

Erratum to: Population fragmentation causes randomly fixed genotypes in populations of *Arabidopsis kamchatica* in the Japanese Archipelago

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In Table 1, errors in the horizontal alignment of column headings were inadvertently introduced during correction. Table 1 should appear as follows:

The online version of the original article can be found under doi:[10.1007/s10265-011-0436-8](https://doi.org/10.1007/s10265-011-0436-8).

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Table 1 CpDNA haplotype composition of each population

cpDNA haplotype			Type 1	Type 2	Type 3	Type 4	Type 5	Type 6	Type 7	Type 8	Type 9	Type 10	Type 11
<i>trnS-G</i> subtype			a	b	c	d	d	d	d	e	d	b	d
<i>trnL-F</i> subtype			c	c	c	a	b	d	f	e	c	c2	b2
No.	Location	Region											
1	Rishirizan	A	7	1									
2	Taisetsuzan	A		8									
3	Rausudake	A						8					
4	Bandaisan	B						8					
5	Shiroumadake	C					1	7					
6	Tateyama	C						3	5				
7	Norikuradake	C								8			
8	Yatsugatake	C		5			2	1					
9	Kitadake	C						8					
10	Fujisan	C						8					
11	Toyama	D		8									
12	Makino	D				7	1						
13	Satsuma	D				1	7						
14	Hamawake	D				7	1						
15	Omimaiko	D					8						
16	Sabae	D					8						
17	Mie	D					8						
18	Kochi	D					8						
19	Tsurugisan	E			8								
20	Daisen	E						8					
Outside the Japanese Archipelago													
	Anchorage (Alaska)			1									
	Paramushir Island (Kuril)										1		
	Nantou (Taiwan)											1	
	Taroko N. P. (Taiwan)												1

The populations in Lake Biwa is Makino (12), Satsuma (13), Hamawake (14), Omimaiko (15) and Sabae (16)

A, Hokkaido; B, Northern Honshu; C, Central Honshu; D, lowland in western Japan (subsp. *kawasakiana*); E, Western Japan