

Supplementary Table S1. Environmental conditions near the rice fields including monthly total rainfall, rainy day number, mean relative humidity (RH), minimum temperature (T_{\min}) and maximum temperature (T_{\max}) during January to December 2021 recorded by Upper Northeastern Meteorological Center, Mueang Khon Kaen District, Khon Kaen, Thailand. The electrical conductivity (EC) of soil and water of the rice fields including non-saline, semi-saline and heavy saline were measured during July to Nov 2021 (gray area).

Month	Total of rainfall (mm)	Rainy day number (day)	RH (%)	T_{\min} (°C)	T_{\max} (°C)	Electrical conductivity (dS m ⁻¹) in the fields					
						Soil			Water		
						Non-saline	Semi-saline	Heavy-saline	Non-saline	Semi-saline	Heavy-saline
January	0.0	0	58.64 ± 4.2	15.81 ± 3.0	28.86 ± 3.0	—	—	—	—	—	—
February	21.4	2	58.46 ± 8.5	19.01 ± 1.6	31.92 ± 3.3	—	—	—	—	—	—
March	92.3	6	66.48 ± 7.6	23.73 ± 1.9	34.61 ± 2.9	0.86 ± 0.11	13.45 ± 5.8	45.27 ± 13.07	—	—	—
April	135.8	9	71.06 ± 9.1	24.87 ± 1.3	34.57 ± 3.2	—	—	—	—	—	—
May	75.9	11	72.58 ± 7.0	25.69 ± 1.1	35.52 ± 2.0	0.12 ± 0.01	2.87 ± 0.02	2.53 ± 0.01	—	—	—
June	142.3	13	74.06 ± 9.1	25.47 ± 1.2	34.48 ± 2.9	—	—	—	—	—	—
July	114.8	15	78.16 ± 8.4	25.08 ± 1.0	33.40 ± 2.4	0.06 ± 0.01	1.08 ± 0.09	1.12 ± 0.01	0.06 ± 0.01	2.26 ± 0.22	2.97 ± 0.11
August	141.7	17	78.19 ± 7.1	25.04 ± 1.0	33.80 ± 1.8	0.04 ± 0.01	1.21 ± 0.14	1.32 ± 0.40	0.21 ± 0.25	1.78 ± 0.07	2.07 ± 0.05
September	271.5	23	86.83 ± 4.5	24.21 ± 0.8	31.58 ± 1.7	0.04 ± 0.03	1.05 ± 0.16	1.33 ± 0.29	0.06 ± 0.01	1.31 ± 0.28	1.48 ± 0.17
October	136.7	14	81.22 ± 5.5	23.44 ± 1.6	31.30 ± 2.2	0.48 ± 0.31	2.86 ± 1.58	4.11 ± 1.95	dry	dry	dry
November	0.0	0	69.93 ± 5.4	21.40 ± 2.0	31.37 ± 1.6	0.87 ± 0.14	4.05 ± 1.22	4.47 ± 1.51	dry	dry	dry
December	0.0	0	60.41 ± 5.0	17.78 ± 2.0	29.40 ± 1.5	—	—	—	—	—	—

Supplementary Table S2. SPAD values and photosynthetic parameters including net photosynthesis (P_n), stomatal conductance (gs), transpiration rate (T_r), water use efficiency (WUE), effective quantum yield of PSII photochemistry (Φ_{PSII}), maximum quantum yield efficiency of PSII (F_v/F_m') and electron transport rate (ETR) of five rice genotypes including Pokkali, RD73, CSSL8-94, TSKC1-144 and KDM105 growing under non-saline, semi-saline and heavy-saline field. The data were measured at six rice growth stages including early vegetative, stem elongation, early booting, flowering, milky and mature. Means which are significantly different ($p < 0.05$ and $p < 0.01$) among genotypes for each saline field and field across genotypes are denoted by different lower-case letters. For each genotype means which are significantly different ($p < 0.05$) among growth stage are denoted with capital letters. Data shows mean of four replicates ± standard error (SE).

Parameter	Saline field	Rice genotype	Early vegetative	Stem elongation	Early booting	Flowering	Milky	Mature	² Critical- p value	
									¹ Critical-p value	¹ Critical-p value
SPAD value	Non-saline	Pokkali	31.40 ± 0.60 bB	36.42 ± 0.33 bA	36.90 ± 0.48 A	36.55 ± 0.65 A	36.30 ± 0.80 bA	27.55 ± 1.27 bC	$p < 0.01$	
		RD73	34.47 ± 0.61 aDE	38.45 ± 0.57 aBC	36.32 ± 0.85 CD	41.00 ± 1.75 AB	42.20 ± 0.75 aA	32.30 ± 0.90 aE	$p < 0.01$	
		CSSL8-94	34.47 ± 0.40 aC	37.25 ± 0.29 abB	36.02 ± 0.39 BC	40.77 ± 1.34 A	41.35 ± 1.01 aA	30.52 ± 0.26 abD	$p < 0.01$	
		TSKC1-144	33.30 ± 0.33 abB	38.07 ± 0.41 aA	34.72 ± 1.12 B	40.40 ± 1.11 A	33.80 ± 1.06 cB	28.45 ± 0.65 bC	$p < 0.01$	
		KDM105	34.22 ± 0.58 aCD	37.47 ± 0.42 abB	35.07 ± 0.60 C	39.76 ± 0.25 A	41.56 ± 0.64 aA	32.75 ± 1.25 aD	$p < 0.01$	
	Semi-saline	¹ Critical-p value	$p < 0.01$	$p < 0.05$	ns	ns	$p < 0.01$	$p < 0.01$	$p < 0.01$	
		mean	33.57 ± 0.34 cD	37.53 ± 0.23 bB	35.81 ± 0.35 bC	39.69 ± 0.59 bA	39.04 ± 0.85 bA	30.31 ± 0.60 cE	$p < 0.01$	
		Pokkali	43.80 ± 1.13 A	42.22 ± 1.05 AB	39.92 ± 0.76 bB	42.30 ± 0.76 bcAB	43.45 ± 1.45 A	33.22 ± 2.32 C	$p < 0.01$	
		RD73	43.57 ± 0.90	40.50 ± 2.41	46.62 ± 1.90 a	48.65 ± 1.83 a	48.42 ± 1.89	43.37 ± 3.11	ns	
		CSSL8-94	43.10 ± 1.16 A	37.70 ± 1.48 C	42.60 ± 1.69 abAB	41.60 ± 0.97 cABC	45.30 ± 1.07 A	38.85 ± 2.18 BC	$p < 0.05$	
	Heavy-saline	TSKC1-144	45.82 ± 1.69	39.75 ± 0.85	40.22 ± 1.87 b	42.65 ± 1.96 bc	43.37 ± 1.65	37.75 ± 2.37	ns	
		KDM105	44.20 ± 0.74 A	39.27 ± 1.80 B	46.27 ± 1.01 aA	46.87 ± 1.19 abA	47.42 ± 1.18 A	37.05 ± 3.06 B	$p < 0.01$	

Parameter	Saline field	Rice genotype	Early vegetative	Stem elongation	Early booting	Flowering	Milky	Mature	² Critical- p value
									p value
P_n	Heavy-saline	¹ Critical-p value mean	ns	ns	p < 0.05	p < 0.05	ns	ns	² Critical- p value
		Pokkali	44.10 ± 0.51 aAB	39.89 ± 0.73 aC	43.13 ± 0.89 aB	44.41 ± 0.86 aAB	45.59 ± 0.75 aA	38.05 ± 1.29 aC	p < 0.01
		RD73	40.85 ± 1.55 B	46.46 ± 1.28 aA	39.87 ± 1.24 B	40.52 ± 1.04 bB	47.52 ± 0.35 aA	28.40 ± 1.23 bC	p < 0.01
		CSSL8-94	41.60 ± 1.27 B	40.57 ± 1.04 bB	44.02 ± 1.95 AB	45.45 ± 0.20 aA	47.20 ± 2.31 aA	33.60 ± 1.91 abC	p < 0.01
		TSKC1-144	42.65 ± 0.56 AB	37.87 ± 1.72 bC	41.65 ± 1.51 ABC	44.67 ± 1.61 aA	43.72 ± 0.63 abA	38.92 ± 1.91 aBC	p < 0.05
	Non-saline	KDML105	41.42 ± 1.47 AB	37.42 ± 0.41 bBC	43.35 ± 2.28 A	39.30 ± 1.56 bAB	40.50 ± 1.40 bAB	33.07 ± 2.89 bC	p < 0.05
		¹ Critical-p value mean	41.80 ± 0.88 B	40.90 ± 0.73 bB	43.50 ± 0.83 AB	46.02 ± 1.07 aA	45.45 ± 0.66 aA	32.10 ± 1.86 bC	p < 0.01
		Pokkali	ns	p < 0.01	ns	p < 0.01	p < 0.01	p < 0.05	
		RD73	ns	40.64 ± 0.87 aC	42.48 ± 0.74 aBC	43.19 ± 0.79 aAB	44.88 ± 0.78 aA	33.22 ± 1.12 bD	p < 0.01
		CSSL8-94	ns	23.13 ± 0.64 A	23.84 ± 0.79 A	22.26 ± 1.41 bA	19.40 ± 0.55 bB	18.29 ± 0.43 cB	p < 0.01
gs	Semi-saline	TSKC1-144	41.42 ± 1.47 AB	37.42 ± 0.41 bBC	43.35 ± 2.28 A	39.30 ± 1.56 bAB	40.50 ± 1.40 bAB	33.07 ± 2.89 bC	p < 0.05
		KDML105	41.80 ± 0.88 B	40.90 ± 0.73 bB	43.50 ± 0.83 AB	46.02 ± 1.07 aA	45.45 ± 0.66 aA	32.10 ± 1.86 bC	p < 0.01
		¹ Critical-p value mean	ns	ns	p < 0.05	p < 0.01	p < 0.01	ns	
		Pokkali	23.29 ± 0.38 bC	24.57 ± 0.45 ABC	25.79 ± 0.71 aA	24.81 ± 0.74 aAB	23.75 ± 0.77 aBC	14.08 ± 0.41 bD	p < 0.01
		RD73	23.29 ± 1.25 B	25.58 ± 1.22 AB	27.98 ± 1.07 aA	26.58 ± 0.82 aA	25.63 ± 0.54 aAB	13.90 ± 1.09 C	p < 0.01
	Heavy-saline	CSSL8-94	24.21 ± 0.38 A	24.44 ± 1.56 A	27.45 ± 1.23 aA	24.86 ± 0.50 aAB	23.03 ± 0.82 bA	15.13 ± 0.97 C	p < 0.01
		TSKC1-144	22.98 ± 0.86 B	23.98 ± 0.88 AB	24.58 ± 1.14 abAB	27.06 ± 0.96 aA	26.07 ± 1.33 aAB	14.45 ± 1.36 C	p < 0.01
		KDML105	22.82 ± 1.11 B	24.98 ± 0.62 AB	26.65 ± 1.59 aA	26.18 ± 1.50 aA	25.72 ± 1.12 aA	13.59 ± 0.54 C	p < 0.01
		¹ Critical-p value mean	ns	ns	p < 0.05	p < 0.01	p < 0.01	ns	
		Pokkali	27.54 ± 0.52 abA (+19%)	24.56 ± 1.47 AB (+3%)	23.97 ± 0.67 B (+8%)	22.77 ± 0.86 BC (+17%)	20.82 ± 1.41 CD (+14%)	17.96 ± 0.53 aD (+35%)	p < 0.01
gs	Non-saline	RD73	28.89 ± 0.42 aA (+24%)	25.25 ± 1.16 B (-1%)	21.22 ± 0.38 C (-24%)	23.63 ± 0.97 BC (-11%)	24.98 ± 1.37 B (-3%)	17.53 ± 1.86 aD (+26%)	p < 0.01
		CSSL8-94	25.85 ± 0.61 bA (-7%)	25.82 ± 0.65 A (+6%)	21.41 ± 1.19 B (-22%)	22.66 ± 0.60 B (-9%)	23.69 ± 1.15 AB (+3%)	15.90 ± 0.85 abC (-5%)	p < 0.01
		TSKC1-144	28.92 ± 0.86 aA (+26%)	25.95 ± 0.95 AB (+8%)	22.48 ± 1.12 CD (-9%)	21.69 ± 1.20 D (-20%)	25.61 ± 0.53 BC (-2%)	12.88 ± 1.23 bE (-11%)	p < 0.01
		KDML105	29.39 ± 0.32 aA (+29%)	25.41 ± 1.17 BC (+2%)	20.68 ± 0.46 D (-22%)	22.26 ± 0.64 CD (-15%)	25.65 ± 2.15 B (0%)	15.47 ± 0.21 abE (+14%)	p < 0.01
		¹ Critical-p value mean	p < 0.01	ns	ns	ns	ns	p < 0.05	
	Semi-saline	Pokkali	28.12 ± 0.37 aA	25.40 ± 0.46 B	21.95 ± 0.43 bC	22.60 ± 0.38 bC	24.15 ± 0.70 aB	15.95 ± 0.60 aD	p < 0.01
		RD73	25.93 ± 1.05 cAB (+12%)	22.83 ± 1.54 BC (-4%)	26.01 ± 1.28 A (+16%)	23.09 ± 0.99 ABC (+19%)	20.36 ± 1.15 C (+11%)	12.55 ± 0.96 D (-6%)	p < 0.01
		CSSL8-94	30.98 ± 0.68 abA (+33%)	28.86 ± 1.53 AB (+13%)	24.82 ± 0.82 BC (-11%)	23.37 ± 0.34 BC (-12%)	19.15 ± 3.46 C (-25%)	12.08 ± 2.42 D (-13%)	p < 0.01
		TSKC1-144	31.64 ± 1.20 aA (+31%)	26.63 ± 1.16 B (+9%)	24.37 ± 1.20 B (-11%)	24.90 ± 0.64 B (0%)	23.04 ± 2.08 B (0%)	11.06 ± 2.01 C (-27%)	p < 0.01
		KDML105	26.43 ± 0.91 cA (+15%)	25.00 ± 1.04 A (+4%)	23.22 ± 0.80 A (-6%)	24.37 ± 0.47 A (-10%)	17.12 ± 1.57 B (-34%)	9.04 ± 1.54 C (-37%)	p < 0.01
	Heavy-saline	¹ Critical-p value mean	p < 0.01	ns	ns	ns	ns	ns	
		Pokkali	28.65 ± 0.67 aA	25.91 ± 0.75 B	24.79 ± 0.50 aB	24.49 ± 0.45 aB	20.52 ± 0.96 bC	11.12 ± 0.73 cD	p < 0.01
		RD73	ns	ns	ns	p < 0.05	p < 0.01	p < 0.01	
		CSSL8-94	ns	ns	p < 0.01	p < 0.05	p < 0.01	p < 0.01	
		³ Critical-p value mean	p < 0.01	ns	ns	p < 0.01	p < 0.01	p < 0.01	
P_m	Non-saline	Pokkali	0.15 ± 0.01 aC	0.57 ± 0.05 A	0.45 ± 0.04 cB	0.39 ± 0.03 B	0.41 ± 0.04 abB	0.14 ± 0.01 C	p < 0.01
		RD73	0.11 ± 0.01 abC	0.54 ± 0.02 AB	0.56 ± 0.04 abA	0.55 ± 0.03 AB	0.48 ± 0.03 aB	0.16 ± 0.02 C	p < 0.01
		CSSL8-94	0.09 ± 0.01 bC	0.52 ± 0.03 A	0.52 ± 0.03 abcA	0.40 ± 0.08 B	0.28 ± 0.04 cB	0.16 ± 0.01 C	p < 0.01
		TSKC1-144	0.12 ± 0.02 abC	0.48 ± 0.03 A	0.48 ± 0.02 bcA	0.41 ± 0.05 A	0.29 ± 0.03 cD	0.14 ± 0.03 C	p < 0.01
		KDML105	0.11 ± 0.02 abD	0.54 ± 0.03 AB	0.58 ± 0.02 aA	0.48 ± 0.07 B	0.37 ± 0.03 bcC	0.14 ± 0.02 D	p < 0.01
	Semi-saline	¹ Critical-p value mean	p < 0.05	ns	p < 0.05	ns	p < 0.01	ns	
		Pokkali	0.12 ± 0.01 bD	0.53 ± 0.01 aA	0.52 ± 0.02 aA	0.44 ± 0.03 B	0.37 ± 0.02 aC	0.15 ± 0.01 aD	p < 0.01
		RD73	0.49 ± 0.01 A	0.16 ± 0.01 C	0.14 ± 0.02 bC	0.34 ± 0.11 AB	0.49 ± 0.03 A	0.18 ± 0.03 BC	p < 0.01
		CSSL8-94	0.50 ± 0.03 AB	0.16 ± 0.02 C	0.51 ± 0.03 aAB	0.56 ± 0.01 A	0.44 ± 0.02 B	0.19 ± 0.04 C	p < 0.01
		³ Critical-p value mean	0.41 ± 0.02 BC	0.17 ± 0.01 D	0.52 ± 0.01 aA	0.49 ± 0.01 AB	0.38 ± 0.06 C	0.17 ± 0.01 D	p < 0.01

Parameter	Saline field	Rice genotype	Early vegetative	Stem elongation	Early booting	Flowering	Milky	Mature	² Critical- p value
Heavy-saline	TSKC1-144	TSKC1-144	0.48 ± 0.05 A	0.14 ± 0.01 C	0.52 ± 0.06 aA	0.49 ± 0.03 A	0.31 ± 0.03 B	0.10 ± 0.01 C	p < 0.01
		KDML105	0.50 ± 0.04 A	0.16 ± 0.01 C	0.49 ± 0.01 aAB	0.46 ± 0.05 AB	0.38 ± 0.07 B	0.12 ± 0.01 C	p < 0.01
		¹ Critical-p value	ns	ns	p < 0.01	ns	ns	ns	
		mean	0.48 ± 0.02 aA	0.16 ± 0.01 bC	0.44 ± 0.04 bAB	0.47 ± 0.03 A	0.40 ± 0.02 aB	0.15 ± 0.01 aC	p < 0.01
		Pokkali	0.39 ± 0.09 A	0.10 ± 0.01 cB	0.11 ± 0.01 cB	0.27 ± 0.10 bAB	0.42 ± 0.04 abA	0.12 ± 0.02 B	p < 0.01
	CSSL8-94	RD73	0.53 ± 0.07 A	0.13 ± 0.01 aBC	0.47 ± 0.02 abA	0.45 ± 0.02 aA	0.23 ± 0.06 abB	0.10 ± 0.02 C	p < 0.01
		CSSL8-94	0.62 ± 0.01 A	0.12 ± 0.01 abD	0.50 ± 0.04 aB	0.49 ± 0.02 aB	0.30 ± 0.06 aC	0.11 ± 0.02 D	p < 0.01
		TSKC1-144	0.38 ± 0.04 B	0.11 ± 0.01 bcCD	0.41 ± 0.01 bAB	0.43 ± 0.02 aA	0.15 ± 0.01 bB	0.08 ± 0.02 C	p < 0.01
		KDML105	0.45 ± 0.07 A	0.12 ± 0.01 abc	0.48 ± 0.03 aA	0.49 ± 0.02 aA	0.27 ± 0.02 aB	0.10 ± 0.02 C	p < 0.01
		¹ Critical-p value	ns	p < 0.05	p < 0.01	p < 0.05	p < 0.01	ns	
		mean	0.47 ± 0.03 aA	0.12 ± 0.01 cD	0.39 ± 0.03 bB	0.43 ± 0.03 AB	0.28 ± 0.03 bC	0.10 ± 0.01 bD	p < 0.01
	³ Critical-p value	p < 0.01	p < 0.01	p < 0.05	ns	p < 0.01	p < 0.01	p < 0.01	
Non-saline	TSKC1-144	Pokkali	6.69 ± 0.50 aBC	10.41 ± 0.27 A	7.13 ± 0.37 B	5.72 ± 0.55 CD	5.06 ± 0.27 cD	3.35 ± 0.23 aE	p < 0.01
		RD73	4.70 ± 0.20 bC	10.28 ± 0.42 A	8.01 ± 0.51 B	7.65 ± 0.56 B	7.37 ± 0.35 aB	7.00 ± 0.52 bB	p < 0.01
		CSSL8-94	4.31 ± 0.37 bD	10.36 ± 0.56 A	7.51 ± 0.39 B	6.72 ± 0.70 BC	5.65 ± 0.55 bcCD	7.08 ± 0.51 bBC	p < 0.01
		KDML105	5.07 ± 0.79 abBC	9.12 ± 0.50 A	7.21 ± 0.14 B	6.68 ± 0.51 BC	5.58 ± 0.52 bcC	6.34 ± 0.94 bBC	p < 0.01
		¹ Critical-p value	p < 0.01	ns	p < 0.05	ns	p < 0.01	p < 0.01	
		mean	5.24 ± 0.27 bE	10.16 ± 0.21 aA	7.71 ± 0.19 aB	6.91 ± 0.29 aC	6.06 ± 0.25 D	6.05 ± 0.40 aD	p < 0.05
	³ Critical-p value	p < 0.01	p < 0.01	p < 0.05	ns	p < 0.01	p < 0.01	p < 0.01	
	TSKC1-144	Pokkali	8.79 ± 0.31 A	4.48 ± 0.26 C	4.02 ± 0.31 bC	5.50 ± 0.85 BC	6.40 ± 0.33 B	4.90 ± 0.88 cBC	p < 0.01
		RD73	9.19 ± 0.22 A	4.64 ± 0.52 C	6.63 ± 0.10 aB	6.95 ± 0.15 B	7.02 ± 0.24 B	7.83 ± 1.16 aC	p < 0.01
		CSSL8-94	8.09 ± 0.21 A	4.58 ± 0.19 C	6.63 ± 0.16 aB	6.56 ± 0.07 B	6.30 ± 0.67 B	7.18 ± 0.43 abAB	p < 0.01
		KDML105	9.07 ± 0.54 A	4.01 ± 0.20 D	6.66 ± 0.27 aB	6.58 ± 0.20 B	5.81 ± 0.31 BC	4.75 ± 0.52 cCD	p < 0.01
		¹ Critical-p value	ns	ns	p < 0.01	ns	ns	p < 0.05	
		mean	8.88 ± 0.16 aA	4.51 ± 0.15 bC	6.11 ± 0.25 bB	6.37 ± 0.20 aB	6.37 ± 0.21 B	6.08 ± 0.40 aB	p < 0.01
Tr	Semi-saline	Pokkali	7.23 ± 0.97 A	3.04 ± 0.11 cC	3.16 ± 0.11 bC	4.41 ± 0.85 bBC	5.58 ± 0.35 AB	3.31 ± 0.33 C	p < 0.01
		RD73	9.01 ± 0.51 A	3.58 ± 0.10 aC	5.97 ± 0.15 aB	5.88 ± 0.15 aB	5.24 ± 1.15 BC	4.86 ± 0.92 BC	p < 0.01
		CSSL8-94	9.81 ± 0.13 A	3.42 ± 0.15 abD	6.23 ± 0.22 aB	6.16 ± 0.22 aB	6.45 ± 0.41 B	4.91 ± 0.86 C	p < 0.01
		TSKC1-144	7.51 ± 0.48 A	3.13 ± 0.06 bcC	5.67 ± 0.09 aB	5.81 ± 0.07 aB	3.83 ± 0.27 C	4.02 ± 0.78 C	p < 0.01
		KDML105	8.45 ± 0.73 A	3.43 ± 0.14 abC	6.16 ± 0.34 aB	6.27 ± 0.22 aB	6.18 ± 0.56 B	4.88 ± 0.91 BC	p < 0.01
	¹ Critical-p value	ns	p < 0.05	p < 0.01	p < 0.05	ns	ns	ns	
		mean	8.40 ± 0.33 aA	3.32 ± 0.07 cD	5.44 ± 0.28 cB	5.71 ± 0.22 bB	5.45 ± 0.33 B	4.40 ± 0.35 bC	p < 0.01
		³ Critical-p value	p < 0.01	p < 0.01	p < 0.01	p < 0.01	ns	p < 0.01	
	TSKC1-144	Pokkali	3.53 ± 0.35 bAB	2.29 ± 0.04 cC	3.11 ± 0.06 B	3.46 ± 0.27 AB	3.64 ± 0.19 bAB	4.03 ± 0.35 aA	p < 0.01
		RD73	4.95 ± 0.18 abA	2.49 ± 0.08 abC	3.54 ± 0.31 B	3.53 ± 0.32 B	3.49 ± 0.16 bB	1.99 ± 0.14 bC	p < 0.01
		CSSL8-94	5.72 ± 0.46 abA	2.35 ± 0.04 bcC	3.67 ± 0.20 B	3.80 ± 0.35 B	4.14 ± 0.28 abB	2.14 ± 0.09 bC	p < 0.01
		Non-saline	3.96 ± 0.29 abAB	2.63 ± 0.05 aCD	3.40 ± 0.09 BC	4.13 ± 0.40 AB	4.73 ± 0.29 aA	2.33 ± 0.14 bD	p < 0.01
		KDML105	5.30 ± 0.93 abA	2.34 ± 0.01 bcC	3.07 ± 0.20 BC	3.46 ± 0.44 B	3.88 ± 0.23 bB	2.14 ± 0.17 bC	p < 0.01
WUE	Non-saline	¹ Critical-p value	p < 0.05	p < 0.01	ns	ns	p < 0.05	p < 0.01	
		mean	4.69 ± 0.28 aA	2.42 ± 0.03 cD	3.36 ± 0.09 bC	3.68 ± 0.16 bBC	3.98 ± 0.14 B	2.53 ± 0.19 D	p < 0.05
		³ Critical-p value	p < 0.01	p < 0.01	p < 0.01	p < 0.01	ns	p < 0.01	
		Semi-saline	Pokkali	3.14 ± 0.13 C	5.52 ± 0.40 AB	6.06 ± 0.48 aA	4.49 ± 0.81 BC	3.24 ± 0.12 cC	3.94 ± 0.55 aC

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Heavy-saline	Pokkali	RD73	3.14 ± 0.03 B	5.57 ± 0.41 A	3.20 ± 0.09 bB	3.39 ± 0.07 B	3.57 ± 0.26 bcB	2.29 ± 0.19 bC	p < 0.01
		CSSL8-94	3.20 ± 0.16 C	5.66 ± 0.27 A	3.22 ± 0.15 bBC	3.45 ± 0.10 BC	3.85 ± 0.33 bB	2.22 ± 0.07 bD	p < 0.01
		TSKC1-144	3.21 ± 0.19 C	6.54 ± 0.57 A	3.37 ± 0.06 bC	3.28 ± 0.13 C	4.44 ± 0.26 aB	2.74 ± 0.15 bC	p < 0.01
		KDML105	3.17 ± 0.03 CD	5.36 ± 0.40 A	3.12 ± 0.07 bCD	3.59 ± 0.25 BC	4.09 ± 0.10 abB	2.70 ± 0.15 bD	p < 0.01
		¹ Critical-p value	ns	ns	p < 0.01	ns	p < 0.05	p < 0.01	
		mean	3.17 ± 0.05 bCD	5.73 ± 0.19 bA	3.79 ± 0.28 bB	3.64 ± 0.18 bBC	3.84 ± 0.13 B	2.78 ± 0.18 D	p < 0.01
		Pokkali	3.73 ± 0.37 C	7.48 ± 0.37 AB	8.21 ± 0.12 aA	5.92 ± 1.23 B	3.65 ± 0.13 C	3.82 ± 0.18 aC	p < 0.01
		RD73	3.46 ± 0.18 C	8.04 ± 0.33 A	4.15 ± 0.12 bB	3.97 ± 0.06 BC	3.83 ± 0.30	2.55 ± 0.37 bD	p < 0.01
Non-saline	Pokkali	CSSL8-94	3.21 ± 0.08 C	7.81 ± 0.42 A	3.90 ± 0.09 bB	4.05 ± 0.20 B	3.55 ± 0.14 BC	2.25 ± 0.10 bD	p < 0.01
		TSKC1-144	3.54 ± 0.17 C	7.97 ± 0.34 A	4.09 ± 0.09 bBC	4.19 ± 0.05 B	4.44 ± 0.19 B	2.27 ± 0.16 bD	p < 0.01
		KDML105	3.41 ± 0.28 C	7.60 ± 0.31 A	4.14 ± 0.11 bBC	4.26 ± 0.17 B	3.77 ± 0.25 BC	2.38 ± 0.32 bD	p < 0.01
		¹ Critical-p value	ns	ns	p < 0.01	ns	ns	p < 0.05	
		mean	3.47 ± 0.10 bC	7.78 ± 0.15 aA	4.90 ± 0.38 aB	4.48 ± 0.28 aB	3.85 ± 0.11 C	2.65 ± 0.17 D	p < 0.01
		³ Critical-p value	p < 0.01	p < 0.01	p < 0.01	p < 0.05	ns	ns	
		Pokkali	0.32 ± 0.01 A	0.24 ± 0.01 B	0.24 ± 0.01 B	0.25 ± 0.01 B	0.25 ± 0.01 abB	0.15 ± 0.01 C	p < 0.01
		RD73	0.32 ± 0.01 A	0.26 ± 0.01 BC	0.31 ± 0.02 AB	0.25 ± 0.03 C	0.23 ± 0.01 bC	0.16 ± 0.01 D	p < 0.01
ΦPSII	Semi-saline	CSSL8-94	0.30 ± 0.01 A	0.27 ± 0.02 AB	0.31 ± 0.02 A	0.23 ± 0.03 B	0.17 ± 0.01 cC	0.17 ± 0.01 C	p < 0.01
		TSKC1-144	0.32 ± 0.01 A	0.27 ± 0.01 BC	0.30 ± 0.01 AB	0.29 ± 0.01 AB	0.25 ± 0.01 abC	0.16 ± 0.02 D	p < 0.01
		KDML105	0.30 ± 0.01 A	0.27 ± 0.01 A	0.27 ± 0.02 A	0.28 ± 0.02 A	0.27 ± 0.01 aA	0.19 ± 0.01 B	p < 0.01
		¹ Critical-p value	ns	ns	ns	ns	p < 0.01	ns	
		mean	0.31 ± 0.01 aA	0.26 ± 0.01 bC	0.29 ± 0.01 aB	0.26 ± 0.01 abC	0.23 ± 0.01 D	0.16 ± 0.01 aE	p < 0.01
		Pokkali	0.29 ± 0.01 A	0.25 ± 0.02 A	0.28 ± 0.03 A	0.24 ± 0.02 A	0.24 ± 0.02 A	0.16 ± 0.01 B	p < 0.01
		RD73	0.31 ± 0.01 A	0.26 ± 0.02 B	0.22 ± 0.01 B	0.25 ± 0.02 B	0.24 ± 0.02 B	0.16 ± 0.01 C	p < 0.01
		CSSL8-94	0.28 ± 0.01 A	0.27 ± 0.01 AB	0.24 ± 0.01 BC	0.24 ± 0.01 BC	0.23 ± 0.01 C	0.15 ± 0.01 D	p < 0.01
Heavy-saline	Pokkali	TSKC1-144	0.31 ± 0.01 A	0.28 ± 0.02 AB	0.24 ± 0.01 B	0.23 ± 0.01 B	0.27 ± 0.02 AB	0.12 ± 0.01 C	p < 0.01
		KDML105	0.30 ± 0.01 A	0.28 ± 0.02 AB	0.22 ± 0.01 C	0.24 ± 0.01 BC	0.27 ± 0.02 AB	0.17 ± 0.02 D	p < 0.01
		¹ Critical-p value	ns	ns	ns	ns	ns	ns	
		mean	0.30 ± 0.01 abA	0.27 ± 0.01 bB	0.24 ± 0.01 bC	0.24 ± 0.01 bC	0.25 ± 0.01 BC	0.15 ± 0.01 abD	p < 0.01
		Pokkali	0.26 ± 0.01 bCD	0.30 ± 0.01 AB	0.34 ± 0.01 aA	0.28 ± 0.03 BC	0.23 ± 0.01 D	0.14 ± 0.01 E	p < 0.01
		RD73	0.30 ± 0.01 aAB	0.34 ± 0.02 A	0.29 ± 0.01 bB	0.27 ± 0.01 B	0.22 ± 0.03 C	0.15 ± 0.02 D	p < 0.01
		CSSL8-94	0.31 ± 0.01 aAB	0.32 ± 0.01 A	0.26 ± 0.01 bBC	0.27 ± 0.01 BC	0.25 ± 0.03 C	0.15 ± 0.01 D	p < 0.01
		TSKC1-144	0.28 ± 0.01 abB	0.34 ± 0.02 A	0.27 ± 0.01 bB	0.28 ± 0.01 B	0.21 ± 0.02 C	0.11 ± 0.01 D	p < 0.01
Fv'/Fm'	Non-saline	KDML105	0.30 ± 0.01 aA	0.33 ± 0.02 A	0.29 ± 0.01 bAB	0.29 ± 0.02 AB	0.24 ± 0.01 B	0.15 ± 0.03 C	p < 0.01
		¹ Critical-p value	p < 0.05	ns	p < 0.01	ns	ns	ns	
		mean	0.29 ± 0.01 bB	0.33 ± 0.01 aA	0.29 ± 0.01 aB	0.28 ± 0.01 aB	0.23 ± 0.01 C	0.14 ± 0.01 bD	p < 0.01
		³ Critical-p value	p < 0.05	p < 0.01	p < 0.01	p < 0.01	ns	p < 0.05	
		Pokkali	0.48 ± 0.01 bB	0.53 ± 0.02 a-cA	0.47 ± 0.02 cB	0.46 ± 0.02 cB	0.50 ± 0.01 bcAB	0.34 ± 0.02 C	p < 0.01
		RD73	0.52 ± 0.02 abA	0.58 ± 0.01 aA	0.57 ± 0.01 abA	0.58 ± 0.01 aA	0.56 ± 0.02 aA	0.40 ± 0.04 B	p < 0.01
		CSSL8-94	0.53 ± 0.01 aA	0.54 ± 0.03 abA	0.57 ± 0.01 abA	0.57 ± 0.01 abA	0.55 ± 0.01 abA	0.40 ± 0.03 B	p < 0.01
		TSKC1-144	0.48 ± 0.01 bA	0.49 ± 0.01 cA	0.53 ± 0.02 bA	0.51 ± 0.03 bcA	0.49 ± 0.02 cA	0.33 ± 0.03 B	p < 0.01
Heavy-saline	Pokkali	KDML105	0.49 ± 0.02 abB	0.52 ± 0.01 bcB	0.61 ± 0.02 aA	0.56 ± 0.03 abAB	0.50 ± 0.03 bcB	0.39 ± 0.02 C	p < 0.01
		¹ Critical-p value	p < 0.05	p < 0.05	p < 0.01	p < 0.01	p < 0.05	ns	
		mean	0.50 ± 0.01 cC	0.53 ± 0.01 bAB	0.55 ± 0.01 cA	0.53 ± 0.01 bAB	0.52 ± 0.01 BC	0.37 ± 0.01 cD	p < 0.01

Parameter	Saline field	Rice genotype	Early vegetative	Stem elongation	Early booting	Flowering	Milky	Mature	² Critical- <i>p</i> value
Semi-saline	Pokkali	0.58 ± 0.01 A	0.56 ± 0.01 bcA	0.56 ± 0.02 cA	0.57 ± 0.02 bA	0.58 ± 0.03 A	0.44 ± 0.01 B	<i>p</i> < 0.01	
	RD73	0.57 ± 0.02 BC	0.60 ± 0.02 abABC	0.62 ± 0.02 aAB	0.63 ± 0.01 aA	0.56 ± 0.03 CD	0.52 ± 0.03 D	<i>p</i> < 0.05	
	CSSL8-94	0.57 ± 0.01 AB	0.61 ± 0.02 aA	0.58 ± 0.01 bcAB	0.62 ± 0.01 aA	0.54 ± 0.03 BC	0.50 ± 0.03 C	<i>p</i> < 0.01	
	TSKC1-144	0.58 ± 0.02 A	0.52 ± 0.01 cB	0.62 ± 0.01 abA	0.61 ± 0.01 aA	0.51 ± 0.01 B	0.40 ± 0.03 C	<i>p</i> < 0.01	
	KDML105	0.59 ± 0.01 A	0.60 ± 0.01 abA	0.63 ± 0.01 aA	0.62 ± 0.01 aA	0.53 ± 0.02 B	0.49 ± 0.03 B	<i>p</i> < 0.01	
	¹ Critical- <i>p</i> value	ns	<i>p</i> < 0.01	<i>p</i> < 0.01	<i>p</i> < 0.05	ns	ns		
	mean	0.58 ± 0.01 aB	0.58 ± 0.01 aB	0.60 ± 0.01 aAB	0.61 ± 0.01 aA	0.54 ± 0.01 C	0.47 ± 0.02 aD	<i>p</i> < 0.01	
	Pokkali	0.52 ± 0.01 bcBC	0.49 ± 0.01 cC	0.52 ± 0.01 cBC	0.54 ± 0.02 bAB	0.57 ± 0.01 aA	0.38 ± 0.02 bD	<i>p</i> < 0.01	
	RD73	0.55 ± 0.01 abB	0.57 ± 0.01 aB	0.61 ± 0.01 aA	0.62 ± 0.01 aA	0.50 ± 0.02 bcC	0.50 ± 0.02 aC	<i>p</i> < 0.01	
	CSSL8-94	0.57 ± 0.01 aB	0.54 ± 0.01 abBC	0.61 ± 0.01 aA	0.61 ± 0.01 aA	0.52 ± 0.02 abC	0.40 ± 0.01 bD	<i>p</i> < 0.01	
Heavy-saline	TSKC1-144	0.51 ± 0.02 cBC	0.51 ± 0.01 bcBC	0.56 ± 0.01 bAB	0.58 ± 0.01 aA	0.46 ± 0.02 cC	0.40 ± 0.04 bD	<i>p</i> < 0.01	
	KDML105	0.56 ± 0.02 aBC	0.54 ± 0.01 abBC	0.59 ± 0.01 aAB	0.62 ± 0.01 aA	0.53 ± 0.01 abC	0.45 ± 0.03 abD	<i>p</i> < 0.01	
	¹ Critical- <i>p</i> value	<i>p</i> < 0.05	<i>p</i> < 0.01	<i>p</i> < 0.01	<i>p</i> < 0.01	<i>p</i> < 0.05	<i>p</i> < 0.05		
	mean	0.54 ± 0.01 bB	0.53 ± 0.01 bBC	0.58 ± 0.01 bA	0.59 ± 0.01 aA	0.52 ± 0.01 C	0.43 ± 0.01 bD	<i>p</i> < 0.01	
	³ Critical- <i>p</i> value	<i>p</i> < 0.01	<i>p</i> < 0.01	<i>p</i> < 0.01	<i>p</i> < 0.01	ns	<i>p</i> < 0.01		
	Pokkali	172 ± 1 A	129 ± 8 B	129 ± 6 B	132 ± 7 B	135 ± 7 abB	80 ± 5 C	<i>p</i> < 0.01	
	RD73	172 ± 6 A	138 ± 5 BC	164 ± 12 AB	133 ± 15 C	123 ± 6 bC	85 ± 6 D	<i>p</i> < 0.01	
	CSSL8-94	161 ± 8 A	146 ± 9 AB	162 ± 9 A	122 ± 15 B	90 ± 5 cC	91 ± 5 C	<i>p</i> < 0.01	
	TSKC1-144	169 ± 4 B	146 ± 4 AB	158 ± 4 AB	154 ± 6 A	132 ± 8 abAB	87 ± 8 C	<i>p</i> < 0.01	
	KDML105	160 ± 7 A	146 ± 7 A	146 ± 11 A	147 ± 10 A	143 ± 6 aA	100 ± 2 B	<i>p</i> < 0.01	
Non-saline	¹ Critical- <i>p</i> value	ns	ns	ns	ns	<i>p</i> < 0.01	ns		
	mean	167.28 ± 2.57 aA	141.39 ± 3.19 bC	152.47 ± 4.64 aB	138.18 ± 5.25 abC	125.04 ± 4.93 D	89.10 ± 2.78 aE	<i>p</i> < 0.01	
	Pokkali	155 ± 7 A	132 ± 10 A	148 ± 15 A	130 ± 9 A	127 ± 12 A	87 ± 7 B	<i>p</i> < 0.01	
	RD73	167 ± 7 A	136 ± 12 B	116 ± 5 B	133 ± 9 B	126 ± 12 B	87 ± 6 C	<i>p</i> < 0.01	
	CSSL8-94	151 ± 5 A	144 ± 6 AB	128 ± 4 BC	129 ± 6 BC	121 ± 8 C	82 ± 8 D	<i>p</i> < 0.01	
	TSKC1-144	164 ± 8 A	149 ± 10 AB	130 ± 6 B	125 ± 7 B	145 ± 9 AB	66 ± 7 C	<i>p</i> < 0.01	
	KDML105	160 ± 2 A	149 ± 13 AB	117 ± 4 C	126 ± 7 BC	143 ± 9 AB	91 ± 9 D	<i>p</i> < 0.01	
	¹ Critical- <i>p</i> value	ns	ns	ns	ns	ns	ns		
	mean	159.76 ± 2.85 abA	142.52 ± 4.42 bB	128.38 ± 4.10 bC	129.08 ± 3.08 bC	132.90 ± 4.53 BC	83.04 ± 3.55 abD	<i>p</i> < 0.01	
	Pokkali	138 ± 5 bCD	161 ± 8 AB	180 ± 5 aA	150 ± 14 BC	124 ± 4 D	78 ± 2 E	<i>p</i> < 0.01	
ETR (μmol e ⁻ m ⁻² s ⁻¹)	Semi-saline	RD73	162 ± 5 aAB	182 ± 8 A	155 ± 4 bB	146 ± 4 B	117 ± 15 C	81 ± 9 D	<i>p</i> < 0.01
		CSSL8-94	165 ± 6 aAB	173 ± 7 A	140 ± 7 bBC	144 ± 7 BC	135 ± 14 C	80 ± 7 D	<i>p</i> < 0.01
		TSKC1-144	151 ± 5 abB	183 ± 8 A	146 ± 7 bB	148 ± 5 B	111 ± 9 C	58 ± 5 D	<i>p</i> < 0.01
		KDML105	157 ± 4 aA	173 ± 9 A	157 ± 6 bAB	153 ± 10 AB	130 ± 6 B	80 ± 13 C	<i>p</i> < 0.01
		¹ Critical- <i>p</i> value	<i>p</i> < 0.05	ns	<i>p</i> < 0.01	ns	ns		
		mean	155.17 ± 2.94 bB	174.87 ± 3.68 aA	156.09 ± 3.89 aB	148.59 ± 3.49 aB	123.62 ± 4.54 C	75.85 ± 3.82 bD	<i>p</i> < 0.01
		³ Critical- <i>p</i> value	<i>p</i> < 0.05	<i>p</i> < 0.01	<i>p</i> < 0.01	<i>p</i> < 0.01	ns	<i>p</i> < 0.05	

¹ Critical-*p* value for testing each trait among genotypes within growth stage (the same column). ² Critical-*p* value for testing each trait among growth stages of each genotype (the same row) and

³ critical-*p* value for testing each trait among salinity fields within growth stage. [Values in the brackets indicated percentage increase (+) or decrease (-) in the saline fields compared with that in the non-saline field]

Supplementary Table S3. Plant height and biomass of leaf, stem, root and total biomass at maturity of five rice genotypes including Pokkali, RD73, CSSL8-94, TSKC1-144 and KDM105 after grew under non-saline (control), semi-saline and heavy-saline field measured at four growth stage including early vegetative, stem elongation, flowering and mature stage. Means which are significantly different ($p < 0.05$) among genotypes for each saline field and field across genotypes are denoted by different lower-case letters. For each genotype means which are significantly different ($p < 0.05$) among growth stage are denoted with capital letters. Data shows mean of four replicates \pm standard error (SE).

Parameter	Saline field	Rice genotype	Early vegetative	Stem elongation	Flowering	Mature	² Critical- <i>p</i> value
Plant height (cm)	Non-saline	Pokkali	40 \pm 1 aD	137 \pm 3 aC	155 \pm 2 bB	178 \pm 2 aA	$p > 0.01$
		RD73	34 \pm 1 bD	106 \pm 3 bcC	151 \pm 3 bcB	163 \pm 1 bA	$p > 0.01$
		CSSL8-94	40 \pm 1 aC	102 \pm 4 cB	152 \pm 2 bcA	156 \pm 1 bA	$p > 0.01$
		TSKC1-144	39 \pm 1 aC	117 \pm 3 bB	170 \pm 1 aA	176 \pm 2 aA	$p > 0.01$
		KDM105	39 \pm 1 aC	102 \pm 7 cB	147 \pm 1 cA	156 \pm 4 bA	$p > 0.01$
		¹ Critical- <i>p</i> value	$p > 0.05$	$p > 0.01$	$p > 0.01$	$p > 0.01$	
		mean	38.925 D	113.15 aC	155.35 aB	166.2 aA	$p > 0.01$
	Semi-saline	Pokkali	35 \pm 1 cC	123 \pm 6 aB	159 \pm 7 aA	152 \pm 1 abA	$p > 0.01$
		RD73	35 \pm 1 cC	104 \pm 1 bB	138 \pm 3 bA	136 \pm 8 cA	$p > 0.01$
		CSSL8-94	42 \pm 1 aC	107 \pm 2 bB	152 \pm 3 abA	147 \pm 5 bcA	$p > 0.01$
		TSKC1-144	39 \pm 1 abcC	102 \pm 3 bB	167 \pm 3 aA	167 \pm 3 aA	$p > 0.01$
		KDM105	37 \pm 1 bccC	104 \pm 4 bB	140 \pm 1 bA	132 \pm 3 cA	$p > 0.01$
		¹ Critical- <i>p</i> value	$p > 0.05$	$p > 0.05$	$p > 0.01$	$p > 0.01$	
		mean	38.05 C	108.55 aB	151.5 aA	146.95 bA	$p > 0.01$
	Heavy-saline	Pokkali	33 \pm 2 C	116 \pm 3 aB	150 \pm 3 aA	163 \pm 6 aA	$p > 0.01$
		RD73	35 \pm 1 C	80 \pm 11 bcB	121 \pm 10 bA	120 \pm 17 cA	$p > 0.01$
		CSSL8-94	57 \pm 15 C	77 \pm 3 cB	125 \pm 2 bA	121 \pm 3 cA	$p > 0.01$
		TSKC1-144	38 \pm 1 C	101 \pm 11 abB	148 \pm 10 aA	148 \pm 9 abA	$p > 0.05$
		KDM105	38 \pm 1 C	77 \pm 6 cB	131 \pm 3 bA	127 \pm 1 bcA	$p > 0.01$
		¹ Critical- <i>p</i> value	ns	$p > 0.01$	$p > 0.01$	$p > 0.01$	
		mean	40.69 C	90.50 bB	135.15 bA	136.15 bA	$p > 0.01$
		³ Critical- <i>p</i> value	ns	$p > 0.01$	$p > 0.01$	$p > 0.01$	
Leaf biomass (g _{DW} plant ⁻¹)	Non-saline	Pokkali	0.09 \pm 0.01 ab	14.42 \pm 1.93 bA	16.06 \pm 1.58 bA	13.26 \pm 0.69 cA	$p > 0.01$
		RD73	0.05 \pm 0.01 cC	24.93 \pm 2.51 aA	20.28 \pm 1.19 bB	18.19 \pm 0.70 bB	$p > 0.01$
		CSSL8-94	0.09 \pm 0.01 abC	25.54 \pm 1.63 aA	20.03 \pm 2.33 bB	21.40 \pm 1.35 abAB	$p > 0.01$
		TSKC1-144	0.07 \pm 0.01 bcB	24.35 \pm 2.04 aA	30.34 \pm 5.19 aA	23.44 \pm 1.01 aA	$p > 0.01$
		KDM105	0.06 \pm 0.01 cC	24.40 \pm 0.98 aB	30.37 \pm 2.46 aA	21.79 \pm 1.75 aB	$p > 0.01$
		¹ Critical- <i>p</i> value	$p > 0.01$	$p > 0.05$	$p > 0.01$	$p > 0.01$	
		mean	0.07 C	22.73 aAB	23.41 aA	19.62 aB	$p > 0.01$
	Semi-saline	Pokkali	0.06 \pm 0.01 B	15.87 \pm 2.10 A	21.89 \pm 4.00 A	6.26 \pm 0.76 cB	$p > 0.01$
		RD73	0.06 \pm 0.01 D	9.22 \pm 0.94 C	18.47 \pm 1.13 A	12.16 \pm 0.79 bcB	$p > 0.01$
		CSSL8-94	0.08 \pm 0.01 C	15.13 \pm 2.29 B	21.06 \pm 1.49 A	23.89 \pm 2.44 aA	$p > 0.01$
		TSKC1-144	0.08 \pm 0.01 B	15.67 \pm 2.76 AB	28.42 \pm 9.81 A	14.76 \pm 1.85 bAB	$p > 0.05$
		KDM105	0.07 \pm 0.01 C	11.62 \pm 1.29 B	25.73 \pm 3.31 A	26.07 \pm 2.91 aA	$p > 0.01$

Parameter	Saline field	Rice genotype	Early vegetative	Stem elongation	Flowering	Mature	² Critical- <i>p</i> value
Stem biomass (gDW plant ⁻¹)	Heavy-saline	¹ Critical- <i>p</i> value mean	ns 0.07 C	ns 13.50 bB	ns 23.11 aA	<i>p</i> > 0.01 16.63 aB	<i>p</i> > 0.01
		Pokkali	0.03 ± 0.01 B	16.44 ± 2.16 A	17.20 ± 1.87 A	13.43 ± 1.99 aA	<i>p</i> > 0.01
		RD73	0.06 ± 0.01 C	8.31 ± 2.41 B	14.60 ± 2.44 A	3.81 ± 0.54 b BC	<i>p</i> > 0.01
		CSSL8-94	0.08 ± 0.01 C	10.34 ± 1.58 B	18.54 ± 3.32 A	12.55 ± 1.33 aAB	<i>p</i> > 0.01
		TSKC1-144	0.07 ± 0.02 B	12.08 ± 1.68 A	19.71 ± 5.90 A	15.78 ± 3.58 aA	<i>p</i> > 0.01
		KDML105	0.07 ± 0.01 C	9.05 ± 0.80 B	16.28 ± 1.97 A	12.31 ± 0.36 aB	<i>p</i> > 0.01
	Non-saline	¹ Critical- <i>p</i> value mean	ns 0.06 C	ns 11.24 bB	ns 17.26 bA	<i>p</i> > 0.01 11.58 bB	<i>p</i> > 0.01
		Pokkali	0.03 ± 0.01 C	12.34 ± 1.33 B	23.07 ± 3.53 cA	18.99 ± 1.20 cA	<i>p</i> > 0.01
		RD73	0.02 ± 0.01 C	18.53 ± 3.20 B	35.59 ± 3.04 bA	20.00 ± 1.12 cB	<i>p</i> > 0.01
		CSSL8-94	0.04 ± 0.01 C	19.09 ± 2.46 B	30.30 ± 3.16 bcA	28.33 ± 1.94 bA	<i>p</i> > 0.01
		TSKC1-144	0.03 ± 0.01 D	17.52 ± 3.29 C	54.01 ± 4.17 aA	33.57 ± 3.35 aB	<i>p</i> > 0.01
Root biomass (gDW plant ⁻¹)	Semi-saline	KDML105	0.03 ± 0.01 D	19.05 ± 1.60 C	52.91 ± 2.63 aA	26.12 ± 2.15 bB	<i>p</i> > 0.01
		¹ Critical- <i>p</i> value mean	ns 0.034 D	ns 17.30 aC	<i>p</i> > 0.01 39.17 aA	<i>p</i> > 0.01 25.40 bB	<i>p</i> > 0.01
		Pokkali	0.01 ± 0.01 cB	18.45 ± 5.53 A	21.30 ± 8.25 bA	29.11 ± 0.93 bA	<i>p</i> > 0.01
		RD73	0.04 ± 0.01 bC	4.05 ± 0.46 C	22.56 ± 2.48 bB	30.38 ± 2.39 bA	<i>p</i> > 0.01
		CSSL8-94	0.05 ± 0.01 aC	13.95 ± 3.68 B	26.16 ± 1.80 bA	24.16 ± 1.52 bA	<i>p</i> > 0.01
		TSKC1-144	0.04 ± 0.01 bB	11.22 ± 1.70 B	46.62 ± 6.99 aA	48.20 ± 3.52 aA	<i>p</i> > 0.01
		KDML105	0.03 ± 0.01 bC	8.89 ± 1.36 B	22.36 ± 3.49 bA	26.07 ± 3.23 bA	<i>p</i> > 0.01
	Heavy-saline	¹ Critical- <i>p</i> value mean	<i>p</i> > 0.01 0.03 C	ns 11.31 bB	<i>p</i> > 0.05 27.80 bA	<i>p</i> > 0.01 31.59 aA	<i>p</i> > 0.01
		Pokkali	0.02 ± 0.01 bB	21.17 ± 5.01 aA	17.99 ± 3.36 A	16.94 ± 3.08 A	<i>p</i> > 0.01
		RD73	0.03 ± 0.01 aC	5.85 ± 2.67 bBC	15.97 ± 5.04 AB	18.61 ± 5.49 A	<i>p</i> > 0.05
		CSSL8-94	0.03 ± 0.01 aD	5.80 ± 1.51 bC	21.44 ± 2.78 A	11.15 ± 0.57 B	<i>p</i> > 0.01
		TSKC1-144	0.03 ± 0.01 abC	10.70 ± 1.80 bBC	29.58 ± 6.76 A	16.42 ± 4.50 B	<i>p</i> > 0.01
Shoot biomass (gDW plant ⁻¹)	Non-saline	KDML105	0.02 ± 0.01 abD	4.36 ± 1.16 bC	15.90 ± 2.10 BA	11.10 ± 1.15 B	<i>p</i> > 0.01
		¹ Critical- <i>p</i> value mean	<i>p</i> > 0.05 0.03 D	<i>p</i> > 0.05 9.58 bC	ns 20.18 bA	ns 14.84 cB	<i>p</i> > 0.01
		³ Critical- <i>p</i> value	ns 0.03 C	<i>p</i> > 0.01 19.44 aA	<i>p</i> > 0.01 15.53 aA	<i>p</i> > 0.01 8.83 bB	<i>p</i> > 0.01
		Pokkali	0.03 ± 0.01 abC	26.44 ± 3.08 aA	10.71 ± 1.55 B	8.44 ± 1.54 B	<i>p</i> > 0.01
		RD73	0.02 ± 0.01 bC	9.24 ± 1.81 bB	15.46 ± 1.11 A	13.47 ± 0.22 A	<i>p</i> > 0.01
	Semi-saline	CSSL8-94	0.05 ± 0.01 aC	14.67 ± 3.61 bAB	20.13 ± 1.75 A	8.24 ± 2.86 B	<i>p</i> > 0.01
		TSKC1-144	0.03 ± 0.01 bB	15.41 ± 3.61 bA	17.29 ± 5.51 A	4.54 ± 0.83 B	<i>p</i> > 0.01
		KDML105	0.04 ± 0.01 abC	31.46 ± 3.47 aA	14.07 ± 4.95 B	9.45 ± 2.96 BC	<i>p</i> > 0.01
		¹ Critical- <i>p</i> value mean	<i>p</i> > 0.05 0.03 C	<i>p</i> > 0.01 19.44 aA	ns 15.53 aA	ns 8.83 bB	<i>p</i> > 0.01
		Pokkali	0.03 ± 0.01 C	8.82 ± 1.06 B	16.02 ± 2.87 A	15.85 ± 2.13 A	<i>p</i> > 0.01
Root-shoot ratio	Semi-saline	RD73	0.06 ± 0.02 B	10.10 ± 5.12 A	10.13 ± 2.30 A	15.25 ± 2.71 A	<i>p</i> > 0.05
		CSSL8-94	0.03 ± 0.01 C	6.47 ± 0.51 B	9.36 ± 0.89 B	20.55 ± 2.56 A	<i>p</i> > 0.01
		TSKC1-144	0.06 ± 0.01 B	7.01 ± 1.71 B	24.79 ± 9.32 A	15.12 ± 3.02 AB	<i>p</i> > 0.05
		KDML105	0.03 ± 0.01 C	5.75 ± 1.43 B	20.22 ± 3.01 A	18.33 ± 1.56 A	<i>p</i> > 0.01

Parameter	Saline field	Rice genotype	Early vegetative	Stem elongation	Flowering	Mature	² Critical- <i>p</i> value
							¹ Critical- <i>p</i> value mean
Total biomass (g _{dw} plant ⁻¹)	Heavy-saline	Pokkali	0.01 ± 0.01 bB	9.56 ± 1.81 A	15.92 ± 4.22 A	10.86 ± 1.97 A	<i>p</i> > 0.01
		RD73	0.04 ± 0.01 aB	3.34 ± 1.53 AB	6.59 ± 2.89 AB	9.91 ± 3.70 A	<i>p</i> > 0.05
		CSSL8-94	0.03 ± 0.01 abC	4.18 ± 0.76 BC	7.87 ± 2.30 AB	11.90 ± 2.19 A	<i>p</i> > 0.01
		TSKC1-144	0.04 ± 0.01 aB	6.39 ± 1.49 AB	8.85 ± 3.32 A	14.18 ± 4.19 A	<i>p</i> > 0.05
		KDML105	0.03 ± 0.01 aC	4.89 ± 2.79 B	3.73 ± 0.59 BC	10.29 ± 0.98 A	<i>p</i> > 0.01
		¹ Critical- <i>p</i> value mean	<i>p</i> > 0.05	ns	ns	ns	
		³ Critical- <i>p</i> value	0.03 C	5.67 bB	8.59 bAB	11.43 bA	<i>p</i> > 0.01
		Pokkali	0.17 ± 0.02 abC	53.20 ± 6.32 B	49.85 ± 5.01 cB	92.78 ± 7.82 bA	<i>p</i> > 0.01
		RD73	0.09 ± 0.01 cD	52.71 ± 4.72 C	71.33 ± 4.81 bB	139.67 ± 6.75 aA	<i>p</i> > 0.01
		CSSL8-94	0.19 ± 0.02 aC	59.30 ± 2.00 B	70.46 ± 4.80 bB	136.26 ± 6.34 aA	<i>p</i> > 0.01
	Non-saline	TSKC1-144	0.13 ± 0.02 bcD	57.29 ± 6.06 C	101.64 ± 9.26 aB	134.81 9.18 aA	<i>p</i> > 0.01
		KDML105	0.13 ± 0.02 bcD	74.91 ± 5.10 C	97.36 ± 4.82 aB	150.76 ± 9.06 aA	<i>p</i> > 0.01
		¹ Critical- <i>p</i> value mean	<i>p</i> > 0.05	ns	<i>p</i> > 0.01	<i>p</i> > 0.01	
		³ Critical- <i>p</i> value	0.14 D	59.48 aC	78.13 aB	130.86 bA	<i>p</i> > 0.01
		Pokkali	0.12 ± 0.01 C (-29%)	43.14 ± 8.48 B (-19%)	59.22 ± 9.67 B (+19%)	139.98 ± 6.07 A (+51%)	<i>p</i> > 0.01
Root length (mm)	Semi-saline	RD73	0.16 ± 0.03 D (+78%)	23.37 ± 5.89 C (-56%)	51.16 ± 4.87 B (-28%)	173.94 ± 10.20 A (+24%)	<i>p</i> > 0.01
		CSSL8-94	0.17 ± 0.01 D (-11%)	35.56 ± 6.07 C (-40%)	56.60 ± 0.31 B (-20%)	157.53 ± 7.25 A (+15%)	<i>p</i> > 0.01
		TSKC1-144	0.18 ± 0.02 C (+38%)	33.91 ± 5.75 C (-41%)	99.83 ± 24.98 B (-2%)	159.53 ± 10.60 A (+18%)	<i>p</i> > 0.01
		KDML105	0.13 ± 0.01 C (0%)	26.27 ± 3.78 C (-65%)	68.32 ± 8.80 B (-30%)	167.89 ± 17.08 A (+11%)	<i>p</i> > 0.01
		¹ Critical- <i>p</i> value mean	ns	ns	ns	ns	
		³ Critical- <i>p</i> value	0.15 D	32.45 bC	67.02 aB	159.78 aA	<i>p</i> > 0.01
		Pokkali	0.07 ± 0.01 bC (-59%)	47.17 ± 8.51 aB (-11%)	51.12 ± 9.27 B (+3%)	117.08 ± 15.94 A (+26%)	<i>p</i> > 0.01
		RD73	0.14 ± 0.01 aB (+56%)	17.51 ± 6.54 bB (-67%)	37.17 ± 9.56 AB (-48%)	71.19 ± 26.41 A (-48%)	<i>p</i> > 0.05
		CSSL8-94	0.14 ± 0.01 aD (-26%)	20.33 ± 3.63 bC (-66%)	47.86 ± 7.83 B (-32%)	84.29 ± 6.72 A (-38%)	<i>p</i> > 0.01
		TSKC1-144	0.15 ± 0.03 aC (+15%)	29.18 ± 3.7 bBC (-49%)	58.16 ± 14.29 AB (-43%)	85.03 ± 21.95 A (-36%)	<i>p</i> > 0.01
Root length (mm)	Heavy-saline	KDML105	0.13 ± 0.01 aC (0%)	18.31 ± 4.3 bBC (-76%)	35.92 ± 3.92 B (-63%)	82.10 ± 12.35 A (-45%)	<i>p</i> > 0.01
		¹ Critical- <i>p</i> value mean	<i>p</i> > 0.05	<i>p</i> > 0.05	ns	ns	
		³ Critical- <i>p</i> value	0.12 D	26.50 bC	46.04 bB	87.94 cA	<i>p</i> > 0.01
		Pokkali	0.07 ± 0.01 bC (-59%)	47.17 ± 8.51 aB (-11%)	51.12 ± 9.27 B (+3%)	117.08 ± 15.94 A (+26%)	<i>p</i> > 0.01
		RD73	0.14 ± 0.01 aB (+56%)	17.51 ± 6.54 bB (-67%)	37.17 ± 9.56 AB (-48%)	71.19 ± 26.41 A (-48%)	<i>p</i> > 0.05
		CSSL8-94	0.14 ± 0.01 aD (-26%)	20.33 ± 3.63 bC (-66%)	47.86 ± 7.83 B (-32%)	84.29 ± 6.72 A (-38%)	<i>p</i> > 0.01
		TSKC1-144	0.15 ± 0.03 aC (+15%)	29.18 ± 3.7 bBC (-49%)	58.16 ± 14.29 AB (-43%)	85.03 ± 21.95 A (-36%)	<i>p</i> > 0.01
		KDML105	0.13 ± 0.01 aC (0%)	18.31 ± 4.3 bBC (-76%)	35.92 ± 3.92 B (-63%)	82.10 ± 12.35 A (-45%)	<i>p</i> > 0.01
		¹ Critical- <i>p</i> value mean	<i>p</i> > 0.05	<i>p</i> > 0.05	ns	ns	
		³ Critical- <i>p</i> value	0.12 D	26.50 bC	46.04 bB	87.94 cA	<i>p</i> > 0.01

¹ Critical-*p* value for testing each trait among genotypes within growth stage (the same column). ² Critical-*p* value for testing each trait among growth stages of each genotype (the same row) and

³ critical-*p* value for testing each trait among salinity fields within growth stage. [Values in the brackets indicated percentage increase (+) or decrease (-) in the saline fields compared with that in the non-saline field]

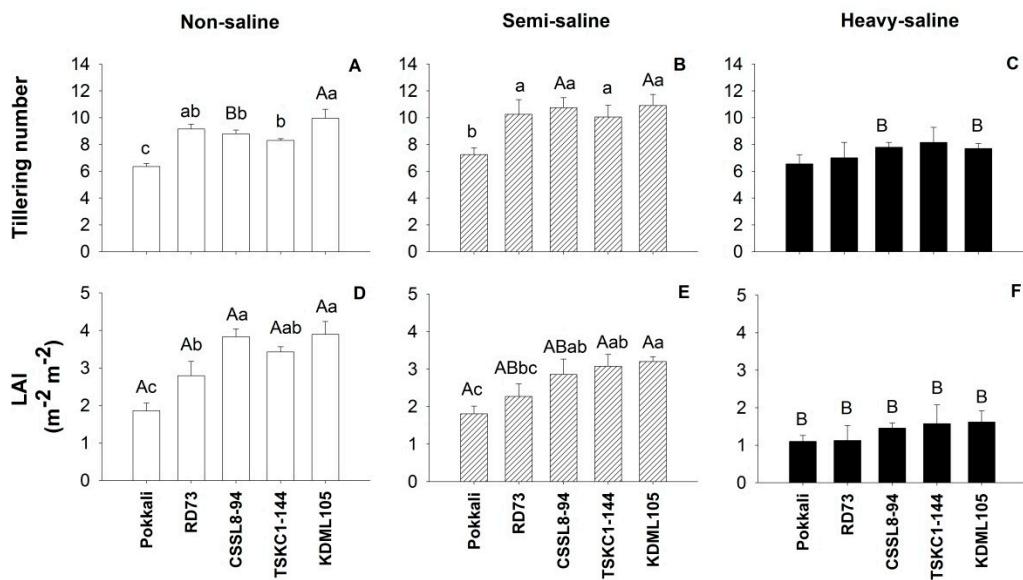


Figure S1. Tiller number and leaf area index (LAI) of five rice genotypes including Pokkali, RD73, CSSL8-94, TSKC1-144 and KDM105 after grew under non-saline (A,D), semi-saline (B,E) and heavy-saline fields (C,F) measured at mature stage. Means which are significantly different ($p < 0.05$) among genotypes for each saline field are denoted by different lower-case letters. For each genotype means which are significantly different ($p < 0.05$) among saline fields are denoted with capital letters. Data shows mean of four replicates \pm standard error (SE) (n=4).

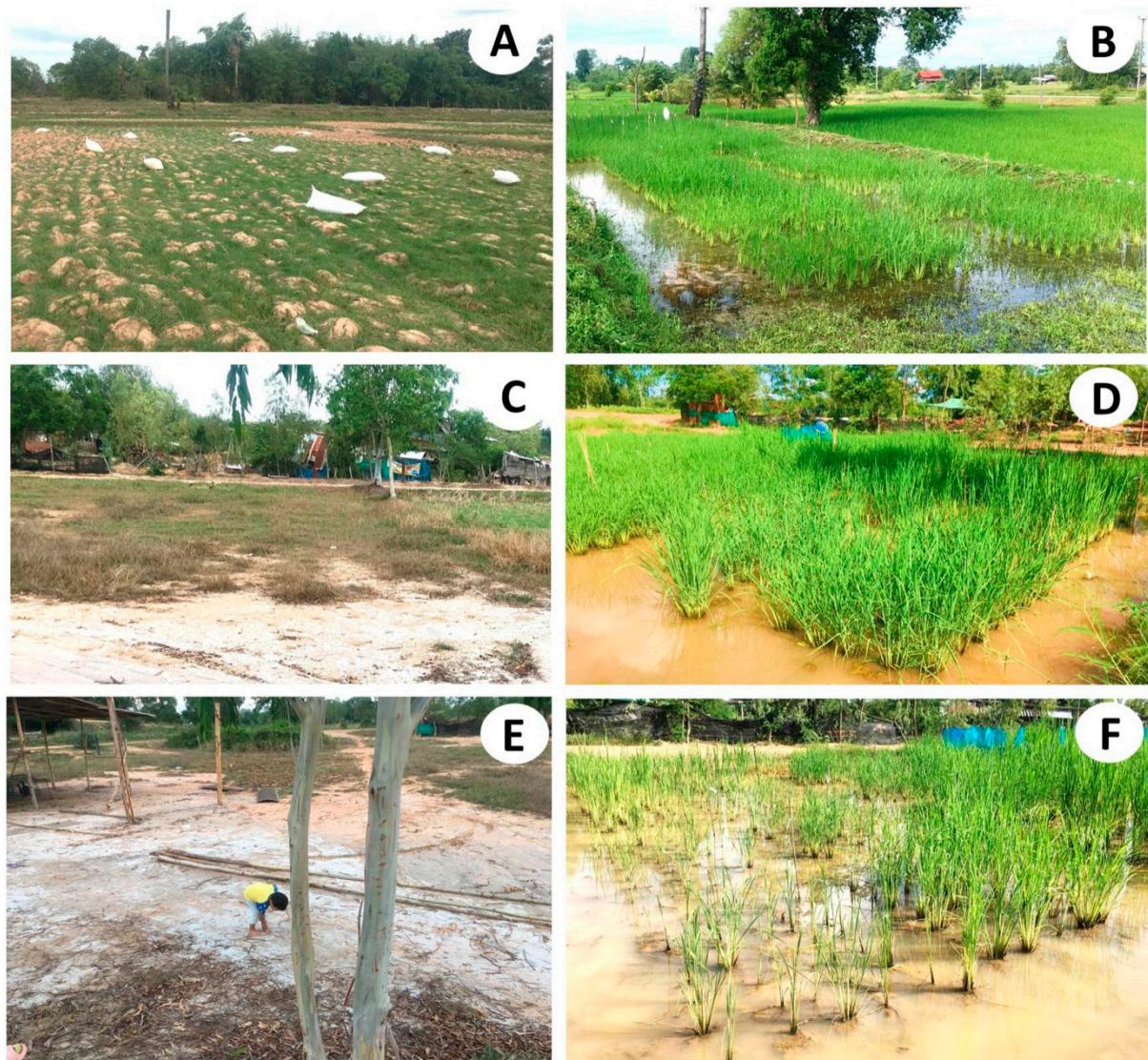


Figure S2. Experimental fields before land preparation; non-saline (**A**), semi-saline (**C**) and heavy-saline (**E**), and rice growing in the non-saline (**B**), semi-saline (**D**) and heavy-saline field (**F**). For the semi-saline and heavy-saline plots, famers avoided growing rice in these areas for many years.