

Supplementary Table S5

Table S5. *trnL-F* sequence matrix. Variable position of the cpDNA sequences (*trnL-F*) of 36 accessions sampled across the distribution range of all three species of *Taxus*.

Species	Country/Locality/Code	Position														
		1	2	2	3	3	3	3	4	5	5	6	6	7	7	7
		0	3	9	1	2	3	4	4	1	4	9	8	5	7	5
		2	8	7	4	8	8	5	8	3	2	0	8	6	0	5
<i>T. contorta</i>	Pakistan: Kurram Valley, KV10	C	A	G	T	T	C	G	G	0	A	C	G	0	C	G
	Pakistan: Tirah, TH15	0	.	.	.	0	.	.
	Pakistan: Chitral, CH5	0	.	.	.	0	.	.
	Pakistan: Palas valley, PV2	0	T	.	.	0	.	.
	Pakistan: Hazara, HZ3	T	.	0	.	.	.	0	.	.
	India: Himanchal Pradesh, FU1	0	.	.	.	0	.	.
	India: Uttranchal, Chamoli, IN74	0	.	.	.	0	.	.
	Nepal: Darchula, DO16	0	.	.	.	0	.	.
	Nepal: Jumla, JD1	0	.	.	.	0	.	.
	Nepal: Manang, MC21	0	.	.	.	0	A	.
	Nepal: Baglung, BB1	0	.	.	.	0	.	.
	China: Tibet, Jilong, GL11	A	0	.	.	.	0	.	.
<i>T. mairei</i>	Nepal: Kavre, KD6	.	.	T	.	G	.	.	.	0	.	A	.	1 ^d	.	.
	Nepal: Sindhuli, SR5	.	.	T	.	G	.	.	.	0	.	A	.	1 ^d	.	.
	Nepal: Sindhuli, SM1	.	.	T	.	G	.	.	.	0	.	A	.	1 ^d	.	.
	Bhutan: Punakha-Tashitang, BT5	.	.	T	.	G	.	.	.	0	.	A	.	1 ^d	.	.
	India: Meghalaya, IN1	.	.	T	.	G	.	.	.	0	.	A	.	1 ^d	.	.
	Myanmar: Hilawng ridge, BU1	.	.	T	.	G	.	.	.	0	.	A	.	1 ^d	.	.
	Vietnam: Lam Dong, VN28	.	.	T	.	G	.	.	.	0	.	A	.	1 ^d	.	.
	China: Jiangxi, JX5	.	.	T	.	G	.	.	.	0	.	A	.	1 ^d	.	.
	China: Guizhou, LS20	.	.	T	.	G	.	.	.	0	.	A	.	1 ^d	.	.
	Nepal: Baglung, BH10	.	.	T	.	G	T	.	.	1 ^b	.	A	T	1 ^e	A	T
	Nepal: Kaski, KC1	.	.	T	G	G	T	.	T	1 ^a	.	A	T	1 ^e	A	T
	Nepal: Gorkha, GK5	.	.	T	.	G	T	.	T	1 ^a	.	A	T	1 ^e	A	T
<i>T. wallichiana</i>	Nepal: Rasuwa, RT8	.	C	T	.	T	T	.	T	1 ^a	.	A	T	1 ^e	A	T
	Nepal: Sagarmatha, ND16	.	.	T	.	G	T	.	T	1 ^a	.	A	T	1 ^e	A	T
	Nepal: Taplejung, TK1	.	.	T	.	G	T	.	T	1 ^a	.	A	T	1 ^e	A	T
	Bhutan: Thimpu, BT1	.	.	T	.	G	T	.	T	1 ^a	.	A	T	1 ^e	A	T
	China: Xizang, Yadong, XY24	.	.	T	.	G	T	.	.	0	.	A	T	1 ^e	A	T
	China: Xizang, Cuona, CN17	.	.	T	.	G	T	.	.	0	.	A	T	1 ^e	A	T
	China: Xizang, Chayu, CY01	.	.	T	.	G	T	.	.	1 ^a	.	A	T	1 ^e	A	T
	China: Yunnan, Gongshan, GS1	.	.	T	.	G	T	.	.	1 ^c	.	A	T	1 ^e	A	T
	China: Yunnan, Lushi, LK01	.	.	T	.	G	T	.	T	1 ^a	.	A	T	1 ^e	A	T
	China: Yunnan, Yingjiang, YJ01	.	.	T	.	G	T	.	T	1 ^a	.	A	T	1 ^e	A	T
Hybrid individual	Nepal: Sindhupalchok, SL18	0	.	.	.	0	.	.
Hybrid individual	Nepal: Sindhupalchok, SL19	0	.	.	.	0	.	.

Character states shown in comparison to accession *T. contorta*, KV10, identical states indicated by "." in other accessions. 0= absent; 1^a= AGAAAGATCAAATATTATTA;

1^b= AGAAAGATCAAATATTATTAAGAAAGATCAAATATTATTA; 1^c= AGAAATATCAAATATTATTA; 1^d=AAAAAA-G; 1^e=AAAAAAAG