

Questions lecture 2: Genomic variation

- 1) A) Define common, versus rare and very rare variants.
B) Are major blood type variants common and why are there so many distinct alleles?
- 2) What is short read alignment and why does it represent an important computational challenge?
- 3) Which genome of which human population (Caucasian, Asian, Yoruban) tends to harbor the most variation, and why?
- 4) How many SNPs are circulating in the “human genome”, how many in each human genome? Are most rare or common, clarify your answer.
- 5) What are LD bins? Explain why the HapMap Project deemed these bins useful for inferring missing genetic information, thereby discussing the implicit trade-off in settling for a threshold of 0.8 or greater linkage.
- 6) What is population stratification? Why is this important when linking allele frequencies to disease prevalence, i.e. in GWAS studies?
- 7) a) What is the major conclusion from the 1000 genome project in terms of how variants impact coding and non-coding function? b) Are we still evolving? Provide an illustrative example that supports your answer
- 8) a) Is the principle of a “meta-genome” still valid and what is mosaicism? Clarify your answer and be able to discuss distinct mutation scenarios. What are the consequences of this finding and what is mosaicism?
- 9) How is the exome typically captured and sequenced?
- 10) Explain at least three disease gene identification strategies using exome sequencing.
- 11) Please explain the difference between the common versus the rare variant contribution to complex disease hypotheses.
- 12) Know how to calculate the odds ratio.
- 13) Why is the term “genome-wide” association studies a misnomer? (see also question 5)
- 14) What’s a Manhattan plot?
- 15) (Patients with = disease) \neq (Patients with = underlying biological disorder). Please explain this statement. How does this relate to personalized medicine?
- 16) A. What is meant with the missing heritability and where could we find it?
B. What is meant by “nature via nurture” or “it takes two to tango”?