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# **Advances in breeding technologies: *an inclusive approach***

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# Presentation Outline

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- Background
  - Beef industry outlook (2018 – 2024)
- Beef industry structure
- Breeding goals
- Genomics
- Concluding remarks



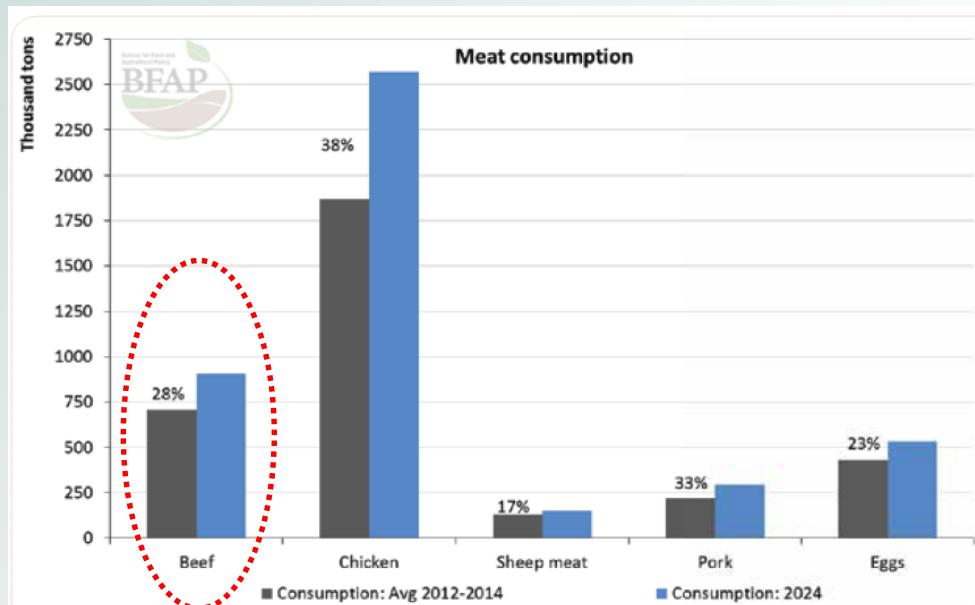
# Beef Industry Outlook – *BFAP 2015 -2024*

- The demand for beef will increase resulting in increase in:
  - Beef consumption (**28%** increase by 2024)
  - Beef auction price (**R57/kg** by 2014) i.e. *annual increase of 5.7%*

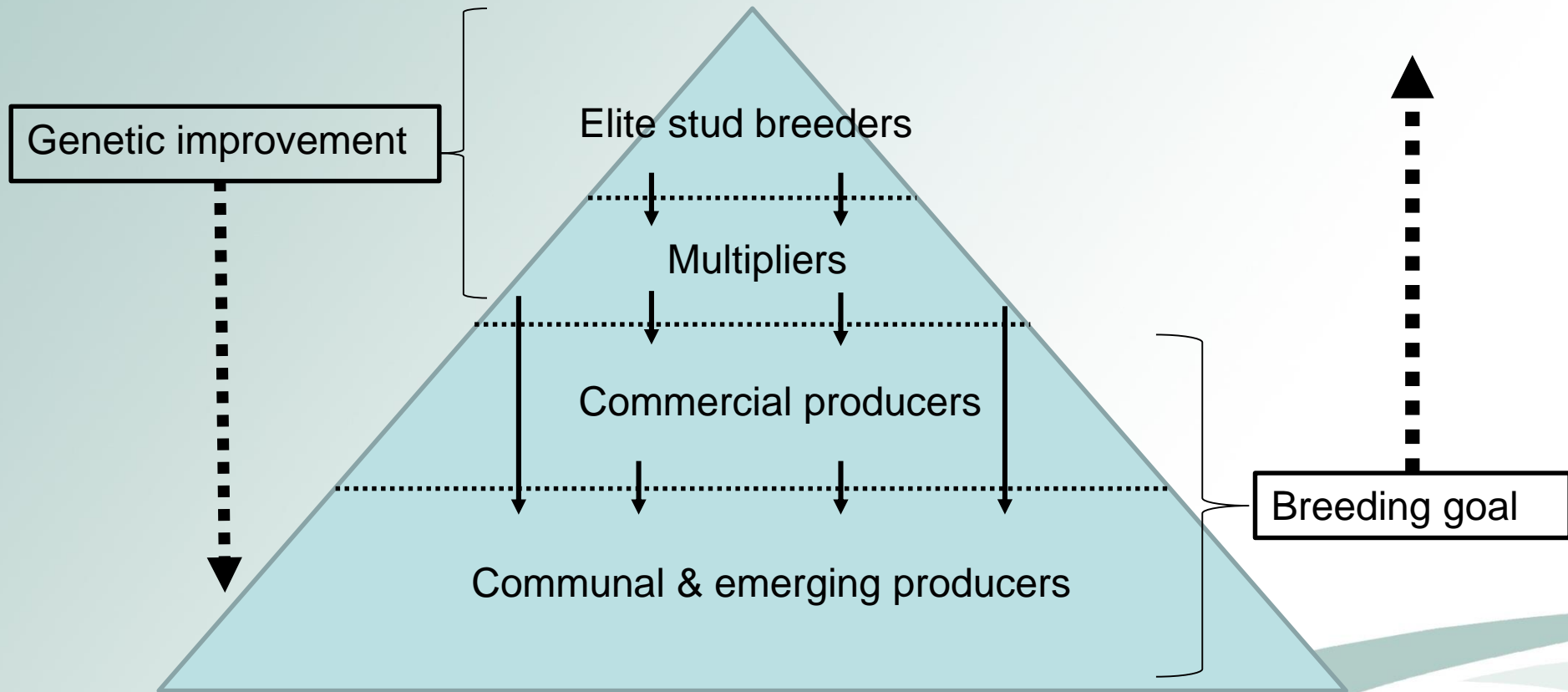
There is a need for «**more beef production**» from the «**same or less**» resource base

Improved efficiency

- Number of cows
- Number of hectares
- Quantity of feed
- etc.

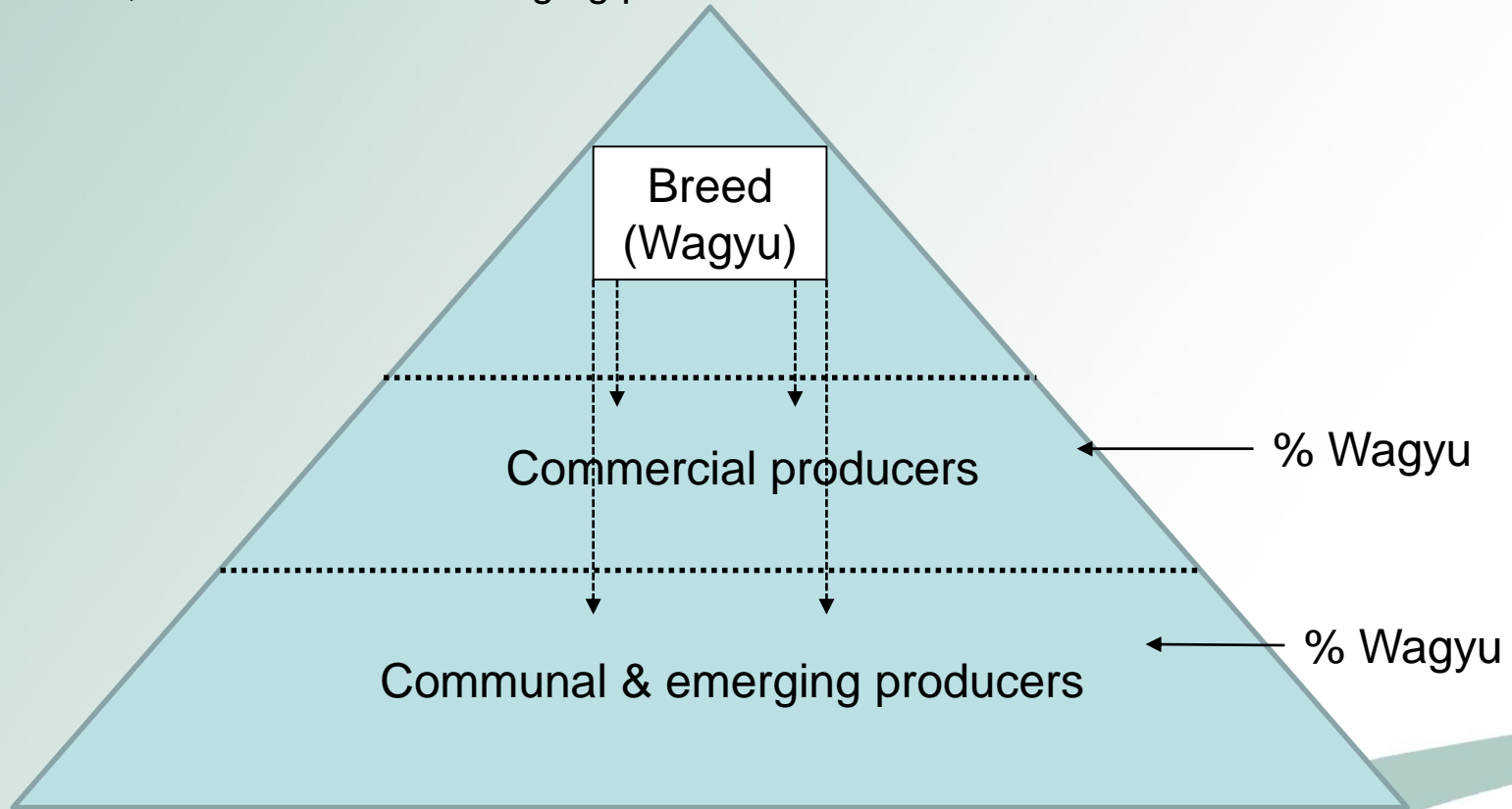


# Beef Production Asset Base



# Breed genetic impact on the national herd

Breed success depends on → compatibility of the **breeding goal** with the **needs** of the commercial, communal and emerging producers



# Needs of commercial LARGE AND SMALL SCALE producers

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- **Desirable «production» cow** (*cow* → *calf*)
  - Produce a «**calf every year**» unassisted
  - Low «**feed requirements**»
  - «**Adapted**» to the production environment i.e. *management and natural environment*
  
- **Desirable product** (*calf* → *beef*)
  - Well «**adapted**» to the production environment
  - Produce more kg of «**high quality beef**» at the «**shortest time possible**» from «**low feed requirements**»

→ **Setting a breeding goal is important**

# Conventional genetic improvement

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- Animal performance depends on:
  - genetics / DNA
  - environment / management
- Genetic evaluation and improvement relied on:
  - Measuring performance (performance testing)
  - Calculate genetic merit (Estimated Breeding Value – EBV) using performance records (*no technology was available to comprehensively read the DNA*)
  - Large number of performance records were required for effective genetic improvement particularly for «lowly heritable traits»

# Bovine Genome Sequencing

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- Bovine genome was successfully sequenced in early 2000s
- A large number of DNA markers called SNPs were discovered throughout the genome
- SNPs allow us to discern differences among animals at DNA level



SNP chip – genome reader



# New genetic improvement paradigm

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- Genetic evaluation and improvement paradigm:
  1. Establish a reference population (>1 000 animals)
    - Animals with performance records and genotyped using SNP chip (genome reader)
  2. Estimate genetic merit (GEBV) using performance and genomic profile
- GEBVs are more accurate particularly for young animals with little or no performance records compared to conventional genetic evaluation
- Implication:
  - Selection can be done earlier in life → reducing selection cycle and increasing the rate of genetic improvement
  - Reduction in the cost of performance recording compared to conventional genetic improvement

# Is genomic selection possible in SA?

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- A National Beef Genomic Project was initiated in 2015 through a partnership between industry and government (including Namibian beef cattle breeds) aimed at:
  - Setting up reference population for different cattle breeds
    - Collecting performance records on traits of economic importance
    - Genotyping animals using SNP chip technology
  - Conducting research in order to build research capacity of genomics nationally
- To date, ~ 3000 animals were genotyped and ~40 animals were sequenced
- The first genomic evaluation in beef cattle was implemented in 2017

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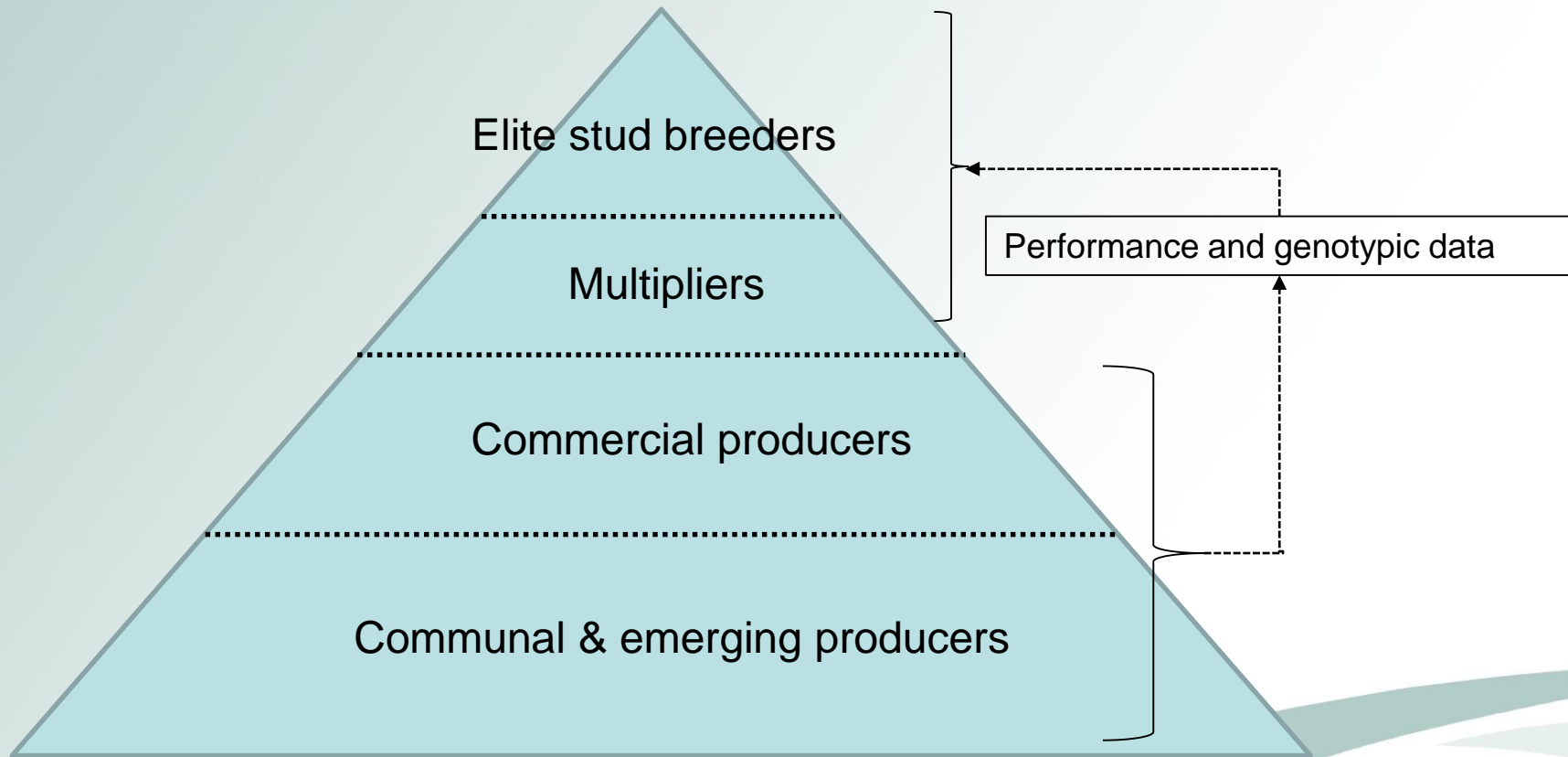
# More benefits of genomic selection

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- Parentage verification
  - Genomic profiles from SNP chip could be used for both parentage verification and genomic selection
- Performance data on commercial animals could be used for genomic selection
  - Carcass data on slaughter progeny of stud bulls
- Identification of genetic defects

# Genomics and Beef Industry Structure

Enhance cooperation between stud breeders and commercial producers



# Summary

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- New breeding technologies (genomics) provides an opportunity to enhance efficiency of production
- Development of breeding objectives is more important in genomic era since in advent mistakes might take longer to fix
- Recording of performance data is key to implementation of genomic selection

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# Thanks for Your Attention

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