

Supplementary material

Table S1. Brief characteristics of climate and soils of the 6 forest natural regions in Lithuania based on Galvonaite et al. [94].

FRN	Name	Briefly	Alt., m a.s.l.	Mean annual T °C	Mean warmest month T °C	Mean coldest month T °C	Annual precipitation, mm	Snow duration, days	Soils
1	Western highland	Wettest, cooler, rich soils	150-200 (234)	6.3-6.7	17.0-17.5	-3.2 to -2.9	810-820	75-90	Medium rich, albeluvisols
2	Midland lowland, north	Drier, rich soils	50-70 (85)	6.5-7.0	17.4-18.1	-3.6 to -3.1	560-700	75-90	Rich, cambisols
3	North-east highland	Coldest, dry, poor	150-200 (289)	6.1-6.7	17.7-18.0	-4.8 to -3.8	610-690	90-105	Poor, albeluvisols
4	Sea-side lowland	Mildest, wet, poor	10-50 (78)	7.4-8.0	17.8-18.4	-1.9 to -1.4	770-800	60	Very poor, arenosols, podsols
5	Midland lowland, south	Mild, dry, rich	50-80 (104)	7.1-7.4	18.0-18.1	-3.6 to -3.1	600-640	65-80	Rich, cambisols, podsols
6	South-east highland	Cold, dry, poor	150-220 (259)	6.8	17.9	-3.7	700	90	Very poor, arenosols

FRN- Forest natural region, Alt. – altitude common range (maximum). Soils – dominant soil type. Data from 1981 to 2010. T is temperature.

Table S2. Molecular characteristics of the nuclear and chloroplast microsatellite loci used for the Scots pine DNA analysis. The repeat motive size is given in bp after the slash. For allele size the range and the most frequent alleles are given.

Locus	Repeat motif, bp	Allele size (most frequent), bp****	Forward primer (5'-3') Reverse Primer (3'-5')	References
Nuclear microsatellites (Psyl= EST-SSR, rest = genomic SSR)				
Psyl16	(AT)7 /2 bp	198-220 (202)	GCTCTGCCCATGCTATCACT TGATGCTACCCAATGAGGTG	Sebastiani et al. [75]*
Psyl18	(GCA)7 /3 bp	291-309 (300)	ACTACCTGGCATTCGTCCTG GGATCTGGTCCATTCTGTGT	Sebastiani et al. [75]
Psyl2	(GC)5 /2 bp	198-224 (208)	TTGCTTTGCAGAACATTG GTCCTGCAGGCAATCAAAAT	Sebastiani et al. [75]
Psyl25	(GCA)5 /3 bp	218-221 (221)	CAGCACCGTTCTTGTATC ACCGTTGCTCGTTGTCTTCT	Sebastiani et al. [75]
Psyl42	(TC)9 /2bp	167-177 (173)	CAACTTCAGCCTTGCAACAA CGACTTCATTGGAACACCA	Sebastiani et al. [75]
Psyl44	(CGG)5 /3 bp	160-177 (171)	TCCAAGTTGGTTCTTGTC GACACGATGGATTCCCTGAT	Sebastiani et al. [75]
Psyl57	(ACC)7 /3 bp	191-209 (200)	CCCCACATCTCTACAGTCAA TGCTCTGGATTGTTGCTG	Sebastiani et al. [75]
PtTX 4001	(GT)15 / 2 bp	201-237 (217)	CTATTGAGTTAAGAAGGGAGTC CTGTGGTAGCATCATC	Auckland et al. [77]
PtTX 4011	(CA)20 / 2 bp	195-283 (261)	GGTAACATTGGAAAACACTCA TTAACCATCTATGCCAATCACTT	Auckland et al. [77]***
SPAC 11.4	(AT)5(GT)19	135-165 (139)	TCACAAAACACGTGATTCA GAAAATAGCCCTGTGTGAGACA	Soranzo et al. [76]**
SPAC 12.5	(GT)20(GA)1 0 / 2 bp	121-193 (155)	CTTCTTCACTAGTTCCCTTGG TTGGTTATAGGCATAGATTGC	Soranzo et al. [76]
SPAC 7.14	(TG)17 (AG)21	181-253 (213)	TTCGTAGGACTAAAAATGTGTG CAAAGTGGATTITGACCG	Soranzo et al. [76]
Chloroplast microsatellites****				
Pt71963	(T)16 (T)14 /	138-146 (143,	TTCATTGGAAATACACTAGCCC	Vendramin et al. [42]
Pt15169	(C)8(T)8A(T)	120-128 (123,124)	CTTGGATGGAATAGCAGCC	Vendramin et al. [42]
Pt30204	(A)12(G)10	138-144 (141)	TCATAGGGGAAGATCCTCTT	Vendramin et al. [42]

*- EST SSR loci developed for *Pinus sylvestris*. **- nuclear SSR loci developed for *Pinus sylvestris*. ***- nuclear SSR loci developed for *Pinus taeda*. ****- cpSSRs for *Pinaceae*, all with 1 bp SSR motive. *****- allele range and most frequent allele from our study.

Table S3. Proportion of mitochondrial DNA mitotypes A and B in natural Scots pine populations within Lithuania.

Pop.No.	Pop. name	Alt. (m)	295 bp (B)	300 bp (A)
1	Plunge	122	3	17
2	Tryskiai	118	7	12
3	Kurtuvenai	108	10	10
4	Varniai	169	6	15
5	Mikieriai	80	5	15
6	Geguzinė	60	9	11
7	Rokiskis	127	10	10
8	Salakas	164	17	3
9	Grazute	183	11	9
10	Labanoras	174	9	12
11	Azvintis	177	12	8
12	Juodkrante	27	3	17
13	Darbenai	40	4	12
14	Sveksna	38	3	17
15	Pagegiai	25	4	16
16	Vaisvydava	87	15	5
17	Braziukai	82	7	13
18	Punia	81	9	11
19	Ancia	122	6	13
20	Veisieiai	138	8	12
21	Trakai	122	7	13
22	Druskininkai	153	11	9
23	Raseiniai	70	0	20
Proportion of A and B mitotypes %			38.6	61.4

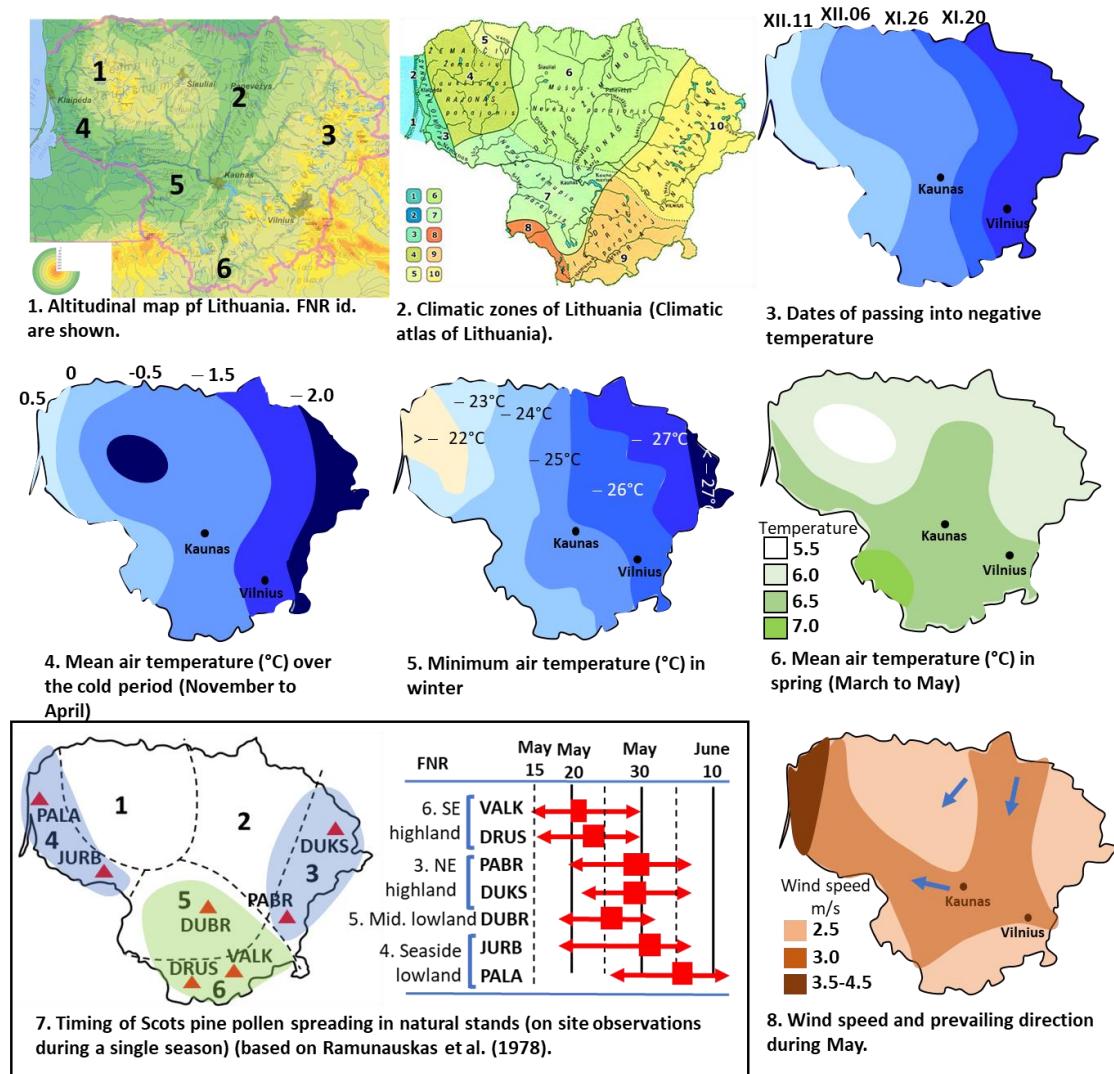


Figure S1. The main climatic characteristics regarding frost hardiness and timing of spring budburst for Scots pine in Lithuania. Maps 1 and 2 taken from Climate atlas of Lithuania [94]. FRN are forest natural regions, i.e., ecoclimatic zones used in forestry and basically reflecting altitudinal variation and continentally gradients similarly as in general climatic zoning in map 2. Maps 3 to 6 and 8 are modified from Galvonaitė et al. [94], where the climatic data averages the period over 1980–2011. The map 7 is modified from an empirical study by Ramanauskas et al [73], where during a single spring onsite observation on budburst timing were carried out simultaneously in different parts of Lithuania (arrows for start and end of pollen spread, the red boxes for mass pollen shed). The map 8 gives prevalent direction and strength of winds during pollination time, which may indicate possible zones of more frequent mating (we outlined these putative zones with different colors in the map 7). Lines on map 7 show forest natural regions.

Pop. 5 MIKE (northern midland lowland, FNR 2)



Pop. 12 JUOD (Nerija spit, FNR 4, sandy dunes)



Pop. 3 KURT, FNR 1 western highland.



Pop. 19, ANCI FNR 6 south-eastern highland



Figure S2. Actual view of the sampled stands in various eco-climatic zones of Lithuania (forest natural regions, FNR). Pop. 5 MIKE typical Scots pine forests in north-eastern highland with pine on the overstory and spruce in the understory. Pop. 12 JUOD is in the forest foots between sandy dunes with accumulations of richer upper soil layers (myrtillous ground vegetation, some oaks in the understudy). Pop. 3. KURT – typical mixed Scots pine

with Norway spruce stands on richer soils in the western highland (717 is a plus tree, FNR 1). Pop. 19 ANCI – Pure Scots pine stands on poorest sandy soils with Juniper in the understory common on the south-eastern highland.

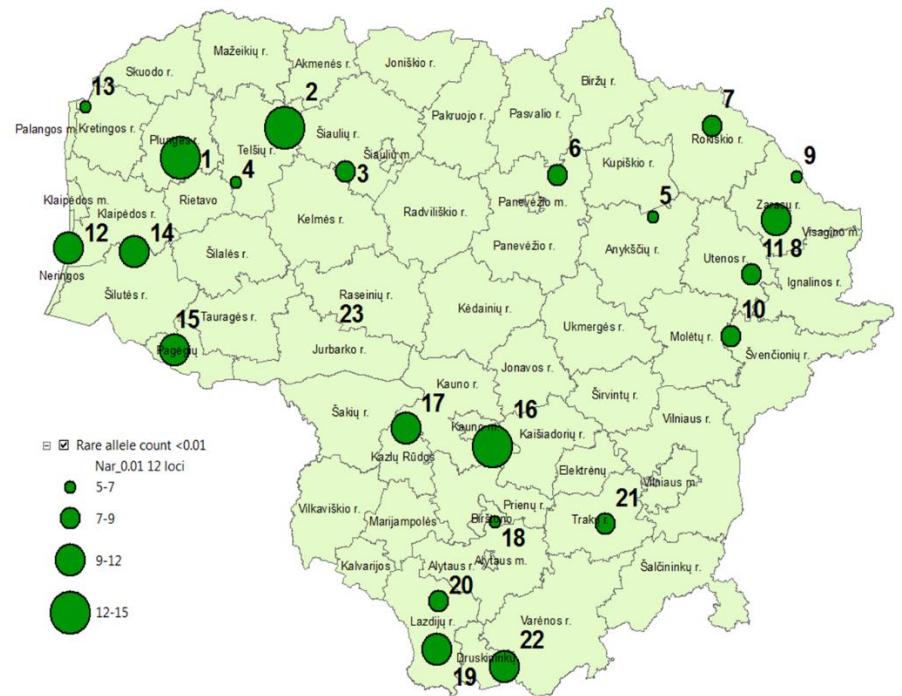


Figure S3. Population frequency of rare alleles (alleles with frequency <0.01 in whole material).

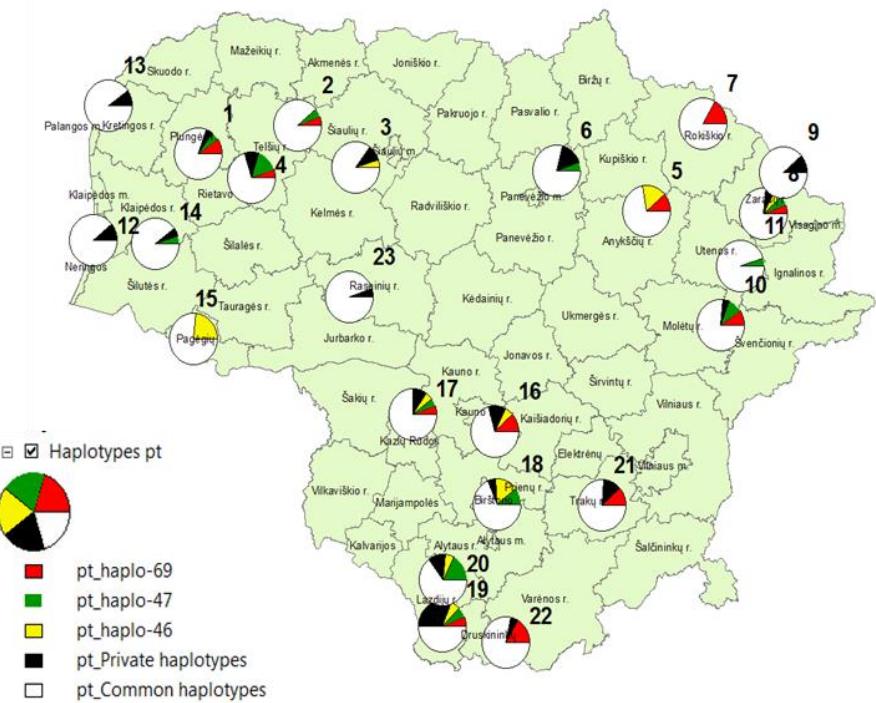


Figure S4. Geographical distribution of four most frequent clusters making up to 61 % of all haplotypes, private and shared haplotypes based on three cpSSR loci of pt series. Paternally inherited haplotypes are highly shared among populations studied (compare common haplotypes in white).