

Priority Setting for Conservation of Animal Genetic Resources

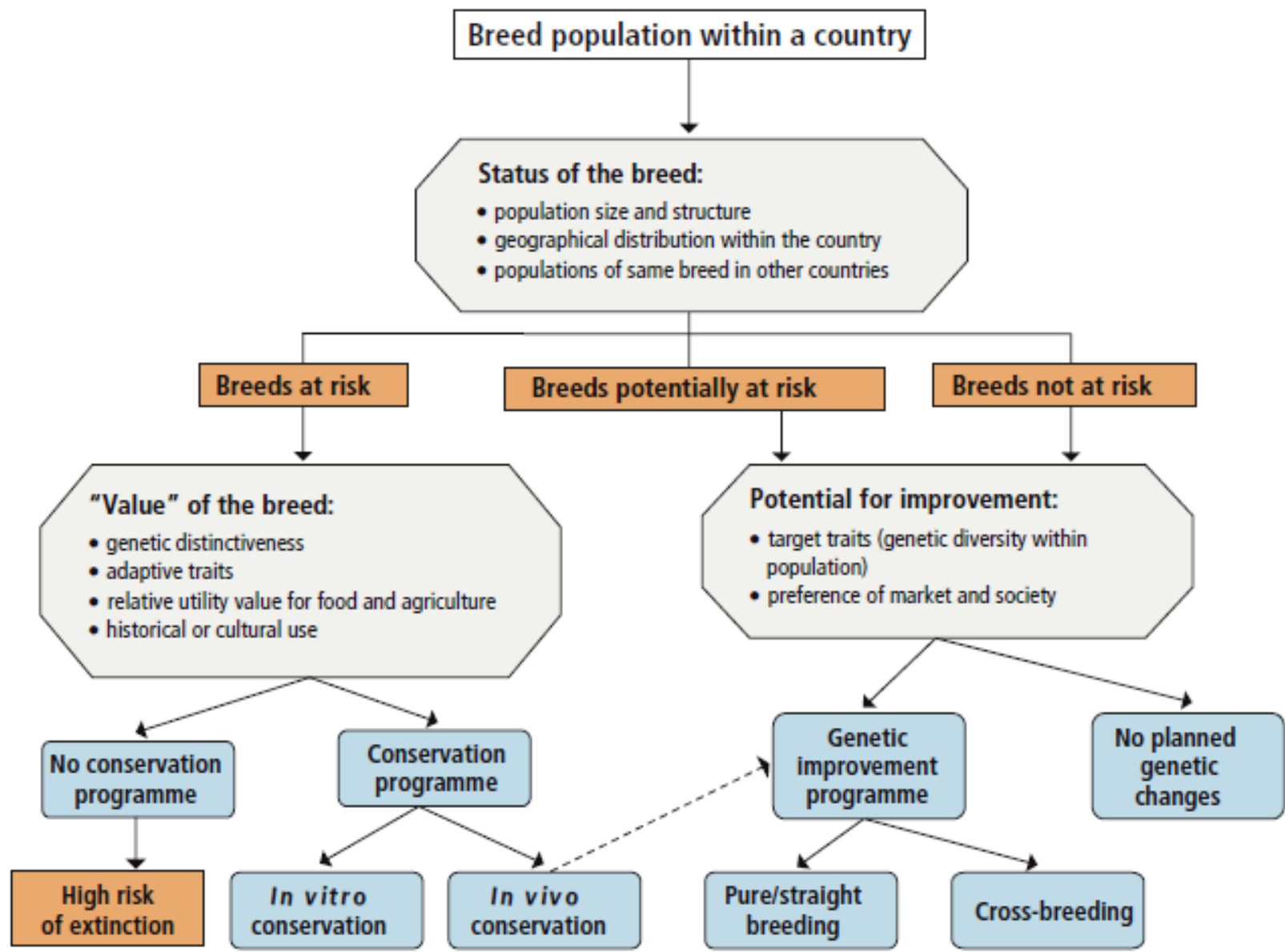
Paul Boettcher
FAO-AGAG
Rome, Italy

Conservation Planning

- For national management of AnGR, comprehensive planning and organization are critical
 - Ensure most valuable AnGR are maintained
 - Increase efficiency
 - Consider interests of all stakeholders
- For conservation, critical step in this process is identifying breeds at risk
 - Based on survey and monitoring

Prioritization

- Once breeds are assigned to risk classes, *prioritization* of breeds should be undertaken
- Conservation of all breeds impossible
 - Many breeds and limited financial resources
 - Food security more important in developing countries
 - Short-term economic return more important in industrialized countries
- Wholesale conservation not scientifically justified
 - Breed may have no apparent short- or long-term value
 - Breeds may be effectively the same genetically



Factors Influencing Conservation Priority

- Risk of extinction
 - Breed Demographics
 - number and distribution
- Genetic variability (seek to maximize)
 - between and within breeds
 - molecular and quantitative genetic
- Phenotypic performance
 - genetic merit for productivity

Factors Influencing Conservation Priority

- Unique traits
 - adaptive traits
 - interaction with environment
- Historical and cultural importance
- Practical considerations
 - species and ease of conservation
 - chance for success

Risk of Extinction

- In most cases, risk of extinction has been used as the primary criterion for conservation
 - Allows for immediate action
- Sometimes, simple qualifiers are used
 - Recognized as a local breed
 - Not available in other countries
- When circumstances (time) allows, consideration of other factors may be more efficient
 - Store the most valuable resources

Accounting for Non-Demographic Factors

- *Action 1. Assign responsibilities and agree on the conservation strategy*
 - National Advisory Committee on AnGR
 - FAO Guidelines on **Preparation of National Strategies and Action Plans for Animal Genetic Resources**
 - <ftp://ftp.fao.org/docrep/fao/012/i0770e/i0770e.pdf>
 - Expert task force
 - Involve key stakeholders

Assessing Conservation Value

- *Action 2. Determine the factors upon which the conservation value will be based*
 - Consider conservation objectives
 - Values of importance
 - direct use and option values most common
 - Factors affecting values
 - Propose strategy (Bennewitz et al., 2007)
 - Maximum risk
 - Extinction risk only
 - Maximum diversity
 - High weight on genetic diversity
 - Maximum utility
 - Multiple factors
 - Determine relative importance of factors
 - Ranking
 - Assign weights

Assessing Conservation Value

- *Action 3. Gather the information necessary to determine the conservation value*
 - Ideally, characterization will have already been completed
 - Undertake characterization
 - Consult stakeholders
 - Productivity, uniqueness, cultural value
 - Genetic variability
 - Molecular markers
 - Pedigree
 - Population history

Assessing Conservation Value

- *Action 4. Discuss and evaluate the advantages and disadvantages of the breeds*
 - National AnGR Advisory Committee
 - SWOT analysis
- *Action 5. Rank breeds for conservation value*
 - Subjective
 - Objective

Subjective Prioritization

- Following the gathering and discussion of information needed to determine the value of breeds, priority order may be determined in a simple approach
- Voting by committee
 - Determine highest priority
 - Continue down list until all breeds are ranked
- Ranking by committee members
 - Order based on average ranking

“Objective” Prioritization: Index

- Assign each breed a value for each factor
- Each value considered should be expressed numerically
 - Breed average for quantitative traits
 - 1/0 for absence presence of special trait
 - Ranking or score for cultural value
 - \uparrow value = \uparrow number
- Calculate mean and SD for each factor to perform standardization
 - For quantitative traits, mean and SD are species specific
- Suggestion: calculate means and SD across risk classes but prioritize within risk class
 - Or include risk as a factor in the index

Conservation Value Index

$$CV_i = w_{F_1} \times (F1_i - \mu_{F_1})/\sigma_{F_1} + w_{F_2} \times (F2_i - \mu_{F_2})/\sigma_{F_2} + \dots + w_{F_n} \times (F_n_i - \mu_{F_n})/\sigma_{F_n}$$

- where, for

$$w_{F_1} \times (F1_i - \mu_{F_1})/\sigma_{F_1}$$

- w_{F_1} = is the weight (i.e. relative importance) of Factor 1 (e.g. milk yield)
- $F1_i$ = is the value for Factor 1 for Breed i
- μ_{F_1} = is the average of all breeds for Factor 1
- σ_{F_1} = is the standard deviation of all breeds for Factor 1

Conservation Value Index

Hypothetical: Factors considered important for conservation and data per breed

	Effective	Genetic	Milk yield	Cultural
	population size	uniqueness	(kg/yr)	importance
Breed 1	60	2	1000	0
Breed 2	100	3	700	0
Breed 3	50	1	500	1
Overall mean	70	2	733.33	0.33
Standard deviation	26.46	1	251.66	0.58
Weight in index	3	1	2	1

Conservation Value Index

Two measures of genetic variability

Hypothetical: Factors considered important for conservation and data per breed

	Effective population size	Genetic uniqueness	Milk yield (kg/yr)	Cultural importance
Breed 1	60	2	1000	0
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Conservation Value Index

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	Effective population size	Genetic uniqueness	Milk yield (kg/yr)	Cultural importance
Breed 1	60	2	1000	0
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Breed 3	50	1	500	1
Measured variables				
Overall mean	70	2	733.33	0.33
Standard deviation	26.46	1	251.66	0.58
Weight in index	3	1	2	1

Conservation Value Index

Hypothetical: Factors considered important for conservation and data per breed

	Effective population size	Genetic uniqueness	Milk yield (kg/yr)	Cultural importance
Breed 1	60	2	1000	0
Breed 2	100	3	700	0
Breed 3	50	1	500	1
		subjective variables		
Overall mean	70	2	733.33	0.33
Standard deviation	26.46	1	251.66	0.58
Weight in index	3	1	2	1

Conservation Value Index

	Breed 1	Breed 2	Breed 3
<i>Standardized values</i>			
Effective Pop'n Size	-0.38	1.13	-0.76
Genetic uniqueness	0	1	-1
Milk yield	1.06	-0.13	-0.93
Cultural importance	-0.58	-0.58	1.15
<i>Weighted values</i>			
Effective Pop'n Size	-1.13	3.40	-2.27
Genetic uniqueness	0	1	-1
Milk yield	2.12	-0.26	-1.85
Cultural importance	-0.58	-0.58	1.15
Conservation value	0.41	3.56	-3.97
Rank	2	1	3

Formal Prioritization

- Various methods have been proposed to consider extinction risk, genetic diversity and possibly other factors together in prioritization
 - “Weitzman” Method (Weitzman, 1992 and 1993)
 - Reviewed by Boettcher et al. (2010)
- Yield a single numerical value as the criterion for prioritization
- Generally consider molecular-based measures of diversity
- Technically sound, but complicated
 - To now, primarily used in research

Formal Prioritization with Molecular Diversity

- Objective
 - Prioritize breeds objectively by considering all factors simultaneously
 - Use molecular data to account for genetic diversity in conservation
- Inputs
 - Conservation objectives to be addressed
 - List of breeds at risk
 - Information about factors that affect conservation value
 - Molecular genetic information
- Outputs
 - Analysis of the genetic diversity of breeds
 - List of breeds prioritized for conservation

Prioritization according to Total Utility

$$U_i = 4 \times (\text{risk}_i \times D_i) + \text{SCV}_i \quad (\text{Simianer et al., 2003})$$

where,

- **U_i** is the total utility for breed I
 - *Criterion for prioritization*
- **4** depends on importance of conservation value
- **risk_i** is the risk of extinction for breed i
- **D_i** is the contribution of breed i to the overall genetic diversity of the collection of breeds
- **SCV_i** is the standardized conservation value of breed i
 - *based on conservation index*
 - *without genetic diversity factors*

Accounting for Extinction Risk

- Option 1: Prioritize within risk category and use an arbitrary constant for risk for all breeds
 - Risk becomes first criterion
 - No refinement within risk category
 - risk with 50 breeding females = risk with 100
- Option 2: Assume risk is equal within risk category and assign reasonable values to each
 - Critical \leftarrow 0.50
 - Endangered \leftarrow 0.25
 - Vulnerable \leftarrow 0.10
- Option 3: Estimate numerically the extinction risk for each breed

Estimation of Extinction Risk

Three Options:

- Base estimates on breed status for various factors related to risk
- Predict extinction based on past data and modeling
- Estimate loss of genetic variation over time

The final two methods require past time-series data

- More difficult in developing countries

Estimating Extinction Risk (Reist-Marti et al. 2003)

- Choose factors affecting risk:
 - 1) population size = p
 - 2) change in population size = c
 - 3) geographic distribution = g
 - 4) presence of breeding programmes = b
 - 5) farmer satisfaction = f

Estimating Extinction Risk (Reist-Marti et al. 2003)

Establish ordered categories for each factor and assign values to breeds according to effect on risk

- $p \sim$ population size
 - $s = 0.0$ if population size is $\geq 100,000$
 - $s = 0.1$ if population size is between 10001 and 100 000
 - $s = 0.2$ if population size is between 1001 and 10000
 - $s = 0.3$ if population size is $< 1\ 000$
- $g \sim$ geographical distribution
 - $g = 0.0$ if the breed is found in locations across the country
 - $g = 0.1$ if animals tend to be found in a single specific area of the country

Estimating Extinction Risk (Reist-Marti et al. 2003)

- Sum values across factors to determine risk for a given breed

$$\text{risk}_i = p_i + c_i + g_i + b_i + f_i + 0.05$$

- Values should be chosen to not exceed 1.00
- 0.05 is to ensure non-zero results

Genetic Diversity

- Different countries may have different objectives for maintenance of genetic diversity
 - Maintain distinct breeds
 - Emphasize distinctiveness
 - Preserve allelic combinations
 - Conserve specific alleles
 - Avoid loss of rare variants
 - Capture maximum diversity
 - Emphasize within-breed genetic variation

Genetic Diversity

- Optimal method to measure diversity will depend on the objective
 - Maintain distinct breeds → genetic distance
 - Conserve specific alleles → allelic diversity
 - Capture maximum diversity → kinships
- Decision can be simplified as a trade-off between within- and across-breed diversity
- In most cases, an intermediate method may be the best choice
 - Piyasatian and Kinghorn (2003) – 5 to 1 ratio
 - Bennewitz and Meuwissen (2005) – 2 to 1 ratio

Genetic Diversity

- “Maximum-Variance-Total” approach to diversity
- Calculate marker-based kinship → matrix **K**
- vector (**c**) of contributions to a “core set”

$$\mathbf{c} = \frac{1}{4} \left(\mathbf{K}^{-1}\mathbf{F} - \frac{\mathbf{1}'_N \mathbf{K}^{-1}\mathbf{F} - 4}{\mathbf{1}'_N \mathbf{K}^{-1}\mathbf{1}_N} \cdot \mathbf{K}^{-1}\mathbf{1}_N \right)$$

Four-Breed Example

$$c = [0.58 \quad 0.13 \quad 0.00 \quad 0.29]$$

- Breed 1 is the most important for genetic diversity
- Breed 3 is not important
 - given that 1, 2 and 4 are present
- In the total utility equation, insert c_i as D_i
 - e.g. for breed 1, $c_1 = D_1 = 0.58$

Assign Conservation Value

- Use Conservation Value Index described earlier
- Standardize to fall between 0.1 and 0.9

$$SCV_i = 0.1 + [0.8 * (CV_i - CV_{min}) / (CV_{max} - CV_{min})]$$

- CV_i = conservation value of breed I
- CV_{min} = value of breed with smallest CV
- CV_{max} = value of breed with largest CV

Prioritization according to Total Utility

$$U_i = 4 \times (\text{risk}_i \times D_i) + \text{SCV}_i \quad (\text{Simianer et al., 2003})$$

where,

- U_i is the total utility for breed i
 - *Criterion for prioritization*
- Breeds are ranked according to total utility and those ranking highest are targeted for conservation

The Final Step: Use the information

- *Action 1. Prepare a report on breed prioritization*
 - Draw up “red list” of endangered breeds
 - Increase public awareness
- *Action 2. Hold meetings with stakeholders*
 - Pass results on to stakeholders
 - government and donors = provide support
 - farmers = design and implement conservation measures
- Implement conservation programmes

Thank You

Current Status of Prioritization

- Extinction risk most often used
- Objective prioritization methods applied to most livestock species on research basis
- Little application in practice
 - Austria
 - ALBC in USA
- Reasons?
 - lack of molecular characterization
 - no clear consensus on optimal approach
 - no simple comprehensive computational tools