Cancer as a system dysfunction

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Extended abstract

In this paper, we describe a system dynamics model that views cancer as a dysfunction of the cellular system analogous to those of human societies. Our experiments with the model replicate the propagation of the ailment and the impacts of the treatments. It also represents work in progress and should be viewed as a proposition that can be further developed to understand cancer and used to create appropriate combinations of interventions for specific situations. Our model represents the interaction between the various types of cells in an organ or a subsystem of the body. It depicts the common structure in the simplest possible terms. It defines a generic system that can be applied to solid as well as blood and bone-related diseases, although we mainly address it to proliferation of sold cancers and their response to treatments in this paper.

Whether viewed in the context of an afflicted organ or a body subsystem, the interacting cell populations can be placed in four broad categories: normal cells, pre-cancer cells, cancer cells and immune system cells. Thus, our model contains four stocks connected by flows shown in Figure 1. Normal cells and Immune system cells have disciplined growth regimes in that they tend to grow to their indicated levels – the former goal is determined by the internal intelligence of the cell, while the later by the surveillance need created by the very existence of the unwanted cells in the body. The cancer cell stock is initially populated by a transformation of precancerous cells meant to reflect the accumulation of tumor-promoting mutations that increase with age. Once transformed, the cancerous cells exhibit unregulated proliferation and thus cancer grows exponentially. The immune response increases to combat the rising cancer cell population, and it is ultimately the imbalance of tumor growth vs immune response that yields a proliferation of cancerous cells.

Model behavior

The model is supplied with an internally consistent set of parameters and initial conditions so under normal conditions, all cell populations except cancer are maintained in a dynamic equilibrium or homeostasis. Cancer population is initialized at zero value. The simulation time is set at 1000 months (83.3 years) that approximates the average life expectancy of a healthy individual. It should be noted that despite the equilibrium, all cell populations constantly turn over rather than remaining constant. The in- and out- flows tied to each stock continue while the stocks remain in a dynamic balance.

When initially populated by the transformation of pre-cancer cells, the cancer cells begin to grow exponentially. Low and intermediate rates of growth of cancer are contained however by a concomitant increase in the immune activity that constantly kills the cancer cells and keeps their population under control. The absolute number of cancer cells is influenced by cancer cell proliferation, as well as the rate at which normal cells acquire pre-cancerous characteristics and

then are ultimately transformed into cancer cells. To model rates of proliferation that may differ between aggressively growing and slow growing tumors, the model allows for modulation of cancer cell proliferation rates.



Figure 1 Key cell population stocks, their connecting flows and rules of conduct in the cellular society.

Conclusion

We have made a preliminary attempt in this paper to model the development of cancer as an interaction between normal, immune system, pre-cancer and cancer cell populations. The model is used to test hypotheses about lifetime risk of cancer and the performance of Cancer treatments, and shows credible results. It is also simulated for experimental treatment options, which reveals interesting contingencies. We recognize the importance of the inclusive nourishment commons of the body embodied in its blood supply and propose extending the model to include this for future work.

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