Part 4.

Genetics of phenotypic variation and disease

In Part 4 we turn our attention to the *genetics of phenotypic variation*. We'll cover three main categories of traits: monogenic diseases, cancer, and complex traits, with particular emphasis on complex traits.

As you read you should pay attention to the themes that repeat, but with key differences, across the different categories:

- The number, allele frequencies and molecular mechanisms of variants;
- The types of selection that are most relevant;
- The study designs used to identify causal genes and variants;
- The main conceptual approaches to data analysis, and major insights.

Specifically, we cover the following:

- *Chapter 4.1: A* **Starter Pack** of trait genetics: an introduction to the topics in this section.
- *Chapter 4.2: The genetics of monogenic diseases: mapping approaches, the major mechanisms, and selection.*
- *Chapter 4.3: The genetics of cancer, emphasizing aspects of this huge field that intersect our main themes including somatic mutation and selection.*
- *Chapter 4.4: Quantitative Genetics: statistical models for the inheritance of polygenic traits, including heritability and artificial selection.*
- *Chapter* 4.5–4.7: *An overview of the main approaches for studying complex traits, and major emerging themes: GWAS; SNP heritability; regulatory genomics and the mechanisms of variant effects; and polygenic scores.*
- *Chapter 4.8: We close with more on the population genetics of complex traits including stabilizing selection and polygenic adaptation.*

I expect to release these chapters in 2024 –JKP.