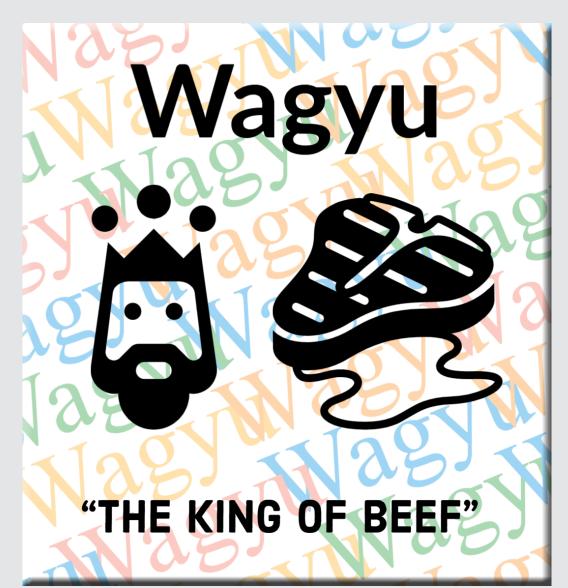
# **Breeding Plan - Fullblood**

## **Breeding Plan**

### **Fullblood herd:**

### 1. Focus 1st on Marbling and Marble Score:

- This is the profit driving trait of Wagyu
- You need a salable carcass that has marbling to generate a premium, other wise you might as well be breeding Angus.



Written and compiled by: Mitchell Ruth

#### 2. Focus 2nd on your choice of a few traits equally justifiable:

#### • Carcass Weight

i. Larger carcass weights allow for maximizing efficiencies of slaughter costs & processing, feed bunk/ feedlot space, and steady beef supply

#### • Rib Eye Area

i. The largest source of revenue from the carcass, drives carcass profit ability

### • Marble Fineness

i. Fineness of marbling drives premiums and appeal to high end chefs and customers

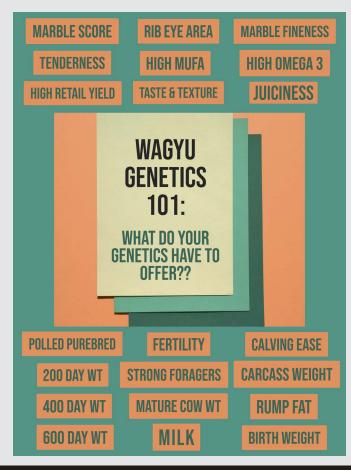
ii. Fineness of marbling drives flavor; the more fine marbling in meat means there is more marbling cell wall which while cooked creates Wagyu's unique flavor.

### • Residual Feed Intake

i. Feed costs commonly account for more than half the cost of produc tion from birth to slaughter.

### • Growth traits, 200 Day, 400 Day, 600 Day Weights

i. Reducing DOF (Days On Feed) is a key factor in remaining profitable and competing with high marbling Angus and other breeds



#### How to accomplish gains in these areas:

#### 1. Use Australian Breedplan

• Largest Wagyu EBV/EPD database outside of Japan

• More than 84,000 dams and 10,000 sires recorded, 6,700 AUS Meat Marble Score recorded, 5,800 Carcass Camera recorded carcasses (MS, MF, REA), 28,000 Weaning Wts, 24,000 400 Day Wts

• Now offers Genomic Testing based on all this data for traits including MS, MF, CW, REA, 200D Wt, 400D Wt, and 600D Wt

• 50K SNP profile based genomics (Cattle Industry Leading Technology)

• Highlights strengths and weaknesses of cattle numerically allowing for easier management and mating

## 2. Use/Invest in high reliability sires backed by actual carcass data (>80% Reli ability for MS)

- Or a variety of sons of elite high reliability bulls
- Young sires should be genomic tested to reduce risk and increase reliability

#### 3. Maintain or invest in a variety of Maternal lines or Cow families

- Use high proven female maternal lines: think Suzutani, Okutani, Yuriko, Chisahime, Hikokura, etc. This minimizes your risk of failure carcasses
- Different maternal lines have different strengths that can be complimenting and successful crosses
- Increases your herds marketability (Seedstock)
- Manages inbreeding



#### Hikokura Maternal Line Sires Sires to Explore

Coates Itoshigenami G113 Mayura Itoshigenami Jr Sumo Cattle Co Michifuku F154 Sumo Cattle Co Michifuku F126 Westholme Fujiteru 3 Sumo Cattle Co Itoshigenami C158 Tamarind T4 Kanadagene 14/2 CHR Takazakura 101L CHR Kitaguni 07K Terutani 40/1 Itomichi 1/2 Mitsuhikokura 149



#### 4. Genomic Test your entire herd of Fullblood/purebred cattle

- Identifies top and bottom of your herd
- Identifies strengths and weaknesses of individual animals
- Adds value by creating higher accuracy EBVs for all cattle tested

## 5. Use Corrective mating, breeding complimentary cattle to one another to eliminate flaws or weaknesses.

- Use carcass bulls on females with size and growth, and maternal/growth bulls on females that need size and milk.
- Evaluate matings individually, often times Wagyu take 2-3 generation of carcass bulls on females that retain size and growth well like the Hikokura maternal line.

## **Common Mating Decision Strategies**

- Random Mating: Turn bulls out breed anyone, no rhyme or reason
- Mating based on inbreeding coefficient
- Corrective Mating:
  Dairy i.e. mating services
- · Linebreeding: Consolidation of desirable traits
- · Blanket AI or Natural Service: One sire on all cows
- Terminal Mating:
  - Sexed male semen
  - Carcass traits emphasized
  - $\cdot\,$  No consideration to maternal traits, etc.
- Replacement AI:
  - Sexed female semen –
  - Focused on key replacement heifer traits
  - Less consideration to carcass traits



#### 6. Make separate matings for Terminal use and Replacement use.

• Terminal matings should focus more on key carcass traits (MS, MF, REA, CWT)

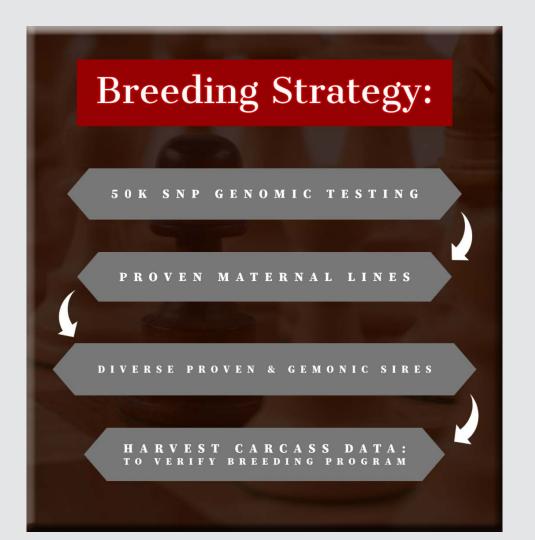
• Replacement matings should focus on growth and maternal traits

### 7. Plan on killing animals to verify the genetics you have and are breeding

• Create sizeable contemporary groups for meaningful data (>10 head, same sex, 2 common reference sires, born in 3 week window, and must be fed and slaughter in the same system for the same period of time)

• Steer 80% of bulls, to prove the value of your top 20% remaining bulls each year

- Creates value thru carcass data
- Shows your cattle's strengths or weaknesses
- Allows for creating better matings in the future
- Submit your carcass data to Breedplan to increase your EBVs accuracy



#### 8. All of this will allow you to share and compare your Wagyu Cattle to other Elite breeders around the USA and Globe.

• Either your cattle will prove themselves to be elite, average, poor

• This allows you cull poor ones, better utilize the average, and flush or market the very elite cattle for more value.

• It also allows for you to identify better or complimenting genetics that can be bought or acquired to improve your herd.

Other Useful Data/Thoughts: What data says I should do all this?

## The Bright Future of Wagyu

- Increasing Global Demand for Premium Beef
- Increasing Buying Power of Global Middle Class
- Genomics/ GEBVs
  - Reducing DOF
  - Continued Progress in Carcass Traits
- Increased Availability of Elite Semen & Genetics
- Strong Diversity of Genetics
- EBVs Make Pedigree Reading Easier

### Proper Strategies for Utilizing Genomics:

#### • Whole Herd:

- Identify the top and bottom of the herd
- Donors & Recipients

#### • Within Herd:

- Individual Selection & Decision Making
- · Comparison of Siblings & Flush mates
- Identify Individual Strengths/Weaknesses

#### National/International Level:

- Top Sire Selection
- Top Females Selectin
- Buy or Acquire new or complimentary genetics



## Mating My Donors

- · What Should I Consider?
  - · Carcass Data (if available): Strengths, Weaknesses
  - · Genomics: Weaknesses, Strengths
  - **Pedigree**: Inbreeding, Complimentary genetics, Linebreeding
  - Phenotype: Strength, Faults/Corrections needed
  - Genotype: Recessives, Exon 5, SCD, etc.

#### Sire Factors:

- Price
- · Availability
- Reliability
- Above factors



Mayura L0010



World K's Michifuku



TF Itohana 2

### "Cheap" or Inferior Genetics

 "Don't be fooled by Cheap or Inferior Genetics, they will have long lasting effects in your herd." – Scott de Bruin 2018 AUS Wagyu Edge Presentation

#### Long Lasting Effects:

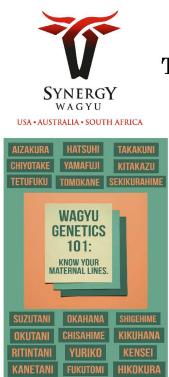
- \* Their Steers 3yrs+ from conception to harvest
- Their Daughters Replacements: 3yrs+ from conception until calving
- \* Their Daughters Daughter's 5 yrs + if retained for breeding
- **Directly Impact Profitability**: Limit the Potential Performance of Animals

### Why Use **Proven** Genetics??

#### • First What is Proven?

- Gold Standard: Performance recorded data, Objective 3<sup>rd</sup> Party Carcass Data, Breedplan data backed proof, etc.
- Next: inhouse performance & carcass data
- Last: Opinion i.e. "I killed some and they looked great"
- Consistency & Reliability of Outcomes:
  - Risk management
  - In reality a son is rarely better than his sire/father!
  - Need a saleable carcass





### The Importance of Maternal Lines

1) Risk Management

2) Females Unique Ability to transmit to offspring

3) Marketability

4) Genetic Merit

### Female Bloodlines are Important!

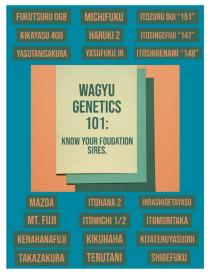
#### • Aka Cow Families or Maternal Lines

- · Female contributes half the DNA, just like the sire
- · Often overlooked in carcass results & data evaluation

\* i.e. Itomichi ½ x Mayura Itoshigenami JNR steers will likely perform far better than Itomichi ½ x World K's Haruki2 steers

- \* May skew your perspective on how good Itomichi  $\frac{1}{2}$  is, Why EBVs are so valuable
- Mitochondrial DNA inheritance only coming from the dam
  https://www.ajas.info/upload/pdf/17\_243.pdf
- Cytoplasmic inheritance theories
  https://www.sciencedirect.com/science/article/pii/S0022030286807731
- · Epigenetics triggered in utero by the dam
  - <u>https://epigeneticsandchromatin.biomedcentral.com/articles/10.1</u> 0081-5

## Wagyu Bloodlines:



**Prefectural Percentages**: 16/16 Analysis: Tajima, Itozakura, Kedaka, etc

-Don't get caught up on these

-There are high marbling Tajima, Shimane, and Kedaka. -There are high growth Tajima, Shimane, and Kedaka.

**Instead classify sires on traits**: Carcass (MS, MF, REA) specialists, Growth & Maternal specialists, or a Balance of both.

Prefectural Percentages are most useful for tracking inbreeding but Inbreeding coefficients do a better job

Essentially they have become obsolete with EBVs

Tajima	Kedaka	Tottori	Itozakura	Shimane	Okayama	Hiroshima	Other	TOTAL
9.4	1.6	0.5	2.8	0.4	1.1	0.3	-	16



# The **Danger** of Breeding with Indexes, Single Traits, Etc.

- Potential Loss of Traits
  Example: Holstein Dairy Cattle
- Potential to Amplify Weaknesses
  Example: Guernsey Dairy Cattle
- No Corrective Mating
- Inbreeding not controlled





1) AS A BREEDER/FEEDER IT IS YOUR JOB/RESPONSIBILITY TO MAKE DECISIONS AND FIGURE IT OUT.

2) AFTER ALL, IT IS YOUR INVESTMENT!

