DEVELOPMENT AND UTILIZATION OF GENOMIC INFORMATION FOR EFFECTIVE PIG PRODUCTION

- RECENT ACHIEVEMENTS AND FUTURE PROSPECTS

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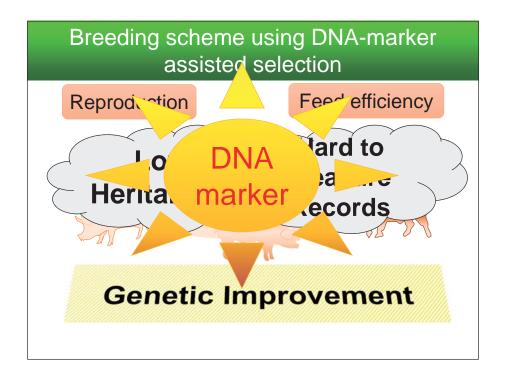
Animal Genome Research Unit

National Agriculture and Food Research Organization



Feed efficiency								
¥70,000				A				
¥60,000				/				
¥50,000		\wedge	~					
¥40,000								
¥30,000		0						
(Price/t)	1990 1992 1994	1996 1998 2000	2002	2006 2008 2010	2012			
	Minis	try of Agriculture	Forestry and Fi	sheries "Index	of agricultural cost"			
Breed -	Average Daily Gain Feed conversion ratio							
Dicca	2010	2020	2010	2020	r ccd books			
Landrace	800	900	3.0	2.9				
Large White	800	910	3.0	2.9	Divid Oolsa			
Duroc	870	1,000	3.1	2.9	Bw 100kg			

Reproduction									
Comparison of reproductive performance between countries									
	JPN	USA	CAN	DNK					
Piglets/litter (B/A)		10.5	10.2	13.0					
Deliveries (A)		2.37	2.32	2.27					
Piglets weaned/year (B)		24.94	23.59	29.62					
JPN: Japan Pork Producers Association (2009) Other: BPEX "2012 Pig Cost of Production in Selected Countries"									
Breeds	Piglet weaned/year 2010 2020								
Landrace Large White	9.9 10.0 8.9		10.8 10.9 9.4						
	ets/litter (B/A) veries (A) ets ned/year (B) Breeds Landrace	pmparison of reproductive per JPN ets/litter (B/A) 9.2 veries (A) 2.2 ets ned/year (B) Preeds Preeds Landrace Large White 1	pmparison of reproductive performance by JPN USA ets/litter (B/A) 9.2 10.5 veries (A) 2.2 2.37 ets ned/year (B) 20.2 24.94 Breeds Piglet weam 2010 Landrace 9.9 Large White 10.0	### Productive performance between concept support of the productive performance between concept support of performance support					



What's Genome? DNA marker?



size **3**Gb = DNA **3 billion**

BRINGING 1% (30 million) Species difference

0.1% (3 million) Individual difference

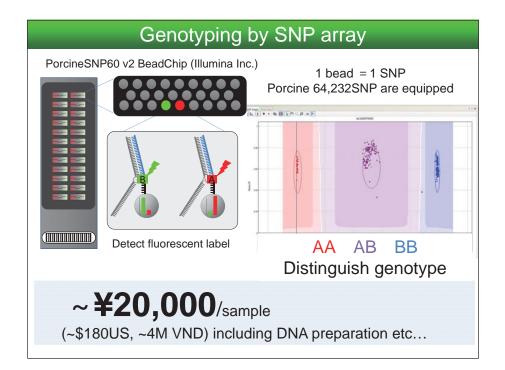
Polymorphisms

Single nucleotide polymorphism

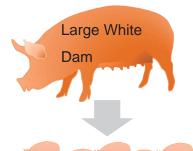
SNP

Development of useful tool to analyze genetic aspect allows to analyze 64,000 SNPs simultaneously

Phenotype SNP genotype SNP genome 3 Gb / 60 K SNP = 50 kb Average interval between 2 SNPs







Dams records

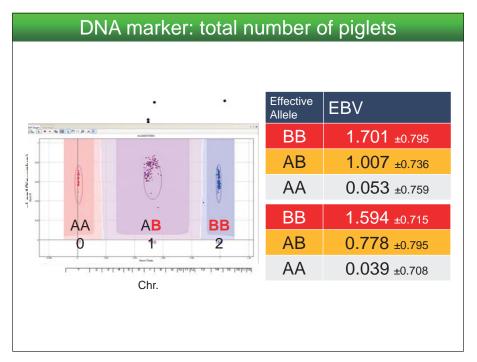
15,121 litters

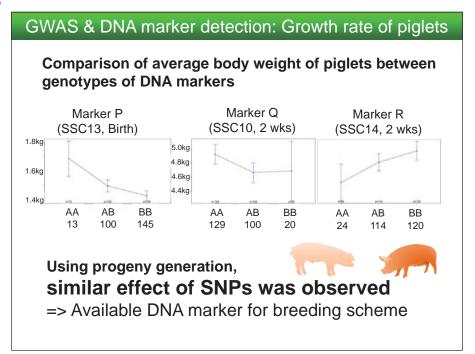
SNP genotyping

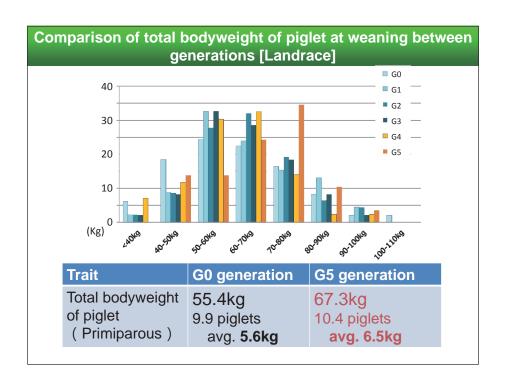
474 heads

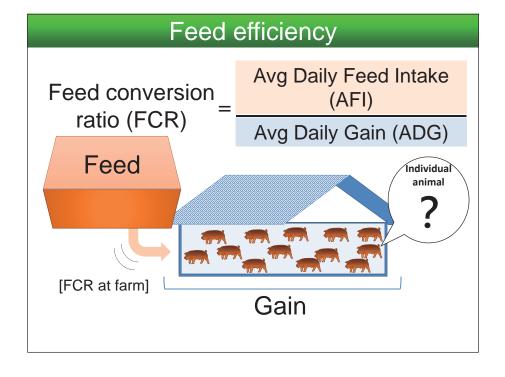


Total number of piglets

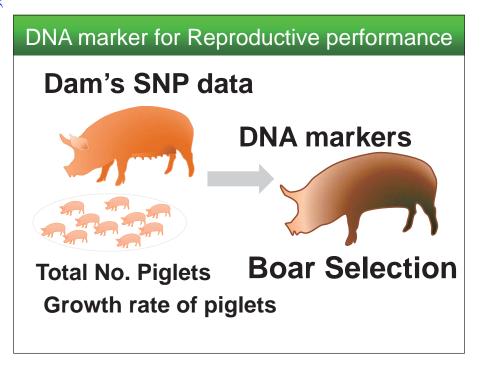


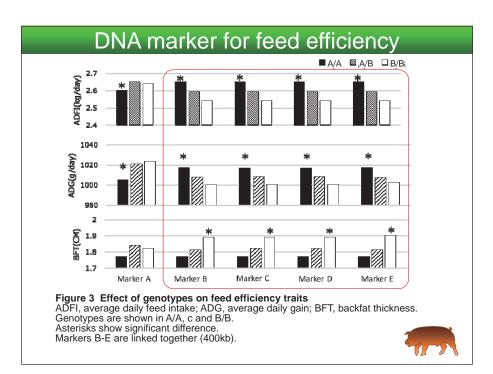


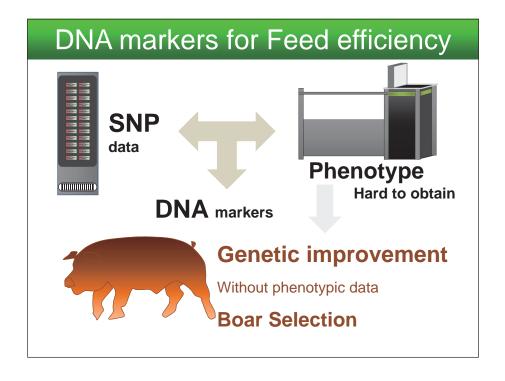




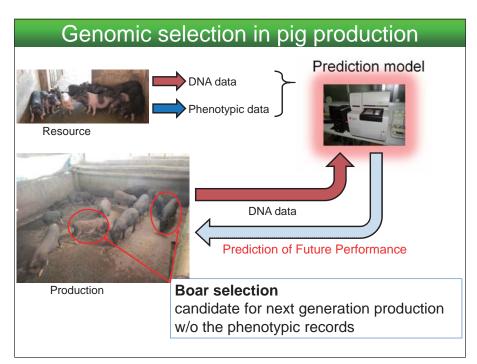
FIRE® system: Automated recording Duroc population 1622 feeding records with SNP genotyping data GWAS Versance explained by 5 adjacents 38º vindou - Trait: 1 Effect: 5 SSC6 SSC10 SSC11 SSC10 SSC11











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