





than 50 billion NOK. Researchers at NIBIO at Ås are using molecular tech-Norway's forest industry accounts yearly for production equivalent to more nologies to characterize current and ancient populations of Norway spruce (1), the most important forest species in Norway.

eastern regions 9000 and 3000 years Norway but his findings (2) were considered as inconclusive evidence for the existence of Norway spruce in Scandinavia during the last Ice or years, plant ecologists assumed that Norway was of vegetation that pine and Norway spruce trees established themselves here from ago, respectively. The Swedish forest researcher Leif Kullman had earlier reported on 11,000 year old remains of Norway spruce near Kongsvinger, during the last Ice Age and T devoid



ments in Trøndelag and Andøya, article (1) was conducted by Mari Mette Tollefsrud at NIBIO together major co-authors at Universities in fromsø, Uppsala and Copenhagen. More than 100 Norway spruce forests across Europe were screened as well The study reported in the Science and international partners, including as ancient DNA extracted from sedinational several Norway (Fig. 1).

versus haplotype B) which were Mitochondrial DNA variation was mh05) gave a clear distinction between 2 haplotypes (haplotype A One of these pairs (designated scored using 11 different PCR pairs.



Fig. 1. Extent of glacial ice coverage in Europe 60,000-15,000 years igo. A and T represent Andøya and Trøndelag sites, respectively, where ancient DNA samples were found. Wikipedia.

identical except for a 21 bp deletion in haplotype A. Haplotype A infound only in frequencies occurring in western highest this genotype may have been reof Norway spruce from refuges after the Norway, suggesting that trees with sponsible for recolonization Scandinavia with the dividuals are last Ice Age. Further evidence for Ice Age refuges of forest trees in Norway are: mitochondrial DNA in 6,500-10,300 year old sediments from Trøndelag DNA and spruce chloroplast DNA in 20,000 year old and 17,700 year old 1) identification of haplotype A and 2) findings of pine chloroplast respectively, sediments,

After its publication, a group of conclusions, saying that inadequate international scientists, including 2 criticized the report's affiliated with the University of protocols were used to exclude contamination in sediment samples Bergen,

independently collected samples, a coincidence they consider to be (3). In answering this criticism, the results to be caused by contamination would require that equipment would have had to be Parducci et al. (4) pointed out that for similarly contaminated in different pieces of

optimistic that individuals with haplotype A might be important for breeding trees that show flexibility to Forest researchers at NIBIO are survive under changing climatic conditions'.

## References:

2. Kullman, L. J. Biogeogr. 29 4. Parducci, L et al.: Science Parducci, L et al.: Science 3. Birks, HH et al.: Science 35 (2012) 1083-1086 338 (2012) 742a (2002) 1117

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