



B+LNZ Genetics + ICAR

# Agenda

- B+LNZ Genetics Background
- New Zealand Livestock Farming
- B+LNZ Genetics & Beef
- B+LNZ Genetics & Sheep
- Why B+LNZ Genetics + ICAR?

Partnership >

Government

+

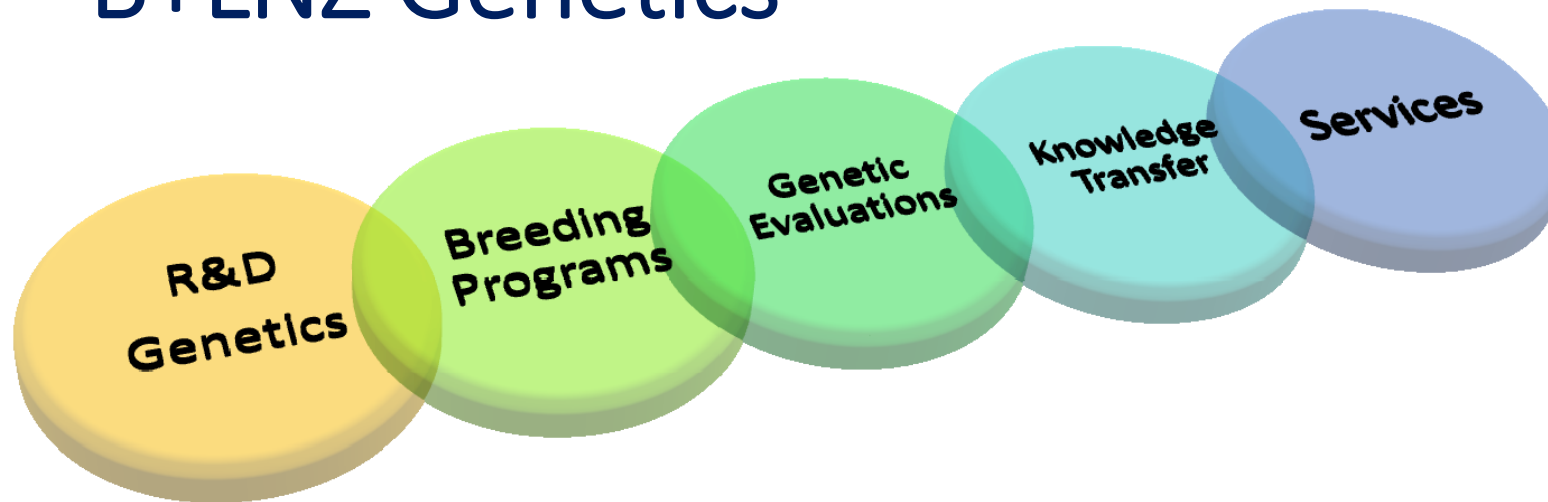
Sheep & Beef Farmers



MINISTRY OF BUSINESS,  
INNOVATION & EMPLOYMENT  
HIKINA WHAKATUTUKI



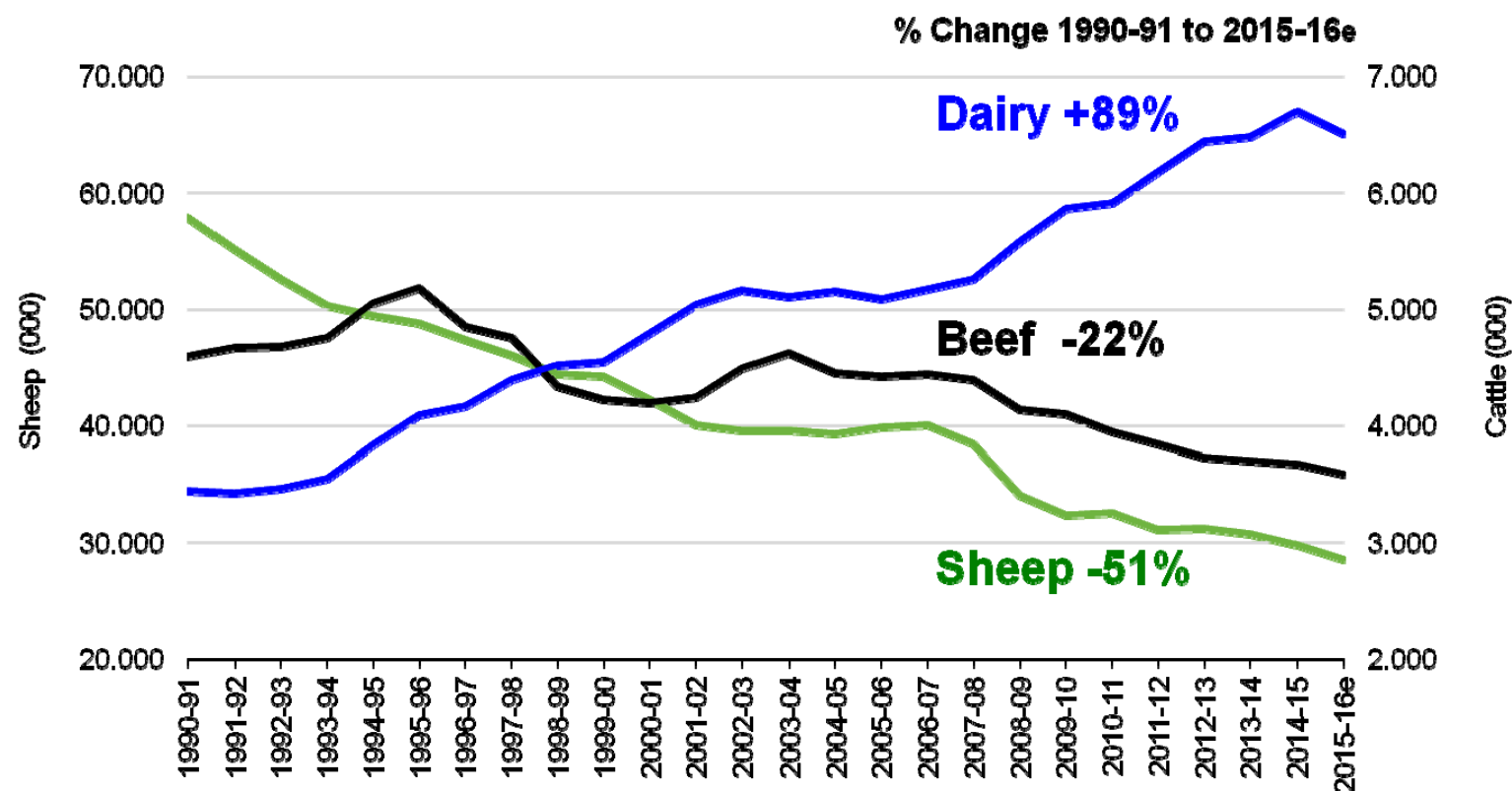
# B+LNZ Genetics



## ***PURPOSE***

***Provide the Information Infrastructure for Breeders, Farmers and Industry to make profitable breeding choices***

# NZ Sheep, Beef & Dairy Numbers



Source: Beef + Lamb New Zealand Economic Service  
Statistics New Zealand

# NZ Production Changes

1990-91 to 2012-13e



MORE  
DAIRY

+175%



MORE BEEF  
AND VEAL

+19%



LESS  
LAMB

-7%

But -51% fewer Sheep



## B+LNZ Genetics: Beef



# Challenges & Opportunities

- Sheep & Beef Farming System
  - Beef provide more value than a lawn mower?
  - Describe & Select Genotypes specifically for NZ?





# Challenges & Opportunities

- NZ Dairy Farming No 1 source of NZ Beef
  - Describe specific Genotypes for Dairy-Beef versus traditional Beef System?



# NZ Beef Genetic Evaluations > Australia





## B+LNZ Genetics: Sheep



# NZ Genetic Evaluation: Scale



	Within-Flock	Across-Flock
Unique animals	14 million	8.3 million
All flocks	1,135	
Current active flocks <sup>‡</sup>	564	
2015 born (NAI*)	330,649	
eBV's stored	22 billion	416 million

*\*NAI = new animal indicators*

*‡Flocks selling rams*

*- Base Year 1995*

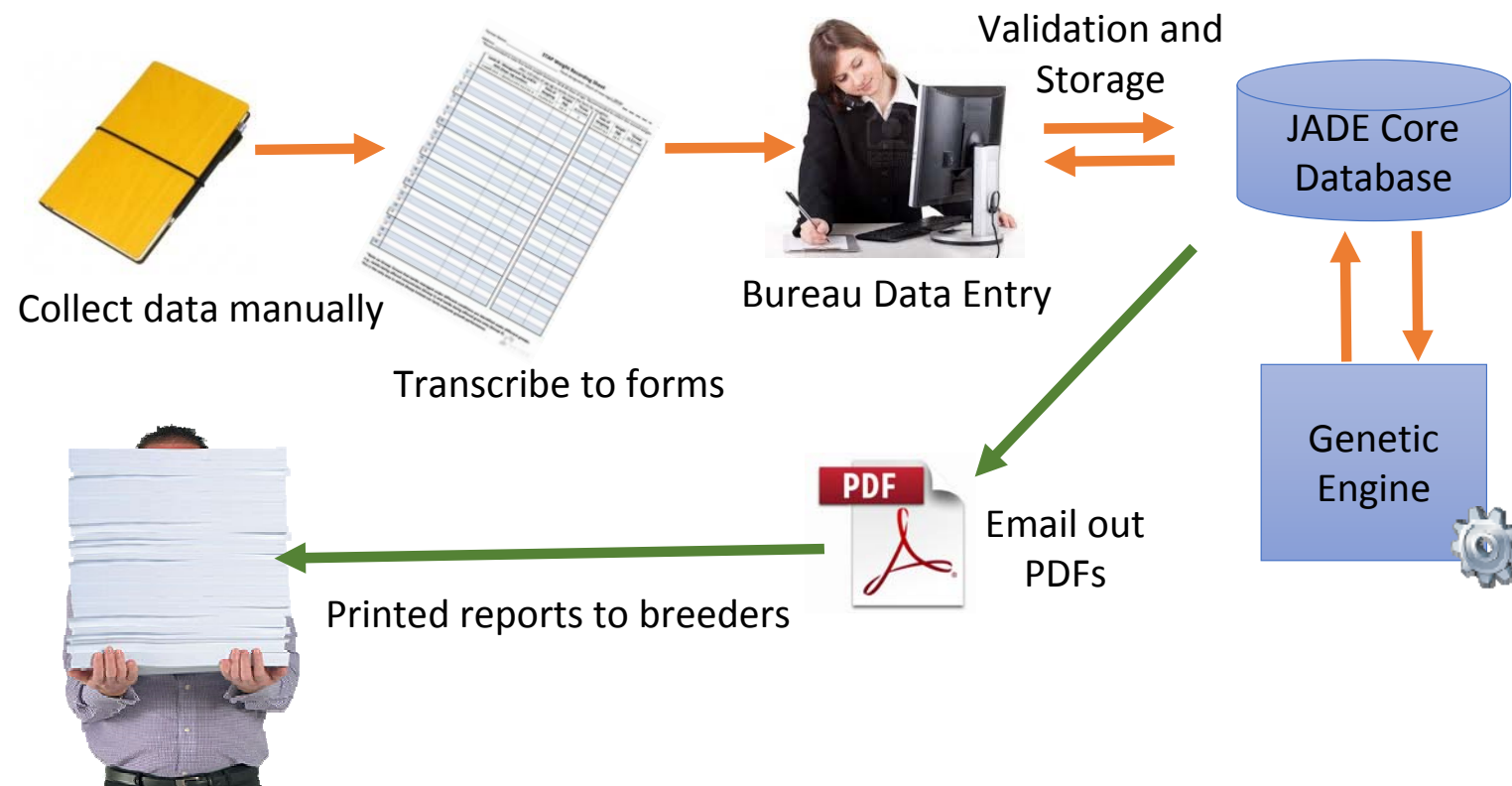
*Figures from Aug 2016*

## 2016: Genetic Engine Upgrade (SIL)

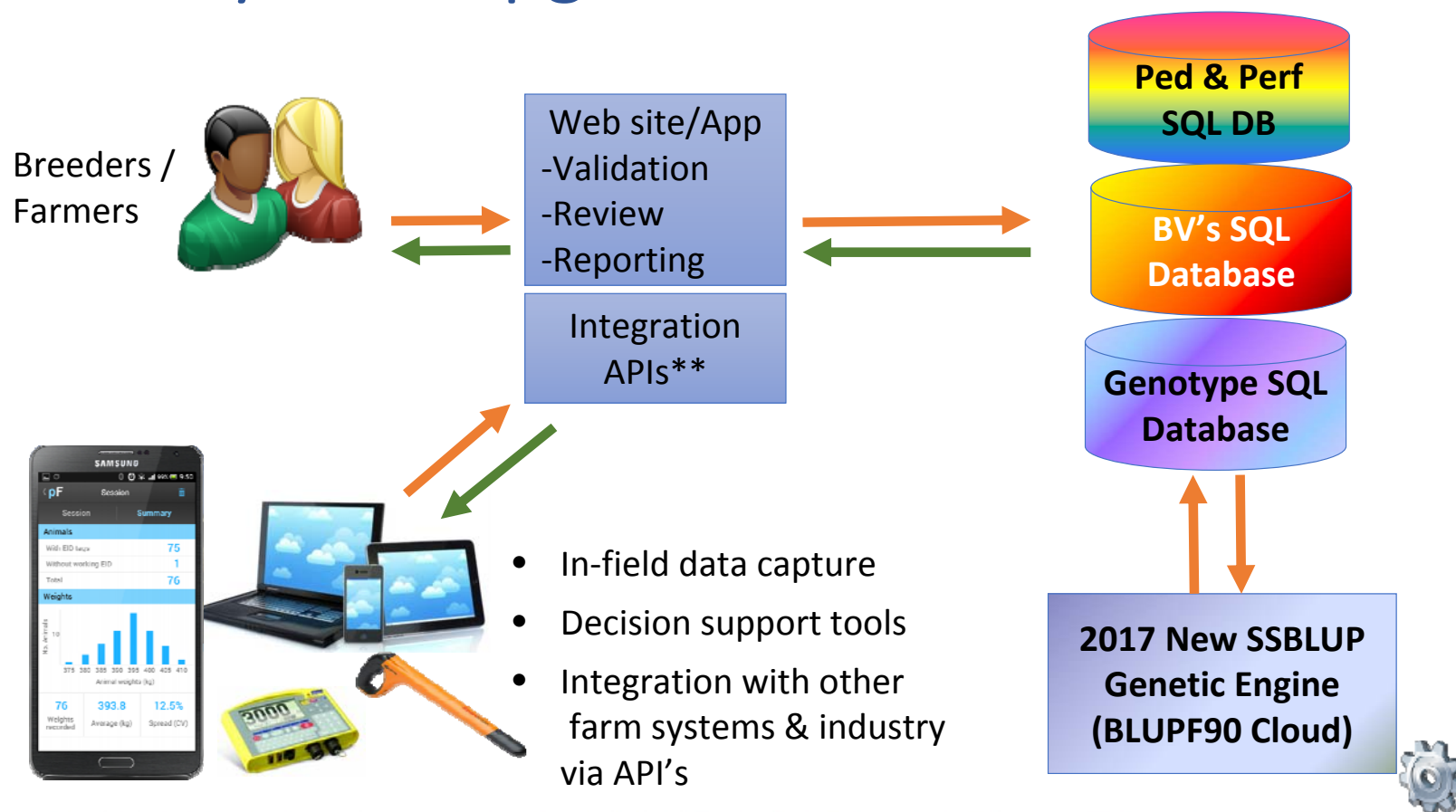
Analysis	Flocks	Animals	ASREMEL	MIX99
Perendale	57	653,826	33 hours	1½ hours
Texel	79	491,988	15 hours	0.5 hours
Coopworth	101	1,612,649	48 hours	4 hours
Multi-Breed Across flock	456	5,348,205	>1 week •multiple computers •simplified models	26 hours
NZGE (Weekly)	1,135	14,387,346	Not Possible	31.5 hours



# GE System Upgrade : Pre – 2016



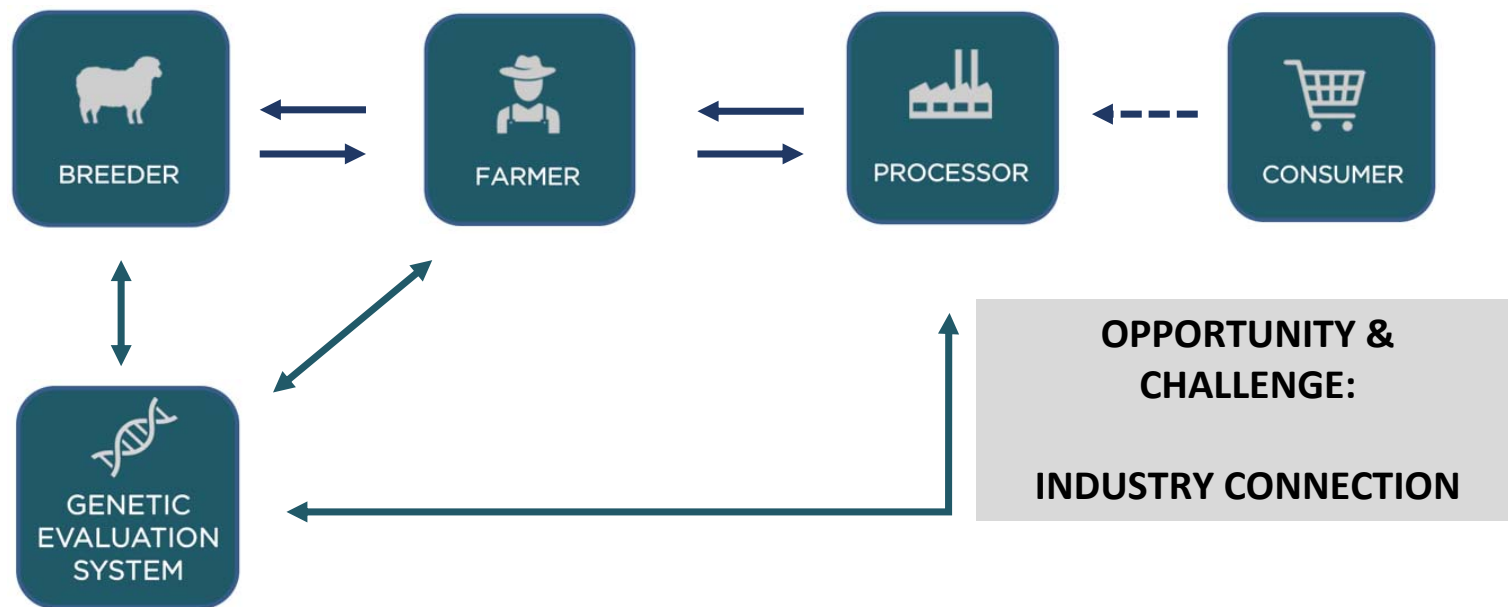
# GE System Upgrade : 2016 -2017



**\*\*Application programming interface (API)**

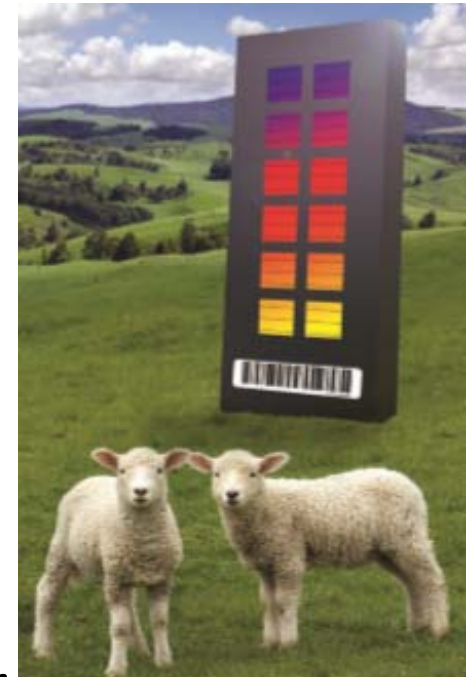
# Genetic Evaluation Data Flow

## CURRENT FLOW OF INFORMATION



# Sheep Genotyping & Genomics

- SNP Parentage
  - Current 80,000 animals / year & growing
  - € 13.00 Euro
- Genomics
  - 36,170 in training: (50k & HD) & 10,000 p.a. genotyped LD
  - € 40.00 Euro
- **Challenge & Opportunities**
  - **Parentage: < € 5.00 Euro**
  - **Genomics: < € 15.00 Euro**
  - **Transition from totally Parentage > Genomics**



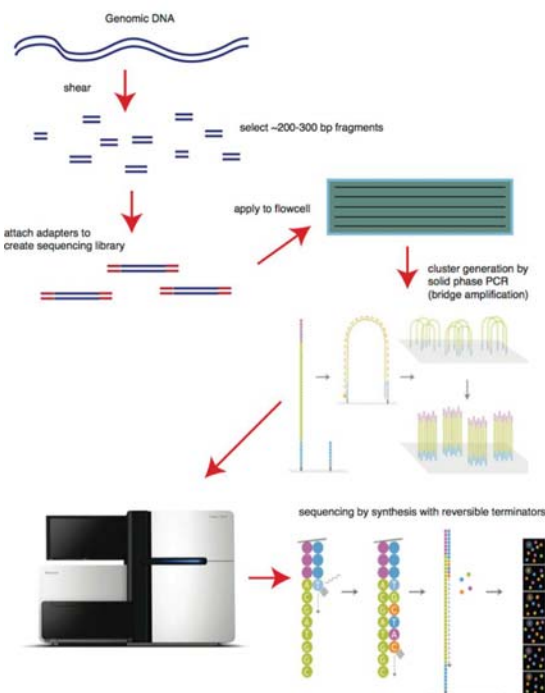
# Sheep Genomic Predictions

Trait		Trait	Romney 2016	Coopworth	Perendale	Composite
<b>Production</b>	Number of Lambs Born	NLB	64%	54%	43%	47%
	Lamb Weaning Weight	WWT	63%	67%	60%	45%
	Weaning Weight Maternal	WWTM	47%	46%	41%	40%
	Live Weight 8 months	LW8	61%	61%	53%	45%
	Live Weight 12 months	LW12	58%	53%	51%	49%
	Carcase Weight	CW	58%	60%	46%	43%
	Ewe Live Weight	EWT	51%	55%	42%	45%
	Eye Muscle Area	EMAc	57%	59%	49%	39%
<b>Meat Yield</b>	Fat Yield	FATY	47%	67%	40%	43%
	Hind Qtr Yield	HQLY	45%	62%	42%	50%
	Loin Lean Yield	LNLY	44%	62%	42%	49%
	Shoulder Lean Yield	SHLY	50%	62%	41%	47%
	Lean Yield	LEANY	47%	62%	42%	49%
<b>Health</b>	Facial Eczema	GGT21	63%		46%	
	Lamb Dag Score	LDAG	48%	62%		59%
	Adult Dag Score	ADAG	52%	58%		53%
	Feacal Egg Count	FEC1	61%	68%	53%	61%
	Feacal Egg Count	FEC2	52%	50%	41%	44%
	Adult Ewe Faecal Egg Count	AFEC	46%	45%	34%	39%
<b>Wool</b>	Fleece Weight 12m	FW12	51%	69%	50%	54%
	Lamb Fleece Weight	LFW	34%	31%	28%	31%
	Ewe Fleece Weight	EFW	42%	26%	25%	27%



# BLG Sheep Genomic Pipeline

1. HD Genotype key Sires with good phenotypes
2. Impute to Sequence
3. GWAS: Causative Mutations & QTL
4. Add SNPs to Panel to improve accuracy for Genomic Selection



**Challenge: ROI on GWAS vs. Phenotypes & Genotypes**

# *Main Areas Sheep Research*

## **Feed Efficacy (RFI)**



## **Meat Yield & Shape**



## **BCS**



## **Maternal Ewe**



## **Meat /Eating Quality**





# Challenge> Knowledge Transfer (KT)

1. KT of R&D outcomes onto Farms
2. Assist Seed Stock producers to increase Genetic Merit of flock/herd



# B+LNZ Genetics + ICAR > Sheep



# Why BLG + ICAR?

- Identifying /implementing key traits of economic value across countries
- Dialog on standardisation / guidelines /codes of practice for the recording of these traits
- The standardisation between countries of nomenclature (IDs, names, units and abbreviations)



# Why BLG + ICAR?

- Sharing/access to hard/expensive to record phenotypes (e.g. RFI)
- Opportunities stimulate across country evaluations/progeny tests and exchange of germplasm
- Exchange and use of data including genomic data for gene discovery and evaluations



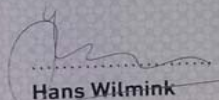
THE GLOBAL STANDARD  
FOR LIVESTOCK DATA

Via Savoia 78, 00198, Rome, Italy

## CERTIFICATE OF QUALITY

### Beef+Lamb New Zealand Genetics

for Identification and production recording in dairy cattle; Laboratory analysis  
(milk samples); Data processing

  
**Hans Wilmink**  
President

Rome, 10 October 2016  
Certificate number: 2016/10  
Valid up-to: April 2018





**The End**