



**Figure S1.** Pedigree of the varieties, which harbor *e1-ret/ghd7-0ret* allele. Bold with underline indicates that the varieties harbor *e1-ret/ghd7-0ret* allele. Gray indicates varieties whose allele is unknown. Other indicates that the varieties harbor *e1/ghd7-0a* allele.

<sup>1)</sup> "Aikoku" includes multiple varieties with the mixture of the insertions.

Table S1. Frequency distributions of days to heading in F<sub>3</sub> lines (T65 × T65w)

Parents and F <sub>3</sub> lines	Genotype of parents and F <sub>2</sub> plants <sup>a</sup>	Days to heading																								Total	No. of F <sub>3</sub> lines observed <sup>b</sup>								
		95	96	97	98	99	100	101	102	103	104	105	106	107	108	109	110	111	112	113	114	115	116	117	118			119	120	121	122	123	124	125	
T65	<i>E1E1</i>	7	5	7	1	1																													21
T65w	<i>E1-rE1-r</i>																			2	3	2	2	3	4	5	1	2						24	
1						2	1	2	2	7	2	1																							18
5							1		3	3	8	7	1	1	1																				25
6					2		1	1	2	3	2	3	3																						17
7				1		1	5	2	3	6		1	2	1	2																				24
12							3	1	3	7		3	3	2																					23
14										1	1	4	1		1																				8
16	<i>E1E1</i>								1	3	5	5	4	2	1	1																			22
17					1	3	1	6	4	3	2	1	1																						22
18								2		2	3	3	2	2																					13
22									3	3	4	5	5	1																					22
23						1		2	2	3	2																								11
25							1	1	1	4	5	3	2	4	2																				22
26							1	1	3	7	3	3	1																						20
2								1		1		1	1	1	1	2	1	4	1	2		3		1	1									21	
4												3				2	1	6	2		1	3	2											20	
8										1		3	2	1	3	2	1	2	2		2	2		1										22	
9													2		5								2	1		1	2	1						17	
10								1	1		2	3				2	2	5	1	3	1		1	1	1									24	
11										1		1	2	3	1	1	2	2	5	2	1	2	1	1	1									25	
13	<i>E1E1-r</i>								2		1		1	3	1	2	2	1	1	3		1	4	1										23	
19										1	1	1	1			4	2	2				1	1											14	
24										2	2	1	2	1		1	3	2	5	1	1					2								23	
29								3		1		1				2	5	2	1	2	1		1											19	
30													2			1	4	2	4	2	4			1	1									21	
33											1	2		1	1	2	1	3	2					5	2	1	1	1						23	
37					1		1	1	3	1	1				3	1	2	1	2				1	2	2									22	
39										1	1		2	3	1	1	2	3		1	4	1	2	2										24	
3																						4	7	8	2	2	1							24	
20																				1	1		2	3	5	4	3	1						20	
21																				2		2	3	5	4	1	3		1					20	
27															1	3	1	3	1	5	2	2	2	1		1								20	
28																			3		4	3	7	6	1									24	
31	<i>E1-rE1-r</i>																		1	2		1	4	7	5	2	1							23	
32																			1	1		3	2	6	7	3								23	
34																						1		4	4	4	3	5	1	2				24	
35																1			2		5	3	2	4	1									18	
36																			1	1	1	1	5	8	2	3								22	
40																							1	1	1	3	6	6	5	1					24

<sup>a</sup> *E1* and *E1-r* are early and late heading alleles of *E1* locus, respectively.

<sup>b</sup> *E1E1* : *E1E1-r* : *E1-rE1-r* = 13:14:11,  $\chi^2 = 8.904$ ,  $P = 0.012$

Table S2. Frequency distributions of days to heading in F<sub>3</sub> lines (T65m × T65w)

Parents and F <sub>3</sub> lines	Genotype of parents and F <sub>2</sub> plants <sup>a</sup>	Days to heading																									Total	No. of F <sub>3</sub> lines observed <sup>b</sup>						
		95	96	97	98	99	100	101	102	103	104	105	106	107	108	109	110	111	112	113	114	115	116	117	118	119			120	121	122	123	124	125
T65m	<i>e1e1</i>				1		8	5	5																									19
T65w	<i>E1-rE1-r</i>																			2	3	2	2	3	4	5	1	2				24		
8			1		2	2	1	6	7	4																							23	
18			1			4	6	2	7	2					2																		24	
20					1	1	3	6	4	1	2	1	1	1																			20	
21			3	2	3	4	2	2	4	1																							21	
28	<i>e1e1</i>				2	4	4	5	4	1																							20	
29					3	5		7	6	1				1																			23	
34					4	2	1	4	8	2				1																			22	
35		1	1		5	1	1																										9	
40										2																							3	
52			1		1		4	5	6	1	2				1																		20	
3								1					1		3	2	2	3	3			3	1	1	2				1			23		
4									2	4	1	1	1			1		3	4	1	1	1	2		1			3				25		
7								2				1		1		1		3	1	4	2	2	2			2						21		
9				1		1			1	4	1				2		3	3	1	3	3	1	1										25	
10							1	2	2						2	2	3	2	1		5	1	1										22	
11								1			1				1	1	4	1	1	2	1	1			1								15	
12						1		2	1	4					1	1	1	2	4					2			1						20	
15					1			3	1					1		2	1	1	1	1	1							1					12	
16									2		2			1		2	1	2	2	2		1	1	1	1	4							22	
19							1	1						1		2	2		1		1	1	1	1									11	
23	<i>e1E1-r</i>							2	1	1		1			2	1	2	2	3	2		3	1	1									22	
24					1	1	1	1	1	1					2	1	2	3	5	1		1	1	2		1							22	
27				2		2									3	1	4	3	1			1	1	1		1							20	
33								1							1	1		1	2				2		1								9	
36				1	1			1	2	2					1	1	3	1	3	1	2	2	1	2	2								24	
37						1	3						1		2	3	1	3	1	2	2	1	1	1	1		1						23	
38							2	1	2	2					1	4	6	1	1	1	1	1	1	1		1							23	
42				1						2	1			1	1				2	5	2	2		3		2	1						25	
51				1			1	1	3						1	3	2	4		1	2	1	1	1	1	1							23	
54				1	1			4	1					1	1	2	2	1	1					1										17
55					1	1	1	1	2				1		2	6	1	3	2				1											21
2																		1	1	3	3	5	6	1	2	3							25	
6																			7	7	3	4	1		1									23
31															1						1		7	7	2	1			2				21	
32																			4	2	1	5	3	3	4									22
39	<i>E1-rE1-r</i>																	5	5	2	4	3	2	2										23
41															1		2	2	2	3	5	2	2	2										21
53																1	4	2	5	6	5													23
56																	1	1	3	5	7	3	4											24
57																					2	5	6	6										19

<sup>a</sup> *e1* and *E1-r* are early and late heading alleles of *E1* locus, respectively.

<sup>b</sup> *e1e1* : *e1E1-r* : *E1-rE1-r* = 9:21:10,  $\chi^2 = 0.150$ ,  $P = 0.928$

**Table S3. Lists of varieties used in this study**

Group	Name	Origin	ID No. <sup>1)</sup>	landrace/improved	Subspecies	Haplotype around <i>E1</i> locus <sup>2)</sup>	Allele of <i>E1/Ghd7</i> locus <sup>2)</sup>	Days to heading <sup>2), 3)</sup>
Hokkaido varieties	Akabozu	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	69.9
	Akage	Hokkaido	-	landrace	Japonica	Hap1	<i>e1/ghd7-0a</i>	68.4
	Akatsuki	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	65.3
	Bozu	Hokkaido	-	landrace	Japonica	Hap1	<i>e1/ghd7-0a</i>	68.4
	Bozu 6	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	67.2
	Dohoku 5	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	65.1
	Eiko	Hokkaido	-	improved	Japonica	Hap3	<i>e1-ret/ghd7-0ret</i>	65.6
	Fukoku	Hokkaido	-	improved	Japonica	Hap3	<i>e1-ret/ghd7-0ret</i>	66.0
	Fukuyuki	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	68.1
	Hashiribozu	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	65.1
	Hayakaze	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	64.5
	Hayakogane	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	67.1
	Hayamasari	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	65.4
	Hokkai 126	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	63.8
	Hokkai 242	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	63.5
	Hokkai 244	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	65.3
	Hokuto	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	64.2
	Iburiwase	Hokkaido	-	landrace	Japonica	Hap1	<i>e1/ghd7-0a</i>	65.3
	Ishikari	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	66.0
	Ishikarisirage	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	67.8
	Jo-iku 100	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	67.8
	Jo-iku 272	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	64.4
	Kirara 397	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	71.3
	Kitaake	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	66.4
	Kitahikari	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	68.3
	Kitakogane	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	66.6
	Kiyokaze	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	65.1
	Kyowa	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	67.7
	Matsumae	Hokkaido	-	improved	Japonica	Hap3	<i>e1-ret/ghd7-0ret</i>	68.3
	Megurosakaemoti	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	71.5
	Norin 11	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	63.0
	Norin 15	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	63.8
	Norin 33	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	64.6
	Sakigake	Hokkaido	-	landrace	Japonica	Hap1	<i>e1/ghd7-0a</i>	65.7
	Sasahonami	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	66.6
	Shimahikari	Hokkaido	-	improved	Japonica	Hap3	<i>e1-ret/ghd7-0ret</i>	69.3
Shinei	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	73.4	
Shiokari	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	65.7	
Sorachi	Hokkaido	-	improved	Japonica	Hap3	<i>e1-ret/ghd7-0ret</i>	73.2	
Tokachikuroge	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	60.8	
Tomomasari	Hokkaido	-	improved	Japonica	Hap3	<i>e1-ret/ghd7-0ret</i>	73.1	
Tomonishiki	Hokkaido	-	improved	Japonica	Hap3	<i>e1-ret/ghd7-0ret</i>	72.4	
Wasebozu	Hokkaido	-	landrace	Japonica	Hap1	<i>e1/ghd7-0a</i>	62.8	
Wasehukoku	Hokkaido	-	improved	Japonica	Hap1	<i>e1/ghd7-0a</i>	66.4	

**Table S3. Lists of varieties used in this study**

Group	Name	Origin	ID No. <sup>1)</sup>	landrace/improved	Subspecies	Haplotype around <i>E1</i> locus <sup>2)</sup>	Allele of <i>E1/Ghd7</i> locus <sup>2)</sup>	Days to heading <sup>2), 3)</sup>
Japanese core collection	Gaisen Mochi	Japan (Unknown)	JRC 01	landrace	Tropical Japonica	N. A.	<i>E1/Ghd7-2</i>	103.5
	Hinode	Kinki	JRC 03	improved	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	86.3
	Senshou	Tokyo	JRC 04	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	87.0
	Yamada Bake	Kagoshima	JRC 05	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	94.5
	Kaneko	Kantotozan	JRC 06	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	84.1
	Iruma Nishiki	Saitama	JRC 07	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	84.9
	Okka Modoshi	Japan (Unknown)	JRC 08	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	101.0
	Hirayama	Tokyo	JRC 10	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	84.3
	Kahei	Kagoshima	JRC 11	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	N. A.
	Oiran	Kumamoto	JRC 12	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	91.8
	Bouzu Mochi	Oita	JRC 13	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	92.8
	Meguro Mochi	Kantotozan	JRC 14	landrace	Tropical Japonica	Hap4	<i>E1/Ghd7-2</i>	103.0
	Akage	Hokkaido	JRC 17	landrace	Japonica	Hap1	<i>e1/ghd7-0a</i>	N. A.
	Hassokuho	Japan (Unknown)	JRC 18	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	93.0
	Wataribune	Shiga	JRC 19	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	109.4
	Hosogara	Aomori	JRC 20	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	78.5
	Akamai	Kochi	JRC 21	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	96.4
	Mansaku	Nagano	JRC 22	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	89.3
	Ishijiro	Toyama	JRC 23	landrace	Japonica	Hap2	<i>E1/Ghd7-2</i>	98.8
	Joushuu	Yamagata	JRC 24	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	92.7
	Dango	Japan (Unknown)	JRC 25	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	109.6
	Aikoku	Fukui	JRC 26	landrace	Japonica	Hap3	<i>e1-ret/ghd7-0ret</i>	91.3
	Ginbouzu	Ishikawa	JRC 27	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	102.3
	Shinriki Mochi	Kumamoto	JRC 28	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	111.3
	Shichimenchou Mochi	Japan (Unknown)	JRC 29	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	107.4
	Morita Wase	Yamagata	JRC 30	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	82.0
	Kameji	Shimane	JRC 31	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	106.5
	Omachi	Okayama	JRC 32	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	114.0
	Shinriki	Hyogo	JRC 33	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	110.0
	Kyoutoasahi	Kyoto	JRC 34	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	110.4
	Kabashiko	Miyazaki	JRC 35	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	106.9
	Sekiyama	Aomori	JRC 36	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	78.8
	Shinyamadaho 2	Hyogo	JRC 37	landrace	Japonica	Hap2	<i>E1/Ghd7-2</i>	108.8
	Nagoya Shiro	Akita	JRC 38	landrace	Japonica	Hap5	<i>E1/Ghd7-2</i>	91.7
	Shiroine (Kemomi)	Tokushima	JRC 39	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	109.8
	Akamai	Nagasaki	JRC 40	landrace	Indica	Hap2	<i>E1/Ghd7-2</i>	116.4
	Akamai	Tokushima	JRC 41	landrace	Indica	Hap2	<i>E1/Ghd7-2</i>	110.9
	Touboshi	Kagoshima	JRC 42	landrace	Indica	N. A.	N. A.	N. A.
	Akamai	Kantotozan	JRC 43	landrace	Indica	Hap4	<i>E1/Ghd7-2</i>	98.6
	Karahoushi	Kagoshima	JRC 44	landrace	Indica	N. A.	N. A.	N. A.
	Hiyadachitou	Yamagata	JRC 45	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	78.6
	Fukoku	Hokkaido	JRC 46	landrace	Japonica	Hap3	<i>e1-ret/ghd7-0ret</i>	N. A.
	Okabo	Japan (Unknown)	JRC 47	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	106.4
	Hakamuri (Yokoyama)	Kagoshima	JRC 48	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	96.0
	Rikutou Rikuu 2	Japan (Unknown)	JRC 49	landrace	Japonica	N. A.	<i>E1/Ghd7-2</i>	87.7
	Himenomochi	Akita	JRC 50	improved	Japonica	Hap2	<i>E1/Ghd7-2</i>	82.8
	Shinshuu	Nagano	JRC 51	landrace	Japonica	Hap2	<i>E1/Ghd7-2</i>	85.1
	Aichiasahi	Aichi	JRC 52	improved	Japonica	Hap4	<i>E1/Ghd7-2</i>	109.8
	Raiden	Kantotozan	JRC 53	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	113.7
	Houmanshinden Ine	Kagoshima	JRC 54	landrace	Japonica	Hap4	<i>E1/Ghd7-2</i>	120.0

1): ID number of the Japanese core collection

2): N. A. : Not Available data

3): Days to heading is from sowing (April 30, 2010) in Kyoto.

**Table S4. Lists of varieties used in this study**

Name	Origin	NIAS gene bank ID	landrace/improved	Haplotype around E1 locus	Allele of E1/Ghd7 locus
Riku Aikoku	Kantotozan	4510	landrace	Hap4	<i>E1/Ghd7-2</i>
Riku Aikoku	Kantotozan	4713	landrace	Hap4	<i>E1/Ghd7-2</i>
Hakubou Aikoku	Gunma	4774	landrace	Hap4	<i>E1/Ghd7-2</i>
Aikoku	Gunma	4982	landrace	Hap4	<i>E1/Ghd7-2</i>
A Aikoku 2	Tohoku	5247	improved	Hap4	<i>E1/Ghd7-2</i>
Aikoku	Akita	5419	landrace	Hap5	<i>E1/Ghd7-2</i>
Gokuwaseaikoku	Akita	5777	improved	Hap4	<i>E1/Ghd7-2</i>
Aikokudaikoku	Akita	5792	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokudaikokumochi A	Akita	5793	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokudaikokumochi B	Akita	5794	improved	Hap4	<i>E1/Ghd7-2</i>
Aikokutarehatou	Akita	5795	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokuwaitou	Akita	5796	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokumitsuryuutou	Akita	5810	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Nakatemubouaikoku	Nagano	5998	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Nakateaikoku 3	Ibaraki	5999	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Waitouaikoku	Tohoku	6044	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokutarehatou	Tohoku	6047	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Bouzuaikoku	Yamagata	6133	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Rikuuaiikoku 20	Akita	6158	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku	Fukui	6365	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku	Fukui	6369	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku 70	Niigata	6442	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku	Miyagi	6730	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Nakateaikoku	Yamagata	6744	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Nakateshinaikoku	Akita	6752	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Waseaikoku	Yamagata	6753	landrace	Hap5	<i>E1/Ghd7-2</i>
Aikoku 1	Miyagi	6757	improved	Hap4	<i>E1/Ghd7-2</i>
Aikoku	Miyagi	6768	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokuibaraki 1	Ibaraki	6771	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokuibaraki 2	Ibaraki	6822	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Akitawaseaikoku 5	Akita	6915	improved	Hap4	<i>E1/Ghd7-2</i>
Waseaikoku	Miyagi	6927	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Waseaikoku 2	Miyagi	6928	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokuho	Miyagi	7013	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Kairyuaikoku 17	Osaka	7053	improved	Hap4	<i>E1/Ghd7-2</i>
Waseaikoku 96	Chiba	7054	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Nakateaikoku 90	Chiba	7055	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Toukyoumubouaikoku	Tokyo	7058	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku 3	Tochigi	7084	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokusai 1	Saitama	7085	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku 20	Akita	7095	improved	Hap4	<i>E1/Ghd7-2</i>
Aikoku 6	Gunma	7096	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Waseaikoku 30	Gunma	7097	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Shizuokaikoku 1	Shizuoka	7189	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku 5	Fukushima	7198	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
A Aikoku 1	Miyagi	7228	improved	Hap2	<i>E1/Ghd7-2</i>
Aikoku 3	Tochigi	7332	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku A	Kanagawa	7556	landrace	Hap4	<i>E1/Ghd7-2</i>
Choubouaikoku	Kantotozan	7560	landrace	Hap2	<i>E1/Ghd7-2</i>
Shinkane/Aikoku	Kanagawa	7576	improved	Hap2	<i>E1/Ghd7-2</i>
Kairyuaikokudaikokugata 1	Miyagi	7621	improved	Hap3	<i>e1-ret/ghd7-0ret</i>

Kairyuaikokudaikokugata 2	Miyagi	7622	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Banaikoku	Gifu	9192	landrace	Hap4	<i>E1/Ghd7-2</i>
Aikoku 68	Mie	9277	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku 1 (Kyouto)	Kyoto	9285	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku	Shiga	9310	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikokumochi	Mie	9412	landrace	Hap4	<i>E1/Ghd7-2</i>
Aikoku B	Osaka	9513	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku C	Osaka	9514	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku D	Osaka	9515	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku F	Osaka	9516	improved	Hap2	<i>E1/Ghd7-2</i>
Aikoku G	Osaka	9517	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku	Tokushima	9650	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku 1	Shimane	9823	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku 2	Shimane	9824	improved	Hap3	<i>e1-ret/ghd7-0ret</i>
Aikoku	Kantotozan	10993	landrace	Hap4	<i>E1/Ghd7-2</i>
Aikoku (4X)	Fukuoka	10994	improved	Hap4	<i>E1/Ghd7-2</i>
Mubouaikoku	Ibaraki	14931	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
Kairyuaikoku	Osaka	15007	improved	Hap2	<i>E1/Ghd7-2</i>
Waseaikoku 3	Kanagawa	15051	landrace	Hap3	<i>e1-ret/ghd7-0ret</i>
W (Waseaikoku)	Akita	37731	unknown	Hap3	<i>e1-ret/ghd7-0ret</i>

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**Table S5. Primer and probe sequences and probes ID for quantitative RT-PCR**

Gene	Forward sequence	Reverse sequence	product size	catalog number of probe <sup>1)</sup>	probe sequence
<i>Ghd7</i>	GAGGGAGGCAAGCTGAT	ACCGATTGCTTCTCGTAG	68	#35 468768001	agaagagga
<i>Ubiquitin</i>	CTCCGTGGCGGTATCATC	TCTTCTGCGGCAGTTGAC	124	#124 4693582001	no data

<sup>1)</sup> The catalog number of probe is based on the Universal Probe Library designed with ProbeFinder version 2.45 (Roche; <https://www.roche-applied-science.com/>).



**Table S6. Primers used for haplotype analysis**

Name of primers	Forward primer	Reverse primer	annealing temperture (°C)	Product size (bp)	SNP	Enzyme <sup>1)</sup>
Haplotype_INDEL	CCCCCAAACATTTTCAGAAC	CTGCACCTAATCCGAATTCA	55	234 <sup>2)</sup>	-	-
Haplotype_SNP1	ATGTCATGCGGTGGGATTG	CCGAAGGCCAAACTAGGAAG	62	623	G/T <sup>3)</sup>	Hpy188I
Haplotype_SNP2	GCTTGAAGATTGGGAGCTTG	CACTAGAAGGAGTTCACCTTTTCTTC	55	295	C/T <sup>3)</sup>	Hpy188I

<sup>1)</sup> Enzyme is used in CAPS marker analysis.

<sup>2)</sup> The number indicates PCR product size without the 20bp deletion.

<sup>3)</sup> The nucleotides indicate the Nipponbare and EG5 genome sequence (left) and Kirara397 genome sequence (right).