Schlötterer (2004). The evolution of molecular markers – just a matter of fashion? *Nature Reviews Genetics*, 5:63-

- 1. What type of information does a genetic marker provide? How is it used to address one of the key questions in genetics?
- 2. What are allozymes? Are they useful in the context of genetic association studies? Why or why not?
- 3.In contrast to allozymes, what are DNA-based markers? What are the advantages of using DNA-based markers?
- 4. What are RFLPs? What downplayed their popularity?
- 5. What are minisatellites? How are they similar/dissimilar to RFLPs? What downplayed their popularity?
- 6. What is DNA fingerprinting?
- 7. What does PCR stand for and how did it change the evolution on molecular markers?
- 8. What are micro-satellites? How are they similar/dissimilar to minisatellites? How large is

- a typical repeat region? What is the key reason for them to have gained popularity in mapping endeavors? What is a major drawback of microsatellites, hampering their use in population genetics studies?
- 9. Are AFLPs PCR-based? Do AFLPs require a priori knowledge about primer sequences in the target species?
- 10. What is a shotgun genome sequence?
- 11. SNPs have become one of the most important genetic markers in genetic (association) studies. Despite their success, they suffer from some shortcomings. Name 4.
- 12. Which technique offers the most finegrained genetic information? Hence, Is DNA sequence analysis or comparison an old-fashioned business or will it revive again.
- 13. Link advantages and weaknesses to the appropriate markers.
- 14. Which markers or variation capturing technique is the most optimal in the context of making inferences of demographic processes?

- Can you make the link with the use of DNA sequences and phylogenetics?
- 15. Which markers or variation capturing technique is the most optimal in the context of paternity testing and forensics?
- 16. Which markers or variation capturing technique is the most optimal in the context of linkage analysis?
- 17. Which markers or variation capturing technique is the most optimal in the context of association analysis?