

Fine mapping of rice drought QTL and study on combined effect of QTL for their physiological parameters under moisture stress condition

K. Baghyalakshmi1*, P. Jeyaprakash¹ , S. Ramchander¹ , M. Raveendran² and S. Robin¹

¹Centre for Plant Breeding and Genetics, Tamil Nadu Agricultural University, Coimbatore - 641003 (Tamil Nadu), INDIA

²Department of Plant Biotechnology, Centre for Plant Molecular Biology and Biotechnology, Tamil Nadu Agricultural University, Coimbatore - 641003 (Tamil Nadu), INDIA

*Corresponding author. E-mail: kauverik@gmail.com

Received: February 4, 2016; Revised received: August 4, 2016; Accepted: November 3, 2016

Abstract: The present investigation was undertaken to study the effect of different yield QTL (DTY_{2.2}, DTY_{3.1} and $DY_{8,1}$) under drought and their physiological response to drought stress. Backcross Inbred Lines (BILs) of IR64 (CB -193 and CB-229) along with IR64, APO and the traditional rice variety Norungan were raised in green house condition under water stress and control to evaluate the effect of the QTL on grain yield. The BIL CB-193 recorded higher photosynthetic rate (22.051), transpiration rate (7.152) and Ci/Ca ratio (0.597) whereas the BIL CB-229 recorded high relative water content (80.76%). It was found that the combination of three QTL (CB-229) performed better than the susceptible parent and the line with two QTL (CB-193 Fine-mapping of two QTLs viz., $qDTY_{2,2}$ and $qDTY_{8,1}$, for grain yield (GY) were conducted using backcross derived lines. Composite interval mapping analyses resolved the originally identified $qDTY_{2.2}$ region of 6.7 cM into a segment of 2.1 cM and two sub QTLs at region between RM23132 and RM1578 (75.75 cM- 77.66 cM), RM515 and RM1578 (75.11 cM-77.66 cM) were identified in $qDTY_{81}$ region. However this study provides a unique opportunity to breeders to introgress such regions together as a unit into high-yielding drought-susceptible varieties through MAS.

Keywords: Backcross inbred lines, Drought QTL, Fine mapping, Photosynthetic efficiency, Rice

INTRODUCTION

Rice (*Oryza sativa* L.) is the world's most important food crop and a primary food source for about half of the world's population. Rainfed rice occupies 50% of the total rice area in the world. Over 700 million people depend on rainfed rice grown on 50 million ha in Asia. Rice is semi aquatic plant and hence water loving plant. Drought is the major abiotic stress limiting rice production in these areas. Even a shorter spell of drought during reproductive stage has a greater impact on yield loss. In India, drought is major constraint for rice production and accounts for as much as 15 per cent of yield losses (Markandeya *et al.,* 2005). Varieties with high yield potential and improved drought tolerance could reduce risk and help alleviate poverty in drought-prone rice growing areas. Drought tolerant cultivars such as Apo, Vandana, Way Rarem, Nagina 22, and Adaysel have been shown to out-yield susceptible lines such as IR64 and IR72 by several-fold under upland and lowland drought (Venuprasad *et al*., 2007; Venuprasad *et al.,* 2009b; Bernier *et al*., 2007; Vikram *et al*., 2011; Swamy *et al.,* 2011). Crosses between such tolerant and susceptible lines are useful to identify important QTL underlying variation in grain yield under drought stress.

Molecular tools facilitate the identification and genomic locations of genes controlling traits related to drought tolerance using quantitative trait loci (QTL) analysis. Larger QTL region results in introgression of unwanted segments. Hence to introgress the precise segment the original QTL region has to be fine mapped and size has to be reduced. The present investigation was undertaken to study the effect of different yield QTL (DTY_2 ², DTY_3 ¹ and DTY_8 ¹) derived from Apo under drought and their physiological response to drought stress. The two QTLs namely *DTY2.2* and *DTY8.1* were fine mapped to locate the markers which were close to the QTL region. The original segment of *qDTY3.1* is 1.4cM hence fine mapping was not required (Venuprasad *et al.,* 2009b). Backcross Inbred Lines (BILs) of IR64 (CB-193 and CB-229) along with IR64, APO and the traditional rice variety Norungan were raised in green house condition under water stress and control to evaluate the effect of the QTL on grain yield.

MATERIALS AND METHODS

Two Backcross Inbred Lines (BILs) of IR64 developed from the cross combination of IR64 X APO along with the parents and Norungan a land race of Tamil Nadu tolerant to drought were used in the investigation.

ISSN : 0974-9411 (Print), 2231-5209 (Online) All Rights Reserved © Applied and Natural Science Foundation www.jans.ansfoundation.org

APO, drought tolerant upland variety, developed at IRRI, is recommended for cultivation under aerobic conditions. Owing to its drought tolerance nature and good performance under aerobic conditions, they serve as important source for mining drought tolerance QTLs. IR64 is a medium duration and high yielding variety but highly susceptible to drought. A set of 50 BILs from the cross between IR64 X APO were generated which carried three mega QTL classes namely *DTY2.2, DTY3.1* and *DTY8.1*.

Green house experiment: The two BIL lines of CB-193 (*DTY3.1* and *DTY8.1*), CB-229 (*DTY2.2, DTY3.1* and *DTY8.1*) were raised under control and water stress condition (pot culture experimentation) in four replication during Summer 2015. The seed materials were grown in plastic pots (30 cm height x 30 cm diameter with drainage hole) filled with three parts of coir pith and one part of natural clay loam soil. Three plants per pots were maintained and were grown in green house under natural temperature. The crop was irrigated till 45th day and there after irrigation was stopped for two replication (till the Relative Water Content reached 60%) and the yield parameters were recorded. The positions of the pots were changed frequently to minimize the micro climate effects. Infrared Gas Analyzer (IRGA), a portable photosynthetic system (LICOR- Model LI 6400 version.5) and used for the measurement of different parameters *viz*., photosynthetic rate, stomatal conductance, transpiration rate and Ci/Ca ratio. Relative water content was estimated by Weatherley (1950) method and expressed in percentage. The basic principle (Barrs and Weatherley, 1962) of this technique consists essentially in comparing the water content of leaf

tissue when fresh leaf sampled with the fully turgid water content and expressing the results on percentage basis. Several biometrical traits were also observed in both under control and moisture stress condition.

Fine mapping and genotyping

Generation of genotypic data: Fresh leaves for 50 BILs of the cross IR64xApo were collected and freezedried. DNA was extracted from freeze-dried leaf samples by a modified CTAB (cetyl tri-methyl ammonium bromide) method (Murray and Thompson, 1980) and the DNA was then quantified on 0.8% agrose gel to adjust the concentration to 25 ng/ μL^{-1} . Polymerase chain reaction was performed as described by Panaud *et al.* (1996). Agrose gel was used for size separation of the amplified DNA fragments using electrophoresis unit (BioRad 96). The DNA fragments were then visualized using trans-illuminator.

Genetic analysis: In this study, rice microsatellite (SSR) markers were used to fine map the previously identified QTL region for grain yield under stress. A total of 13 polymorphic markers for *qDTY2.2* and 12 polymorphic markers for *qDTY8.1*were added in the originally identified QTL regions in the BILs for fine mapping (Table 4). The markers were taken based on their physical position (Mb) on the *indica* genome (http:// www.ricetogo.org) and multiplied by 2.7 for approximate estimation of cM distances for analysis. Composite interval mapping (CIM) was performed using Windows QTL Cartographer 2.5.009 (Wang *et al.,* 2011). The LOD threshold value was obtained empirically from 1,000 permutation tests (Churchill and Doerge, 1994). The LOD thresholds obtained correspond to an experimentwise type I error rate of 0.05.

Fig. 1. *Parents and BILs under irrigated (C-control) and under drought condition (S- Stress).*

Fig 2c

Fig (a–c). *A QTL likelihood curves of LOD score for grain yield (GY) showing significant regions under severe stress. Genetic distance in cM between the markers is indicated on X axis. Horizontal lines correspond to critical LOD value. 2a - DTY 2.2 for number of productive tillers under severe drought; 2b-DTY 8.1 for number of productive tillers; 2c-DTY 8.1 for panicle length*

Fig. 3a. *Fine mapped region of qDTY2.2.*

Fig. 3b. *Fine mapped region of qDTY8.1.*

RESULTS AND DISCUSSION

Marker assisted selection using consistent yield QTLs under drought could be an alterative approach for improving rice grain yield under drought situations (Swamy *et al.,* 2011; Vikram *et al.,* 2012). QTLs have been identified in the past few years for different drought-related trait through phenotyping and genotyping of large mapping population. Various physiological parmeters *viz.,* photosynthetic rate, stomatal conductance, transpiration rate, Ci/Ca ratio and relative water content were measured in BILs namely CB-193 (*DTY3.1* and *DTY8.1*), CB-229 (*DTY2.2, DTY3.1* and *DTY8.1*), their parents (IR64 and APO) and check (Norungan) at different intervals (52, 58, 66,72 and 78 days after sowing) and the results were presented in Table 1. During initial stage of moisture reduction in pot culture at 52 DAS (days after sowing), the BIL CB

K. Baghyalakshmi et al. / J. Appl. & Nat. Sci. 8 (4): 1992-1998 (2016)				

Table 1. Estimation of physiological parameters in two Backcross Inbred Lines [CB-193 (*DTY3.1* and *DTY8.1*), CB-229 (*DTY2.2, DTY3.1* and *DTY8.1*)] of IR64 both under control and water stress condition (pot culture).

C – Control, S – Moisture Stress, DAS – days after sowing

-193 recorded higher photosynthetic rate (22.051), stomatal conductance (0.270), transpiration rate (7.152) and Ci/Ca ratio (0.597) whereas the BIL CB-229 recorded high relative water content (%). The plants

started expressing leaf rolling symptoms at 21 days after drought induction (Fig. 1). When moisture stress increased during 58, 66, 72 and 78 DAS, the line CB-229 registered higher gas exchange parameters espe-

			ъ.						
	DF			PH			NOPT		
Genotypes	$\mathbf C$	S	Per cent Reduction over con- trol	$\mathbf C$	S	Per cent Reduction over con- trol	$\mathbf C$	S	Per cent Reduction over con- trol
CB-193 (2QTL)	75	$62*$	17.33	95	$76*$	40.00	24	13	45.83
CB-229 (3QTL)	75	66	12.00	112	95	30.36	23	$15*$	34.78
Norungan	86	75	12.79	154	109	58.44	11	9	18.18
APO	81	73	9.88	110	87	41.82		6	14.29
IR64	$72*$	64	11.11	97	82	30.93	24	11	54.17
MEAN	77.8	68		114	89.8		18	10.8	
SE	2.52	2.55		10.70	5.72		3.70	1.56	

Table 2a. Yield attributing traits of Backcross Inbred Lines [CB-193 ($DTY_{3,l}$ and $DTY_{8,l}$), CB-229 ($DTY_{2,l}$, $DTY_{3,l}$ and $DTY_{8,l}$)] of IR64 both under control and water stress condition (pot culture). Define c and S below each table.

C – Control, S – Moisture Stress

Table 2b. Yield attributing traits of Backcross Inbred Lines [CB-193 (*DTY3.1* and *DTY8.1*), CB-229 (*DTY2.2, DTY3.1* and *DTY8.1*)] of IR64 both under control and water stress condition (pot culture).

C – Control, S – Moisture Stress; DF-days to 50% flowering, PH- plant height, NOPT- number of productive tillers, RL- root length, NFG- number of filled grains, SPF- spiklet fertility, HGW- hundred grain weight, SPY- single plant yield.

cially photosynthetic rate and relative water content as resistant parent APO and check Norungan. This clearly revealed that, the BIL line CB-229 maintained higher internal water status with normal physiological process of photosyntheis, stomatal conductance and transpiration rate and withstand under higher moister stress conditions. Under stress condition, the population varied widely for all the physiological traits studied *viz.,* photosynthetic rate, transpiration rate, stomatal conductance and relative water content. Apo had higher stomatal conductance when compared to IR64 and the BILs also had higher stomatal conductance than IR64. The same kind of results was obtained by Tezera *et al.* (2002) who reported that higher stomatal conductance would result in higher photosynthetic rate and biomass production. Martinez *et al.* (2007) also pointed out that higher stomatal conductance may be an enhanced adaptation of plants to drought environments. Chen *et al*. (1995) observed that elevating photosynthetic rate is

beneficial to dry matter production and yield. Cao *et al*. (2001) reported that photosynthetic rate among rice varieties were significant and suggested the net photosynthetic rate as a selection parameter for drought resistant genotypes. The similar result was obtained from this study also and Apo had more photosynthetic rate along with BILs when compared to IR64. Araus *et al*. (2002) reported that higher yielding genotypes under drought had greater stomatal conductance and transpiration rate. Sikuku *et al*. (2010) observed transpiration rate in NERICA rice varieties generally decreased with increase in soil water deficit.

Fine mapping: Fine mapping of *qDTY2.2* and *qDTY8.1* was done under severe stress wherein the yield traits were recorded. The drought QTL *qDTY3.1* was not fine mapped in this study since the closest linked marker was already reported by Venuprasad *et al.,* 2009a. CIM analysis of markers within *qDTY2.2* showed a region between RM12529 and RM12571 (13.87cM -

K. Baghyalakshmi *et al.* / *J. Appl. & Nat. Sci.* 8 (4): 1992-1998 (2016)

QTL-Fine mapped	Associated trait Marker Inerval		Peak position	Marker closest	${\bf R}^{2}($ %)	$\mathbf{Add}\%$	LOD
region	under stress			to the peak			score
qDTY2.2	Productive tillers	RM12529-	15.9cM	RM12569	19.0	28.5	$\overline{4}$
		RM12571					
qDTY8.1	Productive tillers	RM23132-	77.1cM	RM1597	23.0	15.3	2.9
		RM1578					
qDTY8.1	Panicle length	RM515-	75.8cM	RM6187	43.0	42.8	7.6
		RM1578					

Table 3. Fine-mapped regions within different QTLs for grain yield under stress.

Table 4. List of polymorphic markers used for fine mapping.

Chromosome 2		Chromosome 8			
Markers	Position	Markers	Position		
	(cM)		(cM)		
RM279	10.67	RM339	73.9		
RM12529	13.87	RM515	75.11		
RM1075	14.20	RM23132	75.75		
RM1285	14.21	RM6187	75.83		
RM3703	14.31	RM8265	75.85		
RM12539	14.61	RM4668	76.05		
RM12543	14.98	RM223	76.47		
RM12548	15.05	RM7049	77.07		
RM12551	15.21	RM1597	77.14		
RM12555	15.39	RM1578	77.66		
RM12569	15.86	RM5887	77.67		
RM12571	15.95	RM284	78.29		
RM71	33.0				

15.95cM) having an effect on GY under drought stress condition. The peak was detected at 15.9 cM under stress conditions with RM12569 as the closest marker to the peak at LOD 4 (Fig. 2a and 3a). The QTL explained a phenotypic variance of 19.0% and had an additive effect of 28.5 (Table 3). Similarly analysis of *qDTY8.1* exhibited two sub QTLs at region between RM23132 and RM1578 (75.75 cM- 77.66 cM), RM515 and RM1578 (75.11 cM-77.66 cM) for the traits *viz.,* panicle length and number of productive tillers under drought conditions with RM1597 and RM6187 as the closest markers to the peak respectively (Fig 2b and 2c & 3b). The peak was detected at 77.1 and 75.8 cM under stress conditions with LOD of 2.9 and 7.6 respectively. The QTL explained a phenotypic variance of 23.1% and 43% and had an additive effect of 15.3 and 42.8 (Table 3) under drought condition.

In the present investgation, several biometrical traits were observed in all genotypes under study which exhibited that, the BIL CB-229 had recorded higher number of productive tillers, root length, number of filled grains per panicle, spikelet fertility percentage, hundred grain weight and grain yield under moisture stress condition when compared to the BIL CB-193. This could be due to the presence of two QTL regions together within *qDTY8.1* showing larger effect in drought stress condition. However, *qDTY2.2*, *qDTY3.1* and *qDTY8.1* have shown a high and consistent additive effect under severe drought condition. The *qDTY2.2* exhibited higher effect on increased number of tillers

and *qDTY8.1* had an effect on panicle length and number of productive tillers thus increasing the yield under drought. Since, the QTL *DTY2.2* was responsible for root length (MacMillan *et al.,* 2006; Kamoshita *et al.,* 2002) and root thickness (Champoux *et al.,* 1995, Dixit *et al.,* 2012). In this study also, the BIL CB-229 possessing QTL *DTY8.1* was associated with grain yield under stress condition. The similar result was already reported by Vikram *et al.* (2012). But the significant advantage of CB-193 was earliness in flowering and reduced plant height. Both the BILs of IR64-193 (*DTY3.1* and *DTY8.1*) and IR64-229 (*DTY2.2, DTY3.1* and *DTY8.1*) were earlier in flowering with reduced plant height, which possessed *DTY3.1* responsible for flowering date and plant height (Venuprasad *et al.,* 2009a). Ding *et al.* (2011) reported complex nature of large-effect QTLs for stress tolerance working on a QTL on chromosome 4 controlling root volume per tiller co-segregating with flag -leaf width and spikelet number per panicle.

Conclusion

The present investigation was undertaken to study the effect of different yield QTL (*DTY2.2, DTY3.1* and *DTY8.1*) derived from Apo under drought and their physiological response to drought stress. The *qDTY2.2* showed larger effect under drought but a very low effect under moderate stress. The other two *qDTY* (*3.1* and *8.1*) showed its effect only under severe drought stress. It should be possible to exploit these fine mapped QTL after development of gene-based markers, for improving grain yield of valuable varieties such as IR64 in stress prone environments via marker assisted backcross breeding. In this study an attempt was made to narrow down the originally identified QTL regions so as to introgress precisely the smallest possible segment of these QTLs showing profound effect on grain yield under drought while minimizing the chances of introgression of any undesirable linked trait. The successful introgression of the fine-mapped regions of these QTLs by MAS provides an opportunity to improve the drought tolerance of well-adapted, highyielding but drought-susceptible popular rice varieties.

ACKNOWLEDGEMENTS

Authors profoundly acknowledge The Department of Biotechnology for funding the scheme "Comparative transcriptomics between rice and resurrection plants and exploitation of novel genes for dehydration tolerance for

improvement of drought tolerance in rice (Indo- Australia)" and successful completion of the research work.

REFERENCES

- Araus, J.L., Slafer, G.A., Reynolds, M.P. and Royo, C. (2002). Plant breeding and drought in C3 cereals: what should we breed for. *Ann. Bot.,* 89 : 925–940
- Barrs, H.D. and Weatherley. P.E. (1962). A reexamination of the relative turgidity technique for estimating water deficits in leaves. *Aust. J. Biol. Sci.*, 15:415–428
- Bernier, J., Kumar, A., Ramaiah, V., Spaner, D. and Atlin, G. (2007). A large-effect QTL for grain yield under reproductive-stage drought stress in upland rice. *Crop Sci.*, *47*: 507– 518
- Cao, S.Q., Zhai, H.Q., Yang, T.N., Zhang R.X. and Kuang, T.Y. (2001). Studies on photosynthetic rate and function duration of rice germplasm. *Chinese J. Rice Sci.,* 15(1): 29–34
- Champoux, M.C., Wang, G., Sarkarung, S., Mackill, D.J., O'Toole, J.C., Huang, N., McCouch S.R. (1995). Locating genes associated for root morphology and drought avoidance in rice via linkage to molecular markers. *Theor. Appl. Genet*., *90*: 969–981
- Chen, W.F., Xu, Z.J. and Zhang, B.L. (1995). Physiological bases of super high yield breeding in rice. Liao Ning Science and Technology Publishing Company, Shenyang, China.
- Churchill, G.A. and Doerge, R.W. (1994). Empirical threshold values for quantitative trait mapping. *Genetics.,* 138:963– 971
- Ding, X., Li, X. and Xiong, L. (2011). Evaluation of nearisogenic lines for drought resistance QTL and fine mapping of a locus affecting flag leaf width, spikelet number, and root volume in rice. *Theor. Appl. Genet.,* 123:815–826
- Dixit S., Mallikarjuna Swamy B.P., Prashant Vikram, Ahmed, H.U., Sta Curz, M.T., Modesto Amante, Dinesh Atri, Hei Leung and Aravind Kumar. (2012). Fine mapping of QTLs for rice grain yield under drought reveals sub-QTLs conferring a respose to variable drought severities. *Theor. Appl. Genet.,* 125: 155-169
- Kamoshita, A, Wade, L.J., Ali, M.L., Pathan, M.S., Zhang, J., Sarkarung, S. and Nguyen, H.T. (2002). Mapping QTLs for root morphology of a rice population adapted to rainfed lowland conditions. *Theor. Appl. Genet.,* 104:880–893
- MacMillan, K., Emrich, K., Piepho, H.P., Mullins, C.E. and Price, A.H. (2006). Assessing the importance of genotype 9 environmental interaction for root traits in rice using a mapping population. II.Conventional QTL analysis. *Theor. Appl. Genet.,* 113:953–964
- Markandeya, G., Babu, P.R. and Lachagari, V.B.R. (2005). Functional genomics of drought stress response in rice: transcript mapping of annotated unigenes of an indica rice (Oryza sativa L. cv. Nagina 22). *Curr. Sci., 89:496–514*

Martinez, J.P., Silva, H., Ledent, J.F. and Pinto, M. (2007).

Effect of drought stress on the osmotic adjustment, cell wall elasticity and cell volume of six cultivars of common beans (*Phaseolus vulgaris* L.). *Euro. J. Agro.*, 26: 30–38

- Murray, M.G. and Thompson, W.F. (1980). Rapid isolation of high molecular weight plant DNA. *Nucleic Acid Res*., 8, 4321-4326
- Panaud, O., Chen, X., McCouch, S. (1996). Development of microsatellite markers and characterization of simple sequence length polymorphism (SSLP) in rice (Oryza sativa L.). *Mol. Gen. Genet.,* 252:597–607
- Sikuku, P.A., Netondo, G.W., Onyango, J.C. and Musyimi, D. M. (2010). Effects of water deficit on physiology and Morphology of three varieties of NERICA Rainfed rice (*Oryza sativa* L.). *ARPN Journal of Agricultural and Biological Science,* 5: 1
- Swamy, B.P.M., Vikram, P., Dixit, S., Ahmed, H.U. and Kumar, A. (2011). Metaanalysis of grain yield QTL identified during agricultural drought in grasses showed consensus. *BMC Genomics.,* 12:319
- Tezara, W., Mitchel, V., Driscul, S.P. and Lawlor, D.W. (2002). Effects of water deficit and its interaction with CO2 supply on the biochemistry and physiology of photosynthesis in sunflower. *J. Exp. Bot.*, 53: 1781-1791
- Venuprasad, R., Lafitte, H.R. and Atlin, G.N. (2007). Response to direct selection for grain yield under drought stress in rice. *Crop Sci.,* 47:285–293
- Venuprasad, R., Dalid, C.O., Del Valle, M., Zhao, D., Espiritu, M., Sta Cruz M.T, Amante M, Kumar .A, Atlin G.N. (2009a). Identification and characterization of large-effect quantitative trait loci for grain yield under lowland drought stress in rice using bulk-segregant analysis. *Theor. Appl. Genet.,* 120:177–190
- Venuprasad, R., Bool, M.E., Dalid, C.O., Bernier, J., Kumar, A. and Atlin, G. N. (2009b). Genetic loci responding to two cycles of divergent selection for grain yield under drought stress in a rice breeding population. *Euphytica,* 167:261– 269
- Vikram, P., Mallikarjuna Swamy, B.P, Dixit, S., Ahmed, H.U., Sta Cruz M.T., Singh, A.K and Kumar, A. (2011). qDTY1.1, a major QTL for rice grain yield under reproductive-stage drought stress with a consistent effect in multiple elite genetic backgrounds. *BMC Genet*., doi:10.1186/1471-2156-12-89
- Vikram, P., Swamy, B.P.M., Dixit, S., Ahmed, H.. Sta Cruz, M.T., Singh, A.K., Ye, G. and Kumar, A. (2012). Bulk segregant analysis: "An effective approach for mapping consistent-effect drought grain yield QTLs in rice". *Field Crops Research,* 134: 185-192
- Wang, S., Basten, C.J. and Zeng, Z.B. (2011). Windows QTL cartographer 2.5. department of statistics, North Carolina State University, Raleigh, NC.
- Weatherley, P.E. (1950). Studies in the water relations of the cotton plant. I. The field measurements of water deficits in leaves. *New Phytol,* 49: 81-97