Poster 17

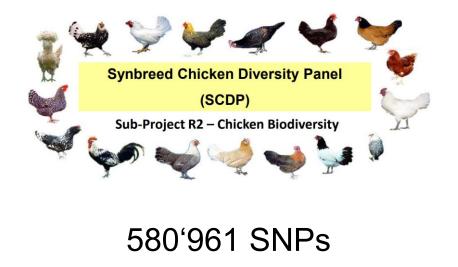
Christian Reimer, Johannes Geibel, Annett Weigend, Steffen Weigend

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





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

