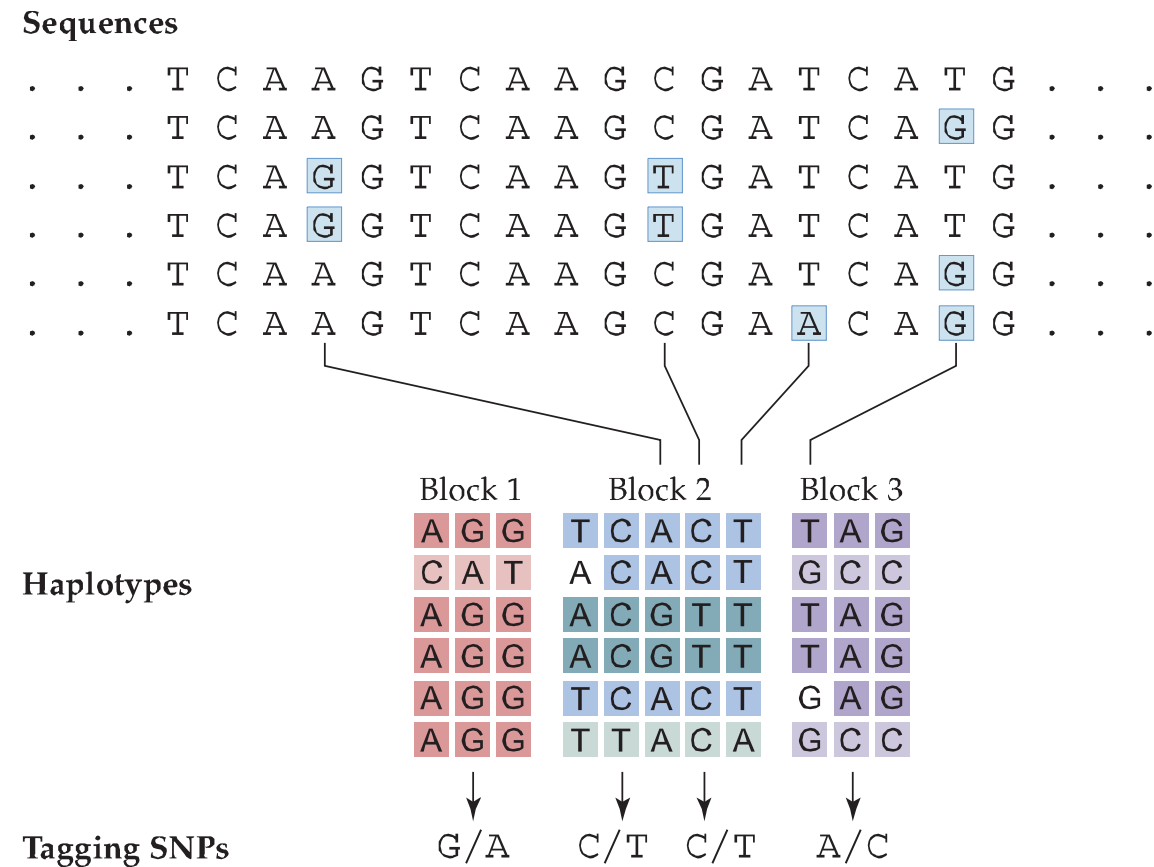


# Not hopeless: SNPs near to one another are correlated...



**Figure 3.5 Tagging SNPs and haplotype blocks.** Extraction of the polymorphisms from a set of sequences typically reveals a blocklike pattern of haplotypes. In this hypothetical example, Block 1 has two classes of haplotypes, one rare and one common; Block 2 has three classes of haplotypes; and Block 3 has two classes of haplotypes. Note that the boundaries between blocks are relatively sharp. The tagging SNPs can be used to define most of the variation in the sample.

## Population Structure can present a problem for GWAS

- What is the potential problem with a GWAS for amylose content?

