Intrapopulational polymorphism of two mtDNA fragments of least cisco from Lake Sobachye

Location	Length of fragment, bp	N	h	S	H_d	π
			COI			
Lake Sobachye	635	9	5	4 (2)	0,81	0,00192
	ND1					
Lake Sobachye	1706	5	4	6 (2)	0,90	0,00164
*Kureyka River	1708	5	5	12 (2)		0,00304
*Data about ND1 sequences' polymorphism of this population are given for						
comparison; information from NCBI were used (GenBank Accession numbers						
JN629025–JN629029). N — sample size: h — number of haplotypes: S — number						

comparison; information from NCBI were used (GenBank Accession numbers JN629025–JN629029). N — sample size; h — number of haplotypes; S — number of polymorphic sites and number (in brackets) of parsimonial-informative sates; H_d — haplotype diversity; π — nucleotide diversity. Length of mtDNA fragments analyzed were depended on quality of sequences obtained; all sequences were aligned according to the smallest sequences.