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MOLECULAR MAPPING AND BREEDING FOR HIGH POPPING VOLUME IN HIGH YIELDING MAIZE LINES

THESIS

SUBMITTED TO ETERNAL UNIVERSITY, BARU SAHIB IN THE PARTIAL FULFILLMENT OF THE REQUIREMENTS FOR THE DEGREE OF

DOCTOR OF PHILOSOPHY

IN

BIOTECHNOLOGY

BY

SHIVANI THAKUR (BS15PSBT007)



DEPARTMENT OF GENETICS-PLANT BREEDING AND BIOTECHNOLOGY

Dr. KHEM SINGH GILL AKAL COLLEGE OF AGRICULTURE ETERNAL UNIVERSITY

BARU SAHIB HIMACHAL PRADESH -173101 (INDIA) OCTOBER-2020



CERTIFICATE-I

This is to certify that the thesis entitled "Molecular Mapping and Breeding for High Popping Volume in High Yielding Maize Lines" submitted in partial fulfillment of the requirements for the degree of Doctor of Philosophy in Biotechnology in the Department of Genetics-Plant Breeding and Biotechnology, Dr. Khem Singh Gill Akal College of Agriculture, Eternal University, Baru Sahib, Himachal Pradesh is a record of bonafide research carried out by Ms. Shivani Thakur Reg. No. (BS15PSBT007) under the supervision of Dr. H. S. Dhaliwal, Professor of Biotechnolgy and no part of the thesis has been submitted for any other degree. The assistance and help received during the course of this investigation have been acknowledged.

The matter presented in this thesis has not been submitted by me for the award of any other degree of this or any other Institute.

Mison thakur (Shivani Thakur)

This is to certify that the above statement made by Ms. Shivani Thakur (BS15PSBT007) is correct to the best of my knowledge.

Date: 14.10.2020 Eternal University, Baru Sahib Fr. Shalitel Dr. H. S. Dhaliwal Major Advisor

The Ph.D. Viva-Voce Examination of Ms. Shivani Thakur has been held on 10.10.2020

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CERTIFICATE-II

We, the undersigned, member of Research Degree Committee of Ms. Shivani Thakur Reg. No. (BS15PSBT007) a candidate for the degree of Doctor of Philosophy in Biotechnology agree that the thesis entitled "Molecular Mapping and Breeding for High Popping Volume in High Yielding Maize Lines" may be submitted in partial fulfilment of the requirement for the degree.

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ABSTRACT

Name: Shivani Thakur Year of admission: March 2016

Subject: Biotechnology

Degree: Doctor of Philosophy

Specialization: Plant Molecular Biology Department: Department of GPB & Biotechnology, Dr. KSG Akal College of Agriculture Thesis Title: Molecular Mapping and breeding for high popping volume in high yielding

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maize lines

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Popcom (Zea mays L.) is a special type of maize that pops up when heated leading to high flake volume. It is being used as popular fiber rich and nutritious snack all over the world. The genetic control of popping rate and flake volume are not understood properly. Most of the popcorn cultivars have low yield and are highly susceptible to various diseases. Southern corn leaf blight (SCLB) caused by Bipolaris maydis and banded leaf sheath blight (BLSB) caused by Rhizoctonia solani cause significant yield losses in maize throughout the world. The present genetics study was carried out to establish the genetic control of popping volume/flake volume using a F2 mapping population. Limited available maize germplasm was screened for resistance against SCLB and BLSB under artificial inoculation field conditions. A teosinte (Zea mexicana L.) race grown for fodder was found to be highly resistant to banded leaf sheath blight under artificial inoculation conditions while most of the maize cultivars were highly susceptible. A commercially used high popping volume (HPV) inbred line with low yield and high susceptibility to SCLB and BLSB was crossed with a low popping volume (LPV) composite, and resistance to SCLB to develop the mapping population. The F₁ HPV x LPV showed low popping volume indicating that the high popping volume is governed by recessive genes. The F2 population of 505 plants showed continuous variation for popping volume indicating that the popping volume is a quantitative trait governed by multiple genes. Bulk segregent analysis (BSA) was carried out for popping volume QTL using 66 maize SSR markers showing polymorphism between HPV and LPV. Genotyping of both positive and negative debulks using the associated markers confirmed the association of four SSR markers with QTL for popping volume. The Single marker analysis (SMA) of the F2 plants showed that the 3 SSR markers bnlg1331, bnlg1520 and bnlg1836 on chromosome 1, 2 and 5 respectively, accounting were closely associated with the QTL for popping volume covering 78% of total phenotypic variance. The QTL for popping volume mapped on chromosome 1, 2 and 5 have been designated as qPVEU-1, qPVEU-2 and qPVEU-5, respectively. As expected using BSA all the three QTLs for popping volume had additive gene action with the positive alleles for popping volume contributed by HPV. The homozygous plants of F4 and BC2F3 backcross population for the three QTL qPVEU-1, qPVEU-2 and qPVEU-5 had high popping volume validating their contribution to popping volume and their utilization for MAS for enhancing popping volume. A number of F4 and BC₂F₃ progenies homozygous for three QTL of high popping volume with resistance to SCLB, have been identified. Only 7 out of 33 BC₂F₃ progenies involving teosinte as the donor for BLSB, were uniformly resistant to BLSB strongly indicating that the BLSB resistance introgressed from teosinte is controlled by a major gene in addition to other minor genes. Selection for high popping volume was done among the BC2F3 progenies homozygous for QTL qPVEU-1, qPVEU-2 and qPVEU-5 and uniformly resistant to BLSB which could be used for mapping the genes for BLSB resistant for MAS of the popping volume in maize lines. The F₄ and BC₂F₃ progenies resistant to SCLB and BLSB diseases, homozygous for high popping volume QTL qPVEU-1, qPVEU-2 and qPVEU-5 with high yield and yield components can be further tested for the traits and used to develop the high yielding and disease resistant popcorn cultivars.

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