

Characterization of Van Well Sweet Cherry Self-Incompatibility (S) Alleles

Sequoia Leuba¹, Ben Kilian² and Amit Dhingra²

¹Yale University, CT ²Department of Horticulture at Washington State University, WA



Introduction

The multi-allelic self-incompatibility (S) locus controls self-incompatibility and incompatibility between cultivars of sweet cherry (*Prunus avium L.*) through a gametophytic self-incompatibility system. If the haploid genome of the pollen and the diploid genome of the style have the same S-allele, pollen tube growth is stopped. Thus, commercially grown cherries are planted with compatible cultivars to ensure adequate fruit set. Van Well nursery sent samples of Sunset Bing, Kimberly, and Glory Cherry sweet cherry cultivars, to determine each of their respective S alleles. Using allele-specific primers, we were able to effectively determine the S genotype of each cultivar. Genetic identification of the S genotype of a specific cultivar is of utmost practical importance to maximize yield for growers and also for breeders to use the genotype to test or modify various fruit characteristics.

Background Information

There are 13 identified S alleles, ranging from S1 to S16 (S8, S11, and S15 are duplicates). Allele specific primers for each allele have been determined and published. By using allele specific primers, one can determine the S allele of a cultivar. The variable cultivars are Sunset Bing (SB), Kimberly (K), and Glory Cherry (GC). We have acquired control samples of cultivars Bing (S3, S4), Chelan (S3, S9), and Skeena (S1, S4), thus having controls for S alleles S1, S3, S4, and S9. We also acquired control primers that anneal to all S alleles to ensure the PCR was successful.

Images of Cherry Cultivars

Bing Cherry



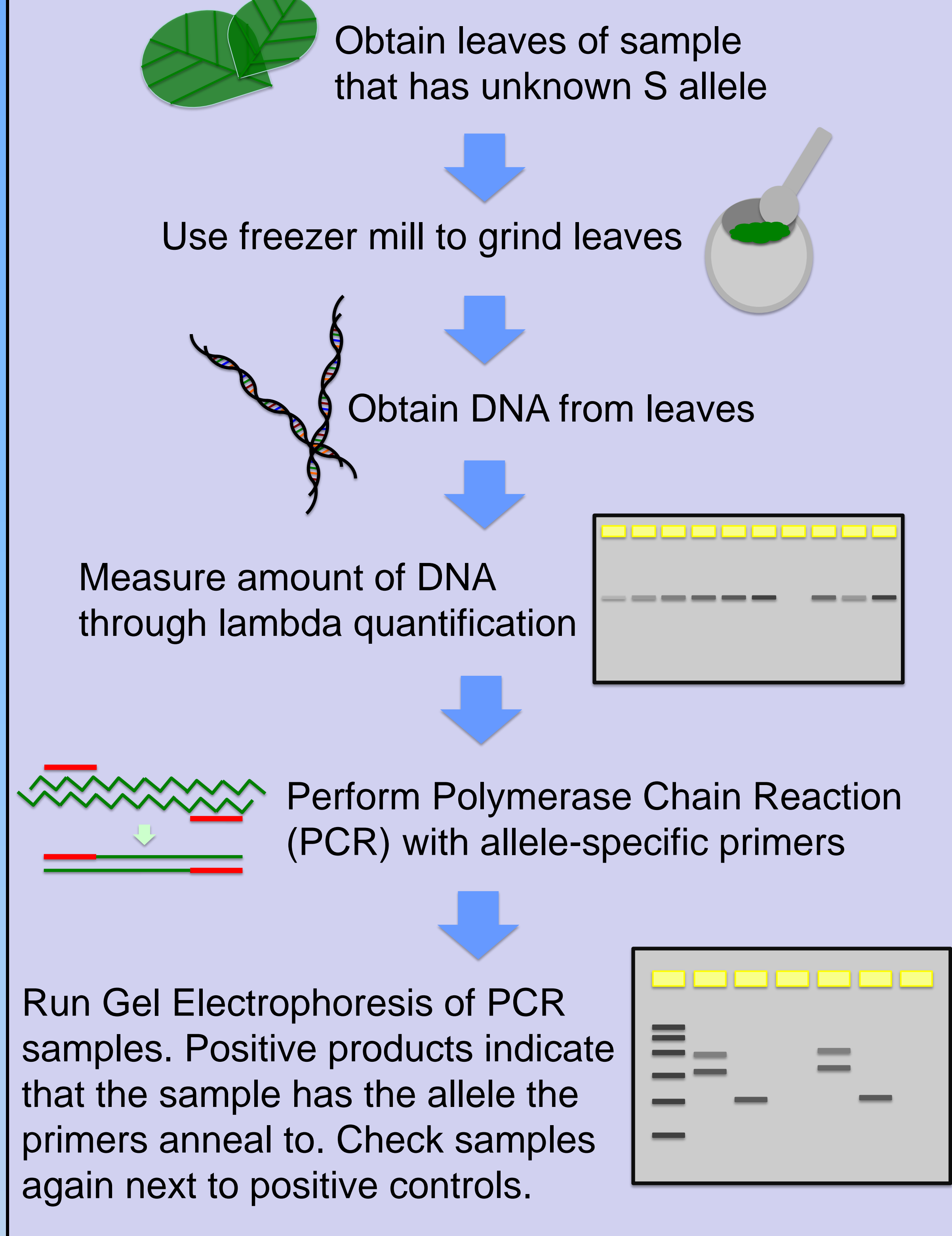
<http://www.f-fruits.com/wp-content/uploads/2011/09/Bing-Cherries1.jpg>

Glory Cherry



<http://images.fruitreeshop.co.uk/images/products/zoom/1313567486-71840900.jpg>

Procedure



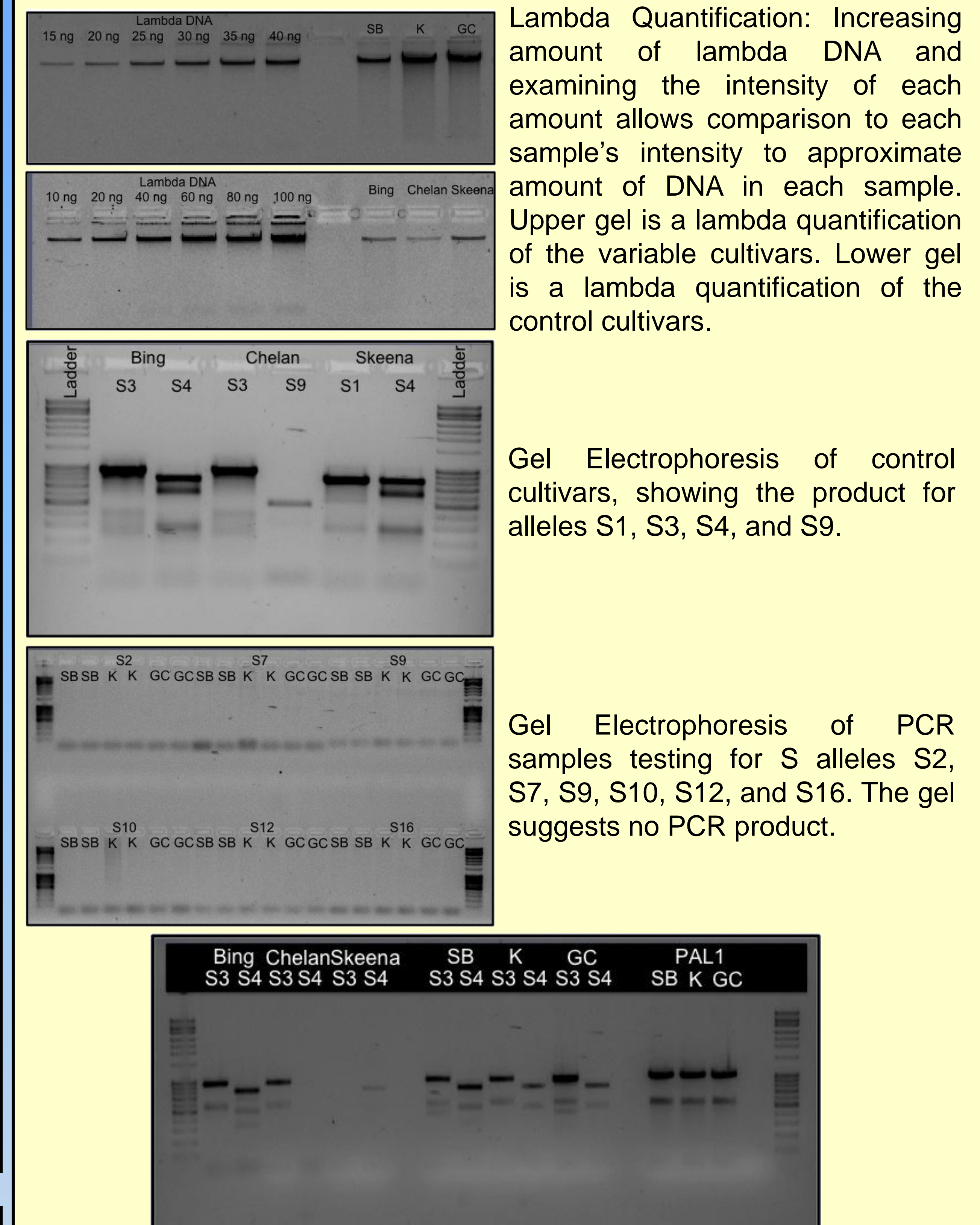
Conclusions

Through our PCR analysis, we were able to determine that Sunset Bing, Kimberly, and Glory Cherry all have the S3 and S4 alleles. With this knowledge, we can determine which cultivars they are able to be crossed with, which is necessary not only for growers to maximize the yield of the fruit set, but also for the breeders in modifying fruit characteristics.

Future Research

More leaves will be obtained to provide positive controls for the other S alleles. In the future, with these positive controls along with the primers for all the S alleles, we will be able to efficiently determine the S allele of any future sweet cherry leaf sample.

Results



Acknowledgements

This work was supported by the National Science Foundation's REU Program under grant number DBI-1156880.

References

- Ipek, A. H. Gulen, M.E. Akcay, M. Ipek, S. Ergin, and A. Eris. "Determination of Self-incompatibility Groups of Sweet Cherry Genotypes from Turkey." *Genetics and Molecular Research* 10.1 (2011): 253-60. Web.
- Sonneveld, T., K. R. Tobutt, and T. P. Robbins. "Allele-specific PCR Detection of Sweet Cherry Self-incompatibility (S) Alleles S1 to S16 Using Consensus and Allele-specific Primers." *TAG Theoretical and Applied Genetics* 107.6 (2003): 1059-070. Print.