Ancient DNA extraction from museum collections

What do Lepidoptera species tell us?

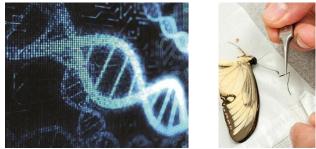
Elsa CALL

Supervisors: Niklas WAHLBERG and Mikael HEDRÉN













BIG4 Midterm Review meeting **Tovetorp** 14/10/2016



Natural history museum collections are abundant

Method

• Use for morphological studies

Projects

- DNA from these specimens
 - Too degraded to be used



• Natural history museum collections are abundant

Method

• Use for morphological studies

Projects

- DNA from these specimens
 - Too degraded to be used
 - Next-Generation Sequencing
 - New possibilities
 - Short fragments
 - Successfully from "fossilized" taxa
 - Neanderthals, mammoths and cave bears



- Complete first draft version of the Neanderthal genome
 - Studies of the recent evolutionary story of modern human
 - Understanding our own genome
 - Finding the "language gene"
 - Diseases (type 2 diabetes, Crohn's disease)



Pääbo et al. 2009

• The main objectives of this PhD

Projects

- Investigate the level of DNA preservation
- Develop protocols to sequence genomic DNA

Method

Target rare and difficult to collect lineages



• The main objectives of this PhD

Projects

- Investigate the level of DNA preservation
- Develop protocols to sequence genomic DNA

Method

- Target rare and difficult to collect lineages
- New advances in the field of "museomics"
- Phylogenetic relationships of major lineages in Lepidoptera



Projects

Method

- Investigating the level of DNA preservation in museum specimens of various ages
 - Are there any general trends in the preservation of DNA over time?



Projects

Method

- Investigating the level of DNA preservation in museum specimens of various ages
 - Are there any general trends in the preservation of DNA over time?
 - Assessment of the state of DNA preservation
 - Investigate the suitability of such samples for Next-Generation Sequencing



 Exploring different possibilities for sequencing DNA from museum samples

• Different methods for sequencing DNA

Projects

Context

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Amplicon sequencing, gene capture or whole genome sequencing

Method

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Projects

• Exploring different possibilities for sequencing DNA from museum samples

- Different methods for sequencing DNA
 - Amplicon sequencing, gene capture or whole genome sequencing

•Which one is the best?

To sequence homologous regions of the genome in different taxa

• Exploring different possibilities for sequencing DNA from museum samples

• Different methods for sequencing DNA

Projects

Context

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Amplicon sequencing, gene capture or whole genome sequencing

•Which one is the best?

To sequence homologous regions of the genome in different taxa

Method

In our case: Gene capture seems more appropriate

• Two projects will concentrate on clades of Lepidoptera to resolve phylogenetic relationships

Method

Geometroidea and Noctuoidea

Projects

- ~35,000 known species each
- Worldwide repartition

Context

• Extensive collections in Copenhagen and Stockholm





- Pieris napi (Pieridae)
- 3017 in the entomological collection's database
- 1551 potentially available for extraction

- Pieris napi (Pieridae)
- 3017 in the entomological collection's database

Method

Conclusion

• 1551 potentially available for extraction

Projects

- 25 selected specimens
- From 1878 to 2008
- 5 differents periods
 - 1880s, 1920s, 1940s, 1960s and 2000s
 - DNA extraction from abdomen

Projects

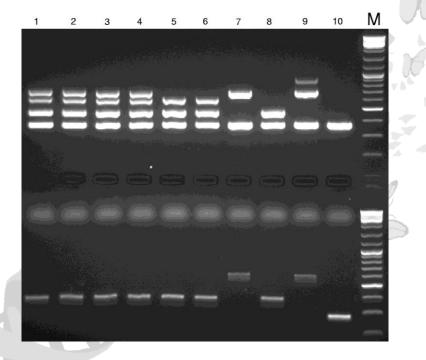
• How the DNA looks like

• DNA is more degraded in older specimens than in fresh ones



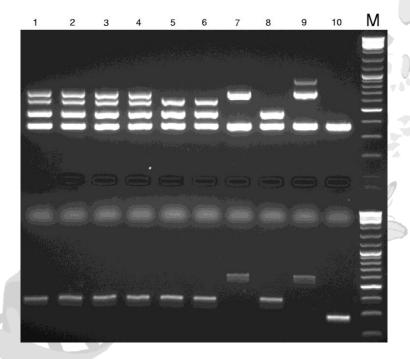
Projects

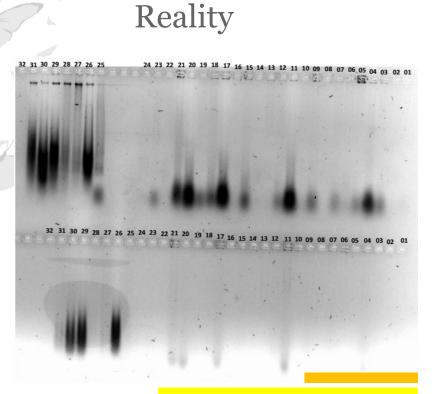
- Attempted method
 - Saturated NaCl
- Expectation



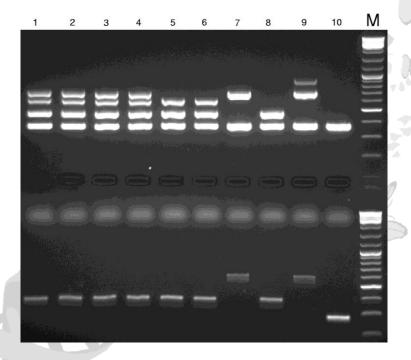
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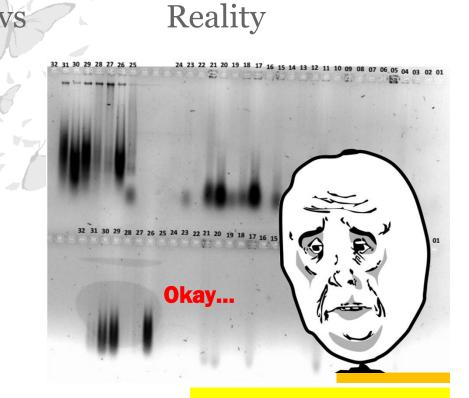
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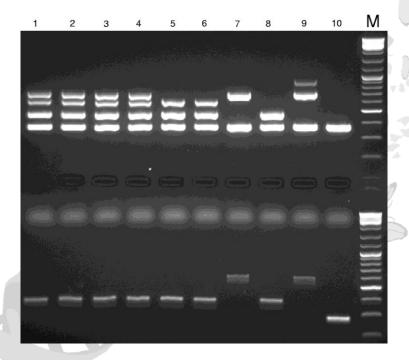
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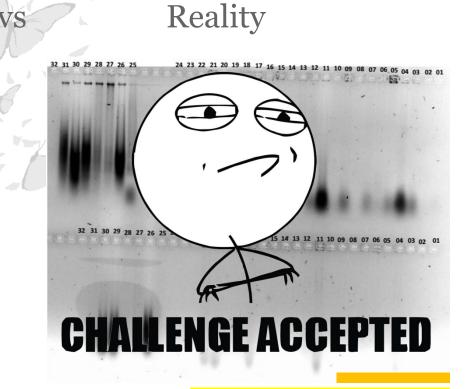




Projects

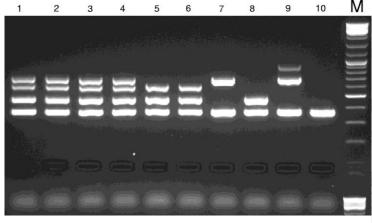
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• First results

- Attempted method
 - Saturated NaCl
- Expectation



Try another method





- Many applications
- But ancient DNA is difficult to sequence
- Several methods to test
- Genomic data can be used to resolve phylogenetic relationships of two clades of Lepidoptera
 - Geometroidea and Noctuoidea



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