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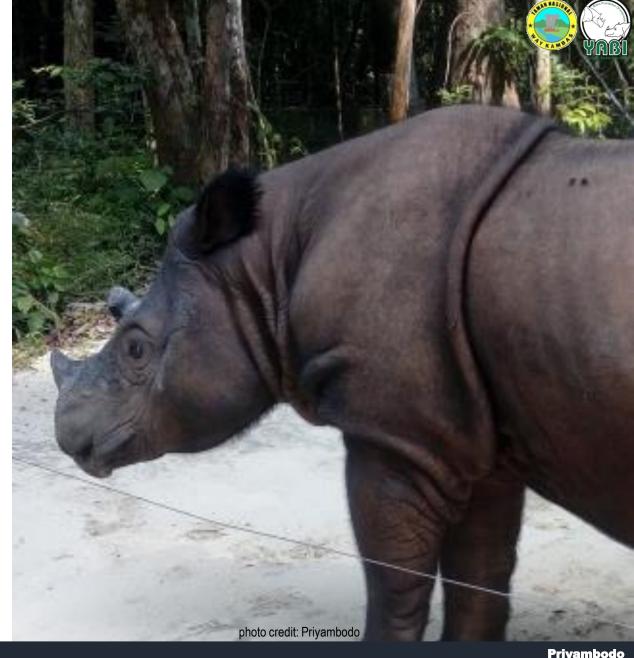
### Sumatran rhino

- There are 7 captive sumatran rhino in semi in situ Sumatran Rhino Sanctuary.
- Phylogenetic study of sumatran rhino based on molecular analysis was unavailable in Sumatran Rhino Sanctuary, Way Kambas National Park.



# Phylogenetic study

 In order to keep up with its existence in the wild, phylogenetic tree will be constructed and individual molecular profiles of its population in the captivity are needed



# Research goal

- This project research was conducted:
  - To collect DNA from each individual captive sumatran rhino by non-invassive method
  - to describe the quality of fecal DNA that collected from time series sampling activity.

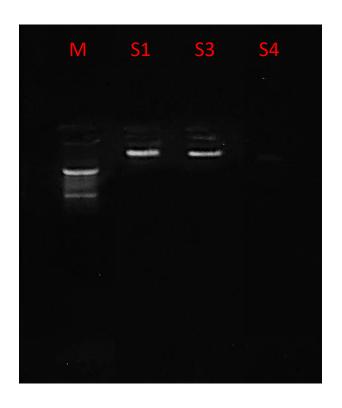


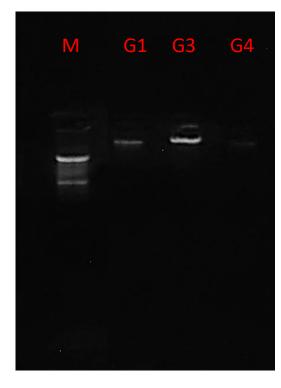


### Research methods

- Six fecal material samples was collected from total 7 individuals.
- Fecal material samples were handled by two different methods for mucus collection, swab and grinding methods.
- Mucus was taken on day 1, day 3, and day 4 for each sample.
- DNA extraction was done by commercial procedure and continued by qualitative analysis with 1% gel horizontal electrophoresis.

## **Qualitative Analysis**





M : marker

S : swab

G : grinding

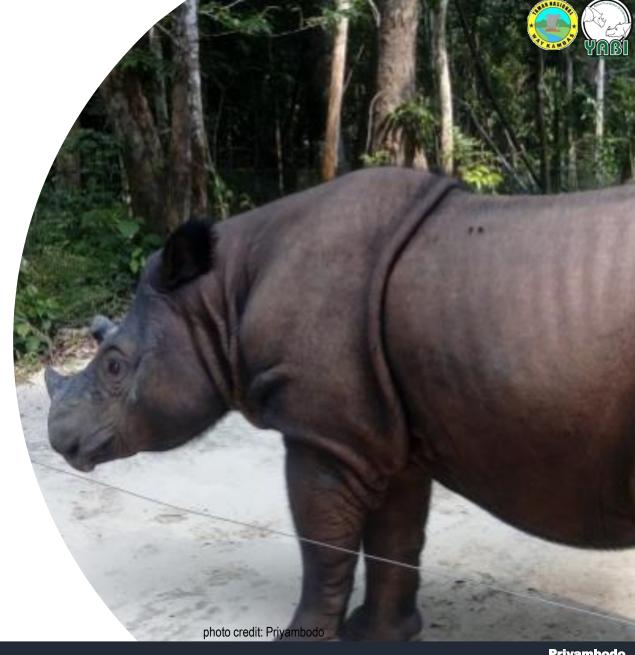
1 : 1<sup>st</sup> day

 $2 : 2^{nd} day$ 

3 : 3<sup>rd</sup> day

#### Discussion

 The absent of DNA band on day 4 caused by decomposition level of fecal samples which loss of mucus as convenient source of DNA.



#### Discussion

- Loss of mucus at feses = loss of source of DNA
- Mucus at feces contain ephitelial tissue from colon



### Discussion

 Decomposition of feses caused by physical factor (humidity, temperature, etc.) and biological factor (microbes activity)



### Conclussion

 Representative DNA from feses could be extracted from fresh feses (day 1 and 2) while mucus present

