

# Population genetics of Rangifer: a circumpolar approach

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#### any partners...



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# RESEARCH QUESTIONS



- How many 'groups' of Rangifer are revealed by genetic clustering analyses?
- Which regions at the circumpolar scale represent a challenge for conservation based on genetic origin and diversity?
- Where are located the most important genetic differences among large regions? What is the connectivity between herds?
- Could we localize Rangifer glacial refugia ?
- What is the temporal dynamic of genetic differentiation between reindeer and caribou?

### SAMPLING / GENOTYPING

> 50 herds over the world

20 individuals / herd

a) 16 microsatellites

b) 470-bp of the control region

## SAMPLING



- 1 NewFounland 28 Chinchaga
- 2 Mealy Mountain 29 Narraway
- 30 A La Peche 3 RedWine
- Caribou Mountain 4 Lake Joseph
- 5 Gaspesie Red Earth
- 7 Portneuf Cold Lake
- 6 Manicouagan 34 Greenland
- 9 Jamésie Iceland
- 36 Svalbard 8 Charlevoix
- 10 Val-d'Or Hardangervidda

Varanger

- 11 La Sarre
- 12 Rivière-George Finland
- 13 Rivière-aux-Feuilles 40 Yamal
- 14 Torngat 41 Taymir
- 15 Southampton 42 Lena-Olenyk
- 16 Beverly 43 Yana Indigurka
- 17 Ahiak 44 Sudrunskaya
- 18 Bathurst
- 19 Dolphin and Union 46 Wrangel
- 20 Bluenose East 47 Qamanirjuaq
- 21 Porcupine
  - 49 Alaska Peninsula
- 22 Central Arctic
  - 50 Interior Alaska

Bathurst Island

Chutkotka

- 23 Teshekpuk 24 Western Arctic
- 61 Northern BC
- 25 Denali
- 52 Nenetsky
- 26 Tay River
- 27 South Nahanni

No samples yet