

MEAL AND OIL QUALITY AMONG GENOTYPES OF INDIAN MUSTARD (*BRASSICA JUNCEA*) VARIES UNDER RECOMMENDED DOSE OF NITROGEN FERTILIZER

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(Received 15th May 2017; accepted 25th Jul 2017)

Abstract. Out of the 24 genotypes of Indian mustard 63 % showed increase in total seed nitrogen content after 80 kg N (N80) application while 37 % showed a decrease compared to control (N0). In some genotypes the increase in total seed nitrogen content under N80 had no influence on total soluble protein. About 88 % genotypes showed reduction in crude fiber by 1.10 % (GM-2) to 35.6 % (DRMR-IJ-31) and in all 24 genotypes soluble sugars also reduced by 8% (Maya) to 80% (HB-9912) suggesting the influence of N application on digestibility of meal. Profiling of seed storage proteins showed differences in banding patterns in treated samples with prominent changes in α and β chains of cruciferin. In case of SFA and $\omega 6:\omega 3$ ratio, 50% of genotypes showed increase while 50 % genotypes showed a decrease in their SFA and $\omega 6:\omega 3$ ratio compared to control. Oil stability index also increased in 63 % of the genotypes under N80. Undesirable Erucic acid was found to increase in 58 % of the genotypes by 0.98% (DHR-991) to 38.31% (HB-9902) and in rest of the 42 % it reduced by 1.14 % (DRMR-IJ31) to 22.55 % (NRCDR-2) over control (N0).

Keywords: total phenol, ascorbic acid, total soluble sugars, crude fiber, seed storage protein

Introduction

Indian mustard is a winter oilseed crop largely grown in semi-arid regions of northern India. It is considered healthy and ideal for cooking because of low saturated fatty acid (SFA), high monounsaturated fatty acid (MUFA), moderate polyunsaturated fatty acid (PUFA) content, ideal $\omega 6:\omega 3$ ratio, low acid value, low saponification value and high smoke point. Besides high quality oil, the seed meal has high protein content which is ideal for animal feed or even fit for human consumption (Wanasundara, 2011). The high protein content also implies a high absorption of nitrogen from the soil which makes nitrogen fertilizers very important for this crop.

Plants take up nitrogen in the form of nitrate, as well as ammonium, this in turn can modulate the uptake of other anions and cations causing a change in the primary and secondary metabolic process and consequently the quality of oil and meal. Optimization of nitrogen application is important for the synthesis of C-based phytochemicals, including natural antioxidants such as phenolic groups (Björkman et al., 2011). Some

reports have shown decrease in phenol content under higher doses of nitrogen fertilizers (Li et al., 2008; Giorgi et al., 2009; Zaghdoud et al., 2016) while Tavarini et al. (2015) have observed increase in phenolic content up to a certain level of nitrogen dose. Likewise, ascorbic acid content is found to vary with nitrogen treatments in different Brassica crops like cabbage (Roni et al., 2015) and cauliflower (Lisiewska and Kmiecik, 1996). Ascorbic acid being an essential vitamin which needs to be obtained through diet is very important as an antioxidant and for synthesis of collagens L-carnitine and conversion of dopamine to norepinephrine.

There is one report of soluble sugars such as stachyose, raffinose and sucrose in diet causing flatulence in animals and digestive problems in poultry (Hartwig et al., 1997). Since seed meal from mustard has been suggested as a protein source for livestock, aquaculture and also for human consumption (Wanasundara, 2011), it is also necessary to look for potential anti nutritional factors in it. Also more emphasis should be given to soluble proteins than total proteins as they add more value to the seed meal. For example it is important to know the amount of storage proteins like albumin and globulin as they can be used for fortification of food products. Low fiber meal is preferred for animal feed because it increases the digestibility and palatability of the feed. The low metabolizable energy of mustard seed meal (2000 kcal/kg) compared to that of soybean (2230 kcal/kg) is one of the reasons for considering mustard seed meal as inferior (Suprianto, 2014).

Increasing population and dwindling natural resources have promoted the excess use of fertilizers. High yielding varieties of crops also demand heavy use of fertilizers which results in irreversible degradation of soil and natural resources. Excessive use of fertilizers also contributes to global warming. There have been numerous studies that relate fertilizer use and its effect on yield and oil content of oilseed Brassica (Joshi et al., 1998; Parmar and Parmar, 2012). However, little is known about the effect of fertilizer application on chemical composition of oil and seed meal.

Keeping this in view, in this study we have tried to see the effect of nitrogen fertilizer on the nutritional quality of oil and seed meal. In many of the mustard growing regions in India, use of 80 kg N/ha has been recommended to enhance the productivity of mustard (Shekhawat et al., 2012). No data to our knowledge are available on the study of nutritional composition of oil and meal in relation with nitrogen fertilizer application in Indian mustard. Such studies can open new avenues for future research and proper documentation pertaining to Indian mustard.

Materials and methods

Experimental site

The experiment was conducted at ICAR-Directorate of Rapeseed Mustard Research (DRMR) Bharatpur (27°11'N Lat. and 77°27'E Long.); India, during 2013-15. The climate of the region is predominantly semi-arid with long hot summers, intermittent short rainy period due to south-west monsoon and almost 5 month long winter when average temperature drops to below 22.5°C. The winter season often coincides with one or two rains due to western disturbances. The soils of the region are mainly coarse to fine loamy, deep, well drained Inceptisols and fallow-mustard is the predominant cropping system of the region. In general, the soils of the experimental site was alkaline (pH 8.1), low in exchangeable salts (EC 0.6 dS/m), organic carbon content (2.6 g/kg), available N (95.4 kg N/ha KMnO₄ oxidisable) and available P (5.4 kg P/ha 0.5N-

NaHCO₃ extractable); and high in available K (267 kg 1N-NH₄OAC exchangeable K₂O/ha). Initially, the field was put under exhaustive pearl millet (*Pennisetum glaucum*) – Indian mustard without any external fertilizer inputs for continuous three pre-experimentation years. *Pennisetum glaucum* was sown in the first week of July every year followed by Indian mustard var. Rohini.

Experimental design

During 2013-14, a replicated trial in split plot design keeping two nitrogen environments (0 and 80 kg N/ha) in main plot and 108 oilseed Brassica (OSB) genotypes in sub-plot was conducted to screen out low nitrogen requiring germplasms. Based on field and yield attributes, 24 genotypes were found promising. These selected 24 genotypes were further evaluated for detailed growth, yield and biochemical studies during 2014-15 under two distinct N environments. The genotypes were sown at 5 cm depth in 30 cm row spacing with the help of manual plough in subplots (4.5 m x 5.0 m) during second fortnight of October 2014. The plant to plant distance was maintained at 10-15 cm by thinning at 20-25 days after sowing (DAS). As per the treatment, half dose of nitrogen as urea, full recommended dose of phosphorus in the form of single super phosphate were applied at the time of sowing, and the remaining nitrogen in the form of urea was top dressed after the first irrigation. At maturity, 5 plants were randomly selected from the second and third row from the left side of the plot for recording growth and yield attributes. Harvesting was done by avoiding boarder area and observation lines manually by cutting at surface level using sickles to estimate biological and economic yields. Further, three samples from each plots were collected for all biochemical analysis. Each biochemical parameters for accuracy of data were done in three technical replicates.

Total nitrogen

The total nitrogen was estimated by miro-kjeldahl (Pro-Nitro A Model of JP Selecta, Spain) method as procedure suggested by AOAC (1955).

Fatty acid profiling

Fatty acid profiling was carried out as described by Paquot and Hautfenne, (1987) with little modification. A Nucon model 5765 gas chromatograph with SP-2300 (2%) + SP-2310 (3%) packed silica column was used. The programme was set at N₂ flow rate: 30 ml/min, H₂ flow rate: 30 ml/min, Zero air flow rate: 300 ml/min, Injector temp: 240 °C and Detector (FID) temp: 250 °C. Individual fatty acids were compared to the retention time of standard methyl esters of fatty acids.

Stability index of oils

Stability index of oils were calculated empirically by the ratio of MUFA: PUFA (Chauhan et al., 2010).

Defattening of seed meal

Oil was extracted from ground seeds after homogenizing with n-hexane using pestle and mortar at room temperature. This was repeated for three to four times to ensure

complete extraction of oil. The seed meal was dried till hexane is completely removed and stored at 4 °C.

Total phenols and crude fiber

Total phenols and crude fiber were estimated by non destructive method using FTNIR-Bruker (model Matrix-1) as described by Bala and Singh (2013).

Vitamin C (Ascorbic Acid)

Ascorbic acid was estimated using the titrimetric method described by Ranganna (1986). 0.1g of the defatted seed meal was homogenized with 3% HPO₃ and final volume made to 10 ml. The supernatant was collected into glass vials after centrifugation for 10 min at 5000 rpm. Aliquot of the extract was taken in a 25-50 ml conical flask and titrated against the standardized dye (2,6 -dichlorophenol indophenol) till persistent pink color was detected that lasted for 5s. Vitamin C was calculated using the following formula.

$$\text{mg of ascorbic acid/100 g} = \frac{\text{Titre} \times \text{Dye Factor} \times \text{Final volume made up}}{\text{Aliquot of extract taken} \times \text{Weight of sample taken}}$$

Total antioxidant capacity

Total antioxidant capacity (TAC) was determined as per Prieto et al. (1999). 100 mg of defatted seed meal was dissolved in 80 % methanol overnight. The supernatant was collected after centrifuging at 4000 rpm for 10 min and final volume was raised to 2 ml. Reduction of Mo(VI) to Mo (V) and the subsequent formation of green colour complex was measured by spectrophotometer (Labomed UV-VIS Double beam UVD-3500) at 695 nm and is expressed as ascorbic acid equivalent (AAE).

Total soluble sugars

Total soluble sugars were estimated as per protocol of Hansen and Møller (1975). 100 mg of defatted seed meal was homogenized with 100% acetone by vortexing. After centrifugation the residue was washed with hot ethanol (80%). This was repeated twice. Supernatant was collected and final volume was raised to 2 ml with 80% ethanol. This extract was used for the estimation of total soluble sugar by Anthrone method whose absorbance was taken at 620 nm.

Total soluble protein

Soluble protein content in the defatted seeds was estimated by the method of Lowry et al. (1951) after precipitation with trichloroacetic acid. Soluble protein was resuspended in laemmili lysis buffer denatured at 95⁰C for 4 min.

Gel electrophoresis

SDS-PAGE of seed storage protein was carried out in a 12 % Polyacrylamide gel in a discontinuous buffer system. 15 µl (20 µg/ µl) was loaded in each well of the stacking gel. Wide range prestained protein ladder was run as reference. The gel was stained with solution containing of 2 % Coomassie Brilliant Blue, followed by destaining after 1 h with mixture of methanol, acetic acid and water (5:1:4).

Statistical analysis

The data obtained under different treatments with respect to various parameters were subjected to Analysis of Variance (ANOVA) using SAS 9.3 software package.

Pair-wise comparisons on the least squares mean (LSMEANS) were performed using the Tukey's honest significant difference (HSD) test.

Results and discussion

Application of nitrogen influences the nutritional quality of meal

Total nitrogen content in seed

Studies on the effect of nitrogen on quality traits have been carried out in leaves of *B. juncea* and *B. rapa* (Falovo et al., 2011), cabbage (Leja et al., 2007), curly kale (Groenbaek et al., 2016) and broccoli (Zaghdoud et al., 2016), but not many reports on effect of applied nitrogen on quality of seed and oil are available. In our study, total seed nitrogen content ranged from 2.86 % (Maya and RL-1359) to 3.95 % (EC399300) at zero nitrogen (N0) application and 3.1% (RL-1359) to 4.3 % (IC212031) at 80 kg N/ha (N80) (Table 1). About 63 % of the genotypes showed increase in total seed nitrogen by 3 % (DRMR-IJ-31) to 34.0 % (NRCDR-2) over N0. The increase in total seed N content was proportional to the increase in total soluble protein which varied from 1% (Maya) to 34 % (NRCDR-2), with a confirmation of a positive correlation ($r=0.33$, $p0.019$) (Table 2.a). Though, exceptions in some genotypes showed reduction in soluble protein by 4 % (78-1-1-1) to 16% (NATP-124 and RGN-55) (Table 1). The decrease in soluble protein, even when there is increase in nitrogen intake, could be due to the partition of nitrogen towards the synthesis of other carbon containing compounds in TCA cycle as it shares common substrates. This is substantiated, by increase in ascorbic acid (14. 29 % to 50 %) and phenol (1.14 % to 14 %) content (Table 1). However, rest of the 37 % genotypes showed decrease in seed N content at N80 by 5.6 % (HB9916) to 20.2 % (EC399300) over N0. These are the genotypes that also showed a reduction in soluble proteins from 5 % (78-1-1-1, NRCHB-101, HB9916) to 16 % (RGN-55) and could be less responsive to nitrogen uptake. According to Chopin et al. (2007) in *Arabidopsis* there are high and low affinity nitrate transporters that regulate the uptake of nitrogen into plants. Our study also found genotypes such as EC399294 (5%), EC399307 (14%) and GM-2 (7%) that showed an increase in soluble proteins in spite of low content nitrogen. This interplay of nitrogen fertilization \times genotype \times quality trait needs more in-depth investigation.

Total phenol content

Natural antioxidants and their use as food additives or supplements has been a hot topic in recent years. Members of Brassica family are known to possess health promoting phytochemicals, especially those with antioxidant properties. Among the phytochemicals, phenolic compounds are very important due to their antioxidant properties (Björkman et al., 2011).

The total phenol content in our study ranged from 1.35 % (RL-1359) to 2.26 % (Rohini) under N0 and from 1.46 % (HB-9902) to 2.18 % (DRMR-IJ-31) under N80 (Table 1). There are reports of decrease in phenol content under higher doses of nitrogen fertilizer (Li et al., 2008, Giorgi et al., 2009; Zaghdoud et al., 2016) which was also

observed in our study as the phenol content reduced by 0.83 % (QM-16) to 21.83 % (Rohini) in about 33% of genotypes (*Table 1*). This reduction in phenol content as reported by Ibrahim et al. (2011) could be due to inhibition of phenylalanine ammonia lyase (PAL) at higher levels of nitrogen or may be due to diversion of metabolites in the direction of protein synthesis. As expected, 63 % of our genotypes increased in phenol content by 1.14 % (GM-2) to 14.01 % (RL-1359) over N0 application (*Table 1*). Our studies is in tune with work by Groenbaek et al. (2016) in traditional kale. Surprisingly, 4 % of our genotype showed no change in phenol content after N80 application. It is interesting to note within the same species the effect of nitrogen can have varied effects on the phenol content.

This discrepancy between different reports could be due to the differences in the form of nitrogen fertilizers (nitrate or ammoniacal) applied. Fallovo et al. (2011) reported increase in phenol content in the leaves of *B. juncea* and *B. rapa* due to forms of N fertilizer. Leja et al. (2007) observed decrease in concentration of phenolic compounds in cabbage head which were fertilized using calcium nitrate, ammonium sulphate, ammonium nitrate and urea by broadcasting. However, Smolen and Sady (2009) observed rise in phenolic concentration in carrots irrespective of the type and mode of nitrogen application. Sousa et al. (2008) have reported the profile of phenolic compounds to have varied even within the plant organ of the same Brassica species.

Our study suggests that variation in total phenol content is due to genetic variability that determines the optimization of nitrogen uptake and in turn alters the biochemical composition. According to Nguyen and Niemeyer (2008) in basil (*Ocimum basilicum* L.) phenolic content is a genotype dependent property. In this study more number of cultivars were analysed unlike the works reported previously in one or two cultivars.

Vitamin C (Ascorbic acid)

Another important natural antioxidant studied was ascorbic acid, an essential nutrient usually obtained from fruits and vegetables. The biological functions of ascorbic acid include radical scavenging, electron transport in plasma membrane and as cofactor in biosynthesis of collagen and conversion of dopamine to norepinephrine. It can scavenge superoxide and hydroxyl radicals besides regenerating α -tocopherol.

Our study on Indian mustard genotypes, showed significant variation in ascorbic acid content from 43mg/100g to 172 mg/100g seed meal (*Table 1*). However, about 58 % of the genotypes showed ascorbic acid content to increase by 14.29% (HB9916) to 50% (EC399307, NATP-124, NRC DR-2, BEC-16), about 21 % genotypes showed decrease in ascorbic acid content by 20 % (QM-16) to 100 % (HB-9902) and the rest 21 % of the genotypes did not show any change. Increase in ascorbic acid content upon nitrogen treatment has also been reported in cabbage (De and Shankar, 1987). Decrease in ascorbic acid in cabbage (Freyman et al., 1991) and cauliflower (Iisiewska and Kmiecik, 1996) have also been reported. The variation in ascorbic acid content could be due to their sensitivity to the enzyme GDP-Mannose pyrophosphorylase, an enzyme essential for the synthesis of metabolite GDP-Mannose, a precursor of ascorbic acid which has been confirmed in model plant to be hypersensitive to NH_4^+ and is a genetic determinant for NH_4^+ sensitivity (Qin et al., 2008). As we have also observed not all genotypes studied had the same levels of nitrogen in seed. This could be the reason for differences in ascorbic acid among the different genotypes. Our study is the first of its kind done in Indian mustard and more enquiries are needed to prove whether there is any dose dependent effect of nitrogen application on ascorbic acid content in seed meal.

Table 1. Biochemical Composition of 24 genotypes of *B. juncea* at N0 and N80 application

Genotypes	Total nitrogen (%)		Total phenols (%)		Ascorbic acid content (mg/100g)		Total antioxidant capacity (mg/g AAE)		Total soluble sugars (mg/g)		Total crude fiber (%)		Total soluble protein (%)	
	N0	N80	N0	N80	N0	N80	N0	N80	N0	N80	N0	N80	N0	N80
DRMR-IJ-31	3.72 ^{abcd}	3.85 ^{abc}	1.93 ^a	2.18 ^a	71.67 ^f	86.00 ^e	39.67 ^{onp}	46.83 ^{fgh}	178.33 ^q	133.33 ^x	8.84 ^{nmopqr}	5.69 ^u	13.97 ^w	16.76 ^{qrst}
Maya	2.86 ^{abcd}	3.58 ^d	1.75 ^a	1.71 ^a	107.50 ^c	86.00 ^e	39.83 ^{onp}	34.33 ^{rq}	243.33 ^g	223.33 ^k	9.24 ^{ijklmnop}	8.99 ^{klmnop}	15.44 ^{tuvw}	15.59 ^{stuvw}
78-1-1-1	3.55 ^{abcd}	3.31 ^{bcd}	1.72 ^a	1.74 ^a	43.00 ⁱ	43.00 ⁱ	45.75 ^{jihg}	40.00 ^{on}	143.33 ^w	131.11 ^y	10.3 ^{defghijklmn}	9.18 ^{klmnopqr}	17.11 ^{mnopqrs}	16.42 ^{qrstu}
DHR-991	3.4 ^{abcd}	3.58 ^{abcd}	1.76 ^a	1.675 ^a	64.50 ^g	86.00 ^e	45.50 ^{jih}	45.00 ^{jik}	546.67 ^a	404.44 ^b	9.85 ^{efghijklmn}	9.34 ^{hijklmnop}	15.67 ^{stuv}	18.43 ^{ijklmn}
EC399294	3.68 ^{abcd}	3.12 ^{bcd}	1.67 ^a	1.83 ^a	43.00 ⁱ	57.33 ^h	39.83 ^{onp}	46.00 ^{fihg}	108.33 ^a	81.11 ^f	8.89 ^{lmnopqr}	10.49 ^{bcddefghijkl}	21.02 ^{edf}	22.09 ^{bdc}
EC399300	3.95 ^{ab}	3.15 ^{bcd}	1.76 ^a	1.84 ^a	57.33 ^h	86.00 ^e	47.33 ^{efg}	51.50 ^c	74.44 ^g	62.22 ⁱ	9.8 ^{efghijklmn}	9.59 ^{ghijklmno}	18.65 ^{hijklm}	21.23 ^{edc}
EC399307	2.94 ^{dc}	3.54 ^{abcd}	1.84 ^a	2.01 ^a	86.00 ^e	172.00 ^a	39.67 ^{onp}	54.33 ^b	143.33 ^w	60.00 ^j	8.76 ^{nmnopqr}	6.22 ^{tu}	18.02 ^{ijklmnopq}	15.92 ^{rstuv}
GM-2	3.85 ^{abc}	3.15 ^{bcd}	1.73 ^a	1.75 ^a	86.00 ^e	86.00 ^e	44.17 ^{jkl}	42.67 ^{ml}	276.67 ^d	66.67 ^h	8.17 ^{opqrs}	8.08 ^{pqrs}	18.22 ^{ijklmnop}	19.47 ^{fghi}
Rohini	3.12 ^{bcd}	3.97 ^{ab}	2.26 ^a	1.855 ^a	71.67 ^f	43.00 ⁱ	48.83 ^{ef}	50.67 ^c	227.78 ^j	165.56 ^s	7.03 ^{stu}	7.53 ^{rst}	17.88 ^{ijklmnopq}	18.30 ^{ijklmno}
HB-207	3.64 ^{abcd}	3.61 ^{abcd}	1.53 ^a	1.65 ^a	172.00 ^a	129.00 ^b	44.50 ^{jik}	47.50 ^{ef}	182.22 ^o	126.67 ^z	9.98 ^{efghijklm}	10.68 ^{abcddefghij}	18.96 ^{hijkl}	21.56 ^{edc}
HB-9902	3.21 ^{bcd}	3.66 ^{abcd}	1.46 ^a	1.46 ^a	86.00 ^e	43.00 ⁱ	45.00 ^{jik}	33.33 ^{rs}	97.22 ^b	60.00 ^j	10.86 ^{abcddefghi}	10.48 ^{bcddefghijkl}	17.43 ^{lmnopqr}	20.66 ^{edfc}
HB-9912	3.92 ^{abc}	3.89 ^{abc}	1.71 ^a	1.76 ^a	43.00 ⁱ	43.00 ⁱ	23.83 ^{vw}	21.83 ^{yx}	173.33 ^r	35.00 ^k	11.07 ^{abcddefg}	9.1 ^{klmnopqr}	19.11 ^{ghijk}	21.27 ^{edc}

HB-9916	3.4 ^{abcd}	3.21 ^{bcd}	1.44 ^a	1.65 ^a	86.00 ^e	100.33 ^d	20.83 ^y	48.67 ^e	277.78 ^d	150.00 ^v	11.64 ^{abc}	10.62 ^{abcdefghij}	18.59 ^{hijklm}	17.60 ^{klmnopq}
IC212031	3.21 ^{bcd}	4.3 ^a	1.66 ^a	1.69 ^a	86.00 ^e	86.00 ^e	33.17 ^{rs}	22.50 ^{wx}	133.33 ^x	90.00 ^d	10.48 ^{bcdefghijkl}	9.95 ^{defghijklm}	17.04 ^{mnpqrst}	18.45 ^{ijklmn}
NATP-124	3.9 ^{abc}	3.94 ^{ab}	1.57 ^a	1.65 ^a	43.00 ⁱ	86.00 ^e	24.33 ^v	31.83 ^{ts}	261.67 ^f	200.00 ^m	11.54 ^{abcd}	10.82 ^{abcdefghi}	24.15 ^a	20.07 ^{efgh}
QM-16	3.36 ^{abcd}	3.48 ^{abcd}	1.82 ^a	1.805 ^a	86.00 ^e	71.67 ^f	39.67 ^{onp}	23.50 ^{vw}	237.78 ^h	92.22 ^c	9.81 ^{efghijklmn}	8.46 ^{nopqrs}	24.25 ^a	22.84 ^{abc}
RGN-55	3.68 ^{abcd}	3.2 ^{7bcd}	1.68 ^a	1.84 ^a	71.67 ^f	86.00 ^e	23.33 ^{vw}	31.17 ^t	263.33 ^e	133.33 ^x	10.5 ^{bcdefghijkl}	10.21 ^{cdefghijklm}	21.01 ^{edf}	17.54 ^{klmnopqr}
NRCHB-101	3.67 ^{abcd}	3.4 ^{abcd}	1.81 ^a	1.80 ^a	57.33 ^h	86.00 ^e	30.83 ^t	28.83 ^u	160.00 ^u	80.00 ^f	9.88 ^{efghijklmn}	9.76 ^{ghijklmno}	16.89 ^{nopqrst}	17.79 ^{ijklmnop}
NRCDR-2	3.19 ^{bcd}	3.88 ^{abc}	1.72 ^a	1.62 ^a	43.00 ⁱ	86.00 ^e	42.50 ^m	43.67 ^{mkl}	297.78 ^c	96.67 ^b	12.2 ^a	11.2 ^{6abcdef}	17.40 ^{lmnopqr}	23.40 ^{ab}
NRCHB506	3.16 ^{bcd}	3.28 ^{bcd}	1.83 ^a	1.64 ^a	86.00 ^e	86.00 ^e	57.17 ^a	39.83 ^{onp}	185.56 ⁿ	162.22 ^t	11.5 ^{4abcd}	10.94 ^{5abcdefgh}	17.87 ^{ijklmnopq}	18.15 ^{ijklmnop}
Pusa Jai kisan	3.16 ^{bcd}	3.17 ^{abcd}	1.56 ^a	1.67 ^a	57.33 ^h	71.67 ^f	23.67 ^{vw}	35.17 ^q	163.33 ^t	88.33 ^e	11.32 ^{a⁴bcdef}	10.52 ^{bcdefghijk}	14.02 ^w	14.51 ^{vw}
NDRE-7	3.04 ^d	3.4b ^{cd}	1.68 ^a	1.58 ^a	57.33 ^h	71.67 ^f	35.17 ^q	40.83 ⁿ	230.00 ⁱ	173.33 ^r	11.39 ^{abcde}	11.33 ^{abcdef}	14.88 ^{uvw}	17.52 ^{klmnopqr}
RL-1359	2.86 ^{bcd}	3.1 ^{bcd}	1.35 ^a	1.57 ^a	43.00 ⁱ	57.33 ^h	38.33 ^p	39.17 ^{qp}	202.22 ^l	180.00 ^p	12.02 ^{ab}	11.1 ^{abcdefg}	15.56 ^{stuvw}	16.64 ^{pqrst}
BEC-16	3.08 ^{bcd}	3.14 ^{bcd}	1.73 ^a	1.92 ^a	43.00 ⁱ	86.00 ^e	38.50 ^{op}	50.33 ^{c^d}	276.67 ^d	243.33 ^g	8.93 ^{klmnopqr}	7.57 ^{rst}	18.18 ^{ijklmnop}	19.20 ^{ghij}

T comparison lines for least Squares means of genotypes*nitrogen
 LS- means with same letters are not significantly different

Total antioxidant capacity

Phenolic compounds and ascorbic acid are known for its antioxidant properties which are highly desirable for improving the keeping quality of meal as well as for providing health benefits. Seed meal extracts of 24 genotypes of Indian mustard showed TAC to range between 20 mg/g AAE (HB-9916) to 48 mg/g AAE (Rohini) under N0 and between 21.83 mg/g AAE (HB9912) to 54 mg/g AAE (EC399307) under N80 application (*Table 1*). We observed, 58 % of genotypes showed TAC to increase and 42 % of genotypes showed reduction in TAC over N0. It may be noted that the genotypes in which the TAC was lowered also showed concomitant reduction in ascorbic acid. According to Nguyen and Niemeyer (2008) reduction in TAC upon treatment with NO_3^- correlated with phenol content. Aires et al. (2011) examined the potential antioxidant activities of ascorbic acid, total flavanoid and total phenol in Brassica vegetables and concluded that among the naturally occurring antioxidants, ascorbic acid contributed the maximum towards antioxidant capacity. Our findings are in agreement with this. Ochoa-Velaso et al. (2016) have reported that TAC was strongly influenced by ascorbic acid and phenolic content in tomatoes. The correlation between TAC with that of phenols and ascorbic acid observed in our study ($r=0.316$, $p=0.028$) is comparatively less but statistically significant (*Table 2a*).

Table 2a. Pearson Correlation between total antioxidant, vitamin C and total phenol

Pearson Correlation Coefficients, N=48 Prob> r under H0: Rho=0				
	Total antioxidant capacity	Phenol	Ascorbic acid	Phenol+ Ascorbic acid
Total antioxidant capacity	1.00000	0.36613 0.0105	0.31511 0.0291	0.31699 0.0281
Phenol	0.36613 0.0105	1.00000	0.12944 0.3806	0.13515 0.3597
Ascorbic acid	0.31511 0.0291	0.12944 0.3806	1.00000	0.99998 <.0001
Phenol+ Ascorbic acid	0.31699 0.0281	0.13515 0.3597	0.99998 <.0001	1.00000

Table 2b. Pearson Correlation between total nitrogen and soluble protein

Pearson Correlation Coefficients, N= 48 Prob > r under H0: Rho=0		
nitrogen	nitrogen	protein
protein	1.00000	0.33743 0.0190

Soluble sugars

Seed meal of *B. juncea* is comprised of about 30% carbohydrates which includes many beneficial as well as harmful components. One report showed soluble sugars such as stachyose and raffinose are harmful as it causes flatulence in animals and digestive problems in poultry (Hartwig et al., 1997). In this study, we evaluated the effect of N80 treatment on soluble sugar content. The total soluble sugar ranged between 74.44 mg/g (EC399300) to 546.71 mg/g (DHR-991) under N0. Under, N80 application it ranged from 60 mg/g (EC399307, HB9902) to 404 mg/g (DHR-991). All genotypes reported reduction in total soluble sugar level ranging from 8 % (Maya) to 80% (HB-9912). Similar observations were made by Smolen and Sady (2009) in carrots (*Daucus carota* L.). Further, Krober and Cartter (1962) had shown reduction in soluble sugar content in four cultivars of soybean at higher nitrogen input with a concomitant increase in protein content. Higher nitrogen levels may divert the biosynthetic process as towards the amino acids and proteins consequently reducing the production of soluble carbohydrates. This may be true in 63% genotypes having inverse relationship between soluble sugars and seed nitrogen content. But, contrasting results was observed in 37 % of genotypes where they experienced reduction in both parameters after N80 treatment.

Cruder fiber

Fiber is primarily derived from cell walls and is made up of cellulose, hemicelluloses and lignin. High fiber in animal's diet is undesirable as it would dilute the availability of energy, proteins and minerals such as Mn and Zn (Suprianto, 2014). As an animal feed, the seed meal should be preferably rich in protein and energy rich carbohydrates rather than crude fiber that only add to the bulk and cause digestive disorders. The crude fiber content in the genotypes under N0 ranged between 7.03 % (Rohini) and 12.02 % (RL-1359 and NRCDR-2) whereas; it ranged from 5.69 % (DRMR-IJ-31) to 11.33% (NDRE-7) at N80 application (*Table 1*). 88 % of the genotypes showed reduction in crude fiber content by 1.10 % (GM-2) to 35 % (DRMR-IJ-31) over N0. One report is in agreement with decrease of crude fiber content in fodder beet but it is also dependent on the type of nitrogen fertilizer (Khogali et al., 2011). Within the same experiment they observed no significant effects in maize. Similarly, in forage kale no effect on fiber content was observed (Chakwizira et al., 2015). It is quite evident that response to nitrogen application is not the same in all crops. In the present experiments about 12 % of genotypes showed increase in crude fiber content by 7 % (Rohini, HB-207) to 17.9 % (EC399294). This suggests that variations in seed nitrogen upon N80 application leads to variation in crude fiber and is genotype dependent.

Total soluble protein and seed storage protein

The two major seed storage proteins of oilseed *Brassica* are the 2S albumin called napin and the 11S or cruciferin a legume type globulin having heteromeric structure. Cruciferin, rich in nitrogen containing amino acids have role in modulating nitric oxide signaling which is potential regulator of nutrient metabolism (Wanasundara, 2011), while albumin rich in sulphur containing amino acids have been annotated with several allergic responses such as celiac disease and Baker's asthma (Wanasundara, 2011). However, not all 2S

albumin causes allergy and it affects only a minority of the population (Wanasundara, 2011). None the less, both storage proteins have found its place in industry as they are highly surface active in monolayers and emulsion formation (Wanasundara, 2011).

Profiling of seed storage proteins with respect to nitrogen treatment has never been done in Indian mustard. We have profiled the seed storage proteins in 24 genotypes of *B. juncea* using SDS–PAGE to see the effect of nitrogen application on the protein profile (*Fig. 1*). Molecular weights of the proteins were compared to previous reports (Wanasundara, 2011, Wanasundara et al., 2012). Our results showed genotypes DRMR-IJ-31, HB-207, HB-9902, NATP-124, RGN-55, NRCHB-506 had low expression of 2S albumin in case of nitrogen treatment (*Fig. 1*). According to Wanasundara et al. (2012) molecular weight of cruciferin ranged between 18.1 kDa and 31.2 kDa in *B. juncea* and that of napin between 6.5 and 12 kDa in reducing environment. But we found the corresponding band between 15 and 16 kDa for napin. This difference in molecular weight may be due to the slight modification in the protocol as we had given heat treatment only for 4 min instead of 15 min as reported earlier. We found genotypic variability in the banding patterns of α and β chains of cruceferin (*Fig 1*). According to Yu (2008) this determines the digestibility of proteins in *B. rapa* and *B. napus*. Since this is the first report in *B. juncea*, it will lead to better ideas on manipulating the meal composition for quality improvement by managing nitrogen fertilization.

Application of nitrogen influences the nutritional quality of oil

Nutritional composition of oil

There are reports on the effect of nitrogen fertilizer on improvement of oil yield and quality (Joshi et al., 1998; Paramar and Paramar, 2012) but, there are no sufficient data to support the role of nitrogen fertilizers in modifying the nutritional composition of oil in *B. juncea* genotypes. Fatty acid profiling in 24 genotypes was done under treated and controlled conditions (*Table 3*). Our results showed, SFA (Palmitic +Stearic acid) content to be within 1.37 % (RGN-55) to 5.76 % (HB-9902) at N0 and between 1.96% (HB-9916) to 4.32 % (NDRE-7) under N80 (*Table 3*). About 50 % of genotypes showed decrease in SFA and 50 % showed increase compared to control, but the rise in SFA levels were still within the permissible limit (< 7%) of healthy edible oil. Paramar and Paramar (2012) have also found a significant decrease in SFA at 50 kg/ha. Reduction in SFA is desirable as it is associated with cardiovascular diseases. Even though, SFA content of Indian mustard is within the optimum range it is possible to manipulate its level by optimizing the nitrogen application as evident from our study.

The level of MUFA (Oleic + Eicosenoic + Erucic) ranged from 50.79% (HB-9902) to 63.51% (NDRE-7) under N0 and from 54.16 % (NDRE-7) to 61.65 % (NATP-124) under N80. The levels of MUFA were found to have increased in about 58 % of genotypes by 1.02% (EC39900) to 20.46 % (HB-9902) over N0, while in the rest 42% of genotypes it decreases by 0.27 % (78-1-1-1) to 14.72 % (NDRE-7) over control.

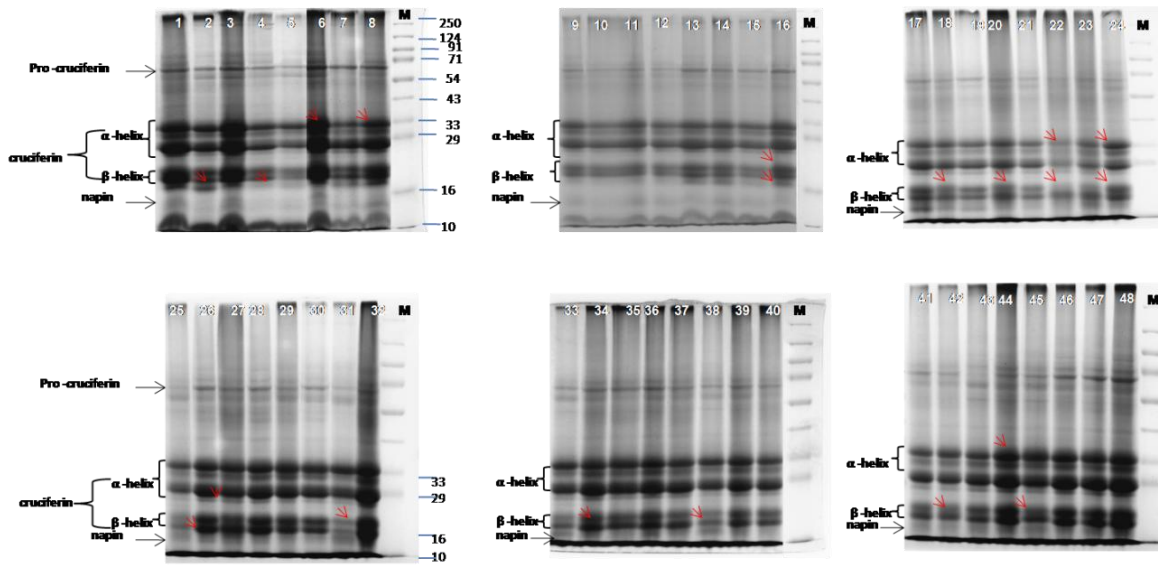


Figure 1. Seed storage protein profiling in 24 genotypes of Indian Mustard under zero and 80kg/ha nitrogen application

Lane 1 and 2 : DRMR II-31 under control and 80kg N/ha; Lane 3 and 4 : Maya under control and 80kg N/ha; Lane 5 and 6: 78-1-1-1 under control and 80kg N/ha; Lane 7 and 8: DHR-991 under control and 80kg N/ha; Lane 9 and 10:GM-2 under control and 80kg N/ha; Lane 11 and 12:EC-399307 under control and 80kg N/ha; Lane 13 and 14: EC-399300 under control and 80kg N/ha; Lane 15 and 16: EC-399294 under control and 80kg N/ha; Lane 17 and 18: Rohini under control and 80kg N/ha. Lane 19 and 20:HB-207 under control and 80kg N/ha; Lane 21 and 22:HB-9902 under control and 80kg N/ha; Lane 23 and 24: HB-9912 under control and 80kg N/ha; Lane 25 and 26:HB-9916 under control and 80kg N/ha; Lane 27 and 28:IC212031 under control and 80kg N/ha; Lane 29 and 30:NATP-124 under control and 80kg N/ha; Lane 31 and 32: QM-16 under control and 80kg N/ha; Lane 33 and 34: RGN-55 under control and 80kg N/ha; Lane 35 and 36: NRCHB-101 under control and 80kg N/ha; Lane 37 and 38: NRCDR-2 under control and 80kg N/ha; Lane 39 and 40: NRCHB-506 under control and 80kg N/ha; Lane 41 and 42:Pusajai kisan under control and 80kg N/ha; Lane 43 and 44: NDRE-7 under control and 80kg N/ha; Lane 45 and 46: RL-1359 under control and 80kg N/ha; Lane 47 and 48 BC-16: under control and 80kg N/ha; Lane M: wide ranged protein ladder (kDa). (→ indicates the change in α and β banding pattern.

Mustard oil contains moderate levels of PUFA that includes linoleic acid and linolenic acid (15-20 %) which is ideal range for healthy oil. They are essential fatty acids and are required for the biosynthesis of eicosapentanoic acid and docosahexanoic acid, which are beneficial to human beings. In this study, linoleic acid (ω_6) ranged from 13.47 % (DHR-991) to 22.94 % (HB-9002) under N0 and 15.20% (NRCHB-506) to 22.38 % (QM-16) for N80. Since, almost all genotypes analyzed in this study are having >15% ω_6 content, they can be used as potential source for extraction (Zambiazzi et al., 2007). It is to be noted, about 42 % of the genotypes showed increase in ω_6 by 1.23% (NRCHB-101) to 29.32 % (DHR-991) over N0. Whereas, 58 % of the genotypes showed reduction in ω_6 by 0.89 % (EC399300) to 18.20 % (HB-9916) over N0 application.

Table 3. Fatty acid profile of 24 genotypes of *B. juncea* at N0 and N80 application

Samples	Palmitic (%)	Stearic (%)	Oleic (%)	Linoleic (%)	Linolenic (%)	Eicosenoic (%)	Erucic (%)	$\omega 6/\omega 3$	SFA (%)	MUFA (%)	PUFA (%)	MUFA:PUFA
DRMRIJ-31 (N₀)	2.83 ^{fg hijklmn}	0.17 ^{qrst}	7.86 ^t	18.7 ^{at}	19.56 ^f	15.42 ^d	35.15 ^x	0.956 ^{uvts}	3 ^k	58.43 ^s	38.26 ^{fg hijklmn}	1.53 ^{bcdefghi}
DRMRIJ-31 (N₈₀)	2.67 ^{fg hijklmn}	0.22 ^{pqr}	8.04 ^s	17.97 ^{yx}	20.36 ^c	15.29 ^e	34.75 ^z	0.883	2.89 ^k	58.08 ^t	38.33 ^{fg hijklmn}	1.52 ^{bcdefg}
Maya (N₀)	3.44 ^{abcdef}	0.53 ^{bcd}	9.99 ^f	20.1 ^{lk}	17.94 ^q	13.26 ^l	34.11 ^b	1.120 ^{ijklmn}	3.97 ^{no}	57.36 ^w	38.04 ^{bcdefg}	1.51 ^{bcdefghi}
Maya (N₈₀)	2.06 ^{ijklmnopq} _r	0.38 ^{hijklm} _n	8.72 ⁿ	17.87 ^z	17.93 ^q	12.2 ^u	39.56 ⁿ	0.997 ^{qrstu}	2.44 ^x	60.48 ^{iw}	35.8 ^{mnpqr}	1.69 ^{bcdefg}
78-1-1-1 (N₀)	2.32 ^{ijklmnop}	0.24 ^{pqr}	10.07 ^f	20.92 ^g	19.17 ⁱ	14.34 ^j	31.75 ^f	1.091 ^{ijklmnop}	2.56 ^g	56.16 ^x	40.09 ^{ijklmnopq}	1.40 ^{fg hi}
78-1-1-1 (N₈₀)	3.5 ^{bcdef}	0.21 ^{qrs}	6.69 ^z	20.54 ⁱ	17.8 ^r	12.8 ^{qp}	36.52 ^u	1.154 ^{ghijk}	3.71 ^k	56.01 ^y	38.34 ^{bcdef}	1.46 ^{bcdefghi}
DHR-991 (N₀)	2.14 ^{lmnopq}	0.41 ^{fghijkl} _m	6.83 ^y	13.47 ^g	13.61 ⁱ	12.99 ⁿ	39.64 ^{nm}	0.990 ^{qrstu}	2.55 ^k	59.46 ^{no}	27.08 ^{lmnopq}	2.20 ^a
DHR-991 (N₈₀)	3.52 ^{bcdef}	0.58	8.42 ^p	17.42 ^b	17.39 ^t	12.01 ^v	40.03 ^k	1.002 ^{qrst}	4.1 ^c	60.46 ⁱ	34.81 ^{bcdef}	1.74 ^{bcdefghi}
EC399294 (N₀)	3.58 ^{bcdef}	0.48 ^{defg}	9.97 ^f	18.83 ^s	16.97 ^w	11.85 ^{wx}	37.63 ^q	1.110 ^{ijklmno}	4.06 ^x	59.45 ^o	35.8 ^{bcdef}	1.66 ^{bcdefghi}
EC399294 (N₈₀)	2.29	0.33 ^{omn}	8.36 ^p	16.06 ^b	15.97 ^{az}	10.75 ^c	41.1 ^g	1.006 ^{pqrst}	2.62 ^g	60.21 ^j	32.03 ^{ijklmnop}	1.88 ^{bcdef}
EC399300 (N₀)	3.41 ^{bcdefg}	0.35 ^{klmn}	9.78 ^g	22.36 ^c	18.45 ^{nm}	14.93 ^g	30.38 ^j	1.212 ^{defgh}	3.76 ^e	55.09 ^{dc}	40.81 ^{bcdefg}	1.35 ^{hi}
EC399300 (N₈₀)	3.02 ^{defghijkl}	0.48 ^{dgef}	11.26 ^c	22.16 ^d	17.79 ^r	12.88 ^{op}	31.51 ^g	1.246 ^{cdef}	3.5 ^h	55.65 ^a	39.95 ^{defghijklm}	1.39 ^{bcdefghi}
EC399307 (N₀)	3.79 ^{bcde}	0.46 ^{defgh}	9.11 ^v	19.61 ^p	16.73 ^x	13.18 ^{ml}	36.46 ^u	1.172 ^{efghij}	4.25 ^v	58.75 ^r	36.34 ^{bcde}	1.62 ^{bcdefghi}
EC399307 (N₈₀)	2.97 ^{defghijkl} _{mn}	0.4 ^{ghijklm}	7.8 ^{tu}	17.61 ^a	15.49 ^e	11.28 ^a	43.25 ^d	1.137 ^{hijklm}	3.37 ^f	62.33 ^c	33.1 ^{defghijklm}	1.88 ^{abcdef}
GM-2 (N₀)	2.97 ^{defghijkk} _{lm}	0.13 ^{ts}	6.48 ^{ao}	18.04 ^x	19.34 ^h	11.94 ^v	41.14 ^{gf}	0.933	3.1 ^t	59.56 ^m	37.38 ^{defghijklm}	1.59 ^{bcdefghi}

GM-2 (N₈₀)	1.89 ^{nopqr}	0.34 ^{lmn}	8.56 ^j	21.47 ^e	15.66 ^c	7.93 ⁱ	43.74 ^c	1.371 ^{ba}	2.23 ^u	60.23 ^{kj}	37.13 ^{nopqr}	1.62 ^{abcdefg}
Rohini(N₀)	1.95 ^{nopqr}	0.19 ^{qrst}	7.52 ^w	18.34 ^v	19.76 ^d	12.12 ^u	39.7 ^{lm}	0.928 ^{uv}	2.14 ^{nm}	59.35 ^p	38.1 ^{nopqr}	1.56 ^{bcdefgh}
Rohini(N₈₀)	3.1 ^{defghijk}	0.25 ^{opq}	9.47 ^j	19.73 ^o	15.57 ^{de}	14.57 ⁱ	37.23 ^r	1.267 ^{cd}	3.35 ^z	61.27 ^f	35.3 ^{defghijk}	1.74 ^{bcdefgh}
HB-207 (N₀)	2.84 ^{efghijklmn}	0.22 ^{pqr}	9.66 ^{ih}	20.86	19.81 ^d	13.16 ^m	32.28 ^e	1.053 ^{mnopq}	3.06 ^f	55.1 ^c	40.67 ^{efghijklmn}	1.35 ^{efghi}
HB-207 (N₈₀)	2.17 ^{klmnop}	0.49 ^{cdef}	10.41 ^e	19.83 ⁿ	15.06 ^g	11.51 ^z	40.2 ^j	1.317 ^{bc}	2.66 ^b	62.12 ^d	34.89 ^{klmnop}	1.78 ^{bcdefgh}
HB-9902 (N₀)	4.8 ^a	0.96 ^a	11.87 ^a	22.94	18.38 ^{npo}	9.16 ^f	29.76 ^k	1.248 ^{cdef}	5.76 ^c	50.79 ⁱ	41.32 ^a	1.23 ⁱ
HB-9902 (N₈₀)	2.43 ^{hijklmnop}	0.53 ^{bcd}	8.23 ^q	19.65 ^{po}	16.02 ^z	11.79 ^{yx}	41.16 ^{ef}	1.227 ^{defg}	2.96 ^{poy}	61.18 ^g	35.67 ^{hijklmnop}	1.72 ^{abcdefg}
HB-9912 (N₀)	4.14 ^{ab}	0.42 ^{efghijkl}	10.4 ^d	20.18 ^k	19.05 ^j	15.13 ^f	29.8 ^k	1.0591 ^{mnopq}	4.56 ⁱ	55.87 ^z	39.23 ^{ba}	1.42 ^{cdefgh}
HB-9912 (N₈₀)	3.51 ^{bcdef}	0.5 ^{bdeg}	9.7 ^{hg}	19.62 ^p	18.36 ^{po}	11.71 ^y	36 ^w	1.069 ^{klmnopq}	4.01 ^{ef}	57.41 ^w	37.98 ^{bcdef}	1.51 ^{a^bcdefgh}
HB-9916 (N₀)	3.23 ^{bcdefghij}	0.53 ^{bcd}	9.14 ^e	19.23 ^r	16.12 ^y	12.75 ^{qr}	38.96 ^p	1.193 ^{defghi}	3.76 ^z	60.85 ^h	35.35 ^{bcdefgh}	1.72 ^{abcdefg}
HB-9916 (N₈₀)	1.6 ^{pqor}	0.36 ^{klmn}	7.74 ^{uv}	15.73 ^e	13.31 ^j	11.24 ^a	41.21 ^f	1.182 ^{defghi}	1.96 ^j	60.19 ^k	29.04 ^{pqor}	2.07 ^{abcd}
IC212031 (N₀)	3.07 ^{defhijkl}	0.37 ^{ijklmn}	9.08 ^l	19.42 ^q	18.48 ^m	14.12 ^k	35.28 ^x	1.051 ^{mnopq}	3.44 ^{pq}	58.48 ^s	37.9 ^{cdefghijk}	1.54 ^{bcdefghi}
IC212031 (N₈₀)	3.13 ^{efghid}	0.3 ^{nop}	8.89 ^m	20.64 ^h	20.49 ^b	15.83 ^c	30.55 ⁱ	1.007 ^{pqrst}	3.43 ^p	55.27 ^b	41.13 ^{defgh}	1.34 ^{defghi}
NATP-124 (N₀)	2.93 ^{cdefghijk}	0.11 ^t	7.79 ^{ut}	18.71 ^t	19.47 ^g	12.7 ^{sr}	37.22 ^r	0.961 ^{rstuv}	3.041 ^m	57.71 ^u	38.18 ^{defghijklm}	1.51 ^{abcdefg}
NATP-124 (N₈₀)	3.18 ^{defghi}	0.36 ^{klmn}	6.98 ^x	17.60 ^a	17.08 ^v	10.24 ^d	44.43 ^a	1.030 ^{opqrs}	3.54 ^p	61.65 ^e	34.68 ^{defghij}	1.7 ^{8abc}
QM-16 (N₀)	2.75 ^{efghijklmn}	0.44 ^{efghi}	10.95 ^d	22.70 ^b	18.48 ^m	12.8 ^{qp}	31.26 ^h	1.228 ^{bfge}	3.19 ^d	55.01 ^{pe}	41.18 ^{efghijklmn}	1.34 ^{efghi}
QM-16 (N₈₀)	3.42 ^{bcdefg}	0.40 ^{ghijkl}	10.49 ^e	22.38 ^c	19.65 ^e	14.78 ^h	28.26 ^l	1.139 ^{hijkl}	3.82 ^b	53.53 ^h	42.03 ^{bcdefg}	1.27 ^{bcdefghi}

RGN-55(N₀)	1.21 ^{qr}	0.16 ^{ts}	6.22 ^b	21.04 ^f	22.61 ^a	17.3 ^a	31.44 ^g	0.931 ^{uv}	1.37 ^a	54.96 ^e	43.65 ^{qr}	1.26 ^{efghi}
RGN-55(N₈₀)	3.61 ^{bcdef}	0.46 ^{defgh}	8.18 ^q	20.04 ^m	15.93 ^a	11.05 ^b	40.31 ⁱ	1.258 ^{dc}	4.07 ^w	59.54 ^{nm}	35.97 ^{bcdef}	1.66 ^{efghi}
NRCHB-101 (N₀)	2.54 ^{ghijklmn} _o	0.42 ^{efghij} _{kl}	9.5 ^j	17.92 ^{zy}	19.44 ^g	14.07 ^k	34.84 ^z	0.922 ^{uv}	2.96 ^t	58.41 ^s	37.36 ^{ghijklmno}	1.56 ^{bcdefghi}
NRCHB-101 (N₈₀)	3.24 ^{bcdefghij}	0.41 ^{fghijkl} _m	8.12 ^{sr}	18.14 ^w	17.73 ^r	12.89 ^{op}	39.28 ^o	1.023 ^{opqrs}	3.65 ^x	60.29 ^j	35.87 ^{bcdefghi}	1.68 ^{bcdefgh}
NRCDR-2 (N₀)	1.52 ^{pqr}	0.45 ^{defghi}	11.38 ^b	18.34 ^v	15.63 ^{dc}	8.87 ^g	43.77 ^{cb}	1.173 ^{efghij}	1.97 ^e	64.02 ^a	33.97 ^{poq}	1.88 ^{ab}
NRCDR-2 (N₈₀)	3.88 ^{abcd}	0.4 ^{ahijklm}	11.32 ^{bc}	20.42 ^j	17.46 ^{ts}	12.33 ^t	33.9 ^c	1.170 ^{fghij}	4.28 ^q	57.55 ^v	37.88 ^{abcd}	1.52 ^{bcdefgh}
NRCHB506 (N₀)	4.13 ^{abc}	0.98 ^a	9.5 ^{l j}	15.20 ^f	15.83 ^b	12.62 ^s	37.09 ^s	0.960 ^{stuv}	5.11 ^h	59.22 ^q	31.03 ^{abc}	1.91 ^{abcdef}
NRCHB506 (N₈₀)	1.7 ^{defghijklm} _{nopqr}	0.57 ^b	7.49 ^w	15.74 ^e	15.16 ^f	10.77 ^c	36.52 ^u	1.038 ^{nopqrs}	2.27 ⁱ	54.78 ^f	30.9 ^{opqr}	1.77 ^{abcdef}
Pusa Jai kisan (N₀)	2.93 ^{defghijk}	0.23 ^{pqr}	8.57 ^o	19.99 ^m	18.96 ^k	14.36 ^j	34.48 ^a	1.054 ^{klmnopq}	3.16 ^j	57.41 ^w	38.95 ^{bcdefghijklm}	1.47 ^{hig}
Pusa Jai kisan (N₈₀)	3.37 ^{bcdefghq}	0.47 ^{defg}	7.74 ^{uv}	18.75 ^{ts}	17.53 ^s	11.49 ^z	40.52 ^h	1.070 ^{klmnopq}	3.84 ^v	59.75 ^l	36.28 ^{dbcdefgh}	1.65 ^{bcdefghi}
NDRE-7 (N₀)	1.18 ^r	0.23 ^{pqr}	6.45 ^a	16.69 ^c	18.35 ^p	13.22 ^{ml}	43.84 ^b	0.91 ^{uv}	1.41 ^a	63.51 ^b	35.04 ^R	1.81 ^{bcdefg}
NDRE-7 (N₈₀)	3.34 ^{bcdefgh}	0.98 ^a	9.47 ^j	20.37 ^j	14.24 ^h	8.45 ^h	36.24 ^v	1.430 ^a	4.32 ^d	54.16 ^g	34.61 ^{bcdefgh}	1.56 ^{bcdefg}
RL-1359 (N₀)	3.25 ^{bcdefghi}	0.13 ^{ts}	7.64 ^v	17.69 ^a	17.25 ^u	11.94 ^{wv}	42.07 ^e	1.026 ^{opqrs}	3.38 ^b	61.65 ^e	34.94 ^{bcdefgh}	1.76 ^{abcde}
RL-1359 (N₈₀)	3.62 ^{bcdef}	0.43 ^{efghij} _k	8.84 ^m	18.46 ^u	18.59 ^l	12.9 ^{on}	36.72 ^t	0.993 ^{qrst}	4.05 ^u	58.46 ^s	37.05 ^{fbec}	1.58 ^{abcdefgh}
BEC-16 (N₀)	2.7 ^{fghijklmn}	0.38 ^{hijklm} _n	9.56 ^{ij}	18.16 ^w	19.44 ^g	9.91 ^e	39.81 ^l	0.934 ^{uv}	3.08 ^s	59.28 ^{qp}	37.6 ^{fghijklm}	1.58 ^{bcdefg}
BEC-16 (N₈₀)	3.37 ^{bcdefgh}	0.4 ^{ghijklm}	9.26 ^k	19.32 ^r	18.44 ^{nmo}	15.95 ^b	33.22 ^d	1.048 ^{nopqr}	3.77 ^r	58.43 ^s	37.76 ^{bcdefg}	1.55 ^{bcdefg}

T comparison lines for least Squares means of genotypes*nitrogen
LS- means with same letters are not significantly different

The ω 3 levels were found to range between 13.62 % (DHR-991) and 22.61 % (RGN-55) under N0 while, under N80 it ranged between 13.31 % (HB-9916) to 20.49 % (IC212031). These levels are high enough (>12%) to be considered as a source for commercial extraction of ω 3 according to Zambiazzi et al. (2007) who reported the bench mark to be 12 %. It has been reported earlier that a dose of 60 kg/ha nitrogen fertilizer was optimum to enhance ω 3 in Indian mustard (Joshi et al., 1998) but it is contradictory to our findings where we found a variation within the same species at N80. 29 % of the genotypes showed an increase in ω 3 by 4.09 % (DRMR-IJ-31) to 27.77 % (DHR-991) over N0 and 71 % of the genotypes reported reduction by 0.06 % (Maya) to 29.54 % (RGN-55) over N0. Previous works have reported the effect of nitrogen only in one genotype and comparative studies have not been explored.

Generally *B. juncea* is known for having all the fatty acids in the ideal range for e.g. high levels of MUFA, low SFA, moderate PUFA, ω 6: ω 3 ratio of 1.2:1 (Chauhan et al., 2010). Under N0, the ω 6: ω 3 ranged from 0.91 (NDRE-7) to 1.23 (QM-16) and under N80, it ranged from 0.88 (DRMR-IJ31) to 1.37 (GM-2) (Table 3). According to various medical researchers (Simopoulos, 2012) there is so no ideal ω 6: ω 3 ratio that would prevent chronological diseases. The optimal ratios vary with the disease under consideration and according to their documentation, ratio beyond 10:1 was observed to have deleterious effects on health (Simopoulos, 2012). However, a low level ω 6: ω 3 is preferable. It was observed that under N80, 50 % of genotypes had increased the ratio by 0.88 % (HB-9912) to 57.28 % (NDRE-7) over N0. While, 50 % genotypes showed reduction in the ratio by 0.33 % (NRCDR-2) to 11.04 % (Maya) over N0.

The MUFA: PUFA ratio depicting oil stability index (OSI) ranged from 1.23 (HB-9902) to 2.20 (DHR-991) under N0 and from 1.27 (QM-16) to 2.07 (HB-9916) in N80 (Table 3). Most of the genotypes (96%) failed to reach the ideal range of 2 (Chauhan et al., 2010). However, under N80 about 63 % of genotypes improved in OSI by 1.81 % (GM-2) to 39.54 % (HB9902) while, the rest 37 % of the genotypes showed reduction by 0.78 % (DRMR-IJ-31) to 20.90 % (DHR-991) over N0 application.

Nutritionally undesirable erucic acid was found to exceed the recommended percentage of 2% at both N0 and N80 application (Table 3). Erucic acid, under N80 observed an increase in 63 % of genotypes by 0.9 % (DHR-991) to 38.31% (HB9902) and 37 % of genotypes showed decrease by 1.14 % (DRMR-IJ 31) to 22.5 % (NRCDR-2) over N0. The variations in nitrogen levels hence contribute to the variation in oil quality as seen in case of composition of individual fatty acids.

Conclusion

This study showed N application can bring about changes in oil and seed meal quality. 63 % of the genotypes showed increase in total N content in seeds by 3 % (DRMR-IJ-31) to 34.0 % (NRCDR-2) and rest 37 % genotypes showed reduction in seed N content by 5.6 % (HB9916) to 20.2 % (EC399300) over control. These genotypes also have positive correlation to the total soluble protein content. However, some genotypes showed an inverse relationship between N and soluble protein which could be due to the flux in N to biosynthesis of other carbon compounds as observed by an increase in the case of ascorbic

acid content and phenol content. Ascorbic acid and total phenolic content showed a positive correlation with TAC ($r=0.316$, $p0.028$). Crude fiber and total soluble sugars were observed to decrease under N80. These reductions are desirable for increasing the digestibility and palatability of meal. Seed storage protein profiled under N80 revealed differences in the banding patterns especially in α and β chain of the cruciferin which determines the digestibility and quality of seed meal. Fatty acid profiling showed reduction of SFA and $\omega6:\omega3$ in 50 % of genotypes and 50 % of genotypes showed increase in their ratios after N treatment. 63% of genotypes showed improvement in OSI after N80 treatment. N80 application caused 63 % of the Indian mustard genotypes to increase in erucic acid. Our findings showed application of nitrogen even with recommended dose can influence biochemical changes within same species. All in all our observations suggest that nitrogen fertilizers can play a role in enhancing nutritive status of oilseed Brassica when applied in appropriate doses. Information on the effect of nitrogen application on nutritional status of oil and meal will be quite useful in quality improvement programmes.

Acknowledgements. The authors are grateful to Indian Council of Agricultural Research for providing all the necessary facilities and funding (DRMR-B7) to carry out this work. The authors are grateful to plant breeders for sharing their materials used in this study.

Conflict of interest statement. The authors do not have any conflict of interest.

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