## Notes

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## Infer mom's genotype by JRI

We have obs. mom and obs. (selfed) kids. We want to know  $P(G|\theta)$ , and  $P(G|\theta) \propto P(\theta|G) \times P(G)$ , where  $\theta$  is observed data. This consists of observed genotypes (G') of both mom and kids. So:  $P(G|\theta) \propto \left(\prod_{i=1}^k P(G'_k|G)\right) \times P(G'_{mom}|G) \times P(G)$  This function is to impute mom's genotype from a progeny array of k kids at a single locus. inferred\_mom=1 -> 00, 2->01, 3->11

## Imputing Founder Genotypes

 $P(G|\theta) \propto P(\theta|G) \times P(G)$ 

$$P(G|\theta) \propto \left(\prod_{i=1}^{k} P(G'_{i}|G)\right) \times \left(\sum_{m=1}^{mom} P(G'_{mom}|G)\right) \times P(G)$$

This function is to impute mom's genotype by finding the maximum likelihood of  $P(G|\theta)$  from a progeny array of k kids at a single locus. - Where  $\theta$  denotes observed data. It consists of observed genotypes (G') of both mom and kids.

- P(G) is the Hardy-Weinberg equilibrium estimated from the population.
- $P(G'_{mom}|G)$  is the error matrix estimated from the data, i.e. homozygote error = 0.02 and heterozygote error = 0.6.
- $P(G'_i|G)$  is the error matrix times Mendelian segregation rate.

## Phasing Founder Genotypes

$$\begin{split} &P(H|\theta) \propto P(\theta|H) \times P(H) \\ &P(H|\theta) \propto \left(\prod_{i=1}^k P(H_k'|H)\right) \times P(H) \\ &P(H|\theta) \propto \left(\prod_{i=1}^k \prod_{l=1}^n P(G_{i,l}'|H)\right) \times P(H) \end{split}$$

- Where  $\theta$  denotes observed data.
- P(H) is the probability of the haplotype for a given window size of n.
- $P(G'_{i,l}|H)$  is the probability of kid i at locus l for a given haplotype H.
- We assume all the possible haplotypes of a given window size are equally likely.