

Supplementary Material

Mild preflowering drought priming improves stress defences, assimilation and sink strength in rice under severe terminal drought

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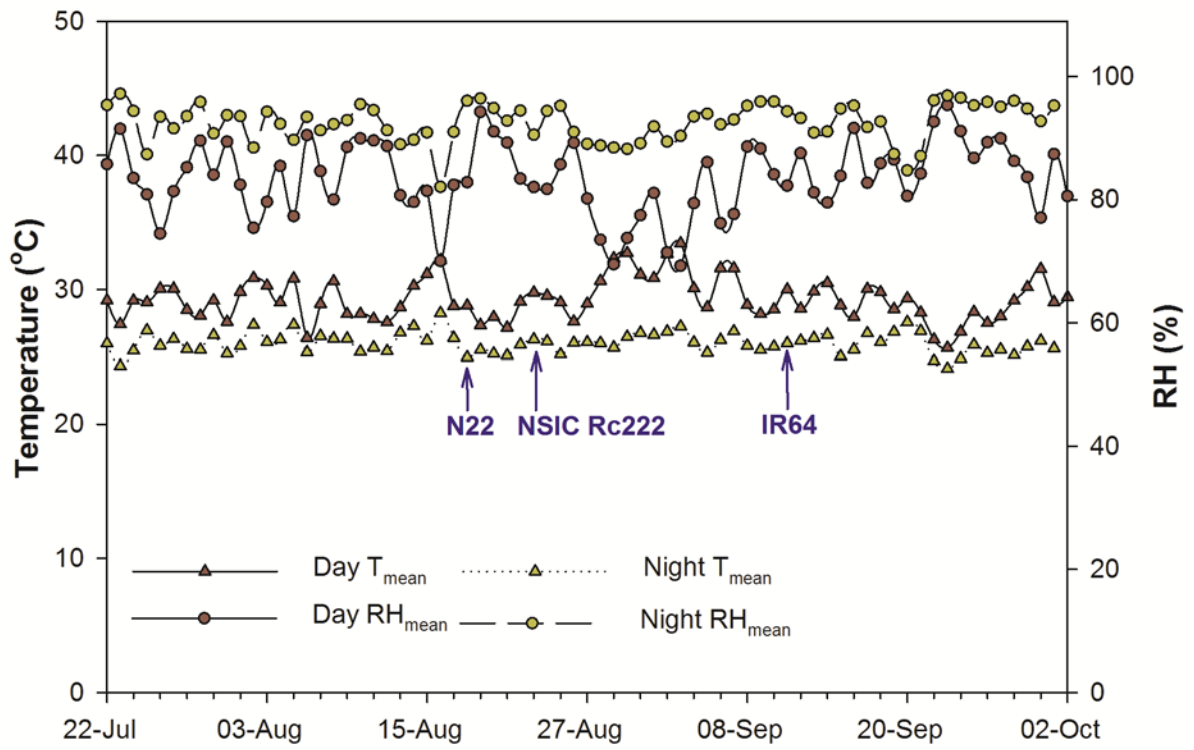


Fig S1

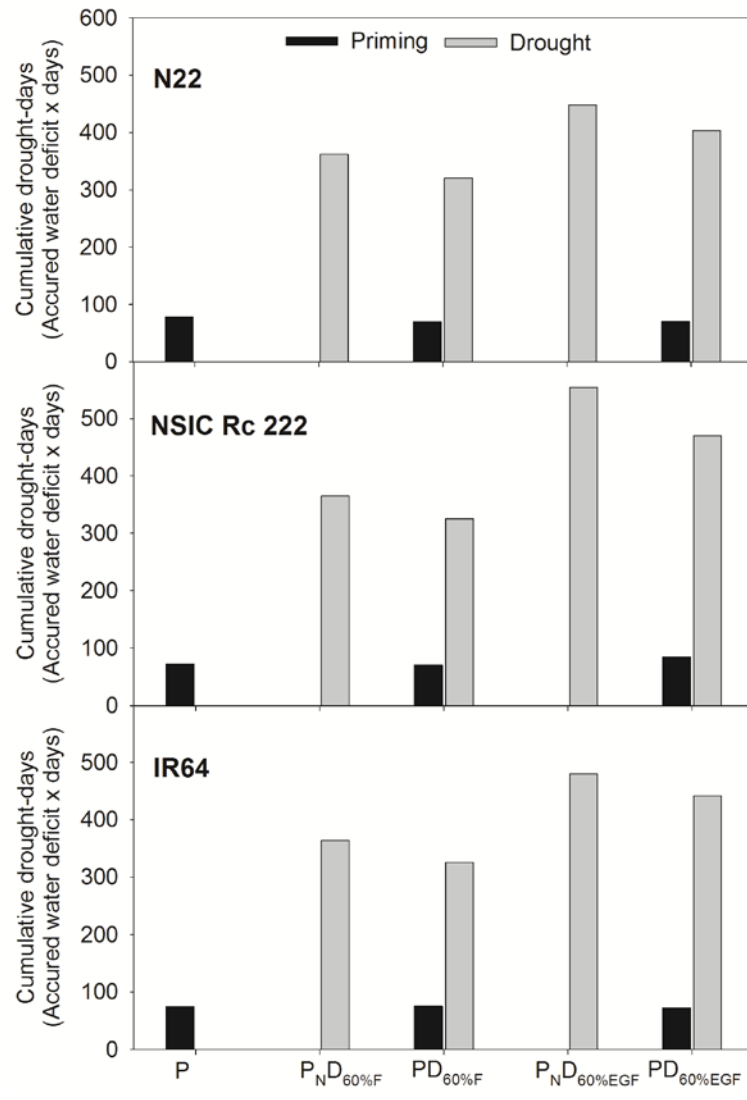


Fig S2

Table S1. ANOVA table for seed set and grain yield of rice cultivars N22, NSICRC222 and IR64

Values represent least significant difference for each trait. Significance level: * $P < 0.05$, *** $P < 0.001$

Interaction	Seed set (%)	Grain yield (g plant⁻¹)
Cultivar (C)	3.95***	2.65***
Treatment (T)	5.59***	3.75***
C x T	9.78*	6.50***

Table S2. ANOVA table for TBARs content analyzed in the flag leaf of rice cultivars N22, NSIC Rc222 and IR64 at flowering and early grain filling stage

Values represent least significant difference for each trait. Significance level: * $P < 0.05$, *** $P < 0.001$

Interaction	TBARs ($\mu\text{g g}^{-1}$ FW)	
	Flowering	Early grain filling
Cultivar (C)	1.73***	1.88***
Treatment (T)	1.73***	1.88***
C x T	2.99*	3.25***

Table S3. ANOVA table for gas exchange traits observed in flag leaf of rice cultivars N22, NSIC Rc222 and IR64 at flowering and early grain filling stage

Values represent least significant difference for each trait. A, CO₂ assimilation rate; g_s, stomatal conductance; E, transpiration rate. Significance level: **P*<0.05; ****P*<0.001; ns, non significant

Interaction	A ($\mu\text{mol m}^{-2} \text{s}^{-1}$)	g _s ($\text{mol m}^{-2} \text{s}^{-1}$)	C _i ($\mu\text{mol mol}^{-1}$)	E ($\text{mmol m}^{-2} \text{s}^{-1}$)
Cultivar (C)	0.90*	ns	9.53***	ns
Treatment (T)	1.27***	0.10***	13.48***	0.80***
C x T	ns	ns	23.35*	ns

Table S4. ANOVA table for enzymes activity in spikelets of rice cultivars N22, NSIC Rc222 and IR64 at early grain filling stage

Values represent least significant difference for each trait. CI, Cytosolic invertase; VI, Vacuolar invertase; CWI, Cell wall invertase; SuSy, Sucrose synthase; SSS, Soluble starch synthase. Significance level: **P*<0.05; ***P*<0.01; ****P*<0.001; ns, non significant

Interaction	CI ($\text{nmol min}^{-1} \text{mg}^{-1} \text{protein}$)	VI ($\text{nmol min}^{-1} \text{mg}^{-1} \text{protein}$)	CWI ($\text{nmol min}^{-1} \text{mg}^{-1} \text{protein}$)	SuSy ($\mu\text{mol h}^{-1} \text{mg}^{-1} \text{protein}$)	SSS ($\text{nmol min}^{-1} \text{mg}^{-1} \text{protein}$)
Cultivar (C)	5.19***	4.38***	2.53***	0.22***	ns
Treatment (T)	5.19*	4.38*	2.53***	0.22***	2.04***
C x T	ns	7.59**	4.38***	0.39***	ns