

## Supplementary material

**Table S1.** Irrigation water salinity and genotype effects on microbial biomass (MBC and MBN).

Factors	MBC <sup>a)</sup> (mg C kg <sup>-1</sup> )				
	6 leaves (Z16)	Tillering (Z30)	Flowering (Z65)	Maturity (Z85)	Harvesting (Z92)
Irrigation water salinity (S)					
0.3 dS m <sup>-1</sup>	292.60 <sup>b) a c)</sup>	341.72 <sup>a</sup>	365.19 <sup>a</sup>	228.00 <sup>a</sup>	122.40 <sup>a</sup>
12 dS m <sup>-1</sup>	250.36 <sup>b</sup> (-14.43%) <sup>d)</sup>	272.29 <sup>b</sup> (-20.31%)	226.28 <sup>b</sup> (-38.03%)	150.45 <sup>b</sup> (-34.01%)	102.40 <sup>b</sup> (-16.34%)
Genotypes (G)					
<i>Agili Glabre</i>	345.85 <sup>a</sup>	391.64 <sup>a</sup>	355.50 <sup>a</sup>	195.65 <sup>ab</sup>	102.05 <sup>bc</sup>
<i>Bayadha</i>	239.34 <sup>b</sup>	272.00 <sup>b</sup>	219.36 <sup>b</sup>	159.76 <sup>b</sup>	136.00 <sup>a</sup>
<i>Maali</i>	268.59 <sup>b</sup>	375.02 <sup>a</sup>	361.23 <sup>a</sup>	212.68 <sup>a</sup>	116.57 <sup>ab</sup>
<i>Razzek</i>	232.14 <sup>b</sup>	189.37 <sup>c</sup>	246.85 <sup>b</sup>	188.80 <sup>ab</sup>	95.97 <sup>c</sup>
ANOVA					
Salinity (S)	***	***	***	***	***
Genotypes (G)	***	***	***	***	***
S × G	***	**	***	***	***
MBN (mg N kg <sup>-1</sup> )					
Irrigation water salinity (S)					
0.3 dS m <sup>-1</sup>	36.23 <sup>a</sup>	39.88 <sup>a</sup>	46.52 <sup>a</sup>	31.51 <sup>a</sup>	17.11 <sup>a</sup>
12 dS m <sup>-1</sup>	29.98 <sup>b</sup> (-17.25%)	40.06 <sup>a</sup> (+0.45%)	29.55 <sup>b</sup> (-36.47%)	22.49 <sup>b</sup> (-28.62%)	14.36 <sup>b</sup> (-16.07%)
Genotypes (G)					
<i>Agili Glabre</i>	35.04 <sup>a</sup>	42.56 <sup>a</sup>	40.38 <sup>a</sup>	32.48 <sup>a</sup>	18.99 <sup>a</sup>
<i>Bayadha</i>	32.56 <sup>b</sup>	38.54 <sup>b</sup>	38.64 <sup>ab</sup>	24.57 <sup>b</sup>	14.13 <sup>b</sup>
<i>Maali</i>	32.85 <sup>b</sup>	38.87 <sup>b</sup>	35.77 <sup>b</sup>	24.82 <sup>b</sup>	15.12 <sup>b</sup>
<i>Razzek</i>	31.98 <sup>ab</sup>	39.92 <sup>ab</sup>	36.96 <sup>b</sup>	26.13 <sup>b</sup>	14.53 <sup>b</sup>
ANOVA					
Salinity (S)	***	NS <sup>e)</sup>	***	***	***
Genotypes (G)	***	***	***	***	***
S × G	***	***	***	***	**

\*\*, \*\*\*Significant at P ≤ 0.01 and P ≤ 0.001, respectively.

a) See Fig. 1 for abbreviations.

b) For each growth stage, irrigation water salinity values are the means of 20 measurements (4 genotypes and 5 replicates), while the genotype values are the means calculated from 10 measurements (2 irrigation water salinity and 5 replicates).

c) Means followed by different letters are significantly different at P ≤ 0.05 (Tukey test).

d) Values in parentheses indicate the rates of increase or decrease of each parameter according to the control treatment (0.3 dS m<sup>-1</sup>).

e) Not significant.

**Table S2. Irrigation water salinity and genotype effects on enzyme activities (D-ase, Alk-ase, and P-ase).**

Factors	D-ase <sup>a)</sup> ( $\mu\text{g INTF g}^{-1} \text{h}^{-1}$ )				
	6 leaves (Z16)	Tillering (Z30)	Flowering (Z65)	Maturity (Z85)	Harvesting (Z92)
Irrigation water salinity (S)					
0.3 dS m <sup>-1</sup>	11.90 <sup>b) a c)</sup>	14.98 <sup>a</sup>	11.21 <sup>a</sup>	10.36 <sup>a</sup>	7.56 <sup>a</sup>
12 dS m <sup>-1</sup>	10.10 <sup>b</sup>	11.90 <sup>b</sup>	8.47 <sup>b</sup>	6.74 <sup>b</sup>	5.12 <sup>b</sup>
	(-15.13%) <sup>d)</sup>	(-20.56%)	(-24.44%)	(-34.94%)	(-32.27%)
Genotypes (G)					
<i>Agili Glabre</i>	12.00 <sup>a</sup>	15.12 <sup>a</sup>	11.97 <sup>a</sup>	10.67 <sup>a</sup>	7.21 <sup>a</sup>
<i>Bayadha</i>	11.52 <sup>a</sup>	10.61 <sup>b</sup>	6.28 <sup>c</sup>	5.48 <sup>b</sup>	4.96 <sup>b</sup>
<i>Maali</i>	10.84 <sup>ab</sup>	16.95 <sup>a</sup>	12.37 <sup>a</sup>	10.80 <sup>a</sup>	8.87 <sup>a</sup>
<i>Razzek</i>	9.64 <sup>b</sup>	11.07 <sup>b</sup>	8.75 <sup>b</sup>	7.25 <sup>b</sup>	4.32 <sup>b</sup>
ANOVA					
Salinity (S)	***	***	***	***	***
Genotypes (G)	***	***	***	***	***
S × G	***	***	***	***	***
Alk-ase ( $\mu\text{g p-NP g}^{-1} \text{h}^{-1}$ )					
Irrigation water salinity (S)					
0.3 dS m <sup>-1</sup>	9.60 <sup>a</sup>	57.31 <sup>a</sup>	64.96 <sup>a</sup>	36.14 <sup>a</sup>	17.34 <sup>a</sup>
12 dS m <sup>-1</sup>	7.27 <sup>b</sup>	40.45 <sup>b</sup>	46.13 <sup>b</sup>	30.20 <sup>b</sup>	16.54 <sup>a</sup>
	(-24.27%)	(-29.42%)	(-28.99%)	(-16.44%)	(-4.61%)
Genotypes (G)					
<i>Agili Glabre</i>	6.18 <sup>b</sup>	66.00 <sup>a</sup>	50.22 <sup>b</sup>	32.99 <sup>b</sup>	21.01 <sup>a</sup>
<i>Bayadha</i>	10.30 <sup>a</sup>	34.89 <sup>b</sup>	54.40 <sup>b</sup>	26.27 <sup>c</sup>	11.19 <sup>c</sup>
<i>Maali</i>	9.44 <sup>ab</sup>	28.66 <sup>b</sup>	73.48 <sup>a</sup>	49.93 <sup>a</sup>	18.89 <sup>ab</sup>
<i>Razzek</i>	7.82 <sup>ab</sup>	65.98 <sup>a</sup>	44.69 <sup>b</sup>	23.50 <sup>c</sup>	16.83 <sup>b</sup>
ANOVA					
Salinity (S)	***	***	***	***	NS <sup>e)</sup>
Genotypes (G)	***	***	***	***	***
S × G	***	***	***	***	***
P-ase ( $\mu\text{g tyr g}^{-1} \text{h}^{-1}$ )					
Irrigation water salinity (S)					
0.3 dS m <sup>-1</sup>	33.77 <sup>a</sup>	37.35 <sup>a</sup>	39.85 <sup>a</sup>	28.52 <sup>a</sup>	15.87 <sup>a</sup>
12 dS m <sup>-1</sup>	31.17 <sup>b</sup>	29.49 <sup>b</sup>	30.71 <sup>b</sup>	21.24 <sup>b</sup>	12.43 <sup>b</sup>
	(-7.70%)	(-21.04%)	(-22.94%)	(-25.52%)	(-21.67%)
Genotypes (G)					
<i>Agili Glabre</i>	32.54 <sup>b</sup>	33.87 <sup>a</sup>	36.25 <sup>a</sup>	25.70 <sup>a</sup>	14.44 <sup>ab</sup>
<i>Bayadha</i>	30.13 <sup>c</sup>	30.91 <sup>a</sup>	34.01 <sup>a</sup>	21.61 <sup>b</sup>	11.86 <sup>c</sup>
<i>Maali</i>	34.89 <sup>a</sup>	35.56 <sup>a</sup>	34.81 <sup>a</sup>	25.91 <sup>a</sup>	16.79 <sup>a</sup>
<i>Razzek</i>	32.33 <sup>b</sup>	33.36 <sup>a</sup>	36.07 <sup>a</sup>	25.30 <sup>a</sup>	13.51 <sup>bc</sup>
ANOVA					
Salinity (S)	***	***	***	***	***
Genotypes (G)	***	NS	NS	***	***
S × G	**	***	NS	***	**

\*\*, \*\*\*Significant at  $P \leq 0.01$  and  $P \leq 0.001$ , respectively.

<sup>a)</sup> See Fig. 2 for abbreviations.

<sup>b)</sup> For each growth stage, irrigation water salinity values are the means of 20 measurements (4 genotypes and 5 replicates), while the genotype values are the means calculated from 10 measurements (2 irrigation water salinity and 5 replicates).

<sup>c)</sup> Means followed by different letters are significantly different at  $P \leq 0.05$  (Tukey test).

<sup>d)</sup> Values in parentheses indicate the rates of increase or decrease of each parameter according to the control treatment (0.3 dS m<sup>-1</sup>).

<sup>e)</sup> Not significant.

**Table S3. Irrigation water salinity and genotype effects on mineral nutrients (available P and N).**

Factors	Available P <sup>a)</sup> (mg kg <sup>-1</sup> )				
	6 leaves (Z16)	Tillering (Z30)	Flowering (Z65)	Maturity (Z85)	Harvesting (Z92)
<b>Irrigation water salinity (S)</b>					
0.3 dS m <sup>-1</sup>	12.11 <sup>b)c e)</sup>	23.97 <sup>a</sup>	17.17 <sup>b</sup>	12.81 <sup>c</sup>	7.50 <sup>d</sup>
12 dS m <sup>-1</sup>	8.48 <sup>c</sup> (-29.96%) <sup>d)</sup>	22.69 <sup>a</sup> (-5.36%)	16.20 <sup>b</sup> (-5.64%)	9.10 <sup>c</sup> (-36.95%)	4.97 <sup>d</sup> (-33.70%)
<b>Genotypes (G)</b>					
<i>Agili Glabre</i>	8.30 <sup>b</sup>	29.85 <sup>a</sup>	13.15 <sup>b</sup>	14.03 <sup>a</sup>	7.96 <sup>a</sup>
<i>Bayadha</i>	15.09 <sup>a</sup>	17.23 <sup>c</sup>	11.44 <sup>b</sup>	8.03 <sup>c</sup>	6.88 <sup>b</sup>
<i>Maali</i>	9.73 <sup>b</sup>	25.49 <sup>ab</sup>	20.98 <sup>a</sup>	10.93 <sup>b</sup>	6.26 <sup>b</sup>
<i>Razzek</i>	8.05 <sup>b</sup>	20.74 <sup>b</sup>	21.18 <sup>a</sup>	10.82 <sup>b</sup>	3.83 <sup>c</sup>
<b>ANOVA</b>					
Salinity (S)	NS <sup>e)</sup>	NS	**	***	***
Genotypes (G)	***	NS	***	***	***
S × G	***	NS	*	***	**
<b>Available N (mg kg<sup>-1</sup>)</b>					
<b>Irrigation water salinity (S)</b>					
0.3 dS m <sup>-1</sup>	35.56 <sup>a</sup>	38.81 <sup>a</sup>	40.92 <sup>a</sup>	29.91 <sup>b</sup>	25.69 <sup>b</sup>
12 dS m <sup>-1</sup>	27.45 <sup>a</sup> (-22.80%)	22.76 <sup>b</sup> (-27.14%)	31.25 <sup>a</sup> (-23.63%)	22.21 <sup>b</sup> (-25.74%)	20.50 <sup>c</sup> (-20.18%)
<b>Genotypes (G)</b>					
<i>Agili Glabre</i>	23.31 <sup>c</sup>	25.50 <sup>c</sup>	39.48 <sup>b</sup>	25.98 <sup>b</sup>	25.35 <sup>ab</sup>
<i>Bayadha</i>	32.22 <sup>b</sup>	22.87 <sup>c</sup>	24.55 <sup>bc</sup>	19.80 <sup>c</sup>	16.34 <sup>b</sup>
<i>Maali</i>	40.77 <sup>a</sup>	47.91 <sup>a</sup>	48.83 <sup>a</sup>	33.45 <sup>a</sup>	29.56 <sup>a</sup>
<i>Razzek</i>	29.77 <sup>ab</sup>	34.89 <sup>b</sup>	31.49 <sup>c</sup>	25.02 <sup>b</sup>	12.16 <sup>c</sup>
<b>ANOVA</b>					
Salinity (S)	***	***	***	***	***
Genotypes (G)	***	***	***	***	***
S × G	NS	***	***	***	***

\* \*\*, \*\*\*Significant at P ≤ 0.05, P ≤ 0.01, and P ≤ 0.001, respectively.

<sup>a)</sup> See Fig. 3 for abbreviations.

<sup>b)</sup> For each growth stage, irrigation water salinity values are the means of 20 measurements (4 genotypes and 5 replicates), while the genotype values are the means calculated from 10 measurements (2 irrigation water salinity and 5 replicates).

<sup>c)</sup> Means followed by different letters are significantly different at P ≤ 0.05 (Tukey test).

<sup>d)</sup> Values in parentheses indicate the rates of increase or decrease of each parameter according to the control treatment (0.3 dS m<sup>-1</sup>).

<sup>e)</sup> Not significant.

**Table S4. Irrigation water salinity and genotype effects on component yields (NG and GY).**

Factors	NG <sup>a)</sup>	GY (g)
Irrigation water salinity (S)		
0.3 dS m <sup>-1</sup>	146.75 <sup>b) a c)</sup>	4.98 <sup>a</sup>
12 dS m <sup>-1</sup>	137.32 <sup>b</sup>	3.67 <sup>b</sup>
	(-6.42%) <sup>d)</sup>	(-26.30%)
Genotypes (G)		
<i>Agili Glabre</i>	134.50 <sup>c</sup>	4.42 <sup>b</sup>
<i>Bayadha</i>	139.80 <sup>b</sup>	4.08 <sup>c</sup>
<i>Maali</i>	149.80 <sup>a</sup>	4.82 <sup>a</sup>
<i>Razzeck</i>	140.10 <sup>b</sup>	3.94 <sup>d</sup>
<b>ANOVA</b>		
Salinity (S)	***	***
Genotypes (G)	***	***
S × G	***	***

\*\*\*Significant at P ≤ 0.001.

<sup>a)</sup> See Fig. 4 for abbreviations.

<sup>b)</sup> Irrigation water salinity values are the means of 20 measurements (4 genotypes and 5 replicates), while the genotype values are the means calculated from 10 measurements (2 irrigation water salinity and 5 replicates).

<sup>c)</sup> Means followed by different letters are significantly different at P ≤ 0.05 (Tukey test).

<sup>d)</sup> Values in parentheses indicate the rates of increase or decrease of each parameter according to the control treatment (0.3 dS m<sup>-1</sup>).