

Capsaicinoid content, pungency and antioxidant potential of Himalayan hot pepper

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ABSTRACT

Hot pepper (*Capsicum annuum* L.) is one of the most important vegetable crops grown worldwide due to its culinary values and potential pharmaceutical applications. Capsaicinoids in hot pepper are the secondary metabolites responsible for its pungency. Since capsaicinoid content depends on the environment, betteradapted genotypes with higher capsaicin content are always required. In the present study, we performed a biochemical screening of twenty-two North-Western Himalayan adapted hot pepper breeding lines. We determined their capsaicin content, dihydrocapsaicin content, total phenolic content, pungency, and antioxidant activity. The HPLC analysis quantified 1000 to 4010 μ g/g capsaicin and 489 to 1863 μ g/g of dihydrocapsaicin, translating to pungency of 16000 to 64160 SHU. The germplasm expressed 28.44% to 78.61% DPPH free radical scavenging activity, 121.29 to 394.294 μ M Fe²⁺/g dry weight (DW) ferric reducing antioxidant power (FRAP) and 4.31 to 8.56 mg/g GAE of total phenolics. Our results identified CITH-HP-91-13 and CITH-HP-92-13 as the breeding lines richest in capsaicinoids and antioxidant activity. The germplasm has been grouped into five clusters based on standardized squared Euclidean distance using Ward's hierarchical clustering and separation based on PCA biplot. Our study revealed significant differences among breeding lines for tested parameters, which indicate the possibility of exploiting them for quality improvement programme of Kashmiri pepper.

Keywords: Capsicum annuum L., Hot pepper, Capsaicinoids, Pungency, Antioxidant potential, Phenolics

INTRODUCTION

Hot pepper or chilli pepper (Capsicum annuum L.) is extensively used as flavouring and colouring spice across the world. Besides culinary value, hot pepper has pharmaceutical application due to its analgesic, neurological, gastroprotective, antiobesity, anti-diabetic and dietetic properties (Basith et al., 2). In modern times, chilli pepper is used in the pharmaceutical industry and as a traditional ethnomedicine (Basith et al., 2). The North-Western Himalayan region is popular for Kashmiri chilli cultivation. Kashmiri chilli is a group of signature cultivars of Kashmiri origin with low pungency and high oleoresins. It imparts a bright red hue to local cuisines without adding too much piquancy. Nonetheless, highly pungent genotypes are also available in the region.

Culinary, disease therapeutic qualities of hot pepper are functions of capsaicinoids whose stable specific levels in a cultivar are imperative for its acceptance by food and pharmaceutical industries. However, environmental conditions significantly influence capsaicinoid content (Loizzo *et al.*, 9). This was also confirmed by our earlier study wherein North-Eastern chilli cultivar '*Bhut Jolokia*' accessions known to possess one of the highest capsaicin contents *i.e.*, ~1 million SHUs (Bosland and Baral, 3), expressed only about one-tenth of Scoville Heat Units (SHU) under North-Western Himalayan climate of Kashmir valley (Indrabi, 7). Hence, selecting highly adapted genotypes stable for capsaicinoid content in a particular geographical area is imperative. These highly adapted hot pepper genotypes with stable capsaicinoids can serve as sources of bioactive compounds and pungency and as breeding material for elite varieties development.

Although there are ample research reports on Capsicum annuum L. for capsaicinoid content and antioxidant potential but genotypes from North-Western Himalayas have not yet been evaluated (Dubey et al., 4). Several hot pepper breeding lines of North-Western Himalayan adaptation are being maintained at ICAR-Central Institute of Temperate Horticulture, Srinagar. This scenario of chilli germplasm availability calls for its biochemical evaluation to authenticate the variation in pungency levels and phytochemical efficacy. The present study was formulated to quantify the variation for biochemical properties such as capsaicin, dihydrocapsaicin content, capsaicinoid-derived antioxidant activity and pungency among Northwestern Himalayan hot pepper breeding lines and

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identification of breeding lines with high pungency in the Kashmir region.

MATERIALS AND METHODS

Twenty-two hot pepper breeding lines developed from collections made from various Kashmir Valley regions were evaluated. These lines were morphologically uniform and stable for economic characteristics like yield and fruit traits. Fresh fruit samples were collected from the plants and dried in the shade. During experimentation and chemical analyses, analytical-grade reagents were used. Extraction was done in a Soxhlet extractor using HPLC-grade methanol as solvent, and a final concentration of 100 mg/ml was prepared.

HPLC analysis for capsaicinoids was conducted in Shimadzu HPLC (Kyoto, Japan). The solution was filtered through 0.2 µm syringe filters. HPLC separation was carried out with an injection volume of 20 µL, a flow rate of 1 ml min⁻¹ with 35-40 minutes of run time. The analyses were done in three replications for each test sample. Both capsaicin and dihydrocapsaicin were detected at 210 nm. Chromatographic separations were performed on C 18 (250 mm × 4.6 mm), 5-µm column using a solvent system of 70% acetonitrile and 30% methanol in an isocratic mode. Prior analysis, the mobile phase was filtered through a 0.45 µm membrane filter (Millipore, Bedford, MA, USA). Class WP v6.1 (Shimadzu, Japan) software package was used for data acquisition and processing. The concentrations were measured with the help of respective peak areas of standards at particular retention time against the concentration and expressed as mg/g of dry sample. Under optimal isocratic conditions, capsaicin and dihydrocapsaicin were separated within 1.74 minutes and 2.3 minutes, respectively. The Scoville Heat Units (SHU) parameter indicates the pungency level. We categorized the pungency of the analysed samples as per Scoville (17).

Capsaicinoid-derived antioxidant potential by means of DPPH and FRAP assays was determined by methods described by Shotorbani *et al.* (18). Fruit samples were shade dried and powdered vigorously in a mortar and pestle. Three grams of fine powder were dissolved in measured levels of methanol. The Whatman filter paper was used to filter the extract. To avoid deterioration of antioxidant activity in the samples, extracts were processed immediately.

The 1,1-Diphenyl-2-picryl-hydrazyl (DPPH) free radical scavenging assay was done using the procedure described by Xu *et al.* (20) with slight modifications. The 15 μ g/ml working solutions of each sample extract were prepared in methanol. To prepare the standard curve, ascorbic acid at 1, 2, 4,

6, 8, 10 and 15 µg/ml concentrations were used. The 1 ml of each sample extract was combined with 1 ml of 0.002% DPPH solution. The solution was mixed well and incubated in the dark for 30 minutes. The absorbance was recorded at 517 nm wavelength. The analyses were done in three replications. Any observed reduction in the absorbance of DPPH caused by the addition of test samples compared with control became the basis for calculating antioxidant activity in terms of percentage inhibition of DPPH radical (% IP). Per cent DPPH inhibition activity was estimated as, where is the absorbance of the control and is the absorbance of the sample. Methanol and DPPH in the ratio of 1:1 mixture was used as blank and standard (ascorbic acid) and DPPH mixture in the ratio of 1:1 was treated as control. The concentration of test sample and standard solution was 15 µg/ml.

The antioxidant capacity of samples was measured in terms of ferric ion reducing antioxidant power (FRAP) also. The FRAP reagent was prepared by mixing sodium acetate buffer (300 mmol/L at pH 3.6), 2,4,6-tripyridyl-s-triazine (TPTZ) solution in 40 mmol/L HCI (10 mmol/L) and FeCl, solution (20 mmol/L) in the ratio of 10:1:1 (v/v), respectively. The FRAP reagent was pre-warmed at 37 °C, and 1.8 ml was used for 200 µl of sample extracts. After 40 minutes, the absorbance of the solution was taken at 593 nm. The calibration curve was prepared based on FeSO, solutions at 0, 40, 80, 160, 320, and 640 µmol/L concentrations. The observations were determined with subsequent formulae determined on calibration curve: 0.998, where stands for absorbance and is µM Fe²⁺/g fresh weight. Antioxidant activities of test sample extracts were finally depicted as µmol Fe2+/g DW of the sample.

The potential of phenolic compounds as antioxidants were estimated by the modified Folin-Ciocalteau method (Omoruyi *et al.*, 11). Five ml of Folin-Ciocalteau reagent (previously diluted with distilled water 1:10 (v/v), and 4 ml (75% w/v) of sodium carbonate (Na₂CO₃) were combined with1 ml of test extract (1 mg/ml) and vortexed for 30 minutes for colour development. The absorbance of the extracts was recorded at 765 nm. A standard curve was made using gallic acid at 0 to 400 mg/L concentrations. Finally, concentrations were expressed as milligram gallic acid equivalent (mg GAE/g DW of sample extract). The formula used for the purpose was 0.997, where is absorbance and is gallic acid equivalent in mg/g.

The existence of significant differences among extracts for capsaicinoids, pungency, antioxidant potential and total phenolics were analysed. For the purpose, one-way analysis of variance (ANOVA) using Duncan's mean range test (DMRT) was carried out. All statistical tests were done using 'SAS[®] Enterprise Guide 4.2' software at 5% significance level. Correlation study was performed using Pearson's test. Principal component analysis (PCA) and cluster analysis were performed using 'SAS[®] Enterprise Guide 4.2 software.

RESULTS AND DISCUSSION

Hot pepper is rich in capsaicin and dihydrocapsaicin, which are natural antioxidants (Ghasemnezhad et al., 5). These capsaicinoids are also determining factors in establishing the commercial quality of hot pepper. In this study, there was significant variability in capsaicin and dihydrocapsaicin contents among tested lines. The capsaicin content ranged from 1000 µg/g (in SEL-1052-11) to 4010 µg/g (in CITH-HP-92-13) with an average of 2434.23 µg/g, while dihydrocapsaicin varied from 489 to 1863 µg/g with an average of 1143.82 µg/g. The highest capsaicin, as well as dihydrocapsaicin contents, was found in line CITH-HP-92-13. Based on capsaicinoid content and SHU were estimated to determine the pungency of germplasm. The highest pungency of 64160 SHU was recorded in CITH-HP-92-13 and the lowest pungency of 16000 SHU was recorded in SEL-1052-11. The average pungency of the germplasm was 38947.64 SHU. According to the Scoville Organoleptic Scale (Scoville, 17), out of the 22 lines, 19 came out to be highly pungent (> 25000 SHU) and 4 moderately pungent (3000-25000 SHU) (Table 1). It means that more than 86% of our tested germplasm is highly pungent, thus a promising genetic stock usable in pungent chilli breeding.

Our study has revealed considerably higher capsaicinoid content and pungency in NW Himalayan adapted chilli germplasm compared to many other germplasms studied worldwide. For example, Othman et al. (13) studied commercial pepper varieties of Riyadh city (KSA), and the mean capsaicin and dihydrocapsaicin contents were 1320 µg/g and 830 µg/g, respectively. Landraces of the Republic of Benin expressed capsaicin content range of 765 to 3070 µg/g (Orobiyi et al., 12). In another study, the local commercial chilli powder products of Visakhapatnam, AP, had capsaicin content of 800 to 1300 µg/g. However, our tested germplasm lines have capsaicin content ranging from 1000 to 4010 µg/g with average of 2434.23 µg/g and dihydrocapsaicin content ranging from 489 to 1863 μ g/g with an average of 1143.82 μ g/g, which are much higher than found in all above studies.

Our germplasm, however, has generally lower capsaicin content than *Capsicum chinense* cv. Habanero, one of the hottest chillies in the world, having up to 200000 SHU (Johnson *et al.*, 8), as

Table 1. Capsaicin and dihydrocapsaicin contents and pungency of hot pepper breeding lines.

S. Line Cap- saicin Dihydro- capsaicin Pungen (SHU) No. saicin capsaicin (SHU) (µg/g) (µg/g) (µg/g) 1) CITH-HP-92-13 4010 ^A 1863 ^A 64160 2) CITH-HP-91-13 3573 ^B 1522 ^B 57168 3) SEL-1065-E 3300 ^C 1450 ^C 52800 4) CITH-HP-1154 3000 ^D 1322 ^D 48000 5) SEL-836-1-2 2937 ^D 1311 ^D 46992
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5) SEL-836-1-2 2937 ^D 1311 ^D 46992
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6) CITH-HP-85-13 2700 ^E 1311 ^D 43200
7) CITH-HP-111-1 2673 ^E 1298 ^D 42768
8) CITH-HP-1154-2-1 2623 ^{EF} 1260 ^{DE} 41968
9) SEL-1050 2601 ^{EF} 1290 ^D 41616 ^D
10) CITH-HP-1154-3-1 2570 EFG 1243 DE 41120 E
11) SEL-136-1-2 2501 ^{FGH} 1200 ^{EF} 40016 ^F
12) CITH-HP-56-13 2493 ^{FGH} 1190 ^{EF} 39888 ^F
13) CITH-HP-111 2483 ^{FGHI} 1185 ^{EF} 39728 ^{FI}
14) SEL-1011-2 (E) 2423 ^{GHI} 1123 ^{FG} 38768 ^G
15) SEL-1055-11 2390 ^{IH} 1150 ^{FG} 38240
16) CITH-HP-1154-1 2323 ^{JJ} 1101 ^G 37168
17) CITH-HP-114-13 2200 ^J 1023 ^H 35200
18) SEL-89 1800 ^K 847 ⁱ 28800
19) KA2-SEL-1 1500 ^L 723 ^J 24000
20) CITH-HP-42-13 1250 ^M 640 ^K 20000 ^H
21) AL-4 1203 ^M 623 ^K 19248
22) SEL-1052-11 1000 ^N 489 ^L 16000
Mean 2434.23 1143.82 38947.6

(Note: Different letters indicate significant differences between means in the column as per Duncan's test)

suggested by a study by Palma-Orozco *et al.* (14). The average capsaicin content of Habanero fruits dried at 40 to 70 °C temperature was 9630 μ g/g. In another study from Czechoslovakia, various Habanero varieties were found to have 1128 to 8215 μ g/g of capsaicin, averaging 3900 μ g/g (Popelka *et al.*, 16). However, in one study on Australian habanero cultivars, capsaicin contents ranged from 1474 to 3916 μ g/g (Johnson *et al.*, 8). Comparing 1000 μ g/g to 4010 μ g/g capsaicin content of our germplasm, it is suggestive that NW Himalayan chilli may be comparable to or more pungent than Australian Habaneros. Hence, extensive studies involving larger sample sizes are needed. Moreover, agroecological conditions may also have a role to play.

Following capsaicinoid concentrations and pungency estimates, we compared the antioxidant

potential of the germplasm via DPPH and FRAP antioxidant assays. DPPH is a very stable free radical and is extensively exploited to estimate the antioxidant potential of biological extracts. In the present study, the free radical scavenging potential of investigated lines estimated with DPPH assay revealed significant differences. The per cent inhibition of DPPH ranged from 28.44% in CITH-HP-91-13 to 78.61% in Sel-836-1-2 (Table 2). Fifteen lines comprising 68% of the tested germplasm elicited more than 50 per cent inhibition. The FRAP of tested germplasm ranged from 121.29 to 394.29 µM Fe²⁺/g DW. Based on the highest FRAP value obtained, SEL-1050 is superior among all lines. It is worth noting that DPPH per cent inhibition and FRAP values did not correspond for the same germplasm. Although DPPH per cent inhibition was highest in SEL-836-1-2 (78.61), the FRAP was highest in SEL-1050 (394.29).

The total phenolic content of twenty-two extracts ranged from 4.31 to 8.56 mg GA/g DW (Table 2).

SEL-1065-E expressed the highest total phenolics (8.56mg GA/g DW). The phenolic contents of chilli lines obtained in this study are similar to those observed in many other studies involving various species of *Capsicum* (Alvarez-Parrilla *et al.*, 1; Sora *et al.*, 19; Hamed *et al.*, 6). The absorbance obtained in test samples over a concentration range of 20-90 mg/L was linear (with 0.997.

To establish the relative competence of different assays and methods of biochemical estimations in chilli, we constructed a correlation matrix for DPPH and FRAP assays and total phenolic, capsaicin and dihydrocapsaicin contents obtained from HPLC (Table 3). Correlation coefficient is a statistic developed for determining the degree of association among parameters. Our results revealed a significant positive correlation among antioxidant assays, total phenolic, and capsaicinoid contents. This observation has strengthened the authenticity of these methods. The positive correlation (P<0.05) among TP-DPPH (r = 0.3098) was significant, whereas it was highly

S. No.	Line	DPPH	Percent Inhibition	FRAP (µM fe2+/g DW)	Total phenols (mg/g GAE)
1)	CITH-HP-92-13	11.51 [⊧]	64.28	122.00 ^L	4.90 ^{JK}
2)	CITH-HP-91-13	1.35 ^к	28.44	302.24 ^{FG}	6.08 FGH
3)	SEL-1065-E	12.67 ^Ĕ	68.37	274.94 ^H	8.56 ^A
4)	CITH-HP-1154	11.89⁵	65.64	317.88 ^{EF}	6.48 ^{EF}
5)	SEL-836-1-2	15.57 ^A	78.61	332.24 ^{DE}	7.38 ^c
6)	CITH-HP-85-13	9.51 ^G	57.22	300.65 ^{FG}	6.12 ^{FGH}
7)	CITH-HP-111-1	3.28 ^J	35.27	341.88 ^{CD}	5.93 ^{GHI}
8)	CITH-HP-1154-2-1	11.54⁵	64.39	315.53 ^{EFG}	4.97 ^J
9)	SEL-1050	14.96 ^{AB}	76.45	394.294 ^A	6.88 ^D
10)	CITH-HP-1154-3-1	3.60 ^J	36.41	363.588 [₿]	8.06 ^B
11)	SEL-136-1-2	8.70 ^H	54.38	121.29 ^L	4.92 ^{JK}
12)	CITH-HP-56-13	13.41 ^D	70.99	356.65 ^{BC}	6.58 ^{DE}
13)	CITH-HP-111	4.86 ⁱ	40.84	314.00 ^{FG}	6.56 DE
14)	SEL-1011-2 (E)	3.25 ^J	35.15	170.00 ^G	5.16 ^J
15)	SEL-1055-11	15.44 [^]	78.16	298.59 ^G	7.53 ^c
16)	CITH-HP-1154-1	14.09 ^c	73.38	357.706 ^{BC}	6.19 ^{EFGH}
17)	CITH-HP-114-13	14.73 [₿]	75.65	346.94 ^{BCD}	5.68 ¹
18)	SEL-89	3.18 ^J	34.93	160.76 ^{јк}	4.55∟
19)	KA2-SEL-1	4.67 '	40.16	148.35 ^ĸ	4.31 [∟]
20)	CITH-HP-42-13	11.28 [⊧]	63.48	229.41 ¹	4.78 ^{JK}
21)	AL-4	9.38 ^G	56.77	121.88 [∟]	5.89 ^{HI}
22)	SEL-1052-11	9.02 ^{GH}	55.52	223.06 ¹	4.36 ^L
	Mean	8.93	57.02	268.81	5.99

Table 2. Antioxidant potential and total phenol content of twenty-two hot pepper breeding lines.

(Note: Different letters indicate significant differences between means in the column as per Duncan's test)

	Total phenols	FRAP	Capsaicin	Dihydro- capsaicin
DPPH	0.3098*	0.3924	0.0782	0.0994
Total phenols		0.5464**	0.4450	0.4420
FRAP			0.2790**	0.2940**
Capsaicin				0.9897

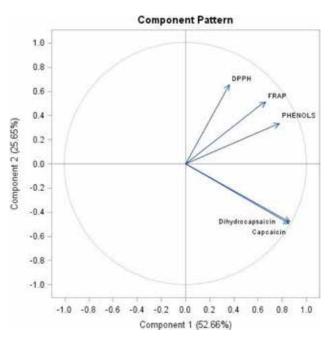
 Table 3. Correlation matrix for five biochemical parameters analysed.

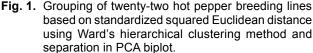
(* Significant at p< 0.05, ** Significant at p< 0.01)

significant (P <0.01) among TP-FRAP (r = 0.5464), capsaicin–FRAP (r = 0.2790) and dihydrocapsaicin-FRAP (r = 0.2940). Observations on the same line have been made in studies on diverse *Capsicum annuum* L., germplasms (Nunez-Ramirez *et al.*, 10).

Principal component analysis (PCA) is done to partition variation and highlight strong patterns in a dataset. We performed PCA using two antioxidant assays (DPPH and FRAP) and three antioxidants (capsaicin, dihydrocapsaicin and total phenolics) which showed that the first two components (PC1 and PC2) explained 78.31% of the total variation. First principal component (PC1) accounted for 52.66% of the total variation and the variables responsible for separation along the PC1 included DPPH (0.22), TP (0.47), capsaicin (0.52) and FRAP (0.40) and DiCAP (0.52). The second principal component (PC2) explained 25.65% of the total variation. The accessions were separated by DPPH (0.58) (Fig. 1). Hierarchical cluster analysis grouped 22 genotypes into two main clusters with cluster-I represented by five genotypes (SEL-89, KA2-Sel-1, SEL-1052-11, AL-4 and CITH-HP-42-13) and cluster-II represented by sixteen genotypes (Fig. 2). Genotype CITH-HP-92-13 does not form the part of any cluster being most superior genotype with respect to capsaicin and dihydrocapsaicin contents. The contribution of individual genotypes to the antioxidant grouping and the relationship between the clusters was assessed by plotting PC1, PC2 and PC3. Expectedly, both PCA and cluster analysis were found equally effective in grouping the breeding lines based on their antioxidant contents (Patras et al., 15).

In conclusion, it was observed that chilli, *Capsicum annuum* L., and breeding lines developed from collections made from North-Western Himalayas have high capsaicin and dihydrocapsaicin contents and antioxidant potential. Two lines *i.e.*, CITH-HP-92-13 and CITH-91-13, expressed the highest values for capsaicin content, dihydrocapsaicin content and FRAP activity. Clustering and PCA for these traits revealed significant differences among lines, which indicates the possibility of exploiting





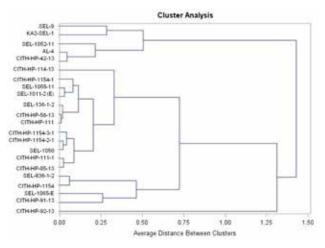


Fig. 2. Dendrogram based on capsaicin, dihydrocapsaicin, total phenols, FRAP and DPPH scavenging activity values obtained for twenty-two hot pepper breeding lines.

them for quality breeding for pharmaceutical and ethnomedical applications. Lines exhibiting high pungency can be used to breed for cuisine-specific targets like high pungency-high oleoresin, for example, combining the trait of high oleoresin (but low pungency) content of Kashmiri mirch with these high pungency breeding lines. These lines can also be used to decipher molecular mechanism and enzyme kinetics of rate limiting factor in capsaicinoids biosynthesis. Most of the genotypes evaluated in this study were highly pungent because of high capsaicin and dihydrocapsaicin content and high content is strongly correlated with significant FRAP activity. Higher FRAP activity of chilli genotypes is due to scavenging potential of capsaicin and dihydrocapsaicin.

AUTHORS' CONTRIBUTION

Conceptualization of research (JIM, DBS, GM, OCS); Designing of the experiments (JIM, DBS); Contribution of experimental materials (DBS, GM); Execution of field/lab experiments and data collection (MAS, AS, RAS); Analysis of data and interpretation (JIM, GM, VD, SY); Preparation of the manuscript (JIM, GM, VD).

DECLARATION

The authors declare that there is no conflict of interest.

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