{a}{2}\right)^{2} \left(\frac{1}{a}\right) dx$$

$$= \int_{0}^{a} \left(x^{2} - ax + \frac{a^{2}}{4}\right) \left(\frac{1}{a}\right) dx$$

$$= \int_{0}^{a} \frac{x^{2}}{a} dx - \int_{0}^{a} x dx + \int_{0}^{a} \frac{a}{4} dx$$

$$= \left[\frac{x^{3}}{3a}\right]_{0}^{a} - \left[\frac{x^{2}}{2}\right]_{0}^{a} + \left[\frac{ax}{4}\right]_{0}^{a}$$

$$= \frac{a^{2}}{3} - \frac{a^{2}}{2} + \frac{a^{2}}{4}$$

$$\sigma^{2} = \frac{a^{2}}{12}$$

Cumulative Distribution Function:

$$F(x) = \int_{-\infty}^{x} f(x')dx'$$
$$= \int_{0}^{x} \frac{1}{a} dx'$$
$$= \frac{1}{a} [x']_{0}^{x}$$
$$= \frac{x}{a}$$

$$F(x) = \begin{cases} 0, & x < 0, x > a \\ \frac{x}{a}, & 0 < x < a \end{cases}$$

b.)

Expected Value/Mean:

$$\mu = \int_{-\infty}^{\infty} x f(x) dx$$

$$= \int_{0}^{\infty} \lambda x e^{-\lambda x} dx$$

$$(\text{let } u = \lambda x, du = \lambda dx, dv = e^{-\lambda x}, v = -e^{-\lambda x}/\lambda)$$

$$= uv|_{0}^{\infty} - \int_{0}^{\infty} v du$$

$$= -xe^{-\lambda x} \Big|_{0}^{\infty} - \int_{0}^{\infty} \frac{-\lambda e^{-\lambda x}}{\lambda} dx$$

$$= 0 + \left[\frac{-e^{-\lambda x}}{\lambda}\right]_{0}^{\infty}$$

$$\mu = \frac{1}{\lambda}$$

Variance:

$$\sigma^2 = E[(X - \mu)^2]$$

$$= E[X^2] - \mu^2$$

$$= \int_0^\infty x^2 f(x) dx - \frac{1}{\lambda^2}$$

$$= \int_0^\infty \lambda x^2 e^{-\lambda x} dx - \frac{1}{\lambda^2}$$
(let $u = \lambda x^2$, $du = 2\lambda x dx$, $dv = e^{-\lambda x}$, $v = -e^{-\lambda x}/\lambda$)
$$= uv|_0^\infty - \int_0^\infty v du - \frac{1}{\lambda^2}$$

$$= \left[-x^2 e^{-\lambda x} \right]_0^\infty + \int_0^\infty 2x e^{-\lambda x} dx - \frac{1}{\lambda^2}$$

$$= 0 + 2\int_0^\infty x e^{-\lambda x} dx - \frac{1}{\lambda^2}$$

$$= \frac{2}{\lambda^2} \left[e^{-\lambda x} (-\lambda x - 1) \right]_0^\infty - \frac{1}{\lambda^2}$$

$$= \frac{2}{\lambda^2} - \frac{1}{\lambda^2}$$

$$\sigma^2 = \frac{1}{\lambda^2}$$

Cumulative Distribution Function:

$$F(x) = \int_{-\infty}^{x} f(x')dx'$$

$$= \int_{0}^{x} \lambda e^{-\lambda x'} dx'$$

$$= \left[-e^{-\lambda x'} \right]_{0}^{x}$$

$$= \left[1 - e^{-\lambda x} \right]$$

$$F(x) = 1 - e^{-\lambda x}$$

Problem 2

(See Jupyter notebook, attached)

Problem 3

a.)

True. A normalized PDF is required so that the inverted CDF, from which we draw our sample, has a domain of 0 to 1. Then, selecting random numbers in this same interval will give us values of the PDF/CDF's independent variable distributed like the original PDF.

b.)

Deterministic Strengths

- 1. Fast execution (all steps contribute to a meaningful solution)
- 2. No statistical uncertainties, solutions are exact (within truncation limits)
- 3. Solutions are global over the problem space

Monte Carlo Weaknesses

- 1. Slow/inefficient (calculations for particles that do not contribute to the tally are essentially wasted)
- 2. Results contain statistical uncertainties
- 3. Solutions are only local (wherever tallies are collected)

Monte Carlo Strengths

- 1. Continuous (in space, energy, angle, etc.)
- 2. Easy to parallelize
- 3. No truncation error, solutions are exact (within uncertainties)

Deterministic Weaknesses

- 1. Difficult to parallelize
- 2. Discretization (energy, angle, space, etc.) limits accuracy
- 3. Solutions contain truncation error (expansions are truncated)

c.)

The relative error of an analog Monte Carlo algorithm is $R = S_{\bar{x}}/\bar{x}$. Since $S_{\bar{x}} = S/\sqrt{N}$, it becomes apparent that $R \propto 1/\sqrt{N}$.

d.)

The central limit theorem allows us to state, specifically for independently and identically distributed random variables, that for many samples of N measurements, the sample means will be distributed according to a normal distribution, with variance σ^2/N (where σ^2 is the true variance of the original variable distribution).

e.)

In a Monte Carlo algorithm, the tracking of particles through a geometry region requires the particle's mean free path in that region. At a boundary, the mean free path must be updated for proper transport through the new material.

f.)

The collision estimator is more likely to be accurate in cases where there are many collisions (such as a large and/or dense geometry region), and the majority of particles passing through the region of interest collide there. In the case that the geometry is optically thin, there will be few collisions in the region, and so a collision estimator will yield poor statistics. Then, a track length estimator will be preferable.

g.)

The Consistent Adjoint Driven Importance Sampling method is termed "consistent" because it applies the appropriate weighting (to the nominal weight) for every particle that is born, according to the importance map defined by the adjoint source.

Problem 4

(See Jupyter notebook, attached)

NE250_HW06_mnegus-prob2

December 1, 2017

1 NE 250 – Homework 6

1.1 Problem 2

12/1/2017

```
In [1]: %matplotlib inline
    import numpy as np
    import seaborn as sns
    from matplotlib import pyplot as plt
```

a.)

To sample from a distribution with an exponentially decaying probability density function

$$p(x) = e^{-x},$$

we must first determine the cumulative distribution function (CDF). This is

$$P(x) = \int_0^x e^{-x'} dx'$$
$$P(x) = \left[-e^{-x'} \right]_0^x$$
$$P(x) = 1 - e^{-x}$$

If we take a random sample from this CDF, ξ , then we can invert the function to find a random x corresponding to ξ .

$$\xi = 1 - e^{-x}$$
$$x = -\ln(1 - \xi)$$

Since ξ is chosen uniformly and ranges from 0 to 1, we can say that $1 - \xi$ is a random number with identical distribution to ξ . Then

$$x = -\ln \xi$$

Now, we want to estimate the mean and variance of samples of size 10, 40, and 160.

```
In [3]: for num_samples in [10,40,160,10_000]:
           x,mu,sig = sample_exp_decay(num_samples)
           mu = round(mu, 4)
           sig = round(sig, 4)
           print('{} samples \t Mean: {} \t Variance: {} '.format(num_samples,mu,
10 samples
                    Mean: 1.1785
                                          Variance: 1.1298
40 samples
                   Mean: 1.0652
                                          Variance: 1.2882
160 samples
                                           Variance: 1.2137
                     Mean: 1.1874
10000 samples
                       Mean: 1.0046
                                             Variance: 0.9755
```

We can see that the values for the required 10, 40, and 160 are reasonably close to the values predicted analytically (We can use Problem 1 part b, with $\lambda=1$ to see that the mean and variance of the distribution should both equal 1). Our precision increases considerably with more histories, and with 10,000 histories, we are significantly more accurate on both mean and variance.

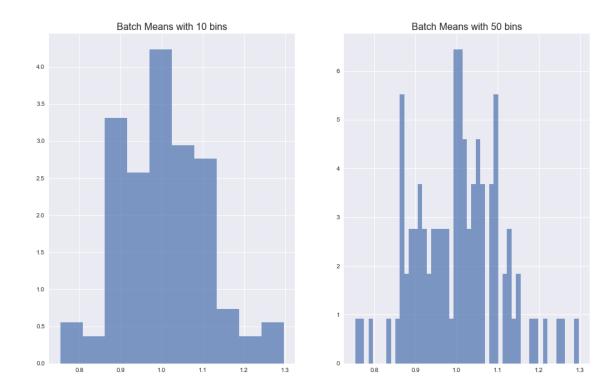
b.)

Next, we will run several batches of runs, each with a given number of histories. To show that the batch average tends to follow the central limit theorem when several batches are considered, we will plot the averages as a histogram

```
In [4]: def batch_sampling(num_batches, num_samples):
            '''Run a given number of batches, each with the given number of samples
            batch_means = [sample_exp_decay(num_samples)[1] for i in range(num_batch_means)
            return batch means
In [5]: def plot_hists(data,title,overlay=False,var=1,n=100):
            '''Plot a histogram with from a given dataset.'''
            binnings = 2
            fig, ax = plt.subplots(ncols=binnings, figsize=(16,10))
            bin_count = [10+40*i for i in range(binnings)]
            data=np.array(data)
            for b in range(binnings):
                ax[b].hist(data,bins=bin_count[b],normed=True,alpha=0.7)
                ax[b].set_title('{} with {} bins'.format(title,bin_count[b]),fonts:
                # Add Gaussian overlay (with mean 1 and variance 1)
                if overlay:
                    x = np.arange(0.5, 1.5, 0.01)
                    g = np.exp(-(x-1)**2/(2*var/n))/np.sqrt(2*np.pi*var/n)
                    ax[b].plot(x,q,'k')
            return ax
```

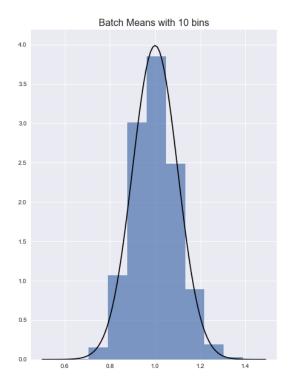
We will first run 100 batches of the sampling algorithm, each batch with 100 histories.

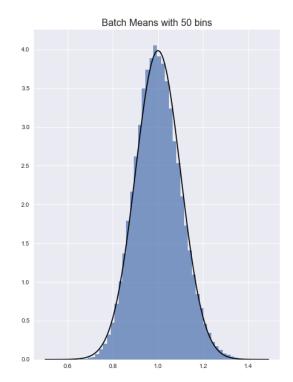
```
In [6]: batch_means = batch_sampling(100,100)
    mean_plots = plot_hists(batch_means, 'Batch Means')
```



The Gaussian behavior is fairly clear in these graphics, but we can make it more apparent if we increase the number of batches. Let's try 100,000 batches instead, but each with still 100 histories. We can overlay the predicted Gaussian distribution (with mean $\mu=1$ and variance $\frac{\sigma^2}{n}=0.01$) given by the central limit theorem on the plots.

```
In [7]: batch_means = batch_sampling(100_000,100)
    mean_plots = plot_hists(batch_means,'Batch Means',overlay=True)
```





NE250_HW06_mnegus-prob4

December 2, 2017

1 NE 250 – Homework 6

1.1 Problem 4

```
12/1/2017
```

a.)

```
In [1]: import numpy as np
    import random
```

We are considering an infinite, steady-state, monoenergetic, two-region Monte Carlo problem with the following characteristics: (note thate we have renamed region 1 and region 2 as region 0 and region 1 respectively; this allows for simpler calculations) * 1-D problem geometry * Region 0 has $\Sigma_s = 0.5 \, \mathrm{cm}^{-1}$ and $\Sigma_t = 1.0 \, \mathrm{cm}^{-1}$ (in both regions the only interactions are scattering or absorption). * Region 1 has $\Sigma_s = 0.75 \, \mathrm{cm}^{-1}$ and $\Sigma_t = 0.9 \, \mathrm{cm}^{-1}$. * Region 0 has $w_{nom} = 1$ and Region 1 has $w_{nom} = 2$. w_{max} and w_{min} can be found as $(w_{nom} \times 2.5)$ or $(w_{nom} / 2.5)$, respectively. * All particles are born in Region 0 with weight 1 at a location that is 1 cm to the left of the interface between Regions 0 and 1. The source is isotropic. * Isotropic scattering.

Problem Statement Write the algorithm for a Monte Carlo code to solve this specific problem. Include the PDFs required for sampling as well as algorithms for conducting sampling. Use a collision estimator to tally the flux. Include implicit capture, rouletting, and splitting.

1.1.1 Underlying parameters

Using the above information, we can create a set of dictionaries to describe our physical situation. Each fundamental parameter gets a dictionary, and each dictionary has entries corresponding to each region.

```
# Max/Min Weights
    w_ext = {region: (w_nom[region] *2.5, w_nom[region]/2.5) for region in w_nom.}
In [3]: w_ext
Out[3]: {0: (2.5, 0.4), 1: (5.0, 0.8)}
```

1.1.2 Tracking the particle

We can track a particle using a particle class, which can model the behavior of each particle as it proceeds through it's lifetime. This particle class will contain methods for each action that the particle will undergo * birth (the class' _init__ method) * transport * boundary encounter * collision * scoring

1.2 Survival Biasing

```
In [4]: class particle:
            11 11 11
            A class to model a single Monte Carlo particle over it's lifetime
            def __init__ (self, verbose=False):
                """The particle is born. Assign a position [cm], angle (cos \theta), end
                self.x = -1
                self.mu = 2 * random.random() - 1
                self.E = 1
                self.w = 1
                self.region = 0
                self.score = np.zeros(2)
                self.verbose = verbose
                if verbose:
                    print('The particle was born at x = -1 cm (region 0), with weight
            def transport(self, sample=True):
                """Transport the particle through the problem geometry"""
                # Sample to find the number of mean free paths that traveled by the
                if sample:
                    xi = random.random()
                    self.mfp_x = -np.log(xi)*self.mu
                    if self.verbose: print('This particle will travel {} MFPs in the
                # Determine the number of mean free paths to a boundary in the curr
                boundary_mfp = -self.x*Sigma_t[self.region]
                if self.verbose: print('The distance to the boundary is {} MFPs.'.:
                # If the particle reaches a boundary before the collision, stop and
                if boundary_mfp == 0: # (the particle was at the boundary, so must
                    if self.verbose: print('(the particle is at the boundary)')
                    self.collision()
                elif self.mfp_x > boundary_mfp:
```

self.boundary(boundary_mfp)

```
else:
        self.collision()
def boundary(self,boundary_mfp):
    """The particle reached a boundary: reevaluate the particle's track
    if self.verbose: print('The particle travels to the boundary.')
    self.region = 1-self.region
    self.mfp_x -= boundary_mfp
    self.update_weight(collision=False)
    self.transport(sample=False)
def collision(self):
    """The particle collided: use survival biasing to continue following
    self.x += self.mfp_x
    if self.verbose: print('The particle collides at x = \{\}'.format(rown)
    self.score_particle()
    self.update_weight(collision=True)
    if self.w == 0:
        if self.verbose: print('\t\t\t\t The particle is killed (w=0).
    self.mu = 2 * random.random() - 1
    if self.verbose: print('The particle scatters, and is now traveling
    self.transport(sample=True)
def update_weight(self,collision=False):
    """Update the particle's weight using survival biasing, splitting,
    if collision: # adjust weight with survival biasing
                          w_{\{i+1\}} = w(1 - \Sigma_a / \Sigma_t)
        # New weight:
        self.w *= (Sigma_t[self.region]-Sigma_s[self.region])/Sigma_t[self.region]
        if self.verbose: print('The particle is survival biased, now was
    # Splitting
    if self.w > w_ext[self.region][0]:
        SR = self.w/w_nom[self.region]
        xi = random.random()
        if xi >= SR - int(SR):
            b = 0
        else:
            b = 1
        n_new_particles = int(SR) + b
        if self.verbose: print('Splitting the particle into {} new part
        global particles
        for i in range(int(SR)):
            particles.append(particle())
            particles[-1].x = self.x
            particles[-1.].w = self.w/n_new_particles
    # Russian Roulette
    if self.w < w_ext[self.region][1]:</pre>
```

```
xi = random.random()
                    RR = self.w/w_nom[self.region]
                    if xi >= RR :
                        self.w = 0
                        if self.verbose: print('\t\t\t ... uh-oh...')
                        self.w = w_nom[self.region]
                        if self.verbose: print('Phew. Particle survived, and now it
            def score_particle(self):
                self.score[self.region] += self.w
                if self.verbose:
                    print('Score! Particle with weight {} added to the tally.'.form
                    print('\tCurrent score: \t Region 0: {} Region 1: {}'.format
In [5]: ## Survival biasing, rouletting, and splitting MC run
       N = 20000
       tally = np.zeros(2)
       particles = [particle() for n in range(N)]
        for p in particles:
           p.transport()
           tally += p.score
       norm_collisions = tally/N
In [6]: print('Average collisions per particle, region 0: {} \t Average collisions
       print('Ratio: ', norm_collisions[0]/norm_collisions[1])
Average collisions per particle, region 0: 1.7315
                                                           Average collisions per p
Ratio: 10.2698695136
```

if self.verbose: print('Rouletting the particles...')

Interestingly, it appears that in our specific problem splitting is never used. This fact could be deduced, however, by noting that splitting only occurs when $w_i > w_{max}$. For both regions, $w_{max} \geq 2.5$. Upon birth in either region, a particle's weight will never be more than 2. Collisions will only trigger a reduction in weight, and rouletting produces particles also with a maximum weight of 2 (roulette products have weight w_{nom}).

1.3 The story of a Monte Carlo Particle

b.)

Let's follow just 1 particle.

```
In [7]: random.seed(1)
        N = 1
        tally = np.zeros(2)
        p = particle(verbose=True)
```

```
p.transport()
        tally += p.score
The particle was born at x = -1 cm (region 0), with weight 1.
This particle will travel -0.121 MFPs in the x-direction.
The distance to the boundary is 1.0 MFPs.
The particle collides at x = -1.121
Score! Particle with weight 1 added to the tally.
                                Region 0: 1.0
        Current score:
                                                Region 1: 0.0
The particle is survival biased, now with weight 0.5
The particle scatters, and is now traveling with mu = 0.528
This particle will travel 0.721 MFPs in the x-direction.
The distance to the boundary is 1.121 MFPs.
The particle collides at x = -0.4
Score! Particle with weight 0.5 added to the tally.
        Current score:
                               Region 0: 1.5
                                                Region 1: 0.0
The particle is survival biased, now with weight 0.25
Rouletting the particles...
                         ... uh-oh...
                                 The particle is killed (w=0).
```

Random seed 4 gives a much more dynamic plot...

```
In [8]: random.seed(4)
       N = 1
        tally = np.zeros(2)
        p = particle(verbose=True)
       p.transport()
        tally += p.score
The particle was born at x = -1 cm (region 0), with weight 1.
This particle will travel -1.199 MFPs in the x-direction.
The distance to the boundary is 1.0 MFPs.
The particle collides at x = -2.199
Score! Particle with weight 1 added to the tally.
        Current score:
                                Region 0: 1.0
                                                Region 1: 0.0
The particle is survival biased, now with weight 0.5
The particle scatters, and is now traveling with mu = -0.208
This particle will travel -0.388 MFPs in the x-direction.
The distance to the boundary is 2.199 MFPs.
The particle collides at x = -2.587
Score! Particle with weight 0.5 added to the tally.
        Current score:
                                Region 0: 1.5
                                                Region 1: 0.0
The particle is survival biased, now with weight 0.25
Rouletting the particles...
Phew. Particle survived, and now it has weight 1
The particle scatters, and is now traveling with mu = -0.197
This particle will travel -0.017 MFPs in the x-direction.
```

The distance to the boundary is 2.587 MFPs.

The particle collides at x = -2.604

Score! Particle with weight 1 added to the tally.

Current score: Region 0: 2.5 Region 1: 0.0

The particle is survival biased, now with weight 0.5

The particle scatters, and is now traveling with mu = 0.601

This particle will travel 0.161 MFPs in the x-direction.

The distance to the boundary is 2.604 MFPs.

The particle collides at x = -2.443

Score! Particle with weight 0.5 added to the tally.

Current score: Region 0: 3.0 Region 1: 0.0

The particle is survival biased, now with weight 0.25

Rouletting the particles...

Phew. Particle survived, and now it has weight 1

The particle scatters, and is now traveling with mu = 0.073

This particle will travel 0.094 MFPs in the x-direction.

The distance to the boundary is 2.443 MFPs.

The particle collides at x = -2.348

Score! Particle with weight 1 added to the tally.

Current score: Region 0: 4.0 Region 1: 0.0

The particle is survival biased, now with weight 0.5

The particle scatters, and is now traveling with mu = -0.655

This particle will travel -1.468 MFPs in the x-direction.

The distance to the boundary is 2.348 MFPs.

The particle collides at x = -3.817

Score! Particle with weight 0.5 added to the tally.

Current score: Region 0: 4.5 Region 1: 0.0

The particle is survival biased, now with weight 0.25

Rouletting the particles...

Phew. Particle survived, and now it has weight 1

The particle scatters, and is now traveling with mu = 0.855

This particle will travel 0.16 MFPs in the x-direction.

The distance to the boundary is 3.817 MFPs.

The particle collides at x = -3.656

Score! Particle with weight 1 added to the tally.

Current score: Region 0: 5.5 Region 1: 0.0

The particle is survival biased, now with weight 0.5

The particle scatters, and is now traveling with mu = 0.613

This particle will travel 0.137 MFPs in the x-direction.

The distance to the boundary is 3.656 MFPs.

The particle collides at x = -3.52

Score! Particle with weight 0.5 added to the tally.

Current score: Region 0: 6.0 Region 1: 0.0

The particle is survival biased, now with weight 0.25

Rouletting the particles...

Phew. Particle survived, and now it has weight 1

The particle scatters, and is now traveling with mu = -0.38

This particle will travel -0.178 MFPs in the x-direction.

The distance to the boundary is 3.52 MFPs.

The particle collides at x = -3.697

Score! Particle with weight 1 added to the tally.

Current score: Region 0: 7.0 Region 1: 0.0

The particle is survival biased, now with weight 0.5

The particle scatters, and is now traveling with mu = 0.464

This particle will travel 0.073 MFPs in the x-direction.

The distance to the boundary is 3.697 MFPs.

The particle collides at x = -3.624

Score! Particle with weight 0.5 added to the tally.

Current score: Region 0: 7.5 Region 1: 0.0

The particle is survival biased, now with weight 0.25 Rouletting the particles...

... uh-oh...

The particle is killed (w=0).