

S7 Table. List of major QTL (PVE > 20%) for seven drought-responsive traits identified in earlier studies along with their linked markers, PVE, mapping population type, and parental genotypes.

S.No.	Trait/QTL	Linked marker	PVE (R^2) ^a	Mapping population; parental genotypes	Favourable allele	Reference
1. Grain yield						
a)	<i>qGYWD.3B.2</i>	<i>Xgpw7774</i>	19.6	RIL; WL711(DS)/C306 (DT)	C306	[1]
b)	<i>4A</i>	<i>Xwmc420</i>	20.0	RIL; Sitta(DS)/Dharwar Dry (DT)	Dharwar Dry	[2]
c)	<i>4A-a</i>	<i>Xgwm397</i>	23.9	RIL; SeriM82/Babax	Babax	[3]
d)	<i>Qyld.csdh.7AL</i>	<i>Xgwm322</i>	20.0*	DH; Chinese Spring (DS)/SQ1(DT)	SQ1	[4]
2. 1000 grain weight						
a)	<i>3B</i>	<i>Xbarc101</i>	45.2	$F_{2:3}$ families; Massara-1 (DS)/ Oste-Gata (DT)	Oste-Gata	[5]
b)	<i>QTgw-7D-b</i>	<i>XC29-P13</i>	21.9	RIL; SeriM82/Babax	Babax	[6]
3. Days to Heading						
a)	<i>QDh-7D.b</i>	<i>XC29-P13</i>	22.7	RIL; SeriM82/Babax	Babax	[6]
b)	<i>QHd.idw-2A.2</i>	<i>Xwmc177</i>	32.2	RIL; Kofa/Svevo	Kofa	[7]
4. Days to Maturity						
a)	<i>QDm-7D.b</i>	<i>X7D-acc/cat-10</i>	22.7	RIL; SeriM82/Babax	Babax	[6]
5. Stem Reserve Mobilization						
a)	<i>QSrm.ipk-2D</i>	<i>Xgwm249a</i>	42.2	RIL; W7984/Opata 85	W7984	[8]
b)	<i>QSrm.ipk-5D</i>	<i>Xfbb238b</i>	37.5	RIL; W7984/Opata 85	W7984	[8]
c)	<i>QSrm.ipk-7D</i>	<i>Xfbb189b</i>	21.0	RIL; W7984/Opata 85	W7984	[8]
6. Water soluble carbohydrate						
a)	<i>QWsc-c.aww-3A</i>	<i>Xwmc0388A</i>	19.0	DH; Kukri (DS)/RAC875 (DT)	RAC875	[9]
7. SPAD/chlorophyll content						
a)	<i>Qchl.ksu-3B</i>	<i>Xbarc68</i>	59.1	RIL; HUW206 (DS)/C306 (DT)	C306	[10]

^aHighest PVE (R^2) values under drought/water stress; * with >20% higher yield per ear, DS, drought sensitive; DT, drought tolerant.

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