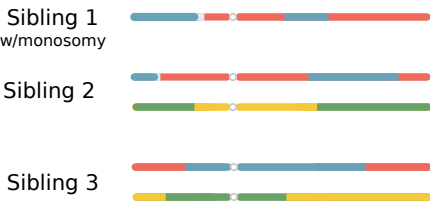


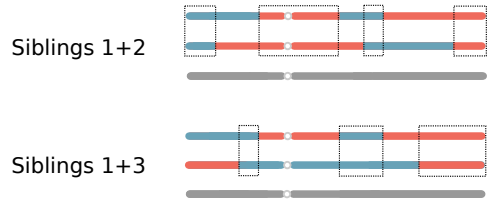
## Chromosome 16



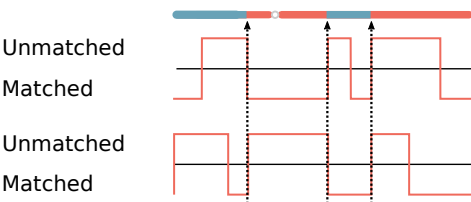
Initially we have no knowledge  
about the haplotypes composition



Our method identifies regions  
with matched haplotypes



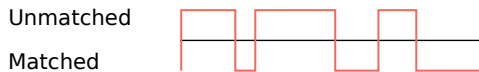
Common crossover positions  
are traced to the monosomy



The output for siblings 1+2



The output for siblings 1+3



The output signal of our method

