

OPEN

# Author Correction: Molecular marker assisted breeding and genome composition analysis of Zhengmai 7698, an elite winter wheat cultivar

Chun-xin Li, Wei-gang Xu, Rui Guo, Jian-zhou Zhang, Xue-li Qi, Lin Hu & Ming-zhong Zhao

Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-017-18726-8>, published online 10 January 2018

This Article contains errors in Table 1. In the HTML and PDF versions of this Article, the denotation of the genes used in F<sub>3</sub>-F<sub>6</sub> generations, and F<sub>7</sub> and subsequent stable generations is incorrect.

Additionally, the legend of Table 1 is incorrect:

“**Table 1.** Names and their primer sequences of markers used in the marker assisted selection of Zhengmai 7698 (the label √ means the existence of that gene).”

should read:

“**Table 1.** Names and their primer sequences of markers used in the marker assisted selection of Zhengmai 7698. \*: genes used in F<sub>3</sub>-F<sub>6</sub> generations; \*\*: genes used in F<sub>7</sub> and subsequent stable generations; √: the existence of that gene. The primers of markers Pm2, Pm4b and Pm8 were presented, respectively, in Mohler et al. *Theor Appl Genet* 93: 1078–1082 (1996), Ma et al. *Theor Appl Genet* 109: 140–145 (2004) and Wang et al. *Acta Genetica Sinica* 28(7), 640–646 (2001).”

The correct Table 1 and its accompanying legend appears below.

This Article contains errors in the Results section under subheading ‘Molecular markers for Zhengmei 7698’.

“In the F<sub>3</sub>–F<sub>6</sub> generations of the selection processes, seven superior genetic markers in the parents were used (genes underlined by straight line in Table 1) to monitor the progress of gene pyramiding in hybrid offspring (e.g., single plant and line), and in the F<sub>7</sub> and subsequent stable generations, and 19 newly developed genetic markers were progressively added (genes underlined by wavy line in Table 1).”

should read:

“In the F<sub>3</sub>–F<sub>6</sub> generations of the selection processes, seven superior genetic markers in the parents were used (genes denoted with an \* in Table 1) to monitor the progress of gene pyramiding in hybrid offspring (e.g., single plant and line), and in the F<sub>7</sub> and subsequent stable generations, and 19 newly developed genetic markers were progressively added (genes denoted with an \*\* in Table 1).”

Furthermore, this Article contains errors in the Reference list. References 15, 18, 22 and 23 are incorrectly given as follows:

‘Geng, H. W., He, Z. H., Zhang, L. P., Qu, Y. Y. & Xia, X. C. Development of functional markers for a lipoxygenase gene on chromosome 4BS in common wheat. *Crop Sci.* 52, 568–576 (2012).’

Traits		Gene	Zhengmai 7698	Parents			Markers and their primers			Reference
Superior traits	High quality gene			4B269	Zhengmai 9405	Zhoumai 16	Marker	Forward primers (5' - 3')	Reverse primers (5' - 3')	
High molecular weight glutenin subunits		Ax1**	✓	✓			UMN19	CGAGACAATATGAGCAGCAAG	CTGCCATGGAGAAGTTGGA	10
		Ax-null**			✓	✓	UMN19	CGAGACAATATGAGCAGCAAG	CTGCCATGGAGAAGTTGGA	10
	Bx7*	Bx7*	✓	✓	✓	✓	Bx7	CACTGAGATGGCTAACGCAGCC	GCCTTGGACGGCACACAGG	11
	By8**	By8**		✓	✓		ZSBy8	TTAGCGCTAACGTGCGCTCT	TTGTCTTATTGCTGCCCTT	12
	By9**	By9**	✓			✓	ZSBy9a	TTCTCTGCATCAGTCAGGA	AGAGAAAGCTGTGTAATGCC	12
	Dx2**					✓	UMN25	GGGACAATACGAGCAGCAA	CTTGTCCGGTTGTTGCCA	10
	Dx5*	Dx5*	✓	✓	✓		Dx5	CGTCCCTATAAAAGCCTAGC	AGTATGAAACCTGCTGCCGAC	13
	Dy10**	Dy10**	✓	✓	✓		UMN26	CGCAAGACAATATGAGCAA	TTGCCTTGTCTGTGTC	10
	Dy12**	Dy12**				✓	UMN26	CGCAAGACAATATGAGCAA	TTGCCTTGTCTGTGTC	10
Grain hardness	Pinb-D1a*	Pinb-D1a*	✓	✓	✓		Pinb-D1a	ATGAAGACCTTATTCCCTCTA	CTCATGCTCACAGGCC	14
	Pinb-D1b*					✓	Pinb-D1b	ATGAAGACCTTATTCCCTCTA	CTCATGCTCACAGGCC	14
Lipoxygenase	TaLox-B1a	TaLox-B1a	✓		✓		LOX16	CCATGACCTGATCCTCCCTT	GCGCGGATAGGGGTGGT	15
	TaLox-B1b	TaLox-B1b		✓		✓	LOX18	ACGATGTGAGTTGACTGTGA	GCGCGGATAGGGGTGC	15
Yellow pigment content	Psy-A1a**					✓	YP7A	GGACCTTGTGATGACCGAG	TGACGGTCTGAAGTGAGAATGA	16
	Psy-A1b**	Psy-A1b**	✓	✓	✓		YP7A	GGACCTTGTGATGACCGAG	TGACGGTCTGAAGTGAGAATGA	16
	Psy-B1a**	Psy-B1a**	✓		✓	✓	YP7B-1	GCCACAACTTGAATGTGAAAC	ACTCTTCCATTGAACCCC	17
	Psy-B1b**	Psy-B1b**	✓				YP7B-1	GCCACAACTTGAATGTGAAAC	ACTCTTCCATTGAACCCC	17
	Psy1-D1g		✓	✓			YP7D-1	TCCGACACCACCAAGTTCC	CGTTGAGTTGTGGGAGT	18
Polyphenol oxidase activity	PPO-A1a**		✓			✓	PPO18	AACTGCTGGCTCTTCTCCA	AAGAAGTTGCCATGTCCGC	19
	PPO-A1b**	PPO-A1b**	✓		✓		PPO18	AACTGCTGGCTCTTCTCCA	AAGAAGTTGCCATGTCCGC	19
	PPO-D1a**	PPO-D1a**	✓			✓	PPO16	TGCTGACCGACCTTGACTCC	CTCGTCACCGTCACCCGTAT	20
	PPO-D1b**		✓	✓			PPO29	TGAAGCTGCCGGTCATCTAC	AAGTTGCCATGTCCCGCC	20
Powdery mildew resistant	Pm2*	Pm2*	✓	✓	✓	✓	Pm2	AGCTGTTGGTACAAGGTG	GCCATGTTTCTACTAG	21
	Pm4b*	Pm4b*	✓		✓	✓	Pm4b	GTGGTGTATCAAATGTCATCA GTACTAC	TCCAGTGACCCATCTGCTCATAC	21
	Pm8*	Pm8*	✓	✓	✓	✓	Pm8	GGAGACATCATGAAACATTG	CTGTTGTTGGCAGAAAG	21
Yellow rust resistant	Yr9**	Yr9**	✓	✓		✓	Xgwm582	AAGCACTACGAAAATATGAC	TCTTAAGGGGTATTACATA	22
	YrZH84**	YrZH84**	✓		✓		Xcfa2040	TCAAATGATTCAGGTAACCACT	TTCTGATCCCCACCAACAT	23
Pre-harvest sprouting resistant	PHS1**	PHS1**	✓	✓		✓	PHS1	GGTGAACAGATGAACTAAAGG/ GGTGAACAGATGAACTAAAGA	GTGAGTGTATATGAAACTAATG ATCCATT	24
	PHS-4AL**	PHS-4AL**	✓	✓			PHS-4AL	TGGAGTCTGAAAGCATTGCA/ TGGAGTCTGAAAGCATTGCG	TCCATGCATCATAGGAAACAA	25

**Table 1.** Names and their primer sequences of markers used in the marker assisted selection of Zhengmai 7698. \*: genes used in F<sub>3</sub>-F<sub>6</sub> generations; \*\*: genes used in F<sub>7</sub> and subsequent stable generations; ✓: the existence of that gene. The primers of markers Pm2, Pm4b and Pm8 were presented, respectively, in Mohler et al. Theor Appl Genet 93: 1078–1082 (1996), Ma et al. Theor Appl Genet 109: 140–145 (2004) and Wang et al. Acta Genetica Sinica 28(7), 640–646 (2001).

‘Wang, L. H. et al. Characterization of low-molecular-weight glutenin subunit Glu-B3 genes and development of STS markers in common wheat (*Triticum aestivum* L.). *Theor. Appl. Genet.* **118**, 525–539 (2009).’

‘Weng, D. X. et al. Microsatellite marker linked with stripe rust resistant gene Yr9 in wheat. *Acta Genetica Sinica* **32**, 937–941 (2009).’

‘Li, Z. F. et al. Molecular tagging of stripe rust resistance gene. YrZH8425B. *Theor. Appl. Genet.* **112**, 1098–1103 (2006).’

The correct references 15, 18, 22 and 23 appear below as references 1–4.

## References

1. Geng, H. W., He, Z. H., Zhang, L. P., Qu, Y. Y. & Xia, X. C. Development of functional markers for a lipoxygenase gene TaLox-B1 on chromosome 4BS in common wheat. *Crop Sci* **52**, 568–576 (2012).
2. Wang, J. W., He, X. Y., He, Z. F., Wang, H. & Xia, X. C. Cloning and phylogenetic analysis of phytoene synthase 1 (Psy1) genes in common wheat and related species. *Hereditas* **146**, 208–256 (2009).
3. Weng, D. X. et al. Microsatellite marker linked with stripe rust resistant gene Yr9 in wheat. *Acta Genetica Sinica* **32**, 937–941 (2005).
4. Li, Z. F. et al. Molecular tagging of stripe rust resistance gene YrZH84 in Chinese wheat line Zhou 8425B. *Theor. Appl. Genet.* **112**, 1098–1103 (2006).



**Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2019