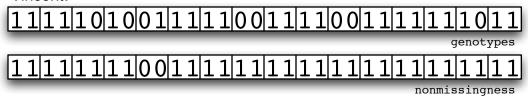
## Identity-by-state and software popcount

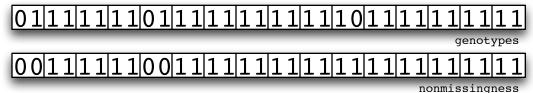
PLINK 1.9's most important optimization is its replacement of slow loops iterating over single genotype calls with bitwise operations on many calls at a time. This document illustrates how identity-by-state (i.e. Hamming distance) between two genomes is computed in this fashion.

Step 1: Transposition and other preprocessing. PLINK's core file format saves genotype calls in a variant-major manner. IBS computation is faster with sample-major data, and its overall time complexity is  $O(mn^2)$  while transposition is just O(mn), so we transpose the data before the main loop. We also assemble a bit array tracking the presence of missing genotype calls.

## Vincent:



## Anton:



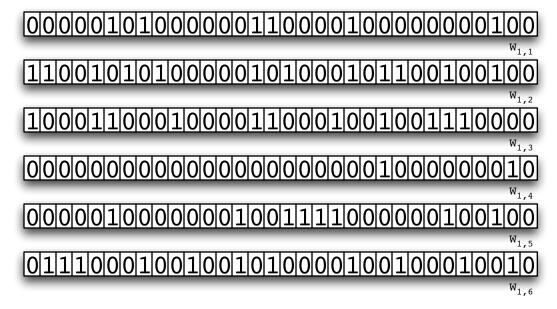
Step 2: XOR-and-mask. PLINK 1 represents homozygous major calls with binary 11, heterozygous calls with 10, and homozygous minor calls with 00.

Conveniently, if you take the exclusive-or of two such values, the number of set bits in the result is the number of differing allele calls; thus, the overall Hamming distance between two genomes in (transposed) PLINK 1 format is the bit population count of their XOR. Excepting missing calls (represented by 01), that is; we "mask" (via AND operations) the final result with both nonmissingness arrays to force those bits to zero. (The red '0' below is due to the mask.)

## 000001010000001100001100000000

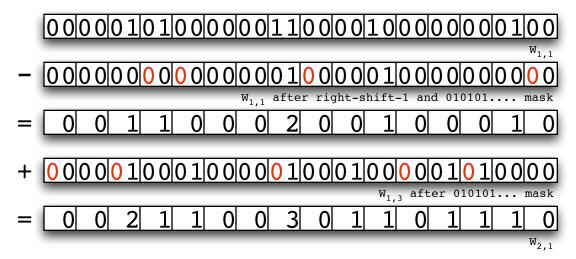
XOR-and-mask result

Step 3: Software popcount. Since PLINK is still used on many machines lacking a hardware popcount instruction, we use SSE2 (in x86-64 builds) or basic word (in 32-bit builds) operations to implement the "bitslice" algorithm discussed by Dalke et al., which is almost as fast when acting on long arrays. For clarity of exposition, we illustrate what happens with six 32-bit words; our SSE2 code applies the same idea to batches of fifteen or thirty 128-bit blocks.



This can be seen as a collection of 192 one-bit values which add up to our desired result.

The bitslice algorithm starts by generating a collection of two-bit partial sums which add up to the same total. Specifically, the partial sums in  $W_{2,1}$  aggregate two bits in  $W_{1,1}$  and an even-position bit in  $W_{1,3}$ ;  $W_{2,2}$  aggregates pairs of bits in  $W_{1,2}$  and odd-position bits in  $W_{1,3}$ ;  $W_{2,3}$  aggregates pairs of bits in  $W_{1,4}$  and even-position bits in  $W_{1,6}$ ; and  $W_{2,4}$  aggregates pairs of bits in  $W_{1,5}$  and odd-position bits in  $W_{1,6}$ . The actual operations are a right-shift-1, a mask with 010101..., a subtraction, a mask (even-position) or right-shift-1-and-mask (odd-position) with 010101..., and an addition.

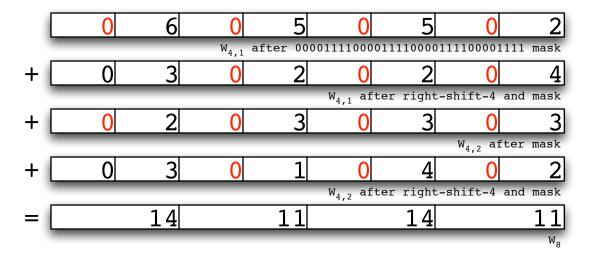


Note that "2" is shorthand for binary 10 and "3" is shorthand for binary 11 here; similar shorthand will be used for four- and eight-bit partial sums on the next page.

The next step is to use these to produce an even smaller collection of four-bit partial sums with the same total. Specifically,  $W_{4,1}$  aggregates two values in  $W_{2,1}$  and two values in  $W_{2,2}$ , while  $W_{4,2}$  aggregates two values in  $W_{2,3}$  and two values in  $W_{2,4}$ .



Then we produce a single word of eight-bit partial sums from  $W_{4,1}$  and  $W_{4,2}$ . (Since none of the four-bit partial sums can be greater than 12, and eight bits can represent values up to 255, we can actually merge up to 10 pairs of partial sums at this stage, rather than just 2; this is done by some of our SSE2 code.)



Finally, we add these eight-bit partial sums: 14 + 11 + 14 + 11 = 50, which is indeed the number of set bits among the original 192.