

**Table S2:** Information on the specimens from different regions used for RAD analyses, number of Illumina reads, unique stacks, polymorphic loci and SNPs. BF=Black Forest, CM=Cantabrian Mountains, MC=Massif Central, PY=Pyrenees. For details see Table S1.

Sample	Species	Region	Barcode	Reads	Unique Stacks	Polymorphic Loci	SNPs found
Est2	<i>T. gallicum</i>	PY	GCCAAT	835701	60897	2934	4483
Ey4	<i>T. gallicum</i>	BF	ATCACG	1337427	141429	4380	6765
Ey8	<i>T. gallicum</i>	BF	TGACCA	1099580	120541	3635	5550
Ey9	<i>T. gallicum</i>	BF	AGTTCC	781254	80096	2878	4422
F3	<i>T. gallicum</i>	MC	GGCTAC	562982	49534	4266	6000
F8	<i>T. gallicum</i>	MC	CGTACG	1004398	92235	8032	10506
Ho2	<i>T. gallicum</i>	BF	GTTTCG	2990589	144491	5188	7620
Ho3	<i>T. gallicum</i>	BF	GTGAAA	1102120	102809	2730	4170
Ho4	<i>T. gallicum</i>	BF	GTGGCC	1532963	115092	3933	5786
Nav14	<i>T. gallicum</i>	CM	CGATGT	3819040	241947	35217	52098
Nav4	<i>T. gallicum</i>	CM	CAGATC	4706880	190410	26704	39563
Pu3	<i>T. gallicum</i>	CM	ACTTGA	1151892	118231	13725	19763
Red8	<i>T. gallicum</i>	PY	AGTCAA	2207080	160655	9460	13925
Roe18	<i>T. gallicum</i>	BF	ACAGTG	667452	85481	1776	2957
Roe3	<i>T. gallicum</i>	BF	GATCAG	304679	31121	809	1540
Roe5	<i>T. gallicum</i>	BF	GGTAGC	6044749	252309	11363	16171
Roe6	<i>T. gallicum</i>	BF	ACTGAT	926705	93497	2163	3489
TtPrad2	<i>T. tellae</i>		ATGAGC	1881539	83496	1718	2126