# *APPLIED BIO-SYSTEMS TECHNOLOGY*

### **Research Article Open Access**

# **Evaluation of the Effect of Morphological Traits on Blister Blight Resistance in Tea Plant (***Camellia sinensis* **L.)**

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### **Abstract**

**Background:** Blister Blight (BB) is a serious leaf disease caused by the fungus *Exobasidium vexans* Masse, damaging Sri Lankan tea plantations.

**Methods:** A morphological trait-based analysis was preformed based on 14 descriptors for Camellia sinensis to differentiate BB resistant and BB susceptible individuals in anF1 population generated by a cross between BB resistant and BB susceptible cultivars (TRI 2043xTRI 2023). The Spearman's correlation analysis, regression modelling, Receiver Operating Characteristic (ROC) and t-test were applied in the analysis of morphological characteristics of the F1 plants.

**Results:** Leaf pubescence (SCC= - 0.530), upper leaf surface (SCC= 0.473) and length of mature leaf petiole denoted significant associations with BB disease index (P<0.05). Threshold values of the developed model to screen vulnerability of tea plant to blister blight were 1.5 for both pubescence of tea leaves and upper leaf surface.

**Conclusions:** Proposed leaf morphology-based thresholds can be successfully applied for preliminary screening of BB susceptibility, prior to further confirmation with more advanced identification techniques.

**Keywords:** Disease Control, Fungal Infection, Marker Assisted Tea Breeding, Non-Alcoholic Beverages, Plant Inherent Resistance

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### **INTRODUCTION**

Tea is one of the most popular non-alcoholic caffeine containing beverages in the world with high amounts of flavonoids and related bioactive compounds [1-2]. Though it remains unclear, it is considered that the tea plant (*Camellia sinensis* L.) was originated from China and spread through the other South East Asian countries as a commercial cultivation [3]. Succulent plant leaves of tea are processed to make three most popular types of tea, green (unfermented), black (fully fermented) and oolong (semi-fermented) [4].

In tea plants, a considerable crop loss results due to nutrient deficiencies, stresses from climatic variations, pests and pathogen attacks. Among diseases, Blister Blight (BB) leaf disease, is caused by the obligatory biotrophic pathogen, *Exobasidium vexans* Massee, which infects only young harvestable succulent leaves, stems and the pericarp of fruits at young stage [5]. The pathogen is spread by windborne basidiospores and infection mostly proceeds through stomata [6]. Mycelium grows intercellularly prior to the formation of basidia fruiting bodies on epidermis. When they grow further, initially small pinhole size spots become visible on young leaves. As the leaves develop the spots become transparent, larger and light brown in colour. They force up and rupture the lower epidermis to form blisters with dark green and water-soaked zones surrounding the blisters [7]. After releasing the fungal spores, the blisters become velvet and white and which subsequently turns into brown in colour [8].

The BB disease causes approximately 25% - 30% crop loss per annum in Sri Lanka. Infected harvestable leaves directly reduce crop yield, not only quantitatively but qualitatively as well, due to the changes of the composition of leaf biochemical compounds such as polyphenols, catechins, enzymes etc*.* [9]. As the environmental conditions such as high humidity and limited sunshine directly fascilitate the infection and development of BB [10] in the field, it is important to control

these microclimatic factors, in which BB may develop.

Morphological characteristics of the infected plants can be used to develop a clear relationship between the morphological characteristics of plants and blister blight resistance using an accurate scoring system for BB susceptibility of plants. Subsequently Quantitative Trait Locus (QTL) mapping can be done for proper identification of the infected plants. Evaluating the agricultural significance of fungal leaf diseases and developing tools that enable rapid recognition of diseases are very important to eliminate these pathogens [11-12].

Various methods are used to identify different pathogens, which cause diseases in crops. Morphological and genetic analysis of infected plants are two main approaches to studyplant pathogen interactions and disease development. Ponmurugan and Baby [13] conducted a study on the morphological, physiological, and biochemical changes in tea plants due to Phomopsis infection. Physiological parameters; photosynthetic and transpiration rates, stomatal conductance, efficiency of water usage and total chlorophyll content were scored both in susceptible TRI-2024 and tolerant TRI-2025 tea cultivars. Plant height, dry weight and plant strength were recorded as morphological characteristics, while total sugar, nitrogen, amino acids, protein, polyphenols and catechins of infected and healthy plants were studied as biochemical parameters. Results revealed all the morphological, physiological and biochemical characteristics tend remain significantly low in infected plants, compared to healthy plants [13].

Growth, photosynthetic and biochemical responses of tea cultivars to BB infection has been studied by Premkumar *et al*. [14], where susceptibility to BB infection has denoted significant strong associations with physical barriers, physiological and biochemical parameters (leaf area, shoot

length and moisture contents etc). Not only the characteristics of the plants, but also the variations in infection causing pathogens (*Exobasidium vexans* Massee) have been studied by Abeysinghe *et al.* [15] using infected tea leaf samples. As the infection has a short (11-28 days), but multiple disease cycles with several generations within a single crop season, it requires repeated applications of fungicides to control the disease [6]. However, continuous application of fungicides can contribute to the development of new strains of *Exobasidium vexans* Massee. Morphological parameters of the pathogen such as colour, length and width of spores and DNA finger printing analysis using RAPD have revealed a high degree of genetic diversity among the samples of the *E. vexans* as an adaptation due to various conditions [16].

Plants utilize structural and chemical characteristics to prevent or reduce the spread of pathogens, which act as their first line of defence against pathogens [17]. Development of plant inherent resistance of tea cultivars against BB is the most suitable solution to control the disease, instead of applying highly toxic fungicides. In this process, marker (morphological, biochemical and molecular) assisted tea breeding is playing a key role. Among those, biochemical markers and molecular markers are more accurate and precise techniques [18-19]. However, the cost and technical requirements of molecular markers and biochemical markers are very high and the process is time consuming. Therefore, developing a simple and rapid assessment approach for early detection of BB resistance is immensely important. Hence, this study aimed to develop an inexpensive, user friendly, accurate and reliable morphological marker for preliminary screening of BB resistance traits in tea cultivars.

# **METHODOLOGY**

# **Materials**

In the present study, 300 individuals of an  $F_1$ segregating population derived from a cross

between TRI 2043 (a tea cultivar resistant to BB disease) and TRI 2023 (a cultivar susceptible to BB) were used in each replicate. Three replicates were grown following Randomized Complete Block Design (RCBD) together with their parents at St. Coombs Estate (Up country wet zone of Sri Lanka), Tea Research Institute, Talawakelle, Sri Lanka.

# **Assessment of Blister Blight Disease Severity of the F<sup>1</sup> Individuals**

Blister Blight Disease Index (BBDI) of each F<sup>1</sup> individual was calculated using the data collected from the field assessments starting from year 2007 to 2010 at one-week intervals based on guidelines given in the BB severity assessment key [19].

# **Morphological Characterization of F<sup>1</sup> Individuals**

Morphological assessment of  $F_1$  individuals was carried out using the descriptors for *Camellia sinensis* L. described by the International Plant Genetic Resources Institute (IPGRI) [20]. Shape of the 5th leaf, size of the 5th leaf, leaf color, apex shape of the 5th leaf, habit of the 5th leaf apex, shape of the 5th leaf base, pubescence of the 1st leaf, leaf venation, leaf vestiture, upper leaf surface, length of the 5th leaf, width of the 5th leaf, length of mature leaf petiole, leaf length to width ratio were the fourteen morphological characteristics scored in this study. Non parametric data were converted into numerical values on the scale mentioned by the IPGRI [20]. The assessment was repeated for each individual established in three different locations and the average of each parameter was used for the statistical analysis.

# **Statistical Analysis**

Offspring of the studied population were separated into two sets of samples based on the seed bearer (mother). The offspring produced from the seeds of TRI 2043 was considered as group 1, while group 2 consisted of the offerings from the seeds of TRI 2023. Spearman's correlation analysis was used to assess the association between different morphological characteristics of the plant and the BBDI. Further, data obtained from morphological characteristics were subjected to regression modelling, after square root transformation in order to develop a model on susceptibility to BB. In addition, Receiver Operating Characteristic (ROC) analysis was used to define the risk thresholds for susceptibility to BB based on the significantly associated morphological characteristics of the  $F_1$  plants.

### **RESULTS AND DISCUSSION**

# **Blister Blight Disease Severity of F1 Individuals**

Individuals with high BB resistivity were grouped on the left-hand side of Figure 1, while high BB susceptible individuals were grouped at the right-hand side. According to the BB severity assessment key, individuals with less than 0.1 BBDI were considered as high BB resistant and individuals with a BBDI higher than 0.5 were considered as highly BB susceptible individuals. According to the analysed data, P219, P58 and P1040 were  $extremely$  resistant  $F_1$  individuals. Meanwhile, P219, P1016 and P1018 were extremely susceptible  $F_1$  individuals.

### **Morphological Characterization of F1 Individuals**

The morphological characteristics of all 300 F<sup>1</sup> individuals were assessed and the results of six  $F_1$  individuals from two extremes with parents are given in Table 1.

**Impact of Mother Plant on the Morphological Characteristics and Incidence of Blister Blight of the Offspring** Among the notable variations in the morphological characteristics of the two offspring groups, only eight morphological characteristics; leaf colour, leaf apex shape, leaf apex habit, leaf vestiture, upper leaf surface, length of mature leaf, width of mature leaf and length of mature leaf petiole, advocated significant differences at 95% level of significance, in accordance with the statistics of the t-test (Table 2). However, the BBDI values of the two test groups did not

show any significant variations (P<0.05).

# **Impact of Leaf Morphology on the Incidence of Blister Blight**

Among the studied leaf morphological characteristics, leaf shape, leaf apex habit, leaf pubescence, leaf vestiture and leaf length to width ratio denoted negative relationships with the BBDI, while rest of the characteristics indicated positive associations. However, only the associations of leaf pubescence, upper leaf surface and length of mature leaf petiole were significant at 95% level of confidence (P<0.05), in accordance with the Spearman's correlation analysis (Table 3).

Leaf pubescence advocated a significant negative moderate relationship with the BBDI (Spearman's Correlation Coefficient [SCC]=- 0.530), while on the other hand, a significant positive moderate association (SCC=0.473) was indicated by the upper leaf surface. Regardless of the significance in correlation, the impact of leaf length to width ratio on BBDI remained to be poor (SCC<0.1).

### **Morphological Characteristics which Affect the Susceptibility to Blister Blight**

The regression analysis based on backward elimination, yielded a simple model for the identification of susceptibility of a tea plant based on the morphological characteristics. As indicated by the model, the susceptibility of the plant remains as a function of Upper Leaf Surface (ULS) and Leaf Pubescence (LP), as indicated by the Equation 1. The model was characterized by a  $\mathbb{R}^2$  value of 0.57, followed by an adjusted  $R^2$  value of 0.51.

Succeptibility to Blister Blight Disease  $\;\;(1)$  $= 0.78 + (0.60 X ULS) - (0.25 X LP)$ 

### **Definition of Risk Thresholds for Blister Blight based on Leaf Morphology**

The Receiver Operating Characteristic (ROC) curve analysis yielded an area coverage of 0.534 and 0.479, for upper leaf surface and leaf pubescence, respectively, while the incidence of BB was defined as BBDI>0.1 (Figure 2).



**Figure 1:** The Bar Chart of 300 F1 Individuals against Blister Blight Disease Index

**Table 1:** Morphological Characteristics of Selected Six F<sub>1</sub> Individuals from the Two Extremes (Highly Resistant and Highly Susceptible) of the BBDI, along with Their Parents



*Note: Sam.: Sample*



**Table 2:** Results of the t-Test for Significant Differences among Leaf Morphological Characteristics between the Two Groups (Highly Resistant and Highly Susceptible) of Off springs

*Note: "\*" in the column indicates significant difference (P<0.05) among the two groups in accordance with the t-test*

**Table 3:** Results of the Correlation Analysis between Different Leaf Morphological Characteristics and Blister Blight Disease Index



*Note: "\*" denotes parameters that indicated a significant correlation with BBDI (P<0.05) at 5% level of significance, while "#" denotes parameters significant at 1% level of significance (P<0.01).*



**Figure 2:** ROC Curve for the Leaf Pubescence and Leaf Vestiture Associated with BBDI

Therefore, it was reassured that both of these leaf characteristics are significantly associated with the incidence of BB. Based on the distribution of the curve and the coordinates of the curve (Figure 2) (sensitivity and 1– specificity), leaf pubescence>1.5 and upper leaf surface>1.5, categories could be considered as risk thresholds, which can symbolize the susceptibility of tea plants to BB incidence. In the analysis of the BB severity of both  $F_1$  individuals derived from TRI 2043 and TRI 2023, resistance and susceptibility characteristics have not shown any maternal segregation as denoted by the statistics of the t test.

As suggested by the overall results of Spearman's correlation analysis, morphological characteristics such as leaf shape, leaf apex habit, leaf pubescence, leaf vestiture and leaf length to width ratio have denoted negative relationships with BBDI. Narrow leaf with notable length has decreased the susceptibility to the BB. The ratio between leaf length and leaf width also have shown negative correlations with BBDI.

Therefore, lanceolate shaped leaf with higher value of ratio between leaf lengths to width can be considered as more resistant to blister blight disease, than ovate, oblong and elliptic leaf shapes. Lanceolate shaped leaves are more resistant to BB, due to the availability of a narrow space for accumulation of BB spores.

In the current study, leaf pubescence has also indicated a significant negative correlation with the BBDI. The leaf pubescence was observed under microscope and categorized as; sparse, intermediate or dense. When pubescence density increases, it can act as a physical barrier for infection [21] limiting the susceptibility to BB. In a study conducted for Uromyces, the presence of dense leaf pubescence has been documented to retard the germination of spores on the surface of bean leaves by trapping the spores [22], thereby reducing the probability of germ tubes reaching the penetration site [24]. A high density of trichomes can also prevent mycelial penetration and infection of other biotrophic fungi [24]. It is reported that an increased number of hydrophobic

pubescence may repel water from the leaf surfaces, thus preventing successful penetration of fungal germ tubes [24]. Alternatively, a high trichome number may simply reduce the frequency of germ tube contact points that can lead to penetration [25]. The straight leaf apex habit was more vulnerable to disease infection rather than down turned leaf, as suggested by the negative correlation between leaf apex habit and BBDI.

On the other hand, several factors such as leaf colour, leaf apex shape, leaf base habit, leaf venation, upper leaf surface, length of mature leaf, length of mature leaf petiole have shown positive correlations with BBDI. All the positively correlated characteristics increase the probability of being infected by pathogen spores, through facilitating the trapping of spores and providing more surface area to interact with the plant leaf. Increase of the leaf length and leaf width may increase the surface area of the leaf and allow the spores of pathogens to increase the chance of contamination [12].

Distinct mid rib and lateral leaf venation system with bullate has also made the leaf more vulnerable to disease infection, than indistinct sunken leaf venation in lamina. It may facilitate the trapping of spores in the wind. When considering the impact of leaf base area, the susceptibility to BB tend to increase from attenuate to blunt shapes, when the leaf base surface area increases. Leaf apex shape denoted a positive correlation with BBDI and therefore the BB severity tend to vary as acute < obtuse < attenuates in shapes, respectively.

Most outstanding leaf morphological characteristics such as upper leaf surface and length of mature leaf petiole were denoting significant positive correlations with BBDI (P<0.05 at 95% level of confidence). A rough upper leaf surface generally leads to high retention of fungal spores, while increasing the length of leaf petiole, may favour the exposure to spores of pathogen [26].

After considering the correlation of all the studied leaf morphological parameters with susceptibility to BB, a simple model to predict the vulnerability of a tea plant to BB (based on morphological features) was developed through step-wise regression analysis. However, the current model only considers the leaf morphological factors with less attention on other external environmental factors such as soil nutrients, light etc. Further, regardless of the combined effect of upper leaf surface and leaf pubescence in terms of BB susceptibility, individual thresholds for each parameter were also developed through a ROC analysis. However, it should be noted that both, upper leaf surface and leaf pubescence are nonparametric morphological parameters and the model was derived with a limited number of samples. Therefore, the current thresholds and leaf morphology-based model is recommended for preliminary screening of BB susceptibility, due to its rapid and limited resource consumptive (labour and cost) nature, prior to further confirmation with more precise molecular markers.

# **CONCLUSIONS**

Constitutive barriers limited or completely inhibited the penetration of tea tissues by pathogenic fungi. The resistant individuals of the analyzed  $F_1$  segregation population were characterized by a significantly higher pubesence density, than susceptible forms. In resistant individuals, upper leaf suface was smooth, which minimise the accumilation of pathogen spores.

Based on the findings, upper leaf surface and leaf pubescence can be used to evaluate the susceptibility to BB incidence in tea plants. The proposed model can be used for preliminary evaluation of BB resistant or BB succeptible traits and it should be validated with more tea cultivars in different ecological regions to enhance the reliability and accuracy.

# **CONFLICTS OF INTEREST**

The authors declare that there are no conflicts

of interest.

# **AUTHORS' CONTRIBUTIONS**

TK: Carried out the investigations, data collection, supported the statistical analysis, and wrote the manuscript; MK and JW: Supervised the study; LU: Analysed data and wrote the manuscript; CP: supervised the study and revised the manuscript; NE: Supported data collection process. All authors read and approved the manuscript.

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