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Interrelationship Among Lipoxygenase Isozymes, Polyunsaturated Fatty Acids and Trypsin Inhibitor During Seed Development in Soybean

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ABSTRACT

Lipoxygenase isozymes, polyunsaturated fatty acids and trypsin inhibitor content were determined in four selected soybean varieties during seed development. A genotypic variation was observed for accumulation pattern of lipoxygenase isozymes and trypsin inhibitor, besides, significant levels of trypsin inhibitor in the early stage of soybean seed development. Linolenic acid was found to be maximum at 30 days after flowering and thereafter decreased continuously at varying rate in all the genotypes till maturity. Lipoxygenase isozymes levels were comparatively low in the early stage of development when linolenic acid was very high. A significant positive correlation of trypsin inhibitor content with lipoxygenase I as observed in present studies suggests a coordinated expression of these biological components during seed development in soybean.

Key words: Soybean, lipoxygenase isozymes, polyunsaturated fatty acids, trypsin inhibitor, seed development

Lipoxygenase (Linoleate: oxygen oxidoreductase, EC 1.13.11.12) and trypsin inhibitor are considered undesirable components in soybean seeds (Rackis *et al.* 1979, Anderson-Hafferman *et al.* 1992). In general, normal soybean seed lipoxygenase (Lox) exists in three isozymic forms namely Lox-I, Lox II and Lox III (Axelrod *et al.* 1981) and constitutes about 1-2 percent of the proteins present in dry seeds (Kitamura 1984). These isozymes catalyse the hydroperoxidation of polyunsaturated fatty

acids (PUFA), linoleic and linolenic acid, containing *cis cis 1,4 pentadiene* moiety and have been categorized into two classes. Class I lipoxygenase (Lox-I) is characterized by high pH optima of around 9.0 and formation of large amounts of 13-hydroperoxides while class II lipoxygenase (II+III) show pH optima of around 7.0 and formation of equal amounts of 9 and 13-hydroperoxides. The hydroperoxidation reaction catalysed by the lipoxygenase isozymes lead to the formation of volatile hexanal compounds

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nsible for the development of products, the prime deterrent in stability by Indian populace. Fur-an seed lipoxygenases have also to cause seed deterioration . During storage and transport, mechanical or bacterial damage, membrane bound and storage lipids are favorable substrates for to catalyse oxidation. Free radicals in the process, set the chain reaction of membrane lipids, ultimately membrane integrity (Vick and 87).

Inhibitor (TI), the protease inhibitor, is responsible for reducing proteins by inhibiting tryptic activity. TI is heat labile, the heat treatment reduces the much-valued proteins and cause loss of essential soy proteins (Rios-Iriarte and

genotypic variability for LoxA and TI has been reported in us (Marczy *et al.* 1995, Yang *et al.* 2001). Recently, variability in characters has been reported in among Indian genotypes (Kumar 2; 2003; Rani *et al.* 2004). The developmental expression of and their substrates, PUFA, and growth is important to view the of immature soybean pods for option and to understand the in among these biological components development. However, the on the expression and inter-ological components during seed e few and scattered (Yao *et al.* *et al.* 1986, Liu and Markakis e present study was undertaken the interrelationship between PUFA and TI in developing seed ndian soybean varieties.

MATERIALS AND METHODS

Four commercial varieties of Indian soybean viz. JS 335, Pb 1, NRC 37 and Shilajeet were sown in 3 meters rows with a spacing of 45 cm in the experimental fields of National Research Centre for Soybean (ICAR), Indore on 27th June 2003. Sufficient number of plants was tagged in each variety on the day of flowering. Hand picking of green immature pods commenced from 30 days after flowering (daf) and continued till harvest maturity with an interval of 5 days. The seeds were removed from picked pods for further analyses.

Extraction and estimation of lipoxygenase isozymes

For determination of lipoxygenase isozymes, fresh green seeds were ground using pestle and mortar in liquid nitrogen. The ground freeze-dried samples were defatted with petroleum ether and air dried to evaporate petroleum ether. The enzyme was extracted with soybean extract with 100 volumes of phosphate buffer (0.2 M, pH 6.8) by agitation in a micro tissue homogenizer for 20 minutes at 0-4°C. The homogenized solution so obtained was centrifuged at 10,000 rpm for 10 minutes at 4°C. The supernatant so obtained was used as the crude extract for the assay of lipoxygenase isozymes following the standard method (Axelrod *et al.* 1981). The reaction mixture for lipoxygenase-I consisted of crude extract as enzyme source, 2.8 ml of boric acid borax buffer (0.2 M, pH 9.0) and 10 mM sodium linoleate as a substrate. Lipoxygenase-II and III were analysed collectively with the reaction mixture consisting of crude extract as enzyme source, 0.2 M phosphate buffer (pH 6.8) and 10 mM sodium linoleate as a substrate. The change in absorbance was recorded in Shimadzu UV-160 spectrophotometer at 234 nm. One unit of enzyme was taken as equivalent to the amount of enzyme that generated an increase in absorb-

per minute due to conjugate diene in
atic hydroperoxidation at 234 nm.

tion and estimation of trypsin inhibitor (TI)

One gram of fresh green pods was ex-
ed in 50 ml of 0.01N NaOH for 4 hours
constant stirring at 125 rpm in an orbital
suspension so obtained was appropriately
ed so that 2 ml of the sample extract inhib-
40-60 percent of the trypsin used as a stand-
n the analysis. TI activity was determined
tandard procedure (Kakade *et al.* 1974) as
ified by Hammerstrand *et al.* (1981). Of the
test tubes taken, 2 ml aliquots of the diluted
ple were added to the four test tubes. A
test tube was prepared for the trypsin stand-
y adding 2 ml of distilled water. To three of
our test tubes containing the sample extract,
of trypsin solution (prepared by dissolving
4 g of the trypsin in 200 ml of 0.001 N HCl)
added and were maintained at a constant
emperature water bath 37°C for 10 minutes.
milliliters of benzoyl DL- arginine para-
anilide hydrochloride (prepared by dissolv-
0.08 g of benzoyl DL arginine paranitroanilide
ochloride in 2 ml of dimethyl sulfoxide and
ed to 200 ml with 50 mM Tris buffer (pH
containing 20 mM calcium chloride and the
ents were warmed to 37°C) was rapidly
d into each tube. The contents were stirred
ediate on a vortex mixture and the tubes
placed in a water bath at 37°C. The reac-
was terminated after exactly 10 minutes by
apid addition of 1 ml of 30 percent glacial
c acid. The fourth tube containing sample
act (sample blank) was prepared by the
procedure except that the trypsin solution
added after the reaction was terminated by
addition of 30 percent glacial acetic acid.
absorbance of each solution was determined
0 nm against the sample blank. Values ob-
d from each of the two sample extracts were

subtracted from trypsin standard. These values
were averaged and the trypsin content was de-
termined as follows:

TI mg /g of defatted sample = Differential
absorbance x Dilution factor / 0.019 x 1000

Percent Inhibition = 100 x Differential absorb-
ance / Absorbance of the standard

Estimation of polyunsaturated fatty acids

Oil was extracted from oven dried shelled
seeds using petroleum ether (bp 40-60°C).
Methyl esters were prepared from the oil by
interesterification in methanol using sodium
methoxide as the catalyst (Luddy *et al.* 1968).
Fatty acid methyl esters (FAMES) prepared
were separated and analyzed in gas liquid
chromatograph (GLC), Shimadzu GC 17A, us-
ing capillary column with length and diameter of
30 meter and 0.32 millimeter, respectively. Oven
temperature of the GLC was programmed at
140°C for 3.6 minutes, and subsequently in-
creased to 170°C at the rate of 13.5°C per minute
and maintained for 3.8 minutes and finally to
182°C at the rate of 5°C per minute for obtain-
ing best resolution of methyl esters. The tem-
peratures of flame ionization detector (FID) and
injector were maintained at 240°C. Nitrogen, the
carrier gas used, was maintained at a flow rate
of 15 ml/minute with column pressure at 90 kpa.
The peaks for individual FAMES were identi-
fied by comparing the retention times with those
of standard methyl esters (procured from Sigma,
USA). The analysis was done in triplicate sam-
ples and the mean values were reported.

Qualitative analysis of Kunitz inhibitor

Kunitz inhibitor from developing seeds was
extracted in Tris-Cl buffer (100 mM, pH 6.8)
containing 0.23 M CaCl₂ and 5 mM phenyl me-
thyl sulfonyl chloride (PMSF) following Kollipara
et al. (1991) and was resolved using non dena-
turing discontinuous polyacrylamide slab gel con-