

**Hidden Markov models**

1. Consider a single intercross individual derived from two inbred mouse strains. Consider  $M$  ordered markers, and let  $G_i \in \{AA, AB, BB\}$  denote the (phase-unknown) genotype of the individual at marker  $i$ . Show that the  $G_i$  form a Markov chain.
2. Fill in the details in the Viterbi algorithm to calculate

$$\hat{g} = \arg \max_{g_1, \dots, g_M} \Pr(G_1 = g_1, \dots, G_M = g_M \mid \mathbf{O})$$