



Genetic structure of five autochthonous and three commercial pig breeds using a high-density SNP chip

María Muñoz, Juan María García-Casco, Ana I. Fernández, Fabián García, Claudia Geraci, Luca Fontanesi, Marjeta Candek-Potokar & Cristina Óvilo



Funded by European Union
Horizon 2020
Grant agreement No 634476

INTRODUCTION



Genetic markers are widely used in the main European pig breeds

- To analyze population structures
- To characterize genetic relationships and diversity

Few studies in autochthonous European breeds

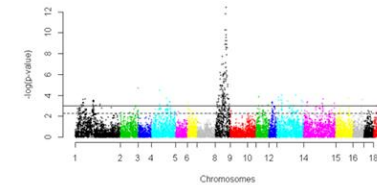


INTRODUCTION

Potential **applications** of genomic characterization

1. Knowledge of potentially negative alleles

2. Genetic basis of specific traits



3. Development of traceability panels



4. *in-situ* management of genetic resources improves utilization of local breeds



OBJECTIVE



Characterization of the genetic relationships and diversity between 20 autochthonous European and five commercial pig breeds



Pilot Study

Iberian, Krskopolje, Casertana, Cinta Senese, Apulo Calabrese, Duroc, Large White and Landrace breeds



Funded by European Union
Horizon 2020
Grant agreement No 634476

GENOTYPING

GeneSeek® Genomic Profiler™ for Porcine HD

Breed	N
Apulo Calabrese	5
Casertana	5
Cinta Senese	5
Duroc	5
Iberian	12
Krskopolje	4
Landrace	4
Large White	4
Total	44



- 68,528 SNPs
- Evenly distributed median 25kb gap spacing
- 43k from the PorcineSNP60 BeadChip

Quality control (QC): plink

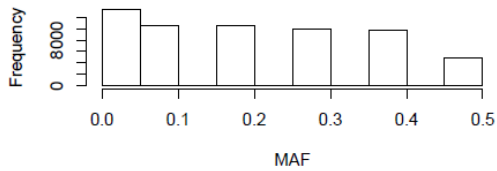
% missing values <2%



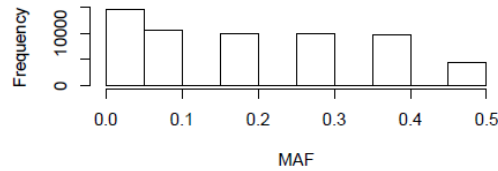
RESULTS

59,193 with % missing values <2%

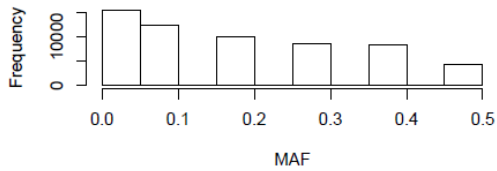
Apulo Calabrese



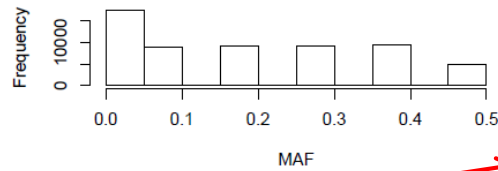
Casertana



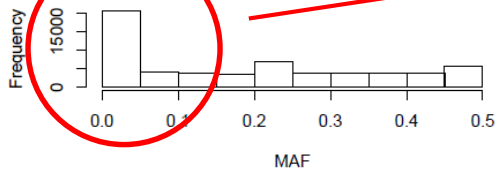
Cinta Senese



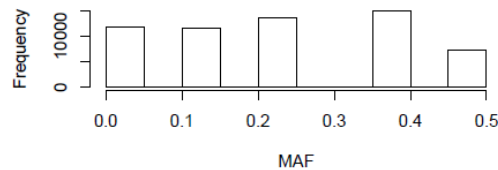
Duroc



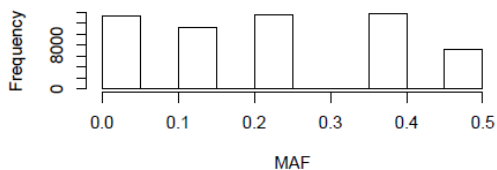
Iberian



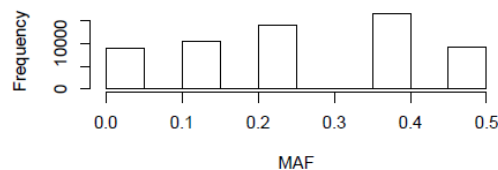
Krskopolje



Landrace



Large White



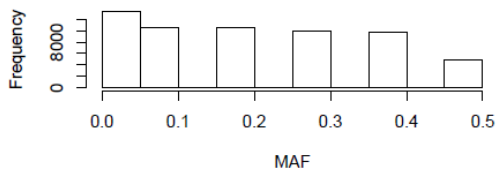
Highest number of monomorphic SNPs
(25.3%)



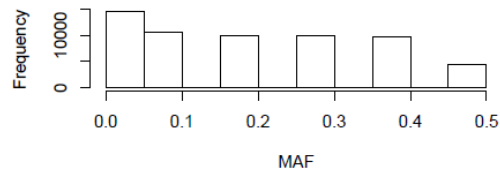
RESULTS

59,193 with % missing values <2%

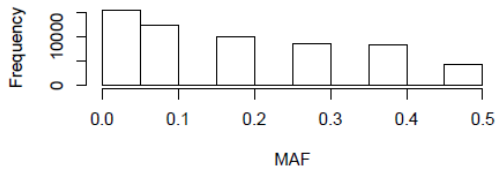
Apulo Calabrese



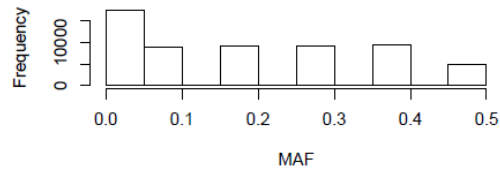
Casertana



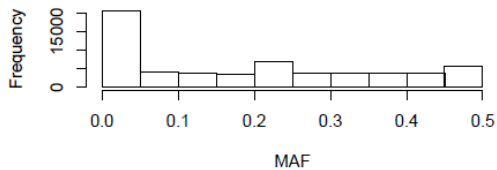
Cinta Senese



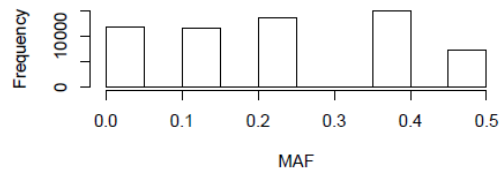
Duroc



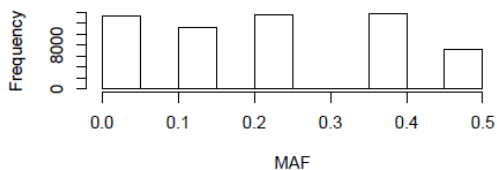
Iberian



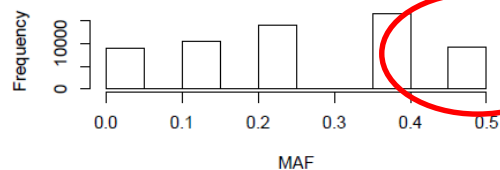
Krskopolje



Landrace



Large White



9,229 SNPs



RESULTS

Library hierfstat (R)

	Ho	Hs	Fis
Apulo Calabrese	0.251	0.319	0.164
Casertana	0.249	0.310	0.147
Cinta Senese	0.289	0.285	-0.026
Duroc	0.287	0.288	-0.012
Iberian	0.230	0.262	0.102
Krskopolje	0.348	0.355	-0.013
Landrace	0.360	0.337	-0.086
Large White	0.320	0.396	0.130

Highest Ho in Landrace & lowest in Iberian

Highest Fis in Apulo Calabrese & lowest in Duroc



RESULTS

Genetic Distance: Ds (Nei's standard genetic distance)

	Calabrese	Casertana	Cinta Senese	Duroc	Iberian	Krskopolje	Landrace	Large White
Calabrese	0							
Casertana	0.15	0						
Cinta Senese	0.15	0.15	0					
Duroc	0.20	0.20	0.19	0				
Iberian	0.13	0.14	0.10	0.18	0			
Krskopolje	0.14	0.15	0.14	0.17	0.13	0		
Landrace	0.15	0.15	0.16	0.22	0.15	0.14	0	
Large White	0.14	0.15	0.14	0.09	0.14	0.13	0.15	0

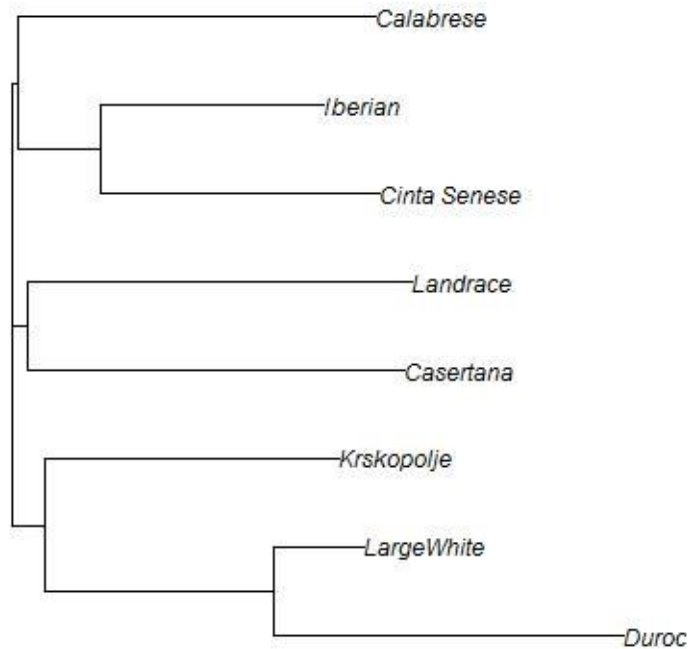
Genetic Distance: Fst

	Calabrese	Casertana	Cinta Senese	Duroc	Iberian	Krskopolje	Landrace	Large White
Calabrese	0							
Casertana	0.27	0						
Cinta Senese	0.27	0.28	0					
Duroc	0.33	0.34	0.33	0				
Iberian	0.26	0.27	0.22	0.32	0			
Krskopolje	0.24	0.25	0.25	0.29	0.24	0		
Landrace	0.27	0.26	0.28	0.34	0.27	0.23	0	
Large White	0.23	0.24	0.24	0.17	0.24	0.20	0.23	0

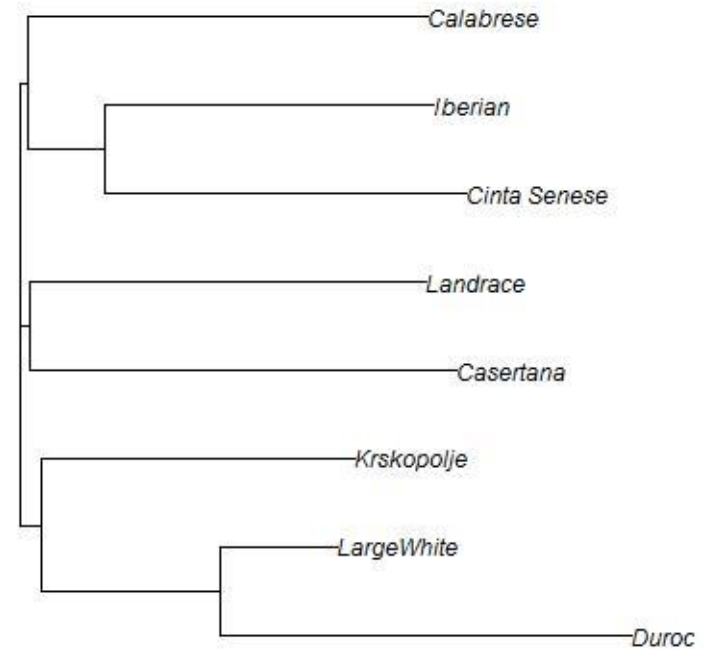


RESULTS

NJ Tree using Ds



NJ Tree using Fst



SOME RESULTS ALL BREEDS

Breeds	Code	Number of succesfully genotyped animals with 70K SNP chip
Alentejano	AL	48
Basque	BA	39
Bisaro	BI	49
Calabrese	CA	53
Casertana	CS	54
Cinta Senese	CI	54
Crna Slavonska	CR	49
Duroc	DU	5
Gascon	GA	48
Iberian	IB	49
Kroskopolje	KR	52
Landrace	LA	4
Large White	LW	4
Lietuvos Vietines	LI	48
Mangulica	MA	50
Mora Romagnola	MR	48
Moravka	MO	50
Nero Siciliano	NS	50
Porc Negre Mallorquí	PN	48
Sarda	SA	48
Schwabisch Hallisches	SH	49
Senojo Tipo Lietuvos Vietines	SE	51
Turopojle	TU	50
Wild Boar	WB	7



SOME RESULTS ALL BREEDS

MAF > 0.01 & missing values < 0.1 51,221 SNPs were located in Ssc11.1 autosomes

Breed	MAF range values									
	0-0.05	0.05-0.10	0.10-0.15	0.15-0.20	0.20-0.25	0.25-0.30	0.30-0.35	0.35-0.40	0.40-0.45	0.45-0.50
AL	18247	5800	4521	4426	4660	3757	4750	5046	4934	4311
BA	20434	6185	5575	4783	4357	4194	3802	3959	3580	3583
BI	4134	4830	5255	5933	6338	6615	6989	7030	7048	6280
CA	10786	6265	5159	6399	4770	4537	5966	5169	5135	6266
CS	8187	4816	5593	5497	6417	5598	6174	6432	5599	6139
CI	8379	7954	8017	5989	6141	4516	4561	5233	4618	5044
CR	5348	5019	5381	5992	6055	6290	6469	6741	6965	6192
DU	16254	9427	177	9567	195	9518	77	10076	0	5161
GA	12192	5098	5574	5328	5182	4345	6206	5917	5533	5077
IB	16255	6189	4581	4566	4671	4704	4847	4871	5133	4635
WB	18233	6326	5939	234	6172	6404	173	6717	6832	3421
KR	4921	4162	4981	5501	6194	6396	6807	7388	7595	6507
LA	11868	0	11782	122	14243	0	123	14580	0	7732
LW	7069	0	10998	151	14779	0	200	17434	0	9820
LI	8726	4567	5135	5732	5656	4923	6239	6657	6700	6117
MA	19086	4917	4787	4540	4537	4578	4777	4673	4415	4142
MR	22286	5509	6264	4933	4217	2713	3803	3723	3677	3327
MO	5226	4849	5734	5757	6062	6248	6834	6732	6931	6079
NS	3604	5057	5854	6419	6331	6499	6562	6821	6938	6367
PN	12377	7783	5741	5273	5067	4078	5094	5266	5095	4678
SA	1542	3500	4988	5929	6718	6786	7724	7906	8069	7290
SH	6718	4552	4857	5314	5782	6305	6392	6652	7030	6850
SE	6761	4688	5085	5631	5937	5179	6873	7043	6928	6327
TU	29743	4809	4619	4418	3141	3344	2988	2521	2438	2431

European Union



SOME RESULTS ALL BREEDS

	HO	HE	FIS
Alentejano	0.248	0.259	0.041
Basque	0.240	0.233	-0.026
Bisaro	0.338	0.355	0.046
Calabrese	0.258	0.305	0.138
Casertana	0.294	0.326	0.087
Cinta	0.299	0.300	0.012
Crna	0.332	0.346	0.040
Duroc	0.300	0.270	-0.109
Gascon	0.299	0.298	-0.005
Iberico	0.251	0.270	0.077
Wild Boar	0.240	0.254	0.041
Krskopolje	0.363	0.361	-0.003
Landrace	0.372	0.310	-0.194
Lietuvos	0.354	0.331	-0.066
Large White	0.336	0.350	0.016
Mangulica	0.257	0.259	0.005
Mora	0.230	0.220	-0.039
Moravka	0.349	0.353	0.012
Nero Siciliano	0.342	0.359	0.050
Porc negre	0.279	0.285	0.015
Sarda	0.360	0.383	0.056
Schwabisch	0.351	0.346	-0.006
Senojo	0.358	0.341	-0.049
Turopolje	0.195	0.187	0.046



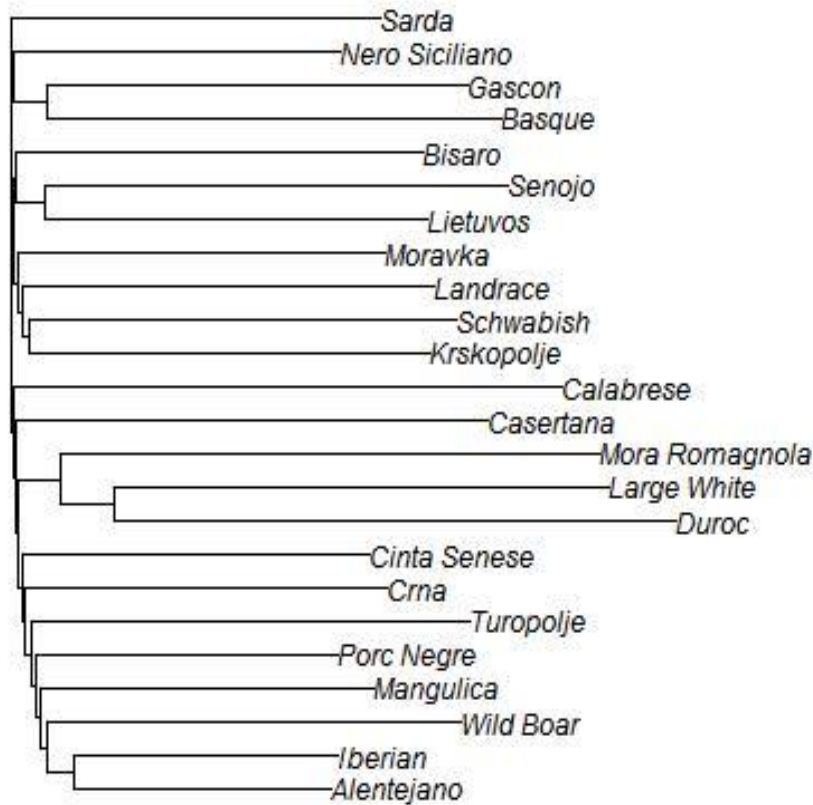
SOME RESULTS ALL BREEDS

	AL	BA	BI	CA	CS	CI	CR	DU	GA	IB	WB	KR	LA	LW	LI	MA	MR	MO	NS	PC	SA	SH	SE
AL																							
BA	0,41																						
BI	0,39	0,48																					
CA	0,46	0,55	0,51																				
CS	0,42	0,51	0,47	0,54																			
CI	0,35	0,45	0,41	0,48	0,44																		
CR	0,36	0,46	0,42	0,49	0,45	0,38																	
DU	0,52	0,61	0,57	0,64	0,60	0,53	0,54																
GA	0,41	0,46	0,46	0,54	0,49	0,43	0,44	0,60															
IB	0,28	0,42	0,39	0,46	0,42	0,35	0,37	0,51	0,41														
WB	0,37	0,49	0,46	0,53	0,49	0,41	0,43	0,59	0,48	0,37													
KR	0,40	0,48	0,44	0,51	0,47	0,41	0,42	0,56	0,46	0,40	0,46												
LA	0,40	0,49	0,43	0,51	0,47	0,42	0,42	0,59	0,46	0,41	0,47	0,44											
LW	0,40	0,48	0,44	0,51	0,47	0,42	0,42	0,58	0,46	0,40	0,47	0,44	0,45										
LI	0,49	0,58	0,53	0,60	0,56	0,50	0,51	0,56	0,55	0,49	0,56	0,53	0,55	0,53									
MA	0,33	0,44	0,41	0,48	0,44	0,37	0,38	0,54	0,43	0,34	0,40	0,42	0,43	0,42	0,51								
MR	0,47	0,58	0,53	0,60	0,56	0,49	0,51	0,60	0,56	0,48	0,54	0,53	0,54	0,53	0,59	0,50							
MO	0,37	0,46	0,42	0,49	0,45	0,39	0,39	0,55	0,44	0,37	0,43	0,42	0,42	0,42	0,51	0,38	0,51						
NS	0,33	0,43	0,40	0,47	0,43	0,36	0,37	0,53	0,41	0,34	0,40	0,40	0,40	0,40	0,49	0,36	0,49	0,37					
PN	0,31	0,42	0,39	0,46	0,42	0,35	0,37	0,52	0,41	0,32	0,38	0,40	0,40	0,40	0,49	0,34	0,48	0,37	0,34				
SA	0,36	0,46	0,42	0,49	0,45	0,39	0,40	0,54	0,44	0,37	0,43	0,42	0,42	0,42	0,51	0,39	0,51	0,40	0,37	0,37			
SH	0,41	0,49	0,45	0,52	0,49	0,43	0,44	0,60	0,48	0,42	0,48	0,44	0,44	0,45	0,55	0,43	0,55	0,43	0,41	0,41	0,43		
SE	0,45	0,53	0,47	0,55	0,52	0,46	0,47	0,63	0,50	0,45	0,52	0,48	0,48	0,45	0,56	0,47	0,58	0,46	0,44	0,45	0,45	0,49	
TU	0,39	0,49	0,46	0,53	0,49	0,42	0,42	0,59	0,48	0,39	0,46	0,47	0,47	0,47	0,56	0,41	0,55	0,44	0,41	0,40	0,44	0,48	0,52



SOME RESULTS ALL BREEDS

NJ Tree using Ds



REMARKS

1. GeneSeek® Genomic Profiler™ for Porcine HD allow genetic characterization of autochthonous European pig breeds
2. Although in the pilot study **Iberian** is the breed with the highest number of **monomorphic** SNPs, **Turopolje** shows the highest value in the whole study
3. Duroc is the breed with the highest genetic distances both in the pilot study and in the whole one.
4. Further analyses (PCAs, STRUCTURE, LD ...) are being carried out to complete the genetic characterization of the autochthonous breeds





Thank you very much!

