

Relatedness among cryo-bank bulls of the Yakutian Cattle breed as estimated with microsatellite data

Ilma Tapio¹, Miika Tapio¹, Meng-Hua Li¹, Ruslan Popov² and <u>Juha Kantanen^{1,3}</u>

¹Biotechnology and Food Research, MTT Agrifood Research Finland, FI-31600 Jokioinen, Finland, ²Yakutian Research Institute of Agriculture, 677002 Yakutsk, Sakha, Russia, ³NordGen – Nordic Genetic Resource Centre, N-1431 Ås, Norway

Semen of 6 Yakutian bulls - Keskil, Moxsogol, Radzu, Erel, Sarial and Alii - is stored in a cryo-bank, but pedigree records of these bulls are not available.
We analysed 30 autosomal microsatellites to clarify genetic relatedness between these bulls and provide recommendations for the use of their semen in conservation and breed management.



Pairwise R

Individual	MER	ML-Relate		
pair	R (SD)	H0	H1	Р
Keskil:Moxsogol	0.233 (0.149)	UR	HS	0.0003
Radzu:Erel	0.163 (0.133)	UR	HS	0.08
		HS	UR	0.184
Radzu:Sarial	0.271 (0.112)	UR	HS	0.032
Erel:Sarial	0.196 (0.144)	UR	HS	0.037
Erel:Alii	0.053 (0.162)	HS	UR	0.042

YAKUTIAN CATTLE are the unique last remnants of the Siberian Turano-Mongolian cattle, with 1200 purebred animals left, and well adapted to the extreme sub-arctic conditions.

Up to 74% of the Yakutian Cattle are distributed in the three northern villages of Dzhargalakh, Kustur and Batagay-Alyta.



Observed mean relatedness (R) [black spots] and standard deviation bars for pairwise R comparisons for Yakutian cryo-bank bulls were computed using MER v.3.0 (*Wang J. 2002.Genetics 160:1203-1215*). The upper dashed line represents the upper limit of the 90% confidence interval for simulated random individuals. Ten out of 15 pairwise R-estimates approached zero or fell below it, suggesting that for those pairs relatedness did not significantly deviate from the average (zero) in the contemporary population.

The R-estimates above zero were tested using a maximum-likelihood estimator using ML-Relate (Kalinowski ST et al. 2006. Molecular Ecology Notes 6:576-579). A statistical test between two *a priori* hypotheses (unrelated (H_0) vs. half-sibs (H_1) was applied.

SD=standard deviation of a relatedness estimate (R) obtained using the MER program UR=unrelated and HS=half-sibs hypothesis tested using ML-Relate, P=significance of test result

• We suggest that Keskil:Moxsogol, Radzu:Sarial, Erel:Sarial and Radzu:Erel can be treated as bulls having up to 25% of their genes in common. We are more concerned about the type II error (related individuals are missclassified as unrelated) than the type I error (unrelated are identified as related).

We suggest dividing the bulls into 3 separate groups in a rotation mating scheme:
 I) Keskil and Moxsogol II) Alii III) Radzu, Erel and Sarial. In the rotation, a female belongs to the same group as the sire and is crossed with a bull from another group.

SAKHA YNAGA – CATTLE OF THE YAKUTS. 2009. Annales Academiae Scientiarum Fennicae. Humaniora No. 355. A new book edited by Leo Granberg, Katrina Soini and Juha Kantanen will tell a story of the endangered Yakutian Cattle, of their important role in the Siberian cultural heritage and of their status for safeguarding the food production in the future. The book is based on multidisciplinary studies and published in autumn 2009. Please make your order on web bookstore www.tidedkrij.af.(j)