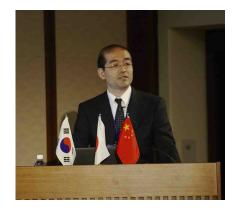
Session 2

Biodiversity Conservation

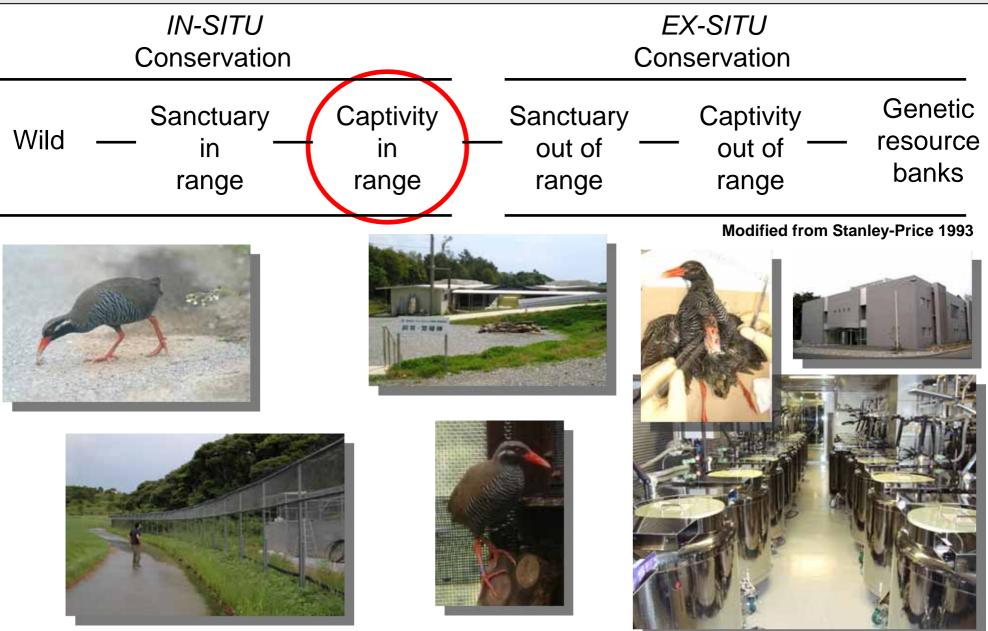


The results of reproductive and genetic research on Okinawa rail (*Gallirallus okinawae*) and its application for a captive breeding program

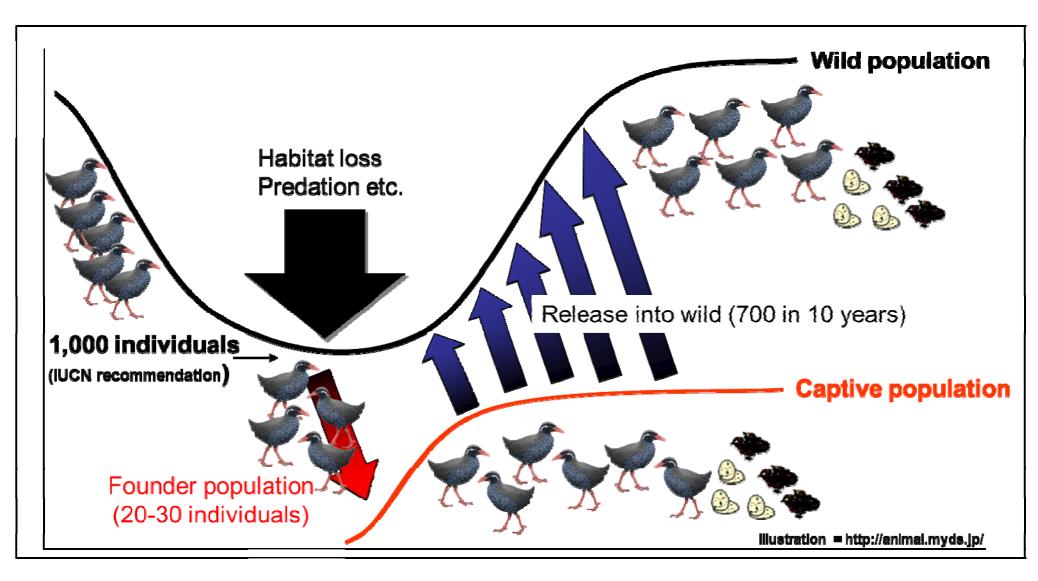


Manabu ONUMA, DVM, PhD Ecological Genetics Analysis Section Center for Environmental Biology and Ecosystem National Institute for Environmental Studies

Options for wildlife conservation



Captive breeding program for Okinawa rail

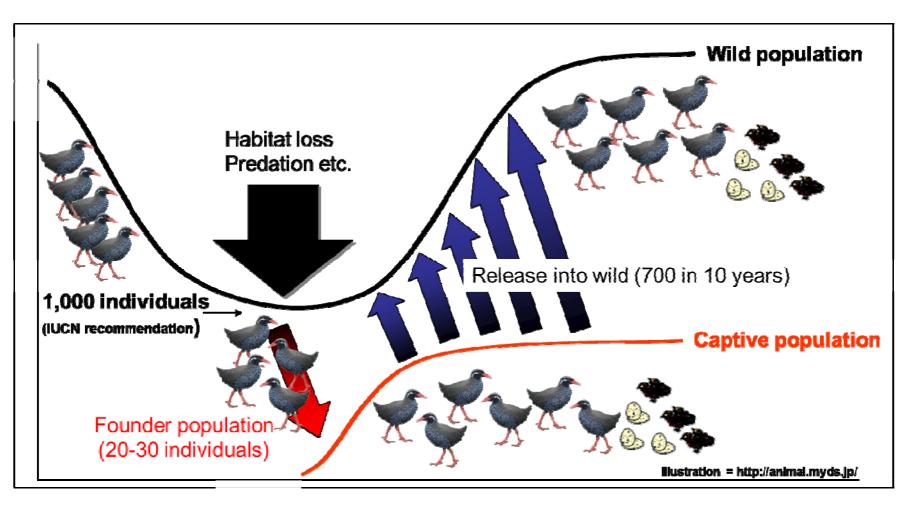


Essential information for successful captive breeding program

1, Efficient and stable reproduction : Breeding season, breeding age

(Onuma et al., 2011)

2, Maintaining genetic diversity : Evaluate genetic diversity



Sample collection

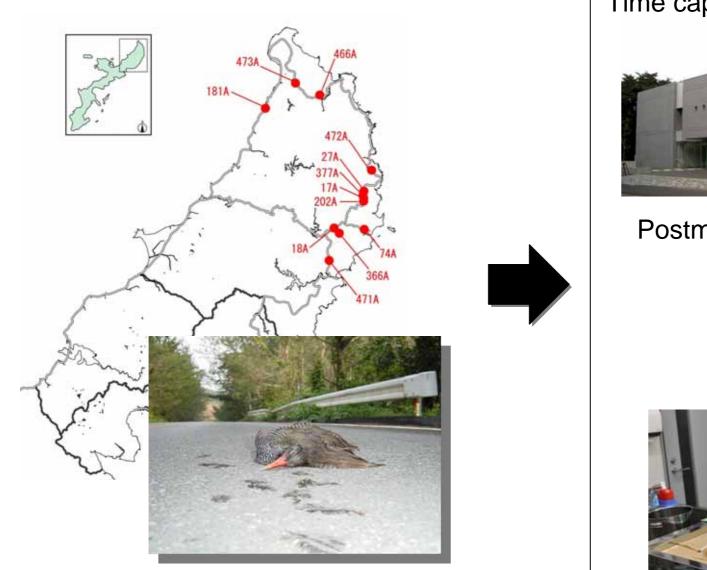
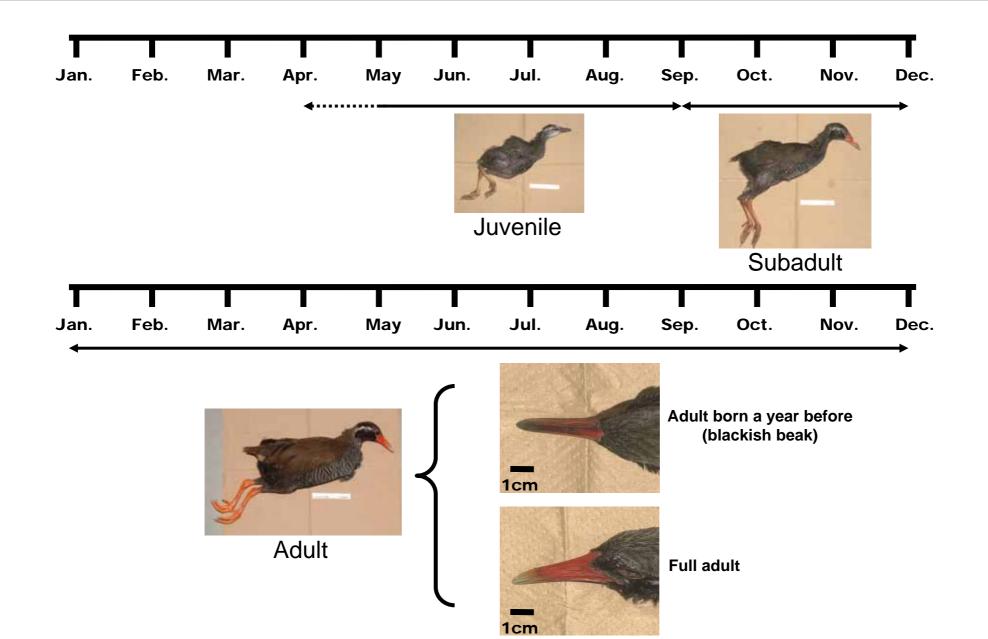


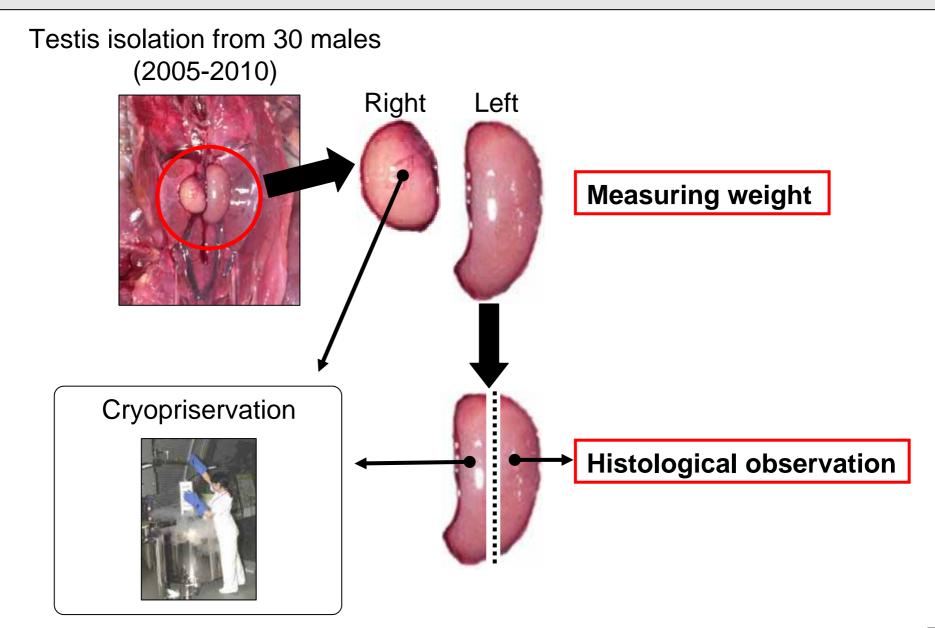
Photo by Ministry of Environment



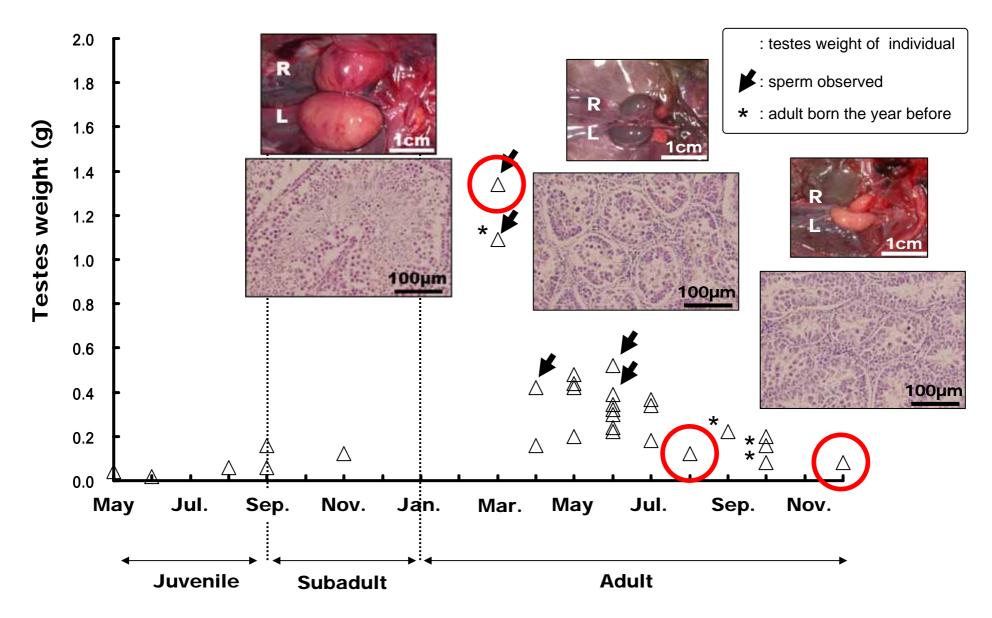
Age determination criteria of the Okinawa rail



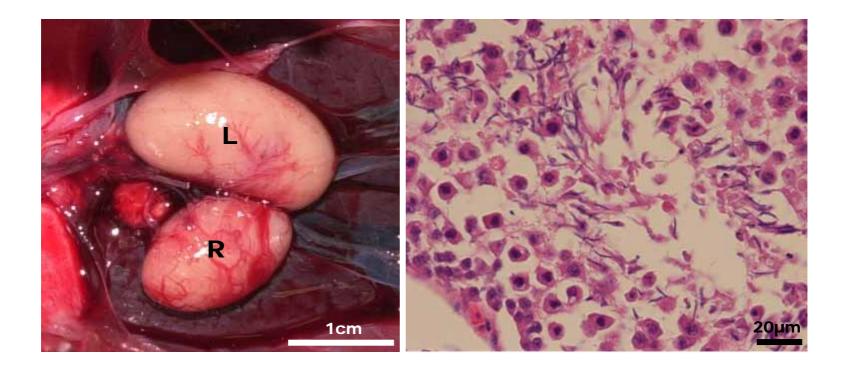
Methods (Testis observation)



Temporal changes in testis weight of Okinawa rail



Gross and histrogical observation of testis in the first breeding season



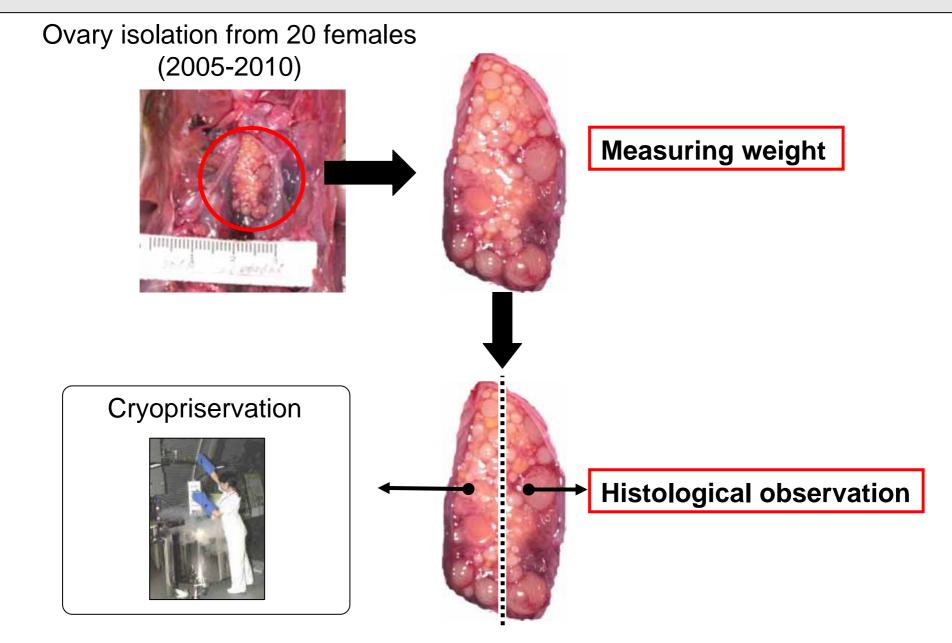
Semen collection of Okinawa rail



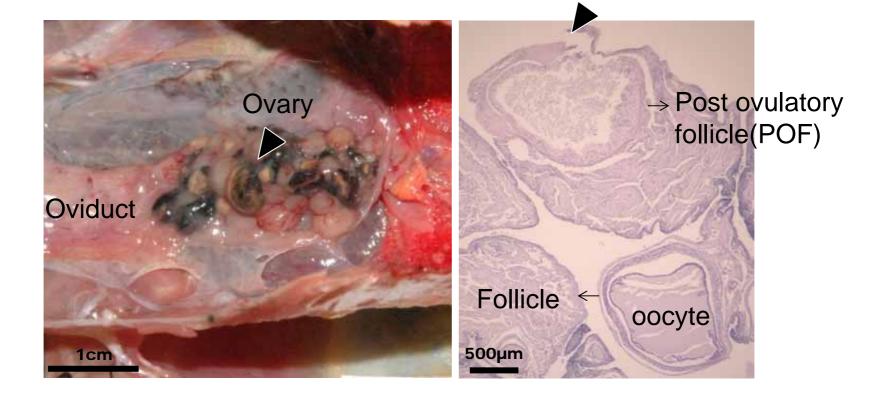




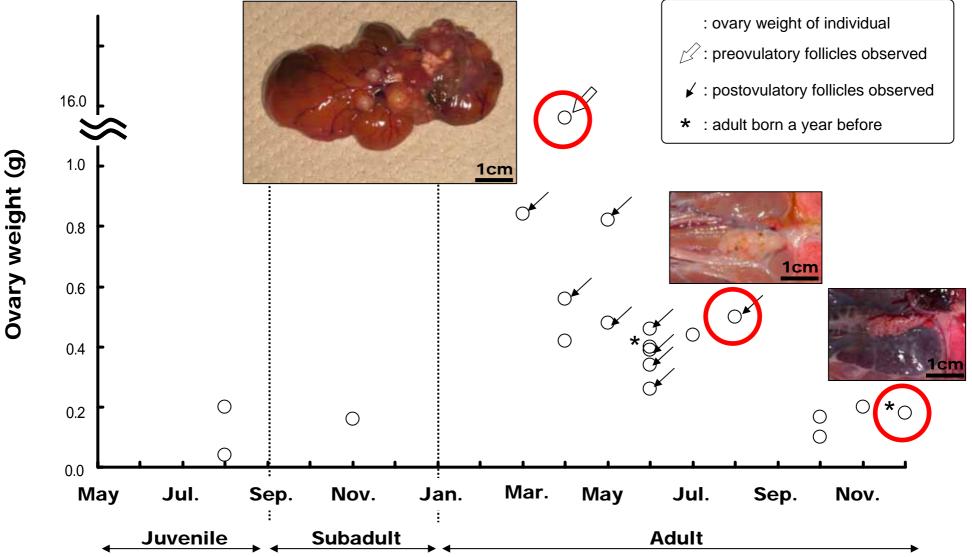
Methods (Ovary observation)



Gross and histological observation of ovary



Temporal changes in ovary weight of Okinawa rail

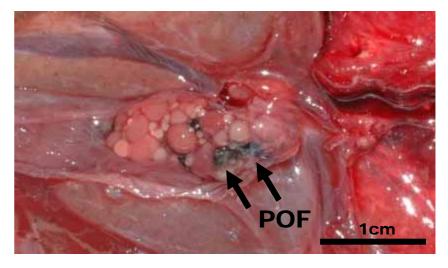


Gross observation of ovary in June

Adult born a year before



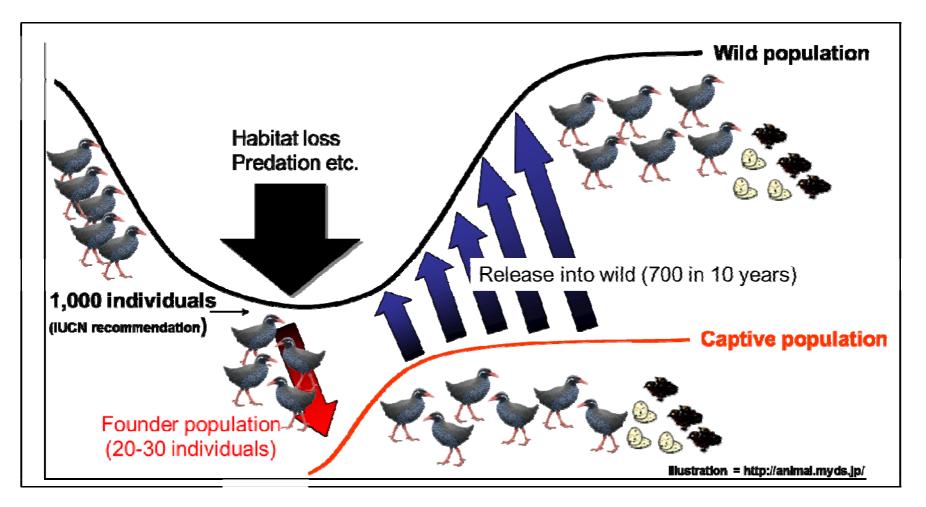




Essential information for successful captive breeding program

1, Efficient and stable reproduction : Breeding season, breeding age

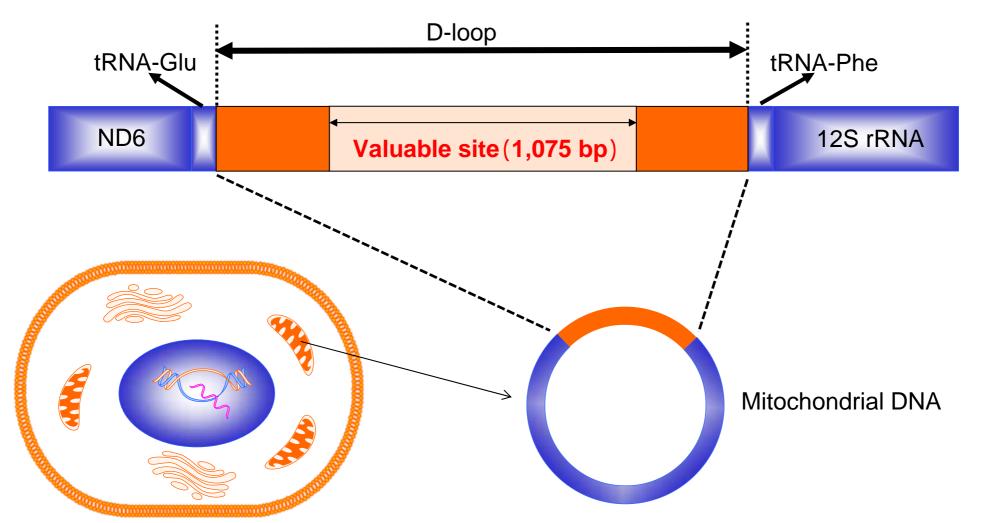
2, Maintaining genetic diversity : Evaluate genetic diversity



Method (Genetic diversity evaluation)

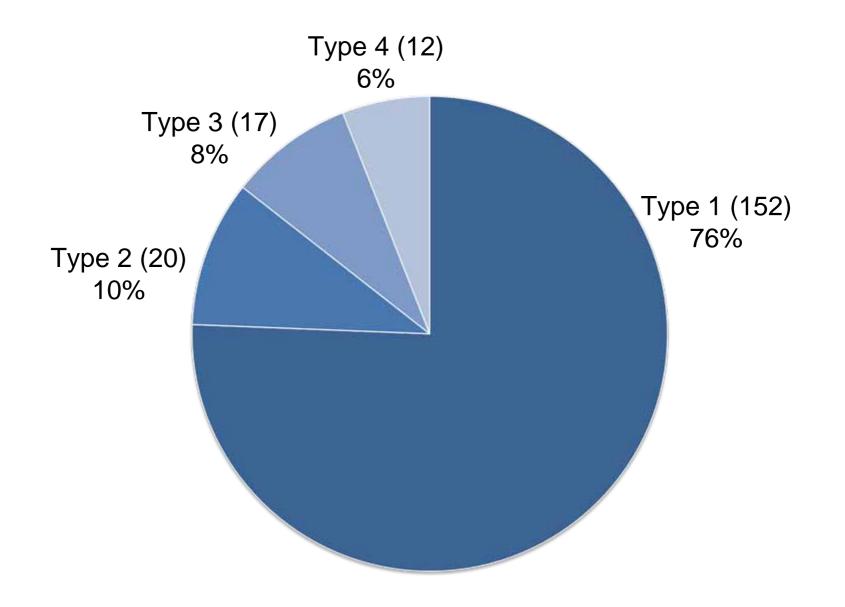
Animal: 201 individuals (male:107, female:94)

Genetic marker: Mitochondrial DNA D-loop region



16

D-loop variations of Okinawa rail



Conclusions

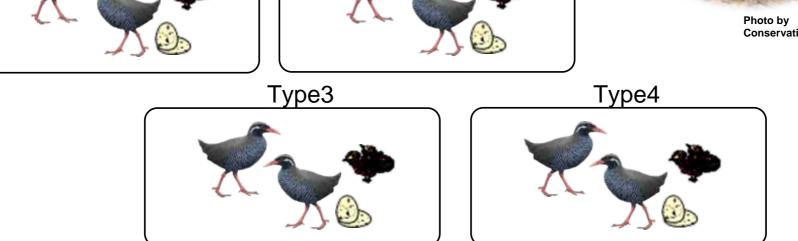
Type2

- Efficient and stable reproduction
 Breeding season: March to June (Seasonal breeder)
 Male: mature in the first breeding season
 Female: mature in the second breeding season
- Maintaining genetic diversity Maintaining 4 haplotypes

Type1







Whole genome sequencing of Okinawa rail in progress



High-throughput sequencing (32GB)

Sequence assembly

AGGACAAGTATTGAGATGATGC IGTATTGATATAAGGACAAGTAT IGATGCGTGGATAAATACTGTAA AATGATGCGTGGATAAATACTGT CAAGTATTGAAATGATGCGTGGA STTCTAAGACATTAATTGTAATG IGATGCGTGGATAAATATTGTAA

1,000 gene sequences will be identified by the end of 2011.

Draft genome sequence and chromosome mapping will complete by 2015.

ACKNOWLEDGMENTS

This research was conducted as part of the environmental specimen time capsule program funded by the Ministry of the Environment of Japan.

Special thanks to Dr. Yasumasa Sawashi, Mr. Yuji Miyake, Mr. Shota Fukuchi, Mr. Hiroshi Shichiri, Ms. Nahoko Eto, Mr. Nakata Katsushi, Mr. Michio Kinjyo, Mr. Makoto Fukuda, Mr. Manabu Nakachi, Mr. Yousuke Amano, Mr. Arata Kurihara, Mr. Sugao Ohshiro and all staffs of the time capsule building.

