## **Supplementary information**

## Cytoplasmic-genetic male sterility gene provides direct evidence for some hybrid rice recently evolving into weedy rice

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Sampling code	Abbreviation Province		Source location (Population site)	
WRLN001-01	LN1	Liaoning	Shenyang	
WRLN002-01	LN2	Liaoning	Shenyang	
WRLN002-01	LN2 Liaoning		Shenyang	
WRLN004-02	LN4	Liaoning	Dandong	
WRLN005-01	LN5	Liaoning	Dandong	
WRLN006-02	LN6	Liaoning	Panjin	
WRLN007-20	LN7	Liaoning	Panjin	
WRLN008-01	LN8	Liaoning	Panjin	
WRLN009-01	LN9	Liaoning	Tieling	
WRLN010-01	LN10	Liaoning	Tieling	
WRLN010-02	LN11	Liaoning	Tieling	
WRLN010-03	LN12	Liaoning	Tieling	
WRJS022-03	JS1	Jiangsu	Yangzhou	
WRJS022-04	JS2	Jiangsu	Yangzhou	
WRJS022-05	JS3	Jiangsu	Yangzhou	
WRJS022-06	JS4	Jiangsu	Yangzhou	
WRJS024-13	JS5	Jiangsu	Yangzhou	
WRJS024-14	JS6	Jiangsu	Yangzhou	
WRJS013-20	JS7	Jiangsu	Taizhou	
WRJS014-16	JS8	Jiangsu	Taizhou	
WRJS038-14	JS9	Jiangsu	Nantong	
WRJS038-15	JS10	Jiangsu	Nantong	
WRJS038-20	JS11	Jiangsu	Nantong	
WRJS039-02	JS12	Jiangsu	Nantong	
WRGD001-01	GD1	Guangdong	Zhanjiang	
WRGD001-02	GD2	Guangdong	Zhanjiang	
WRGD001-13	GD3	Guangdong	Zhanjiang	
WRGD001-14	GD4	Guangdong	Zhanjiang	
WRGD003-09	GD5	Guangdong	Zhanjiang	
WRGD003-13	GD6	Guangdong	Zhanjiang	
WRGD011-02	GD7	Guangdong	Zhanjiang	
WRGD011-03	GD8	Guangdong	Zhanjiang	
WRGD012-06	GD9	Guangdong	Zhanjiang	
WRGD012-07	GD10	Guangdong	Zhanjiang	
WRGD013-03	GD11	Guangdong	Zhanjiang	
WRGD013-04	GD12	Guangdong	Zhanjiang	

**Supplementary Table S1** Source of 36 weedy rice accessions for initial scrutiny of possible origin from commercial hybrid rice.

Population code	Sampling number	Province	Source location	
(number)		L'au	(Population site)	
WRJS007	4	Jiangsu	Lianyungang	
WRJS013	4	Jiangsu	Taizhou	
WRJS014	4	Jiangsu	Taizhou	
WRJS015	4	Jiangsu	Taizhou	
WRJS023	4	Jiangsu	Yangzhou	
WRJS024	4	Jiangsu	Yangzhou	
WRJS025	4	Jiangsu	Yangzhou	
WRJS026	4	Jiangsu	Yangzhou	
WRJS027	4	Jiangsu	Yangzhou	
WRJS035	4	Jiangsu	Taizhou	
WRJS036	4	Jiangsu	Yangzhou	
WRJS037	4	Jiangsu	Nantong	
WRJS038	4	Jiangsu	Nantong	
WRJS044	4	Jiangsu	Changzhou	
WRJS045	4	Jiangsu	Changzhou	
WRJS046	4	Jiangsu	Changzhou	
WRJS047	4	Jiangsu	Yangzhou	
WRJS049	4	Jiangsu	Taizhou	
WRJS050	4	Jiangsu	Nantong	
WRJS051	4	Jiangsu	Changzhou	
WRJS052	4	Jiangsu	Nantong	
WRJS053	4	Jiangsu	Yangzhou	
WRJS054	4	Jiangsu	Yangzhou	
WRJS055	4	Jiangsu	Yancheng	
WRJS056	4	Jiangsu	Suqian	
WRJS057	4	Jiangsu	Yancheng	
WRJS058	4	Jiangsu	Yancheng	
WRJS059	4	Jiangsu	Yangzhou	
WRJS060	4	Jiangsu	Taizhou	
WRJS061	4	Jiangsu	Zhenjiang	
WRJS062	4	Jiangsu	Yangzhou	
WRJS063	4	Jiangsu	Lianyungang	
WRJS064	4	Jiangsu	Lianyungang	
WRJS065	4	Jiangsu	Xuzhou	
WRJS068	4	Jiangsu	Yancheng	
WRJS069	4	Jiangsu	Yancheng	
WRJS070	4	Jiangsu	Huaian	
WRJS072	4	e		
WRJS072 WRJS074	4	e		
WRJS075	4	Jiangsu	Nanjing Nanjing	

Supplementary Table S2 Source of additional 322 weedy rice accessions for scrutiny of possible origin from commercial hybrid rice

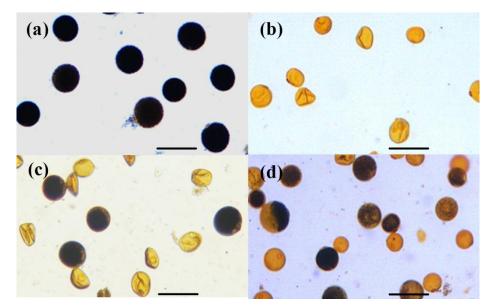
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WRJS076	4	Jiangsu	Changzhou
WRJS077	4	Jiangsu	Suzhou
WRJS078	4	Jiangsu	Wuxi
WRJS079	4	Jiangsu	Nanjing
WRSH002	3	Shanghai	Chongming
WRAH005	3	Anhui	Wuhu
WRAH017	3	Anhui	Maanshan
WRHN001	3	Henan	Xinyang
WRJX005	3	Jiangxi	Shangrao
WRZJ007	3	Zhejiang	Shaoxing
WRHU007	3	Hunan	Changde
WRHU011	3	Hunan	Yiyang
WRYN001	3	Yunnan	Kunming
WRYN007	3	Yunnan	Dali/Dali
WRGZ003	3	Guizhou	Anshun
WRGX001	3	Guangxi	Liuzhou
WRGD001	3	Guangdong	Zhanjiang
WRGD002	3	Guangdong	Zhanjiang
WRGD003	3	Guangdong	Zhanjiang
WRGD004	3	Guangdong	Zhaoqing
WRGD005	3	Guangdong	Zhaoqing
WRGD008	3	Guangdong	Maoming
WRGD009	3	Guangdong	Maoming
WRGD011	3	Guangdong	Zhanjiang
WRGD012	3	Guangdong	Zhanjiang
WRGD013	4	Guangdong	Zhanjiang
WRGD016	3	Guangdong	Zhanjiang
WRGD022	5	Guangdong	Zhanjiang
WRGD023	5	Guangdong	Zhanjiang
WRGD024	3	Guangdong	Yangjiang
WRGD025	3	Guangdong	Zhaoqing
WRGD026	3	Guangdong	Zhanjiang
WRGD027	3	Guangdong	Zhaoqing
WRGD028	3	Guangdong	Zhanjiang
WRGD029	3	Guangdong	Yangjiang
WRGD030	3	Guangdong	Zhanjiang
WRGD031	3	Guangdong	Zhanjiang
WRGD032	3	Guangdong	Zhanjiang
WRGD033	3	Guangdong	Zhaoqing
WRGD034	3	Guangdong	Zhaoqing
WRHA008	3	Hainan	Dongfang
WRHA009	3	Hainan	Dongfang
WRHA010	3	Hainan	Dongfang
WRHA011	3	Hainan	Dongfang
WRHA013	3	Hainan	Lingshui
WRHA014	3	Hainan	Lingshui
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WRHA015	3	Hainan	Lingshui
WRHA016	3	Hainan	Yanzhou
WRHA018	3	Hainan	Qionghai
WRHA019	3	Hainan	Wenchang
WRHA020	3	Hainan	Haikou

Character	Score/character state				
	0	1	2	3	4
Glume hairiness	Short, regular	Short, less regular	Nearly glabrous	Long, irregular	Long, overlapped
Phenol reaction	Dark	Light	Very light	Trace	None
Interval between 1st and 2nd node of panicle axis, cm	1.5	2.0	2.5	3	3.5
Glume color at heading	Greenish white	Whitish green	Yellowish green	Light green	Green
Leaf pubescence	Very dense	Dense	Medium	Few	No
Length-width ratio of spikelets	4.0	3.5	3.0	2.5	2.0

## Supplementary Table S3 The index characters and their scoring standards of the Cheng's Index Method

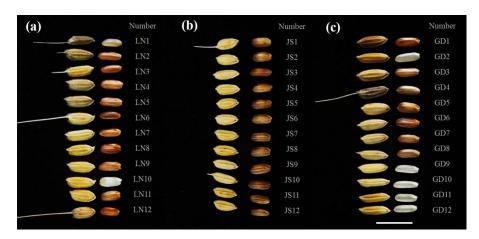
Subspecies type is judged by totaling the scores of the six indexes, 0~8 considered as *indica* type, 9-13 for *indica*-cline type , 14-17 for *japonica*-cline type, 18-24 for *japonica* type.



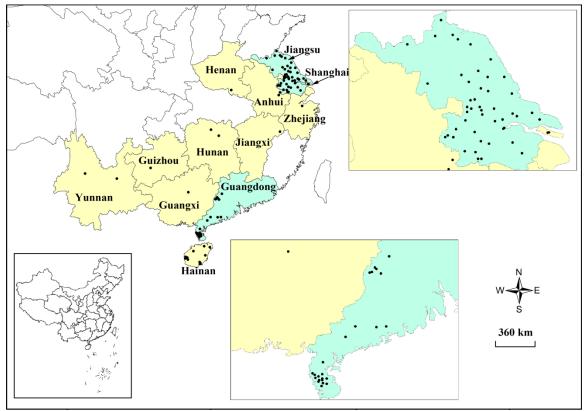
**Supplementary Figure S1** Morphology of pollen grains used to determine their viability. (a) Normal fertile pollen grains stain black and exhibit similar particle size; (b) Aborted pollen grains stain yellow and are deformed; (c) Easily distinguished fertile and aborted pollen grains in hybrid progeny for which LTPA was the female parent and weedy rice was the male parent; (d) Difficult-to-distinguish fertile and aborted pollen grains in hybrid progeny for which S28A was the female parent and weedy rice was the male parent, because the presence of the Boro-type (BT) abortive pollen grains with spherical shape, different particle size and staining. Pollen grains were stained with  $1\% I_2$ -KI solution. Bars,  $50 \mu m$ .



**Supplementary Figure S2** Corroboration of results obtained by *cms* marker using orWA352. Marker orWA352 only amplifies a 1432bp fragment from samples containing CMS-WA. Four wild abortive (WA) type sterile line samples and sixteen weedy rice accessions were confirmed to contain CMS-WA. The results corresponded to those obtained by the *cms* marker. From left to right: Marker (D2000 DNA ladder), Zhenshan97A, Zhenshan97B, Tianfeng A, Tianfeng B, Zhenpin A, Zhenpin B, LTPA, LTPB, WRJS026-01 (Jiangsu), WRHA016-01, WRHA016-03 (Hainan), GD9, GD10, GD11, GD12, WRGD012-04, WRGD013-06, WRGD013-13, WRGD013-17, WRGD022-03, WRGD022-06, WRGD022-09, WRGD022-10, WRGD022-13 (Guangdong). Sequenced amplified products contained the *WA352* gene.



**Supplementary Figure S3** Caryopses of 36 weedy rice populations from three provinces in China. (a) Liaoning; (b) Jiangsu; (c) Guangdong. Bars, 10mm.



**Supplementary Figure S4** Collection sites of 91 weedy rice populations in 12 provinces where hybrid rice is widely planted. Maps generated using ArcGIS 10.0.