University of Hull

Report to Analyse the Simulation of Random Growth of Cells

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Programming Assignment

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**Introduction**

Computational science is the discipline within computer science that advances knowledge in various scientific fields through complex computational analysis, modelling, simulation, and numerical approximation (Plonus, 2007). As a result, it can provide innovative solutions to problems by using powerful computers to replace or complement traditional modelling and experimental methods.

This study aims to understand the different complexities that encapsulate growth simulations and how they can be applied to real-world scenarios. We shall build upon various random grid walk models to understand random movement, which can be further applied to random cell growth, simulating a disease such as a cancer tumour. In addition, the Gompertz model will support this simulation by applying more realistic characteristics, such as time, cell capacity, and growth rate. Combining these methods shall form a basic simulation of how a cancer tumour can start to grow and multiply over time in a living organism. Thus, aims to provide more insight into cell growth that can be applied to real-world situations.

**Task 1.1**

The first steps to create the simulation include creating a random grid walk function. The goal for this task is for a cell to start in the middle of a one hundred by one hundred grid and move randomly: left, right, up and down one grid place for one hundred steps.

This task relies on two random number generators and a simple binary algorithm to decide at random whether the cell should move in any of the four directions. This algorithm can be looped one hundred times to simulate the movement for one hundred steps.

However, one issue with this algorithm is the implementation of two random number generators. As a computer cannot be biologically truly random, we must implement pseudo-random number generators to simulate randomness. A pseudo-random number generator is an algorithm that ‘produces a deterministic stream of seemingly random numbers’ (Reeds & Walkenhorst, 2011). Using this function, a distribution type must be applied to determine the range and type of outputs to expect from the function. The two main types considered for this project include: uniform and normal distribution.

**A comparison of a graph

Description automatically generated with medium confidence**‘Normal distribution is a probability distribution that is symmetric about the mean, showing that data near the mean are more frequent in occurrence than data from the mean’ (Chen, 2024). For example, if a normal distribution function had a mean of 0 and a standard deviation of 1, most results would be in the range -1 and 1, primarily around 0, and values out of this range would rarely be selected in comparison. Resulting in a ‘bell curve’ when graphed, as shown in Figure 1. Normal distributions can be used in several circumstances. Often, normal distribution can be found to occur naturally. ‘For example, the average height of a human is roughly 5’ 9”’(Chen, 2024). Taller and shorter people exist but decrease in frequency within the population. Thus, normal distribution would be helpful in simulations relating to natural trends such as population height.

**Figure 1: Normal Distribution Graph**

**References**

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