

NUR - Hand-in Exercise 2

Martina Cacciola (s4170814)

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Abstract

In this document, I present the results for the Hand-in Exercise 2 for the course Numerical Recipes for Astrophysics.

1 Satellite galaxies around a massive central

The exercise is done in the script

```
1 import time
2 import numpy as np
3 from math import *
4 import matplotlib.pyplot as plt
5
6 '''
7 This script contains the code for Exercise 1.
8 '''
9
10 class CombinedRNG:
11     '''
12     Pseudo-random number generator combining the XOR-shift and Multiply-With-Carry (MWC)
13     methods.
14     We set the seed to the current time in milliseconds, as a source of variability
15     '''
16     def __init__(self, seed = None):
17         # If no seed is provided, use the current time in milliseconds
18         if seed is None:
19             seed = int(time.time() * 1000)
20         # Initialize the states for XOR-shift and MWC methods
21         self.xor_state = seed
22         self.mwc_state = seed
23
24     def xor_shift(self):
25         # XOR-shift method
26         # The seed is converted to a 64-bit unsigned integer
27         # By performing a shift of 21 bits to the left and XORing the result with the
28         # original state
29         self.xor_state ^= (self.xor_state << 21) & 0xFFFFFFFFFFFFFFFF
30         # Then, a shift of 35 bits to the right and again XOR with the previous result
31         self.xor_state ^= (self.xor_state >> 35)
32         # Finally, a shift of 4 bits to the left and XOR with the previous result
33         self.xor_state ^= (self.xor_state << 4) & 0xFFFFFFFFFFFFFFFF
34         return self.xor_state
35
36     def mwc(self):
37         # Multiply-With-Carry (MWC) method
38         # Set the multiplier
39         a = 4294957665
40         # Extract the upper and lower 32 bits of the state
41         mwc_upper = (self.mwc_state & 0xFFFFFFFF00000000) >> 32
42         mwc_lower = self.mwc_state & 0x00000000FFFFFFFF
43         # Calculate the next state by multiplying the lower 32 bits with the multiplier
44         # and adding the upper 32 bits
45         x_next = a * mwc_lower + mwc_upper
46         # Update the state using AND operation with a 64-bit mask
```

```

45     # Only the lower 64 bits are kept, the rest are set to zero
46     self.mwc_state = x_next & 0xFFFFFFFFFFFFFFFF
47     return x_next >> 32
48
49     def combined_rng(self):
50         # Combine the XOR-shift and MWC methods by XORing their outputs
51         return (self.xor_shift() ^ self.mwc()) & 0xFFFFFFFFFFFFFFFF
52
53     def uniform(self, low, high, num_samples):
54         # Generate a list of uniformly distributed random numbers
55         # By scaling the output of the combined RNG to the desired range
56         return [low + (high - low) * self.combined_rng() / 0xFFFFFFFFFFFFFFFF for _ in
57                 range(int(num_samples))]
58
59 # Create an instance of the combined RNG
60 rng = CombinedRNG()
61
62 ## 1a)
63
64 # Define the parameters
65 a = 2.4
66 b = 0.25
67 c = 1.6
68 xmax = 5
69 Nsat = 100
70
71 ## 1a)
72
73 def integrand(x):
74     # Take into account singularity at x=0
75     # Consider the volume element is 4*pi*x^2 (4*pi is taken outside)
76     if x == 0:
77         return 0
78     return Nsat * (x/b)**(a-3) * exp(-(x/b)**c) * x**2
79
80 # Implement a numerical integration
81 # Using the trapezoidal rule
82 def trapezoidal_rule(f, a, b, n):
83     '''
84     Inputs:
85     f: Function to integrate
86     a, b: Integration limits
87     n: Number of intervals
88     It works by approximating the integral of f(x) between a and b
89     by the sum of the areas of trapezoids formed by the function and the x-axis
90     '''
91     # Width of each trapezoid
92     h = (b - a) / n
93     # Initialize the result with the average of the function at the limits
94     result = 0.5 * (f(a) + f(b))
95     for i in range(1, n):
96         # Add the value of the function at each interior point
97         # Each of the function values corresponds to the height of a trapezoid
98         result += f(a + i * h)
99     # Multiply the sum by the width of the trapezoids
100    # This gives the total area under the curve
101    result *= h
102    return result
103
104
105 # Romberg integration
106 def romberg(f, a, b, n):
107     '''
108     Inputs:
109     f: Function to integrate
110     a, b: Integration limits
111     n: Number of rows in the Romberg table
112     '''
113     R = np.zeros((n, n))

```

```

114 # At each iteration , calculates an approximation of the integral
115 # using the trapezoidal rule with 2 ** i intervals
116 # This forms the first column of the Romberg table
117 for i in range(n):
118     R[i, 0] = trapezoidal-rule(f, a, b, 2 ** i)
119 # At each iteration , calculates an improved approximation of the integral
120 # using Richardson extrapolation
121 # Takes a weighted average of the current approximation and the previous one
122 # The weights are chosen to cancel out as much of the error as possible
123 # This fills the rest of the Romberg table
124 for j in range(1, n):
125     for k in range(j, n):
126         R[k, j] = R[k, j - 1] + (R[k, j - 1] - R[k - 1, j - 1]) / (4 ** j - 1)
127 return R[-1, -1] # Return the last element of the last row of the Romberg table
128
129
130 # Apply the method above to the integral
131 n = 10
132 m = 6
133 R = romberg(integrand, 0, xmax, m)
134 result = R * 4 * pi
135 A = Nsat / result
136 with open("normalization.txt", "w") as file:
137     file.write("The result of the numerical integration is: " + str(A))
138
139
140 ## 1b)
141
142 # Define the number density of galaxies
143 # This represents the number of galaxies per unit volume at a given radius x
144 def n(x):
145     return A * Nsat * (x/b)**(a-3) * exp(-(x/b)**c)
146
147 # Define the number of galaxies N(x) in a shell of radius x and thickness dx
148 # We compute N(x) by multiplying the number density n(x) by the volume element 4*pi*x^2
149 def N(x):
150     return 4 * np.pi * x**2 * n(x)
151
152 # Define the probability distribution function p(x) given that p(x)dx = N(x)dx / Nsat
153 # It gives the probability of finding a galaxy at a given radius x
154 def p_x(x):
155     return N(x) / Nsat
156
157
158 # Implement inverse transform sampling to sample from the distribution
159 def inverse_transform_sampling(pdf, n_samples, x_min, x_max):
160     x_values = np.linspace(x_min, x_max, 10000)
161     cdf_values = np.zeros_like(x_values)
162     # Calculate the cumulative distribution function using numerical integration (
163     # trapezoidal rule)
164     # The cdf at a point x is the integral of the pdf from x_min to x
165     for i in range(1, len(x_values)):
166         cdf_values[i] = cdf_values[i-1] + trapezoidal-rule(pdf, x_values[i-1], x_values[
167         i], 1)
168     cdf_values /= cdf_values[-1] # Normalize to ensure CDF ranges from 0 to 1
169
170 # Generate random numbers uniformly distributed between 0 and 1
171 random_numbers = rng.uniform(0, 1, num_samples=n_samples)
172
173 # Apply the inverse CDF to the random numbers
174 # For each random number, find the corresponding value of x
175 # such that the CDF of x is equal to the random number
176 # This is done by finding the first x value for which the CDF is greater than the
177 # random number
178 # The corresponding x value is then stored as a sampled point
179 sampled_points = np.zeros(n_samples)
180 for i in range(n_samples):
181     for j in range(len(cdf_values)):
182         if random_numbers[i] < cdf_values[j]:
183             sampled_points[i] = x_values[j]

```

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181         break
182
183     return sampled_points
184
185 # Generate 10,000 sampled points
186 n_samples = 10000
187 x_min, x_max = 10**-4, 5
188 h = (x_max - x_min) / n_samples
189 sampled_points = inverse_transform_sampling(p_x, n_samples, x_min, x_max)
190
191 # Calculate the bin edges
192 bin_edges = np.logspace(np.log10(x_min), np.log10(x_max), 21)
193
194 # Calculate histogram and divide each bin by its width
195 hist, _ = np.histogram(sampled_points, bins=bin_edges, density=False)
196 bin_widths = np.diff(bin_edges)
197 hist_density = hist / (bin_widths * n_samples / Nsat) # Correcting for the
198               normalization offset
199
200 # Plot the analytical function N(x) and the histogram of the sampled points on a log-log
201       scale
202 x_values = np.logspace(-4, np.log10(x_max), 20)
203 N_x_values = [N(x) for x in x_values]
204
205 plt.figure(figsize=(10, 6))
206 plt.loglog(x_values, N_x_values, label='Analytical N(x)', color='r')
207 plt.bar(bin_edges[:-1], hist_density, width=bin_widths, label='Sampled Points Histogram'
208         )
209 plt.xlabel('Relative Radius x')
210 plt.ylabel('Number of Galaxies N(x)')
211 plt.ylim(10**-3, 10**5)
212 plt.legend()
213 plt.title('Analytical N(x) vs Sampled Points Histogram')
214 plt.grid(True)
215 plt.savefig('my_solution_1b.png', dpi=600)
216
217 ## 1c)
218
219 # Define a selection method following these rules:
220 # Select each galaxy with same probability
221 # Not draw the same galaxy twice
222 # Not reject any drawn galaxy
223 def reservoir_selection(sampled_points, n_galaxies):
224     selected_galaxies = []
225     num_samples = len(sampled_points)
226     for i in range(num_samples):
227         # If we haven't selected n_galaxies yet, add the current galaxy
228         if i < n_galaxies:
229             selected_galaxies.append(sampled_points[i])
230         else:
231             # If we have already selected n_galaxies, decide whether to replace one of
232             them
233             # Generate a random index j between 0 and i
234             j = rng.combined_rng() % (i + 1)
235             # If j is less than n_galaxies, replace the galaxy at index j with the
236             current galaxy
237             if j < n_galaxies:
238                 selected_galaxies[j] = sampled_points[i]
239     return selected_galaxies
240
241 # Define a sorting method (quicksort)
242 def quick_sort(arr):
243     # Base case: if the array is empty or has only one element, it is already sorted
244     if len(arr) <= 1:
245         return arr
246     # Choose the pivot element as the middle element of the array
247     pivot = arr[len(arr) // 2]
248     # Partition the array into three parts:
249     # elements less, equal and greater than the pivot

```

```

246     left = [x for x in arr if x < pivot]
247     middle = [x for x in arr if x == pivot]
248     right = [x for x in arr if x > pivot]
249     # Recursively sort the left and right parts
250     # then concatenate the three parts to produce the final sorted array
251     return quick_sort(left) + middle + quick_sort(right)
252
253 # Select 100 galaxies
254 n_galaxies = 100
255 selected_galaxies = reservoir_selection(sampled_points, n_galaxies)
256
257 # Sort the galaxies using quicksort from smallest to higher radius
258 sorted_galaxies = quick_sort(selected_galaxies)
259
260 # Plot the cumulative number of the chosen galaxies
261 fig1c, ax = plt.subplots()
262 ax.plot(sorted_galaxies, np.arange(n_galaxies))
263 ax.set(xscale='log', xlabel='Relative radius',
264        ylabel='Cumulative number of galaxies',
265        xlim=(x_min, x_max), ylim=(0, n_galaxies))
266 plt.savefig('my_solution_1c.png', dpi=600)
267
268 ## 1d)
269
270 # Define Ridders' method for numerical differentiation
271 def ridders_method(f, x, h, m, tol=1e-10):
272     # Compute first approx with central difference for high h
273     D = [(f(x + h) - f(x - h)) / (2 * h)]
274
275     # Decrease h by a factor of 2 and calculate a new approximation. Repeat until you
276     # have m approximations
277     for i in range(1, m):
278         h /= 2
279         D.append((f(x + h) - f(x - h)) / (2 * h))
280         for j in range(i):
281             D[j] = (4**(j+1) * D[j+1] - D[j]) / (4**(j+1) - 1)
282
283     # Terminate when the improvement over previous best approximation is smaller than
284     # the target error or if error grows
285     best_approximation = None
286     for i in range(1, len(D)):
287         if abs(D[i] - D[i-1]) < tol:
288             best_approximation = D[i]
289             break
290
291     # Terminate early if the error grows and return the best approximation from
292     # before that point.
293     if abs(D[i] - D[i-1]) > abs(D[i-1]):
294         best_approximation = D[i-1]
295         break
296
297     if not best_approximation:
298         print("Tolerance not reached or error grew significantly. Consider increasing m.")
299
300     return best_approximation
301
302 # Calculate the analytical result using derivative function
303 #Following the formal definition of the derivative
304 def derivative_n(f, x, h= 1e-10):
305     return (f(x + h) - f(x)) / h
306
307 # Calculate the numerical result using central difference method
308 numerical_result = ridders_method(n, 1, 0.1, 15, tol=1e-10)
309 analytical_result = derivative_n(n, 1, h=1e-10)
310
311 # Output the results
312 with open('derivative.txt', 'w') as f:
313     f.write("Analytical result: " + format(analytical_result, '.12f') + "\n")

```

312

```
f.write("Numerical result: " + format(numerical_result, '.12f') + "\n")
```

```
sat.py
```

. The necessary explanations of the methods used are in the comments of the code. For question (a), the normalization factor A we obtain is the following:

```
1 The result of the numerical integration is: 9.05216643171278
```

```
normalization.txt
```

. For question (b), we generate 3D satellite positions that statistically follow the satellite profile $n(x)$. We sample them from the probability density function $p(x)$, using the inverse transform sampling. We compare the distribution of the sampled points with $N(x)$, the analytical function describing the number of galaxies in a shell of thickness dx at a given distance x . From Fig.1, we can see that the sampled points match the expected distribution. Nevertheless, we can see how the samples are not present in the first interval of x values. Inverse sampling works by transforming uniform random numbers into samples from the desired distribution, using the cumulative distribution function (CDF) of the PDF. The CDF maps values from the sample space to the interval $[0, 1]$, and its inverse can be used to map values from $[0, 1]$ back to the sample space. In the case of smaller values, a small range of uniform random numbers can map to a large range of small values in the sample space, leading to undersampling in these regions. This might explain why there is a discrepancy at smaller values in our histogram.

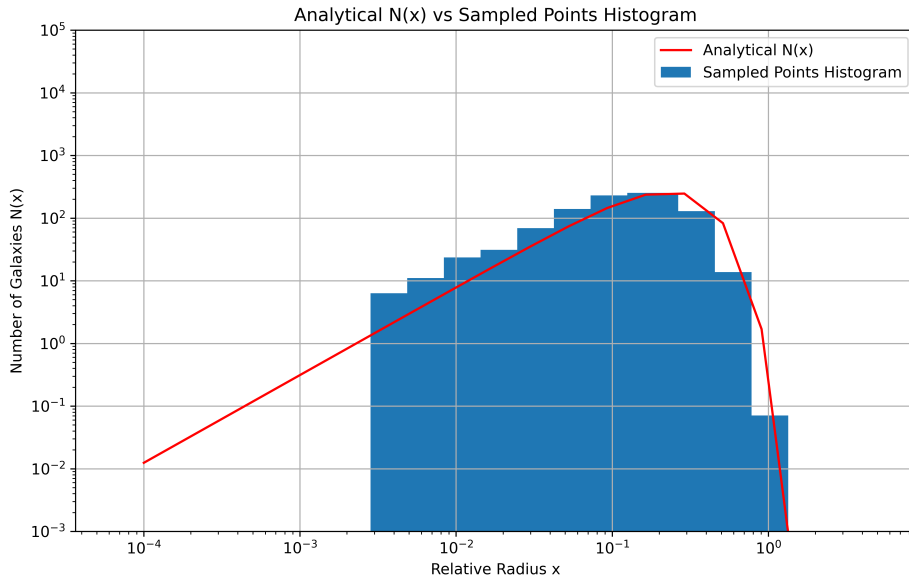


Figure 1: Plot in logarithmic scale showing $N(x)$, function of the number of galaxies at given distance x , and the histogram of the 10000 sampled points. The sampled points match the analytical distribution, but there is a problem of undersampling in the region of smaller radii.

For question (c), we select 100 random galaxies from the ones sampled in point (b), using Reservoir method. It is a sampling algorithm that chooses a random sample, without replacement, of k items from a population of unknown size n in a single pass over the items. The 100 drawn galaxies are ordered from smallest to largest radius. The number of galaxies within a radius x are shown in Fig. 2. The radii selected are starting from higher values because of the previous undersampling. The increasing trend makes sense as the number of galaxies within a given radius should increase as the radius increases.

For question (d), we compute the derivative of the function $n(x)$ using both the Ridders' method and the analytical procedure. The obtained results can be seen in:

```
1 Analytical result: -0.615624912159
```

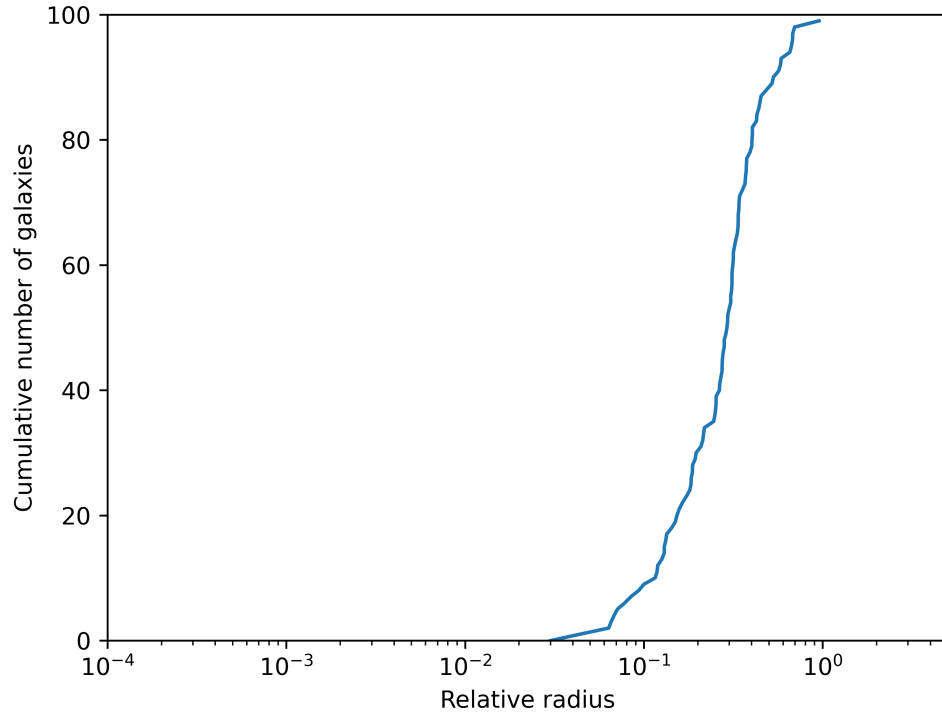


Figure 2: Plot showing the cumulative number of galaxies randomly selected with respect to the relative radius x . The number of galaxies progressively increases with larger distances.

² Numerical result : -0.615624525721

derivative.txt

2 Heating and cooling in HII regions