

# A software system for tobacco germplasm data

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## Abstract

Tobacco germplasm are of different 'types', with large numbers of accessions in each type. In order to effectively manage this tobacco germplasm information, a user-friendly menu-driven software has been developed based on Visual C++ and MS-Access. The system allows the user to store, modify and update germplasm information as and when required, and represents a decision-making tool.

**Keywords:** database; germplasm management; germplasm resources

## Introduction

Accessing the large amounts of plant genetic resource data requires the development of germplasm databases to store and retrieve the information (Fox *et al.*, 1996; Fox and Skovmand, 1996; Bruskiwich *et al.*, 2003; Christopher *et al.*, 2005; Chen and Huang, 2007; Agrawal *et al.*, 2007; Mundankar and Karibasappa, 2008). Some of the current databases are international based, while others are either national or crop based ([www.globalplanofaction.org](http://www.globalplanofaction.org)). Some examples are the AVRDC Vegetable Genetic Resources Information System (AVGRIS; <http://203.64.245.173/avgris/>), the Biodiversity Directory of Germplasm Collections database (<http://www.bioversityinternational.org/index.php?id=168>) and the System-Wide Information Network for Plant Genetic Resources (SINGER; <http://singer.grinfo.net/>). These database systems provide online access to the germplasm data.

Tobacco is grown as a commercial crop in many different countries, and each of these maintains its own germplasm collections. Most of these countries are managing these collections without computer databases. Although a single database in Chinese language is available

(Zhang *et al.*, 1990), its global accessibility is restricted in view of its language.

In India, the Central Tobacco Research Institute takes the responsibility to collect, maintain, evaluate and document tobacco germplasm, to facilitate their improvement for domestic and export purposes (Rao *et al.*, 2005). The collection consists currently of approximately 2400 accessions, including 57 wild *Nicotiana* species, but at the moment lacks a database system. Here, we describe a software system for tobacco germplasm, which has been designed to store, update and access the germplasm data.

## Software system for tobacco germplasm

A datasheet with 100 fields has been prepared based on various parameters related to tobacco germplasm (Fig. 1). Based on this datasheet, a software system for tobacco germplasm data (SSTG) has been developed in English language using Visual C++ (Bennett, 1997; Harvey and Paul, 2000, 2004) as a front-end application and MS-Access (Microsoft corporation, 1994) as a back-end application. A personal computer system with a minimum of 64 MB RAM, preloaded with Visual C++ and MS-Access, a CD-ROM drive for loading the software

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**DATA SHEET**

- **HISTORY:**  
Variety, Donor, Species, Nbpgr, Institute, special Information, Evaluators Name and Address
- **CLASSIFICATION:**  
Curing, Use
- **MORPHOLOGICAL DESCRIPTION:**  
Plant Habit, Height up to Crow Foot, Internodal Length, Plant Width or Spread (cm), Stem Colour, Leaf Characters, Total Leaf Number, Economic Leaf Number, No of Cured leaves / kg, Suckering Habit, Spangling, No. of Days Taken for Flowering, Inflorescence, Flower Colour, Corolla Shape, Capsule Shape, Capsule, Seed, Seed Dormancy, Rooting Habit.
- **REACTION TO BIOTIC & ABIOTIC STRESS:**  
Diseases, Pests, Abiotic Stresses
- **CROP SEASON**  
Irrigated, Rain-fed
- **YIELD DATA (KG/HA)**  
Green Leaf, Cured Leaf, Bright Leaf, Grade Index
- **QUALITY CHARACTERISTICS**  
Chemical, Physical, Cured Leaf Assessment Score Card.

Fig. 1. Datasheet consisting of various fields in the database.

and a Super Video Graphics Adaptor monitor is required for the execution of this software.

SSTG helps to store, update and access the tobacco germplasm data for the optimum management of germplasm resources. The main menu (Fig. 2) of this system consists of five modules. The 'INSERT' module allows the user to enter the entire record of a germplasm accession including history (variety, type, donor, specifics, etc.), classification (curing and purpose), morphological classification (plant habit, height, internodal length, stem width, stem colour, etc.), leaf characters (colour, stalk, shape, base, midrib, surface, maturity, size and leaf thickness), reaction to biotic and abiotic stresses (diseases, pests and abiotic stress), crop season (crop condition, green leaf, cured leaf, bright leaf and grade index) and quality characteristics (chemical, physical, cured leaf, etc.) into the database. The 'DELETE' module helps to delete the entire information of a particular accession. The 'UPDATE' module allows changing the values of a record in the stored database. The 'RECORD REPORTS' module permits the display of

the entire information by selecting an accession name. The 'QUERY' module (Fig. 3) provides an option of selecting the germplasm lines having combinations of various parameters. Multiple queries can be made

**Select upto 10 Fields for View / Printing**

**Fields in Table**

SCORE4  
SCORE5  
SCORE6  
SCORE7  
SCORE8  
SEC\_FLOWER  
SPANGLING  
SPECIES  
SPL\_INFO  
SUK\_O\_TOP  
SUK\_WD\_TOP  
TOT\_LF\_NO  
YD\_BRIGHT  
YD\_GRADE  
YD\_GREEN

>

<

**Select Fields:**

USE  
VARIETY  
YD\_CURED

OK

CANCEL

Fig. 3. Query menu 1 helps the user to select various parameters for accessing the required information from the database.

GERM PLASM

MAIN MENU

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INSERT

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DELETE

---

UPDATE

---

QUERY

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RECORD REPORT

EXIT

Fig. 2. Main menu allows the user to add, delete, update and retrieve the data from the database.

**Enter the Query/Press OK to Default Query**

**Fields in Table**

SCORE4  
SCORE5  
SCORE6  
SCORE7  
SCORE8  
SEC\_FLOWER  
SPANGLING  
SPECIES  
SPL\_INFO  
SUK\_O\_TOP  
SUK\_WD\_TOP  
TOT\_LF\_NO  
YD\_BRIGHT  
YD\_GRADE  
YD\_GREEN

>

<

**Selected Field**

USE  
VARIETY  
YD\_CURED

Relation:

Values

OK

CANCEL

Fig. 4. Query menu 2 helps the user to access the database with multiple queries using relational operators.

using relational operators (Fig. 4) to obtain the required report. A hard copy of the selected report can be obtained using the 'PRINT' option.

## Conclusion

SSTG with user-friendly menus for storing and retrieval of tobacco germplasm data will be of use for tobacco researchers. This software enables the scientists to add new data, delete unwanted data, update existing data and retrieve data on the selected parameters with conditional queries on germplasm accessions. This database can be kept in the web server for online access by the scientists engaged in tobacco research, throughout the world. SSTG can be obtained by contacting hravisankar@india.com.

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