# HIGHLIGHTS

- Farmers wanting to improve rear teat placement can now include this trait in their breeding objective.
- To close the distance between rear teats, select for bulls from the Good Bulls Guide with a Rear Teat Placement ABV more than 100.
- To widen the distance between rear teats, select for bulls from the Good Bulls Guide with a Rear Teat Placement ABV less than 100.

### **Rear teat placement**

Applying a cluster to some cows for milking is difficult because of close or crossed rear teats. In recent years, breed associations have scored the rear teat placement of heifers during classification. This data is used in Rear Teat Placement Australian Breeding Value (ABV) to allow farmers to breed for this trait.

Data for rear teat placement is collected through the classification of first lactation females. Breed association classifiers assess and record rear teat placement on a scale of 1 to 9 based on the position of the rear teats in relation to the centre of the quarter as illustrated in Figure 1.

An analysis of 7,060 Holstein and 1,657 Jersey cows scored in 2010/11 found a phenotypic mean score is 7.1 in Holstein and 6.5 in Jersey, indicating that both breeds tend to be closer to the crossing end of the scale.

# **Rear Teat Placement ABV**

DataGene calculates the Rear Teat Placement ABV from the data collected about an animal's daughters, information from its pedigree and genotype.

Rear Teat Placement ABV is standardised with an average of 100 and standard deviation of 5. The average represents the modern dairy cow.

Rear teat placement is an example of a trait where neither extreme is desirable. Therefore, the direction of expression can be tricky.

The Rear Teat Placement ABV is expressed:

- 1. In the same direction in which the linear trait is measured (higher is closer).
- 2. In the same direction as front teat placement (more than 100 is closer).
- 3. In the same direction as many foreign countries currently express the same trait (more than 100 is closer).

## Breeding for improved rear teat placement

There is a range in rear teat placement, meaning farmers can select for closer or wider rear teats.

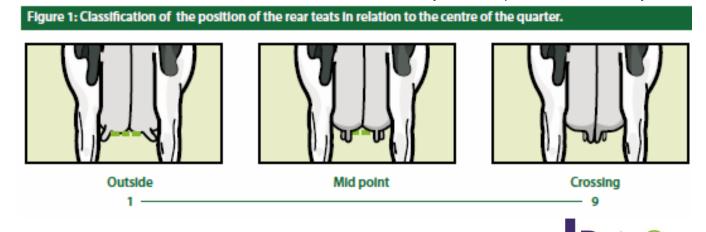
To widen the distance between rear teats, select bulls from the Good Bulls Guide with an ABV less than 100.

An ABV of 95 means 1 standard deviation below average (wider rear teats). About 34% of bulls will have an ABV between 95 and 100 as illustrated in Figure 2.

#### Heritability

Heritability is an estimate of the level of genetic control for a trait. Genetic gain is faster in high heritability traits compared to low heritability traits.

olutions for Herd Development



The heritability of Rear Teat Placement is 0.25, meaning that it is moderately heritable and the pace at which changes can be made is similar to traits such as fore and rear udder attachments.

### Correlations

In dairy cattle breeding, most traits have a relationship with each other. For example, there is a positive relationship between protein kilograms and milk litres. More milk generally means more protein.

These relationships are known as correlations and are reported in a range between -1 to +1. In the case of rear teat placement, the genetic correlation between rear teat placement and front teat placement is strongly positive at 0.71. This means that as you breed for closer front teats there is a strong tendency to have closer rear teats as well, but this is not always the case, as such the Rear Teat Placement ABV will provide direct information to farmers about this trait.

Correlations between rear teat placement and other udder traits are shown in Table 1.

Table 1: Genetic correlations between rear teat   placement and other udder traits		
	Genetic correlation	Description of the relationship
Front Teat Placement	0.71	Strongly positive
Centre Ligament	0.81	Strongly positive
Mammary System	0.61	Strongly positive
Udder Depth	0.07	Very little relationship

Farmers wanting to improve Rear Teat Placement can now include this trait in their breeding objective.

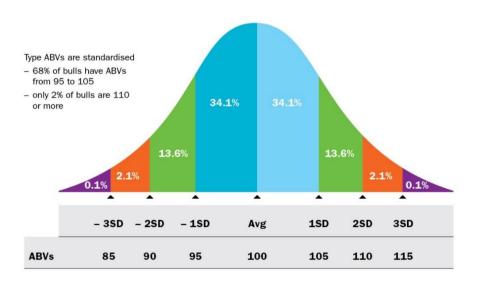
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## For more information

Genetics Backgrounder: Type ABVs explained

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