

## **Analysis of lysophospholipid content in low phytate rice mutants**

Chuan Tong<sup>1,2</sup>, Yaling Chen<sup>1</sup>, Yuanyuan Tan<sup>3</sup>, Lei Liu<sup>2,\*</sup>, Daniel L. E. Waters<sup>2</sup>, Terry J. Rose<sup>2,4</sup>, Qingyao Shu<sup>3</sup>, Jinsong Bao<sup>1,\*</sup>

<sup>1</sup>Institute of Nuclear Agricultural Sciences, College of Agriculture and Biotechnology, Zhejiang University, Huajiachi Campus, Hangzhou, 310029, China.

<sup>2</sup>Southern Cross Plant Science, Southern Cross University, Lismore, NSW, 2480, Australia.

<sup>3</sup>Institute of Crop Sciences, College of Agriculture and Biotechnology, Zhejiang University, Hangzhou, 310029, China.

<sup>4</sup>Southern Cross Geoscience, Southern Cross University, Lismore, NSW, 2480, Australia.

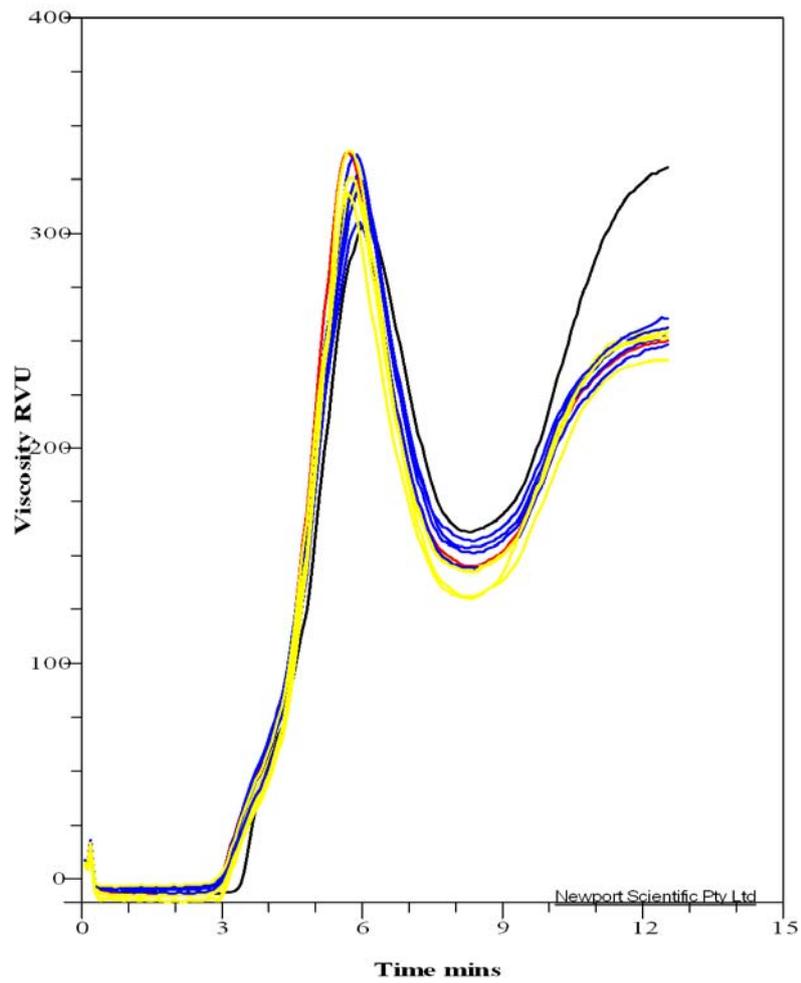
### **\*Corresponding authors:**

**Jinsong Bao**, phone +86-571-86971932, fax +86-571-86971421 and emails: jsbao@zju.edu.cn (J. Bao).

**Lei Liu**, phone +61-02-6622-3211, fax +61-02-6622-3459 and emails: lliu@scu.edu.au (L. Liu)

**Table S1.** Primers used for gel-based genotyping of *Wx* and *lpa1* genes

Markers	Primer sequence(5'-3')	Tm (°C)	Restriction enzyme	Fragment size (bp)	
				WT	MT
W <sub>x</sub> -SSR	P <sub>1</sub> F: CTTTGTCTATCTCAAGACAC	51.5	-	123	123
	P <sub>1</sub> R: TTGCAGATGTTCTTCCTGATG	56.8			
W <sub>x</sub> 484-W2R	P <sub>1</sub> F: CTTTGTCTATCTCAAGACAC	51.5	<i>Acc I</i>	253	126+127
	P <sub>3</sub> R: TTTCCAGCCCAACACCTTAC	58.6			
LPA1-CAPS	P <sub>2</sub> F:AGGGAGAAGTTCCTAAGTGTGT	59.6	<i>Xsp I</i>	184	121+63
	P <sub>2</sub> R:GGTAGCTTTCCATTGGCTGT	58.7			



**Figure S1.** RVA pasting properties of KBNT-*lpa* and Jiahe 218 derived F<sub>7</sub> breeding lines. \*Black is KBNT-*lpa* line; red is Jiahe218; yellow are those wild breeding lines and the blue are those mutant breeding lines.