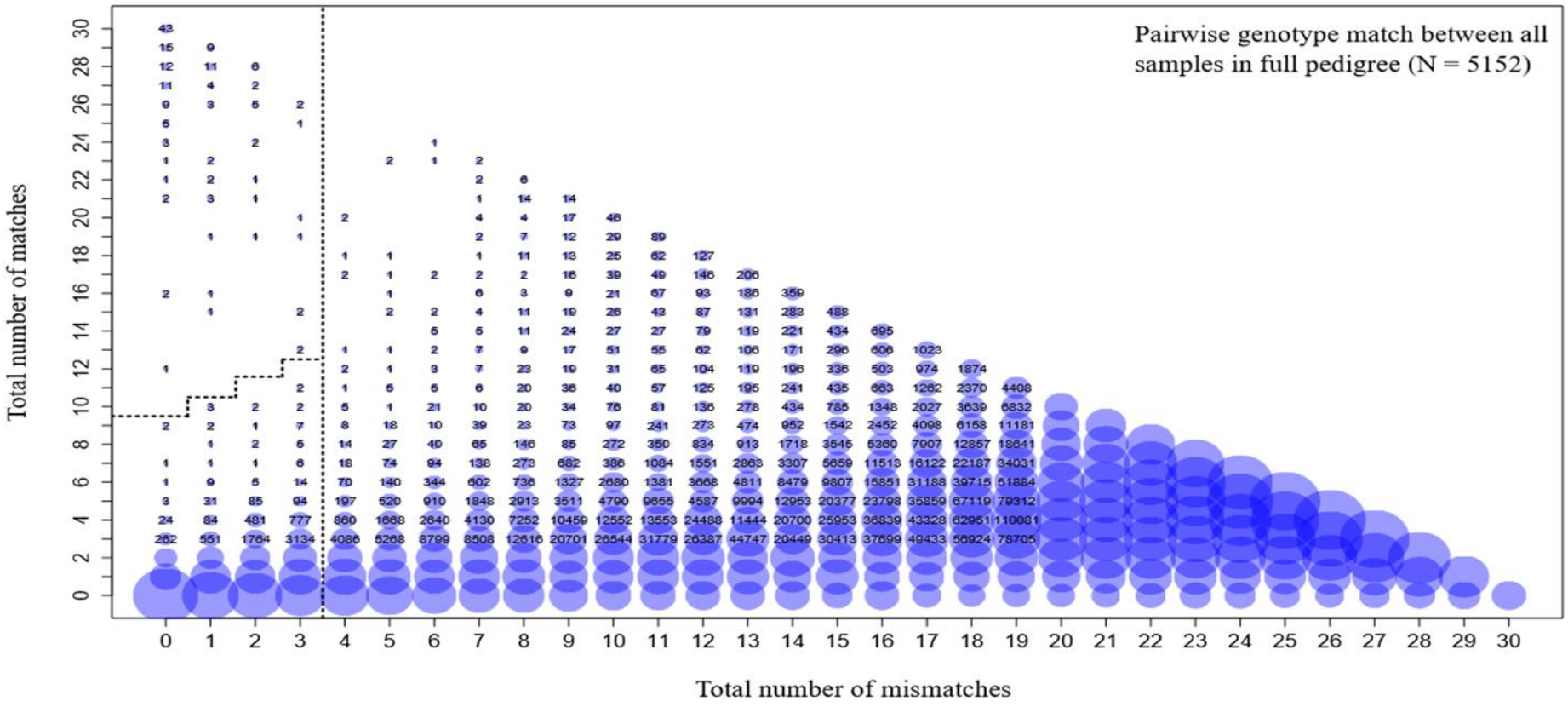

Figure S2: Identity analysis showing the total number of all pairwise genotype matches and mismatches.