

Original Research Article

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Media Standardization for Hydroponic Culture to Screen Wheat Genotypes for Nitrogen Use Efficiency

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ABSTRACT

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Root architecture is critical for efficient nutrient acquisition. This necessitates the investigations on plant root system to identify genotype with high nitrogen uptake efficiency under target environment. A hydroponic culture was employed to standardize the media to be used for screening of wheat genotypes for Nitrogen Use Efficiency (NUE) using proven low nitrogen susceptible and tolerant genotypes. Four different modified Hoagland solutions (NH₄NO₃ media, NO₃ complete, NO₃ half, and N minus media) were used and five wheat genotypes namely, UASBW 11328, GW 322 and PDW 291, NARBADA 4 and C 306 were grown in hydroponic culture for 30 days. Results regarding analysis of variance for interactions of genotypes and nitrogen levels revealed that all the genotypes exhibited the significant difference among themselves for all the four different solutions. Over the different media, low N susceptible genotype C 306 exhibited significant difference for most of the traits. While over the genotypes, both half NO₃ and N-minus recorded significant difference for root and shoot traits. Thus it was concluded that, half NO₃ medium is the best to screen the wheat genotypes for screening for NUE under 50 per cent supply. Further, this method gives the way for quick initial screening of wheat genotypes for NUE.

Introduction

Wheat (*Triticum aestivum* L.) is the main staple food of the world population and second important staple food crop in the country. Most of the agricultural crops, including wheat, are inefficient at uptaking and utilizing applied nitrogen fertilizer (Hitz, 2015). This emphasises breeding of wheat cultivars with improved nitrogen use

efficiency for reducing excessive input of fertilizers along with maintaining an acceptable yield. Plant roots play important role in nutrient uptake and selection for traits related to root architecture are of great importance for yield performance of crop species however, they remain poorly understood (Petarulo *et al.*, 2015). Study of

factors affecting the uptake of nutrients by roots and mechanism of element uptake in the field can be technically difficult (Hirel *et al.*, 2007). However, hydroponically grown plants, which facilitate rapid access to the root system where both physiology and molecular techniques could be, applied together (Garnet *et al.*, 2009). Cultivation of plant in mineral nutrient solution rather than in soil allows us to study the relationship between plant root nutritional uptake mechanism and plant growth (An *et al.*, 2006). Hence, An *et al.*, (2006) studied the effect of N levels on early growth of wheat seedlings using hydroponic culture. Keeping these points in view, the present investigation was conducted by supplying different forms and levels of N to Hoagland medium using the proven genotypes for NUE, by studying the effect of N levels on the growth and development of wheat shoot and root architecture.

Materials and Methods

To explore the objectives outlined above a hydroponic experiment was carried out in the Department of Biotechnology, University of Agricultural Sciences, Dharwad. Five diverse wheat genotypes namely UASBW 11328, GW 322, PDW 291, NARBADA 4 and C 306 were selected on the basis of their yield potential and NUE component traits like harvest index, NUE *etc.* falling in category such as high (UASBW 11328 and GW 322), medium (PDW 291 and NARBADA 4) and low (C 306) NUE genotypes (Satisha, 2016). The genotype UASBW 11328 is high yielding, having maximum biomass and high NUE. GW 322 is having high grain yield, high nitrogen uptake and utilization efficiency coupled with high harvest index and protein content under 50 per cent N. PDW 291 and NARBADA 4 are with high grain protein content and moderate NUE. The genotype C 306 is the tallest genotype, low chlorophyll content during grain filling stage, low grain

yield, harvest index, nitrogen harvest index (NHI) and NUE under normal N condition.

The experiment was conducted in hydroponic culture with modified Hoagland medium. The media was standardized for the hydroponic culture to screen the genotypes for nitrogen use efficiency in bread wheat. Four different modified Hoagland solutions were used *viz.*, media with both ammonical and nitrate form of N (NH₄NO₃ media), NO₃ complete media, NO₃ half media, and N minus media wherein in the N minus media, remaining micro nutrients and iron sources used was same as that of standard Hoagland solution (Hoagland and Arnon, 1938). The compositions of all four media are given in the Table 1, 2 and 3. Five genotypes were grown on the four different medium. Uniform seeds of five genotypes were surface-sterilised in 70 per cent ethanol for 1 min, followed by three rinses with deionised water. The seeds were germinated on filter paper in petri plates at room temperature over two days. The seedlings, with the embryogenic primary root, the two pairs of secondary roots (seminal roots) and the coleoptile (1–2 cm long) were then transplanted into supported hydroponic system with two replications. This system was formed using a polystyrene thermocol substrate that was placed into thermocol tanks that was filled with four different nutrient solutions (Hoagland modified solution) (An *et al.*, 2006; Petrarulo *et al.*, 2015). The nutrient solution in the tanks was renewed every 4-5 days and pH of the solution was adjusted to 6.0 using diluted NaOH and HCl before refreshing and maintained at the temperature of 25±2⁰ C with 75 per cent relative humidity and photoperiod of 16/8 hours. Four weeks after transplanting, the plant along with entire root system was carefully removed from the trays with the thermocol sheet. Data was collected on chlorophyll content, root length, shoot length, root to shoot ratio, number of primary roots, number of leaves, root fresh

and dry weight, shoot fresh and dry weight. After observations, the aerial parts of plants and roots were separated and oven dried for 48 hours at 80 °C and finally recorded the shoot dry weight (SDW) and root dry weight (RDW) (Petrarulo *et al.*, 2015).

Results and Discussion

Analysis of variance for factorial CRD analysis

The mean sum of squares for root and shoot traits under four nitrogen levels *viz.*, NH₄NO₃, NO₃ complete, NO₃ half and N minus media for five genotypes are presented in Table 4.

Analysis of variance showed significant variation among five genotypes at four different levels of nitrogen for root and shoot traits. This indicated existence of differences among genotypes at different levels of N. The significant genotype × nitrogen level interaction was observed for most of the traits except for the traits like, number of leaves and

dry weight of root. This indicated considerable amount of variability present in the material.

Effect of different nitrogen levels on five bread wheat genotypes grown under hydroponic culture

Data obtained from the analysis of interaction between genotypes and different nitrogen levels revealed the existence of sufficient genetic variability among the genotypes. In general, over the different nitrogen levels, the low N susceptible genotype C 306 exhibited significant differences with respect to mean performance of the root and shoot traits (Table 5 and Fig. 1). C 306 genotype which was bred for drought tolerance having good root parameters however, susceptible to low N supply (Renu Munjal and Satyavir Singh Dhanda, 2016). This suggests that root parameters alone may not contribute to NUE however it requires efficient nitrogen uptake and high harvest index.

Table.1 Components for Media with both ammonical and nitrate form of N used for hydroponic study

Sl. No.	Composition	For 1 litre (ml)
1	Calcium dihydrogen phosphate	10
2	Calcium sulphate	2
3	Potassium sulphate	5
4	MgSO ₄	2
5	NH ₃ NO ₄	1
6	Micronutrients	g l ⁻¹
	H ₃ BO ₃	2.86
	ZnSO ₄	0.22
	CuSO ₄	0.051
	MnCl ₂	1.81
	H ₂ MoO ₄ , H ₂ O	0.09
7	Iron EDTA	1.5

Table.2 Components for NO₃ complete media and NO₃ half media used for hydroponic study

Sl. No.	Composition	NO ₃ complete media (ml ⁻¹)	NO ₃ half media (ml ⁻¹)
1	Potassium nitrate (1M)	12.5 ml	6.25 ml
2	Calcium nitrate (1M)	12.5 ml	6.25 ml
3	Potassium dihydrogen phosphate (1M)	2.5 ml	1.25 ml
4	MgSO ₄ (1M)	5 ml	2.5 ml
5	Micronutrients	g ⁻¹	g ⁻¹
	H ₃ BO ₃	2.86	2.43
	ZnSO ₄	0.22	0.11
	CuSO ₄	0.051	0.026
	Mncl ₂	1.81	0.95
	H ₂ MoO ₄ , H ₂ O	0.09	0.045
6	Iron EDTA	3.75 ml	1.87 ml

Table.3 Components for N minus media used for hydroponic study

Sl. No.	Composition	For 1 litre (ml)
1	Calcium dihydrogen phosphate	10
2	Calcium sulphate	2
3	Potassium sulphate	5
4	MgSO ₄	2
5	Micronutrients	g ⁻¹
	H ₃ BO ₃	2.86
	ZnSO ₄	0.22
	CuSO ₄	0.051
	Mncl ₂	1.81
	H ₂ MoO ₄ , H ₂ O	0.09
6	Iron EDTA	1.5

Table.4 Mean performance of the parents for root and shoot traits under half NO₃ media in hydroponic culture

Parents	SPAD	RL	SL	RSR	NPR	NL	FWR	FWS	DWR	DWS
GW 322	30.80	28.50	37.60	0.76	10.38	6.58	0.16	0.93	0.03	0.26
2 WYCYT 34	30.05	30.71	40.53	0.76	11.91	6.00	0.17	0.97	0.04	0.22
K9107	28.65	29.82	39.30	0.76	11.25	5.90	0.17	0.86	0.04	0.19
UAS323	30.45	29.15	38.97	0.75	9.50	6.00	0.14	0.90	0.04	0.20
RAJ4248	28.95	28.05	37.52	0.75	10.53	6.00	0.13	0.86	0.03	0.18
C306	30.35	28.68	39.46	0.73	9.83	6.42	0.17	0.96	0.04	0.27

* and ** indicates significant at 5 (%) and 1 (%) level of significance, respectively.

SPAD = Chlorophyll content, RL = Root length (cm), SL = Shoot length (cm), RSR = Root to shoot ratio, NPR = Number of primary roots, NL = Number of leaves, FWR = Fresh weight of root (g), FWS = Fresh weight of shoot (g), DWR = Dry weight of root (g), DWS = Dry weight of shoot (g).

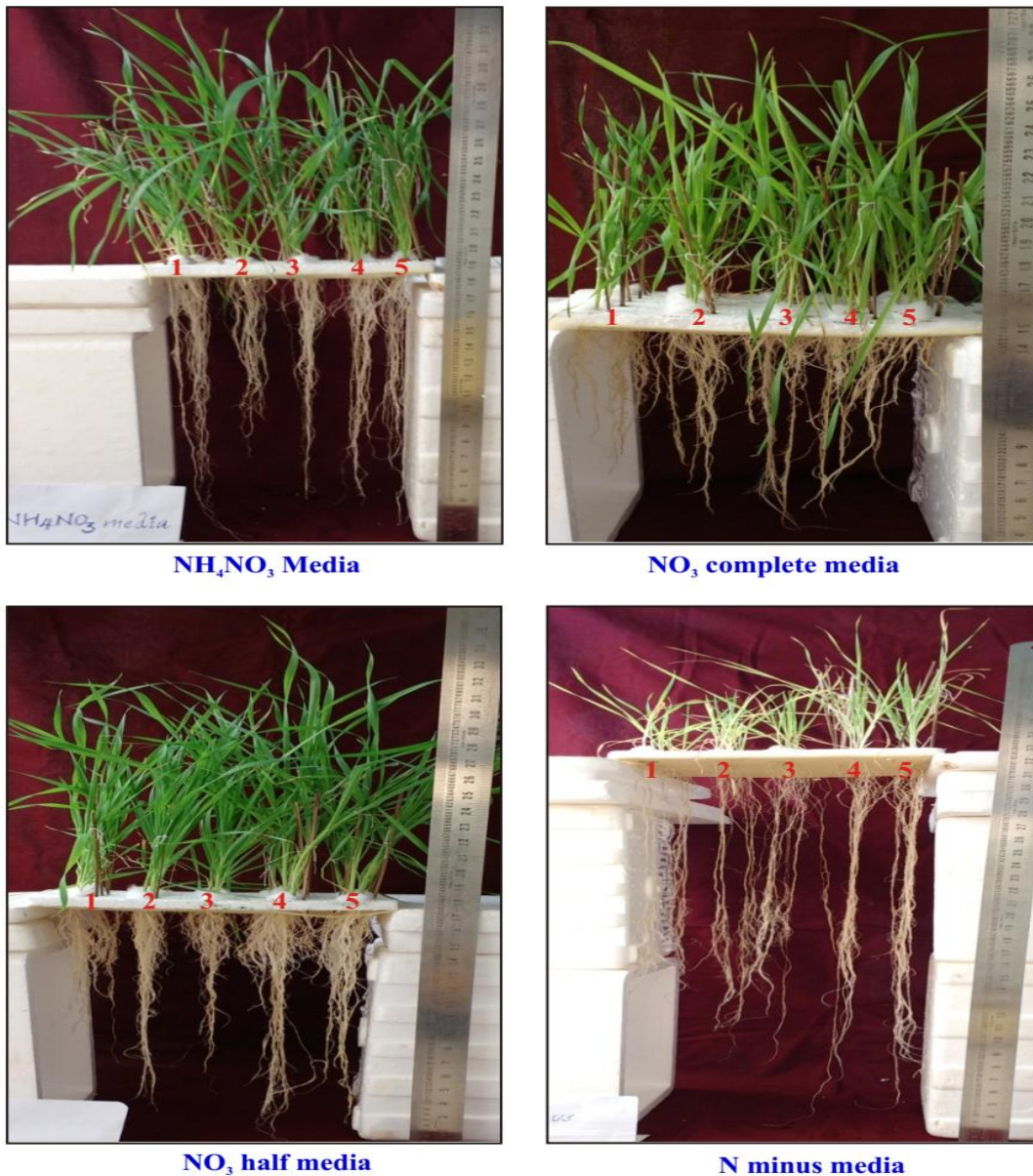
Table.5 Effect of different nitrogen levels on five bread wheat genotypes grown under hydroponic culture

	Treatments	SPAD	RL	SL	RSR	NL	NPR	FWR	FWS	DWR	DWS
Genotypes (G)	G1	30.15**	35.85	33.65	1.15	5.35	7.22	0.08	0.78	0.03	0.12
	G2	27.74	33.67	32.84	1.20**	5.89**	9.03	0.06	0.69	0.03	0.12
	G3	29.73**	36.62	33.06	1.18**	5.18	8.77	0.06	0.65	0.03	0.10
	G4	25.42	41.28**	44.83**	0.93	5.44	10.61	0.09	1.16**	0.05**	0.16**
	G5	28.00	40.13**	40.13**	0.96	5.76**	9.55**	0.10**	0.86	0.05**	0.14**
	Mean	28.21	37.51	36.90	1.09	5.52	9.04	0.08	0.83	0.04	0.13
	S.Em. ±	0.42	0.56	0.52	0.02	0.09	0.31	0.01	0.07	0.00	0.01
	C.D. (1 %)	1.70	2.27	2.11	0.07	0.36	1.24	0.03	0.27	0.01	0.02
Nitrogen level (N)	N1	36.12**	35.62	42.22**	0.82	6.08	9.18	0.07	1.00	0.03	0.16
	N2	29.49**	26.44	35.70	0.73	5.72	10.25	0.08	0.92	0.04	0.13
	N3	27.02	32.66	41.81**	0.79	6.05**	9.89**	0.10**	1.30**	0.06**	0.18**
	N4	19.65	55.32**	27.88	2.00	4.23	6.82	0.06	0.09	0.03	0.05
	Mean	28.07	37.51	36.90	1.09	5.52	9.04	0.08	0.83	0.04	0.13
	S.Em. ±	0.38	0.50	0.47	0.02	0.08	0.28	0.01	0.06	0.00	0.01
	C.D. (1 %)	1.52	2.03	1.89	0.06	0.32	1.11	0.02	0.24	0.01	0.02
Interaction (G x N)	G1N1	37.95**	40.36	39.17**	1.02	5.92**	7.25	0.05	1.02**	0.02	0.17**
	G1N2	31.70**	22.37	33.57	0.64	5.50	9.47	0.09	0.91**	0.02	0.11
	G1N3	29.75**	26.05	37.40**	0.69	5.80**	6.00	0.10	1.11**	0.05	0.16**
	G1N4	21.20	54.64	24.47	2.23**	4.17	6.17	0.07	0.07	0.03	0.05
	G2N1	34.40**	26.77	36.12	0.73	6.50**	8.63	0.04	0.63	0.03	0.14**
	G2N2	30.85**	29.08	32.21	0.92	6.43**	10.00**	0.08	0.83	0.04	0.13
	G2N3	28.75**	37.66	40.08**	0.94	6.45**	11.00**	0.09	1.25**	0.05	0.19**
	G2N4	16.95	41.16	22.97	2.22	4.17	6.50	0.03	0.07	0.04	0.04
	G3N1	39.03**	34.08	39.75**	0.90	6.00**	8.45	0.05	0.98**	0.02	0.17**
	G3N2	29.45**	24.62	35.17	0.72	5.00	9.64	0.06	0.62	0.04	0.12
	G3N3	29.10**	28.10	33.50	0.87	5.55	9.67	0.08	0.91	0.04	0.10
	G3N4	21.35	59.68	23.83	2.24**	4.17	7.33	0.07	0.09	0.03	0.04
	G4N1	32.90**	40.32	53.73**	0.75	6.00**	11.30**	0.10	1.42**	0.03	0.17**
	G4N2	27.25	30.22	36.02	0.70	5.75**	12.17**	0.12	1.23**	0.06	0.18**
	G4N3	22.50	39.34	53.48**	0.74	6.00	11.65**	0.11	1.89**	0.08	0.24**
	G4N4	16.25	55.26	36.10	1.55	4.00	7.33	0.05	0.11	0.04	0.06
	G5N1	36.30**	36.57	42.33**	0.72	6.00**	10.25**	0.12	0.94**	0.04	0.15**
	G5N2	28.20**	25.94	41.52**	0.64	5.91	10.00**	0.07	1.02**	0.03	0.14**
	G5N3	25.00	32.15	44.60**	0.73	6.45**	11.15**	0.13	1.35**	0.08	0.22**
	G5N4	23.61	65.89**	32.06	1.76	4.67	6.80	0.09**	0.12	0.06**	0.07
Mean	28.12	37.51	36.90	1.09	5.52	9.04	0.08	0.83	0.04	0.13	
S.Em. ±	0.85	1.13	1.05	0.03	0.18	0.62	0.01	0.13	0.01	0.01	
C.D. (1 %)	3.41	4.54	4.22	0.14	0.72	2.48	0.05	0.54	0.03	0.05	

* and ** indicates significant at 5 (%) and 1 (%) level of significance, respectively.

Genotypes (G): G1- UAS BW 11328, G2 - GW 322, G3 - PDW291, G4 – NARBADA 4, G5 - C306. Nitrogen level (N) : N1- NH₄NO₃ media, N2- NO₃ complete, N3- NO₃ half, N4- N minus.

Fig.1 Phenotype with respect to root and shoot traits of different bread wheat genotypes under different nitrogen levels



1 - UASBW 11328; 2 - WH 1022; 3 - PDW 291; 4 - NARBADA 4; 5 - C 306

With respect to different nitrogen levels, NO₃ half media exhibited significant differences from other nitrogen levels for the traits shoot length, number of primary roots, number of leaves, fresh weight and dry weight of root and fresh weight of shoot over the genotypes. The results regarding interactions of genotypes with nitrogen levels indicated that

the low NUE genotype C 306 at half NO₃ recorded significant difference for most of the traits and further, low N susceptible and tolerant genotypes exhibited significant differences among themselves for root parameters suggesting suitability of this media for screening of wheat genotypes for NUE.

Hydroponic system was proposed to screen the genotypes for nitrogen use efficiency by studying root and shoot traits. Results regarding analysis of variance for interactions of genotypes and nitrogen levels revealed that all the genotypes exhibit the significant difference among themselves for all the four different solutions and indicated that the low NUE genotype C 306 at half NO₃ medium recorded significant difference for most of the traits and further, it distinguished medium and high nitrogen use efficiency genotypes. Hence, the NO₃ half media can be further utilized to screen genotypes for NUE that are competitive for nitrogen (N) uptake in early vegetative stages of growth. Hence, the half NO₃ medium was further used in the screening of fifteen single cross hybrids and parents for nitrogen use efficiency. The present study gives new method for quick initial screening of wheat genotypes for NUE based on root parameters.

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