



# CALIFORNIA STATE SCIENCE FAIR 2013 PROJECT SUMMARY

<b>Name(s)</b> <b>Andrew Q. Ninh</b>	<b>Project Number</b> <b>S1209</b>
<b>Project Title</b> <b>Two Discrete Stochastic Cellular Automata Models of Cancer Stem Cell Proliferation</b>	
<p style="text-align: center;"><b>Abstract</b></p> <p><b>Objectives/Goals</b> The objective was to try to create a general model (using set parameters) of cancer stem cell (CSC) induced tumor growth by combining discrete mathematical models, automata theory, and principles of cellular automaton to create a Java program. This program would in turn produce both custom mathematical models as well as growth visualizations.</p> <p><b>Methods/Materials</b> The mass-action and spatial discrete mathematical models and CSC automata theory were turned into a Java program (on the BlueJ IDE) which models CSC growth, which was graphed on Mathematica. Visual depictions of the automata arrays of the mass-action and spatial Turing machines were created using Mathematica's ArrayPlot function.</p> <p><b>Results</b> After results were averaged from thousands of trials using the law of large numbers, the differentiated cancer cell populations followed the standard Gompertzian growth curve with the mass-action model reaching a lower carrying capacity at a faster rate while the spatial model reached a higher carrying capacity at a slower rate; the cancer progenitor cells exhibited a gradual Gompertzian growth curve; and the CSCs remained at a lifelike percentage of total cells and exhibited a von Bertalanffy growth curve.</p> <p><b>Conclusions/Discussion</b> Cellular automaton, discrete mathematical models, theoretical computer science, and programming was used in creating mathematical models as well as automaton visualization of the progression of solid CSC-induced tumor growth over time. Automata-based modeling of tumors is useful in that automaton "rules" may be potentially substituted by boolean structures of genes, thus bridging bioinformatics and individualized tumor modeling.</p>	
<b>Summary Statement</b> Cancer stem cell (CSC) induced tumor growth is modeled using theoretical computer science and the tumor growth is visualized using cellular automata, potentially helping with creating individualized models of csc-induced tumors.	
<b>Help Received</b> Professor Komarova of UC Irvine introduced me to the mass-action and spatial models (which I used in a different project) that I applied in my research.	