



**CALIFORNIA STATE SCIENCE FAIR
2008 PROJECT SUMMARY**

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| Name(s) David K. Tang-Quan | Project Number S1416 |
| Project Title Antimicrobial Peptide Susceptibility of Candida albicans Kinase Mutants | |
| Abstract Objectives/Goals Disseminated candidiasis occurs in hospitalized patients when the fungus, <i>Candida albicans</i> , enters the bloodstream and infects almost all organs of the body. The mortality rate of this disease is close to 40%. Immunocompromised patients can also develop oropharyngeal candidiasis, or <i>Candida</i> overgrowth in the mouth. Although the mechanisms behind <i>C. albicans</i> defense against reactive oxygen intermediates have already been discovered, the mechanisms for resistance to antimicrobial peptides are still unknown. Methods/Materials Sixty-four kinase mutants were screened with protamine containing antimicrobial peptides to discover mutant strains that were hyper-susceptible. Serial dilutions of each strain were pipetted onto protamine and YPD plates, starting at 10^8 cells/ml and decreased by a factor of 10 until the final concentration of organisms was 10^3 cells/ml. Hyper-susceptible strains were retested and the validity of results was further confirmed in a trial using a second, independent clone. Results Results from the first 96-well microtiter plate of kinase mutants showed three genes that are responsible for hyper-susceptibility to antimicrobial peptides: PBS2, YCK2, and HST7. Findings from the second 96-well plate of kinase mutants were unconfirmed, but six possible hyper-susceptible strains and some likely hypo-susceptible strains were identified. Conclusions/Discussion It was concluded that the genes PBS2, YCK2, and HST7 play a significant role in <i>C. albicans</i> resistance to antimicrobial peptides. Further research to confirm the validity of these results includes gene complementation and a clean knock out of genes. Pharmacologists and other medical researchers can use these findings to determine which genes need to be inhibited when developing medications for <i>Candida</i> infections. | |
| Summary Statement This study focuses on identifying the genes responsible for <i>Candida albicans</i> resistance to antimicrobial peptides. | |
| Help Received Used lab equipment at Los Angeles Biomedical Research Institute; mentored by Dr. Scott Filler; supervised by Norma Solis. | |