PLSC 731: Paper Reviews

Li et al. – Genetic structure and diversity of cultivated soybean (Glycine max (L.) Merr.) landraces in China

- 1. Describe the origin and beginning of soybean cultivation. (858)
- 2. What are landraces? What are they useful for crop improvement. (858)
- 3. Describe the Chinese soybean gene bank. What is a difficulty in maintaining the collection? (858)
- 4. Describe the use of germplasm evaluation studies. What did previous studies learn about the organization of soybean diversity? (858)
- 5. How can a population for a divergence study be organized? (859)
- 6. Describe several features of the STRUCTURE software. (859)
- 7. Why are SSRs used for diversity studies? (859)
- 8. What is a prerequisite for association mapping? (859)
- 9. What factors affect linkage disequilibrium (LD)? (859)
- 10. Describe the results of previous LD studies in soybean? (859)
- 11. What are the objectives of this study? (859)
- 12. How was a subset of the Chinese soybean genebank selected? (859)
- 13. Why were modern cultivars excluded from this study? Is this a good choice? (859)
- 14. Is this a diverse population? (859-860)
- 15. Describe the selection process for the SSRs used in this study. Was this a good choice of markers? (860)
- 16. What are the assumptions of the STRUCTURE software analysis? (860)
- 17. Describe the allelic variation and distribution of rare alleles. (863)
- 18. Why was k=7 decided upon as the subpopulation number? (863)
- 19. What does Fst measure? What is the overall distribution of Fst in this population? (864)

- 20. Is there a relationship between the STRUCTURE and neighbor joining results. (865)
- 21. Why was a subset of genotypes selected to compare Fst? (865)
- 22. What was the degree of LD within population clusters? (866)
- 23. Why was LD measured within a linkage group? (866)
- 24. What can the degree of rare alleles imply about genetic variation within this population? (867)
- 25. What result supports the concept of conservation of landraces within a genebank? (868)
- 26. Why did the authors conclude that the Yellow River Valley is the center of origin of soybean? (869)