

Joining of genotyping data of diverse sets of chicken samples originating from different arrays can be improved by imputation with sequence data

Problem:

Studies on Chicken Diversity use various genotyping array versions
→ Data merging is difficult



580'961 SNPs



8'583 SNPs

**Overlap only
1'364 SNPs**

Joining of genotyping data of diverse sets of chicken samples originating from different arrays can be improved by imputation with sequence data

