

3 **Differential biochemical response among Banana (*Musa* spp.)**
4 **genotypes against Banana Bunchy Top Virus (BBTV)**
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10 **ABSTRACT**
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Banana bunchy top virus (BBTV) is one of the major viruses causing high yield loss in bananas. The study was carried out to gain a better understanding of the host and virus interaction and to explore the adaptive mechanism and biochemical responses in banana cultivars viz., Rasthali and Grand Naine against the banana bunchy top virus (BBTV). In the leaf samples of BBTV infected Rasthali and Grand Naine, estimated the total chlorophyll, carbohydrates, phenols and enzyme activities such as peroxidase, polyphenol oxidase, catalase, ascorbate peroxidase, guaiacol peroxidase and superoxide dismutase. The virus infected samples of both cultivars showed a significant increase in the defense enzymes over the healthy sample. Higher total phenols in healthy Rasthali plants which further significantly increased after BBTV infection was observed in comparison to Grand Naine. In contrast to Grand Naine, Rasthali showed higher polyphenol oxidase (PPO) activity contributing to increased polyphenol content. Higher superoxide dismutase (SOD) activity in virus infected Rasthali was observed in comparison to Grand Naine. The increased amount of total phenols, polyphenols and SOD activity in Rasthali might have contributed to less susceptibility to bunchy top virus. However, total protein and chlorophyll content were reduced after BBTV infection in both the banana cultivars.

12
13 *Keywords: [Banana Bunchy top virus, Rasthali, Grand Naine, Biochemical changes and*
14 *Defense enzymes]*

15
16 **1. INTRODUCTION**

17 Plants are frequently exposed to infection by a wide array of pathogens that lead to different
18 responses in the host plant. During compatible plant-pathogen interaction, along with the

19 development of visible symptoms, the pathogen adversely affects the growth and
20 development, physiological status and yield of a plant [1]. The constitutive defense
21 mechanism in the plants such as pre-existing physical and chemical barriers, along with
22 inducible defense responses restrict pathogen colonization [2,3,4]. Bananas and plantains
23 are important agricultural produce which are attacked by various pests and pathogens
24 causing major production losses. Grand Naine (AAA subgroup Cavendish) is most
25 commonly cultivated commercial Cavendish cultivar in the world. Rasthali (AAB subgroup
26 Silk) is cultivated mostly in India and is popular in the local and world market as a premium
27 dessert variety similar to Cavendish bananas. Plant viral diseases cause significant losses
28 by reducing plant growth and yield. Banana bunchy top disease (BBTD) is one of the most
29 damaging viral diseases affecting various banana cultivars. There is no resistant germplasm
30 available in bananas and plantains; however, the level of susceptibility varies among the
31 banana cultivars.

32
33 BBTD infects the fruit and foliage. It is caused by a single-strand DNA virus, the banana
34 bunchy top virus (BBTV; Genus *Babuvirus*; Family *Nanoviridae*). The virus colonizes in the
35 phloem tissue and damages the host cells. The etiology of the disease comes from the
36 typical symptoms which occurs in banana plants, in which the newly emerging leaves are
37 narrow, chlorotic and reduced leaf size, which causes a "bunchy" appearance at the top. In
38 addition, few distinctive symptoms are 'morse code streaking,' 'green J hooks' and 'keikis'.
39 Viruses uses the host machinery for their replication and multiplication, which initiate
40 significant changes in their usual physiological processes such as loss of pigment contents,
41 increasing respiration rates, soluble sugar and starch accumulation and production of higher
42 levels of enzymatic antioxidants. Due to viral infection, changes occur in the host plants at
43 the molecular level, thereby leading to biological and physiological changes. Hence, it is of
44 value to estimate the physiological and biochemical changes in banana cultivars Rasthali
45 and Grand Naine and to measure biochemical changes occurring due to BBTV infection.
46 The present investigation will lead to better understanding of the defense mechanism in two
47 banana cultivars, which will be useful for adopting suitable control strategy against bunchy
48 top disease in banana.

49 2. MATERIALS AND METHOD

50 2.1 Plant material and source of infection

51 Banana cultivars Rasthali and Grand Naine were used in the present investigation. Leaf
52 samples were collected from BBTV infected plants in Orchard of Tamil Nadu Agricultural

53 University, Coimbatore district, Tamil Nadu, India. Leaf samples from healthy plants of each
54 cultivar were taken as control.

55 **2.2 PCR confirmation of BBTV presence in the infected banana samples**

56 The plant genomic DNA was isolated from 100 mg leaf samples of healthy and infected
57 (showing characteristic symptoms of BBTV) samples of Rasthali and Grand Naine using the
58 cetyl trimethyl ammonium bromide (CTAB) method with some modification as described by
59 Doyle and Doyle [5] and subjected to PCR using the BBTV specific primers designed for
60 Replicase gene F-5' ACGACAGAATGGCGCA3' and R-
61 5'TCAGCAAGAAACCAACTTTATTC3'. The PCR products were resolved on 1 % agarose
62 gel, electrophoresed at 70 V for one h and the amplicons were assessed with 1.0 kb DNA
63 ladder.

64 **2.3. Leaf samples for biochemical analyses**

65 BBTV infected samples were collected from the most recent fully expanded leaf for
66 biochemical analysis. All biochemical parameters were measured using a spectrophotometer
67 (Jasco V- 730 BIO spectrophotometer, USA).

68 **2.4 Estimation of photosynthetic pigments:**

69 The photosynthetic pigments such as chlorophyll 'a', chlorophyll 'b' and total chlorophyll
70 content of healthy and infected leaves were estimated according to the non-destructive
71 DMSO method as explained by Hiscox and Israelstam [6]. The absorbance was recorded at
72 663 and 645 nm, respectively in a spectrophotometer. Chlorophyll a, b and total chlorophyll
73 were calculated by the following formulas:

$$74 \text{ Chlorophyll a (mg g}^{-1} \text{ tissue)} = \frac{[12.7(\text{OD}_{663}) - 2.69(\text{OD}_{645})] \times V}{1000} \times W$$

$$76 \text{ Chlorophyll b (mg g}^{-1} \text{ tissue)} = \frac{[22.9(\text{OD}_{645}) - 4.68(\text{OD}_{663})] \times V}{1000} \times W$$

$$78 \text{ Total Chlorophyll (mg g}^{-1} \text{ tissue)} = \frac{[8.02(\text{OD}_{663}) + 20.20(\text{OD}_{645})] \times V}{1000} \times W$$

80 Where OD, Optical density at respective nm, V, Final volume of chlorophyll extract, W, Fresh
81 weight of the tissue extracted.

82 **2.5 Total sugars and starch content:**

83 Total reducing sugars were calculated according to the method described by Dubios *et al.*,
84 [7] and the total starch content as suggested by McCready *et al.*, [8]. The absorbance of the
85 samples was recorded at 625 nm in a spectrophotometer along with the blank sample. The
86 amount of total sugars and total starch was estimated by using a standard curve prepared

87 for D-glucose. The content of reducing sugar and total starch was expressed as mg g⁻¹ fresh
88 weight.
89

90 **2.6 Phenolic content:**

91 Phenol content was measured using Folin-Ciocalteu reagent and estimated using the
92 method described by Folin and Ciocalteu [9]. The absorbance of the samples was recorded
93 at wavelength 660 nm against a reagent blank. Using pyrocatechol as standard, a standard
94 curve was generated to determine the concentration of total phenols in the leaf extract.

95 **2.7 Measurement of total protein content**

96 Total protein was estimated by using the Bradford method [10] and absorbance was
97 recorded at 595 nm. Bovine serum albumin was used as a standard. Protein contents in leaf
98 samples were recorded as µg of protein per gram of leaf tissue.

99 **2.8 Preparation of enzyme extract:**

100 To obtain the total enzyme extract, a one-gram leaf sample was homogenized at 4°C in 1 ml
101 of extraction buffer [50 mM potassium phosphate buffer (pH 7.0), 1 % Triton X-100 and 7
102 mM 2-mercaptoethanol]. The obtained homogenate was then centrifuged at 12000 rpm for
103 20 min at 4°C. The resulting supernatant was used for analysis of enzymes.

104

105 **2.8.1 Peroxidase activity**

106 POX activity was assessed following the oxidation of pyrogallol according to the method
107 given by Malick and Singh [11]. The absorbance of the reaction mixture of the sample was
108 recorded at 430 nm at 30-sec intervals up to 3 min. The specific activity of the enzyme was
109 expressed as micromoles pyrogallol oxidized per minute per milligram protein.

110

111 **2.8.2 Polyphenol oxidase activity**

112 PPO activity was determined according to the method described by Ngadze *et al.*, [12]. The
113 activity was measured by monitoring the increase in absorbance for 3 min at 410 nm. The
114 specific activity of the enzyme was expressed as micromoles catechol oxidized per minute
115 per milligram protein.

116

117 **2.8.3 Catalase activity**

118 Catalase activity was calculated by measuring the rate of disappearance of H₂O₂ using the
119 method followed by Maechlay and Chance [13]. The decrease in H₂O₂ was followed as a

120 decline in absorbance at 240 nm. Catalase activity was expressed as micromoles of H₂O₂
121 oxidized per minute per milligram protein.

122

123 **2.8.4 Ascorbate peroxidase activity**

124 APX activity was determined using the method described by **Chen and Asada** [14]. The
125 oxidation of ascorbate was followed by a decrease in the absorbance at 240 nm. The
126 enzyme-specific activity is expressed as micromoles ascorbate oxidized per minute per
127 milligram protein.

128

129 **2.8.5 Guaiacol peroxidase activity**

130 GPX activity was calculated using the method described by **Upadhyaya et al.** [15]. The
131 increase in absorbance at 420 nm was recorded for 1 min. The enzyme-specific activity is
132 expressed as micromoles guaiacol oxidized per minute per milligram protein.

133

134 **2.8.6 Superoxide dismutase activity**

135 SOD was assayed by measuring its ability to inhibit the photochemical reduction of nitroblue
136 tetrazolium (NBT) using the method described by **Dhindsa et al.**, [16]. The absorbance of the
137 reaction mixture was recorded spectrophotometrically at 560 nm.

138

139 **2.9 Statistical analysis:**

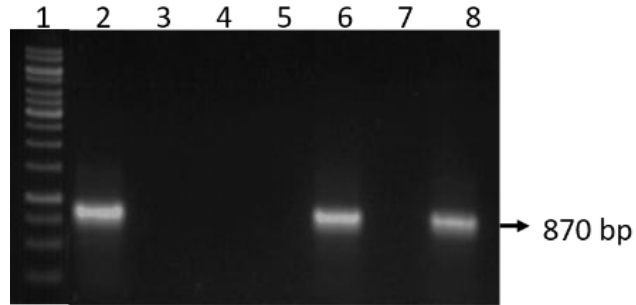
140 All the experiments were performed in two duplicates (n=4). The significance of differences
141 between healthy and infected samples was determined by using one-way analysis of
142 variance (*ANOVA*) and means and standard errors were calculated. Differences in means
143 were considered significant when the *P*-value was <0.05.

144

145 **3. RESULTS AND DISCUSSION**

146 **3.1 PCR based confirmation of BBTV**

147 The presence of BBTV in symptomatic leaves of Rasthali and Grand Naine was confirmed
148 by PCR amplification of 870 bp BBTV Rep gene using designed gene specific primers
149 (Fig.1.)



150

151 **Fig. 1. PCR amplification of BBTV Rep gene in symptomatic Rasthali and Grand Naine**
152 **plants**

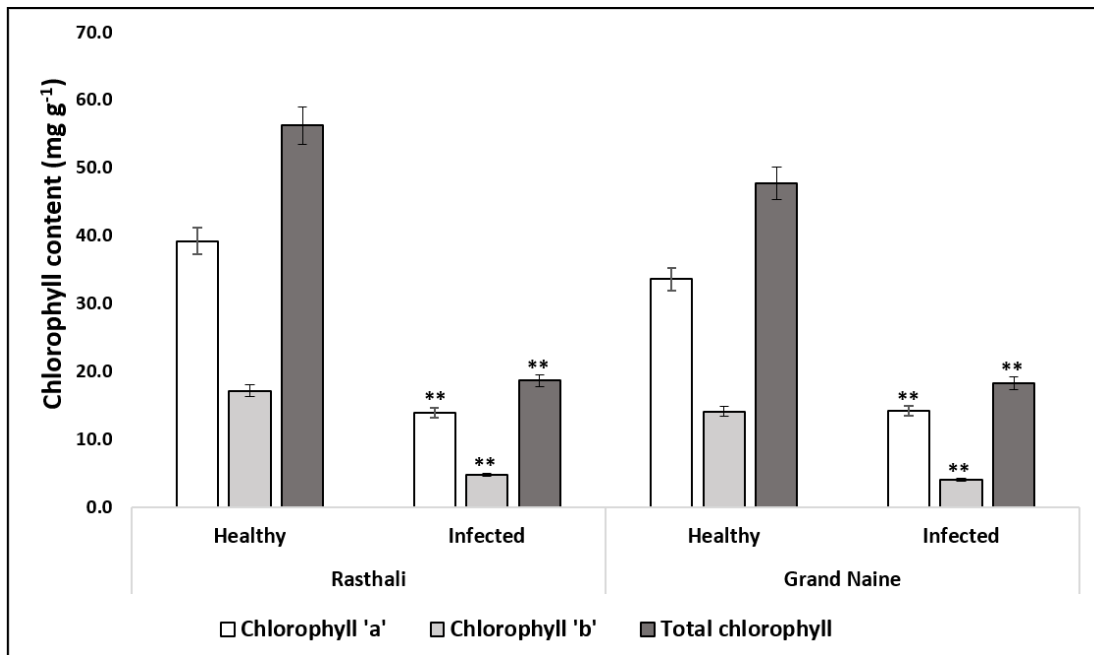
153 *Lane 1, 1 kb ladder; 2, positive control; 3, negative control; 4, water control; 5-6, Rasthali healthy and*
154 *infected sample; 7-8, Grand Naine healthy and infected sample.*

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156 **3.2 Effect of BBTV incidence on the photosynthetic pigment**

157 The BBTV infected plants exhibited a two- fold reduction in photosynthetic pigment contents
158 (chlorophyll a, chlorophyll b and total chlorophyll) compared to healthy plants (Fig. 2). The
159 decrease in the chlorophyll content in the infected plant reduces the photosynthetic capacity
160 and plant growth resulting in the symptoms such as stunting and chlorosis. This change in
161 the chlorophyll content can be due to the stimulation of enzymes like chlorophyllase that
162 degrades chlorophyll [17], or it may be due to the effect of the virus on pigment synthesis
163 [18, 19]. A recent study suggests the possibility of BBTV utilizing the chloroplast for the
164 synthesis of viral proteins [20]. They found during BBTV infection, outer membranes of
165 chloroplasts are **disrupted** and crystalline aggregation of virus- like particles accumulate in it.

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167

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Fig. 2. Chlorophyll content of healthy and infected Rasthali and Grand Naine plants.

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Data represent the mean \pm standard error of mean of four independent replications. Significant differences in healthy and infected from each cultivar analysed by Student's t test (* $P < 0.05$, ** $P < 0.01$) are shown.

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3.3 Carbohydrates

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The total sugars and starch were significantly higher in infected plants compared to healthy

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in both the cultivars of banana (Fig. 3 a-b). Our study suggests that sugars increase during

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BBTV infection may control photo inhibitory processes and produce symptoms. The

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carbohydrate content reported by [21] was similar to the findings of our study where changes

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in the sugar and starch content were the same in all the banana cultivars viz., Virupakshi,

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Grand Naine and Rasthali. Viruses appear to alter both their rate of synthesis and rate of

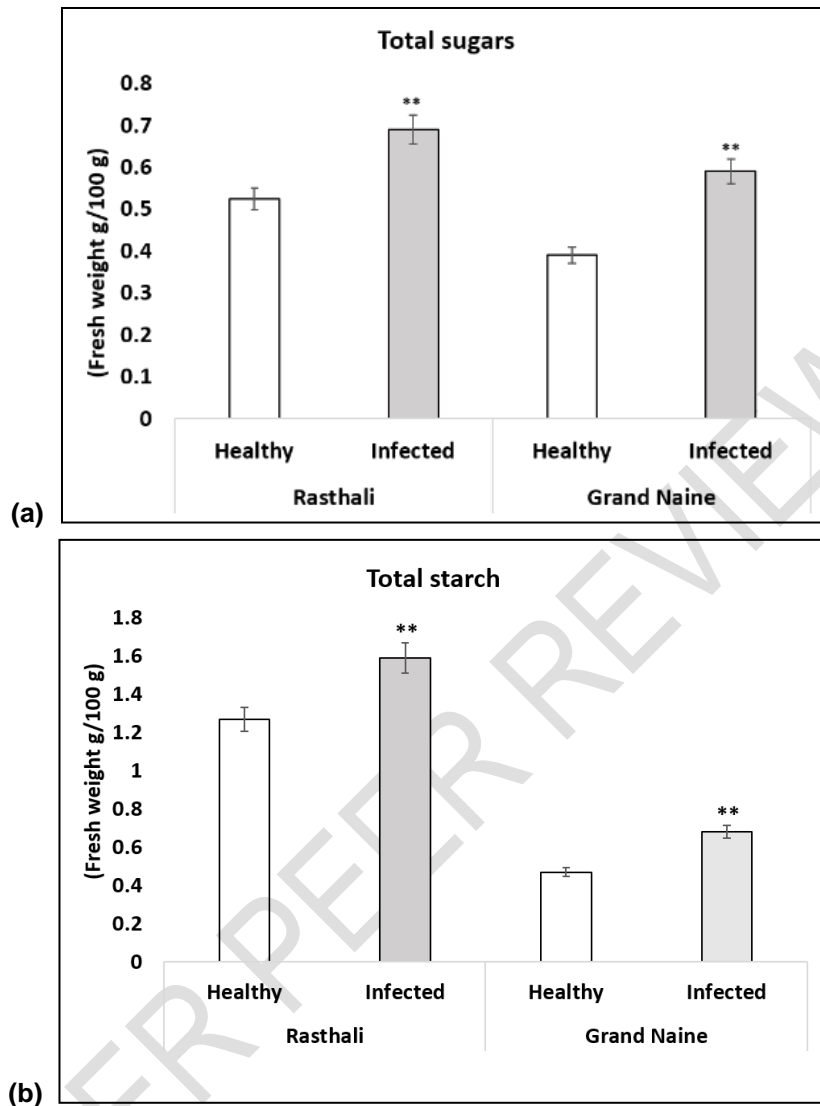
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translocation and have little effect on carbohydrates which affects the overall growth of the

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plant [22].

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183

184 **Fig. 3. Carbohydrate content (a) Total sugars and (b) Starch of healthy and infected**
185 **Rasthali and Grand Naine plants.**

186 *Data represents the mean \pm standard error of mean of four independent replications. Significant*
187 *differences in healthy and infected from each cultivar analysed by Student's t test (* P <0.05, ** P <0.01)*
188 *are shown.*

189 3.4 Total Phenol

190 In the present investigation a significant variation in the total phenolic compounds of banana
191 cultivars in response to infection with BBTv was apparent (Fig. 4a). The total phenol content
192 was significantly higher in virus-infected leaves in both the cultivars tested. Increased
193 quantities of phenols might be attributed to a defense mechanism where plant polyphenols
194 act as secondary metabolites. It has been reported that resistance to disease caused by
195 pathogens can be attributed to the presence of a high amount of phenol [11, 19, 23, 24, 25].

196 It has been reported that Rasthali viral **infection occurs** later than in the Grand Naine cultivar
197 because of the difference in the genomes [26]. Although no Musa genotype is known to be
198 resistant to BBTv, cultivars in the AA and AAA genomic groups are highly susceptible,
199 whereas cultivars containing the B genome are regarded as less susceptible. The less BBTv
200 susceptible Rasthali had higher total phenol content in healthy plants which further increased
201 >2 fold after BBTv infection. This is in contrast to Grand Naine displaying lower total phenol
202 content in healthy plants which increased, but to a lower level after BBTv infection. Hence,
203 the increased phenolics in the infected plant may be contributing to the resistance against
204 the infection of viral pathogens [27].

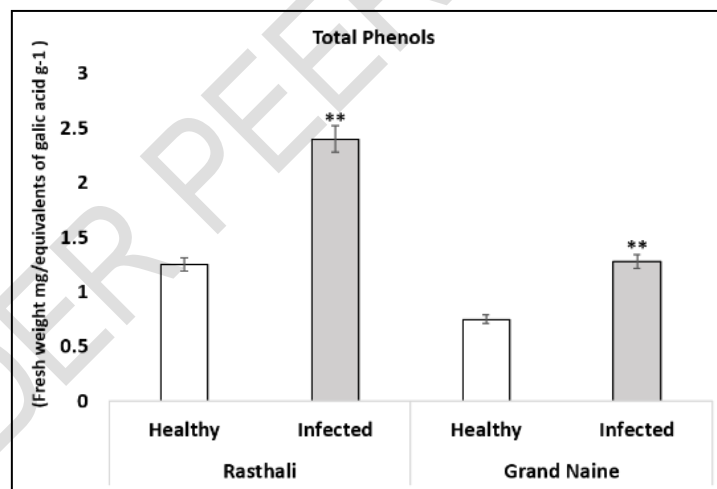
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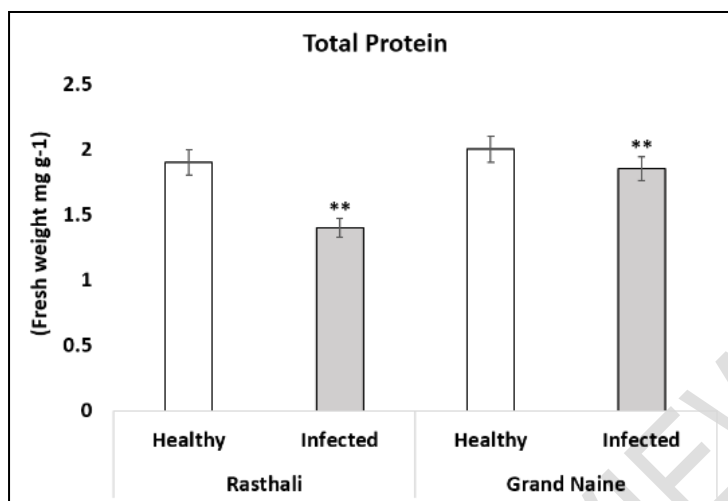
206 3.5 Total Protein

207 Protein content was found to decrease significantly in the BBTv infected plants of both
208 cultivars (Fig. 4b). The involvement of host proteins in disease resistance has been
209 demonstrated in various plant pathogenic interactions [15, 28]. The results obtained are in
210 accordance with the results of Tobacco mosaic virus-infected tobacco plants [29], Tomato
211 yellow leaf curl virus-infected tomato plants [30], Banana bunchy top virus-infected cultivars
212 of banana [31], geminivirus infected *Capsicum annum* [32] and cotton with CLCuBuV [33].

213

(a)





214

(b)

215 **Fig. 4. (a) Total phenol content and (b) Total protein of healthy and infected Rasthali**
 216 **and Grand Naine plants.**

217 *Data represents the mean \pm standard error of mean of four independent replications. Significant*
 218 *differences in healthy and infected from each cultivar analysed by Student's t test (* P <0.05, ** P <0.01)*
 219 *are shown.*

220 **3.6 Enzyme activities**

221 **3.6.1 Peroxidase**

222 **Peroxidase** activity was increased significantly in BBTv infected plants of both cultivars, in
 223 comparison to healthy (Fig. 6). In an earlier report, a similar increase in the activity of POX
 224 was observed in Virupakshi and Grand Naine cultivar [21]. The peroxidases are enzymes
 225 whose primary function is to oxidize hydrogen donors at the expense of peroxides which is
 226 known to be involved in oxidative damage in response to stress to the plant. POX activity
 227 was found to be increased in chilli against chilli leaf curl virus as reported by [34].

228

229 **3.6.2 Poly Phenol Oxidase**

230 Polyphenol oxidase is involved in the formation of insoluble polyphenols in plants by the
 231 oxidation of soluble phenols. Higher poly phenol oxidase activity was observed in Rasthali in
 232 healthy plants which marginally reduced after BBTv infection, in contrast, Grand Naine
 233 showed lower activity in healthy plants which increased during BBTv infection (Fig. 6).
 234 Higher total soluble phenols, together with higher PPO **have been demonstrated to** play a
 235 role in resistance to viral pathogens [12,21].

236

237 **3.6.3 Catalase**

238 A significant elevation was observed in the CAT activity of BBTv-infected samples in both
 239 banana cultivars tested (Fig. 5). Changes in catalase activity have been found to be a

240 significant monitoring index for plant responses under abiotic or biotic conditions. An
241 elevation in the CAT activity was reported in leaves of *Arachis hypogaea* infected with
242 Peanut mottle virus [35] and cotton plants infected with the Cotton leaf curl burewala virus
243 [33].

244 **3.6.4 Ascorbate Peroxidase**

245 The activity of ascorbate peroxidase was significantly higher in BBTV infected plants of both
246 cultivars when compared to the healthy (Fig. 5). Peroxidases acts as an antioxidant
247 response activated by the increasing presence of H₂O₂ within cells. Among the major
248 peroxide detoxifying systems in plant cells ascorbate-glutathione cycle, is the one in which
249 ascorbate peroxidase enzyme plays a key role catalyzing the conversion of H₂O₂ into H₂O.
250 The increase in APX activity in BBTV infected banana was similar to reports of *Hibiscus*
251 *cannabinus* infected with begomovirus *Nicotiana benthamiana* infected with Pepper mild
252 mottle virus [36] and sunflower infected with sunflower chlorotic mottle virus [37].

253 **3.6.5 Guaiacol Peroxidase**

254 Guaiacol peroxidase (GPX) activity was recorded to be significantly higher in BBTV infected
255 cultivars when compared with healthy (Fig. 5 and 6). GPX is an essential group from
256 peroxidase enzymes, which oxidize guaiacol and are found in cellular cytoplasm and
257 apoplasm fractions. It is involved in a range of processes related to plant growth and
258 development. GPX activity was found be higher in mesta plants infected with yellow vein
259 mosaic virus as reported by [38]

260 **3.6.6 Superoxide dismutase**

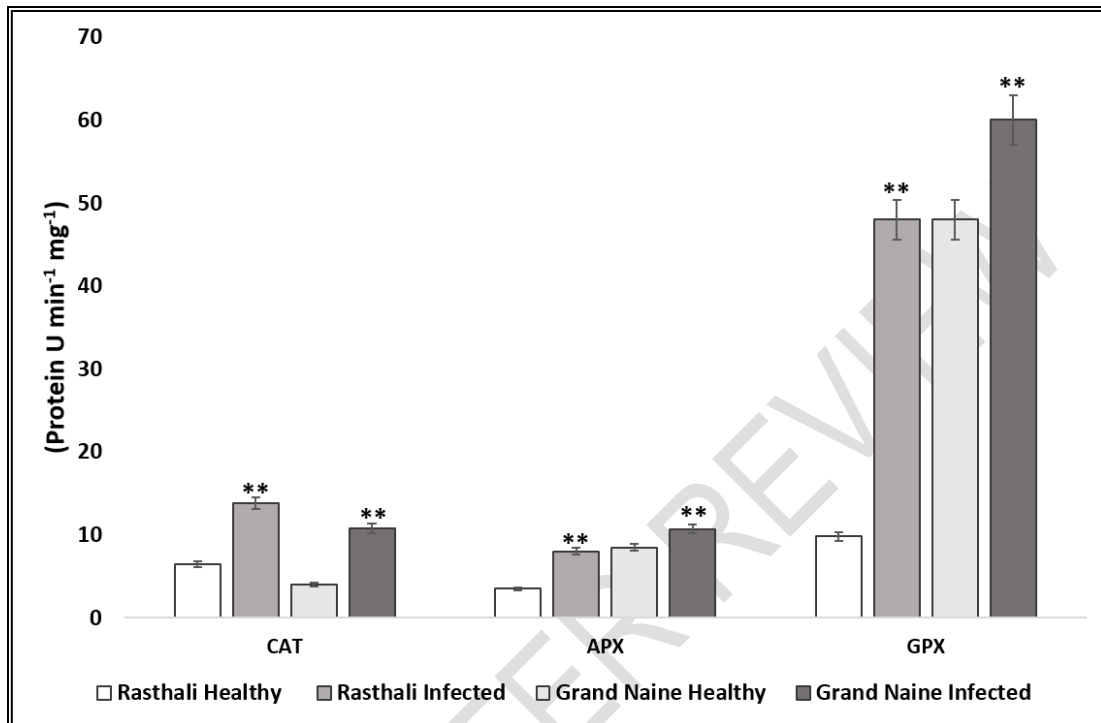
261 Superoxide dismutase (SOD) is an enzyme that breaks down superoxide radical generated
262 during stress into molecular oxygen or hydrogen peroxide thereby preventing cell damage.
263 SOD activity was significantly higher in the leaves of healthy plants of Rasthali in contrast to
264 healthy plants of Grand Naine (Fig. 6). Upon BBTV infection, Rasthali showed increased
265 SOD activity, whereas, Grand Naine showed decrease in SOD activity. In contrast, early
266 reports showed an increase in SOD activity upon BBTV infection in Grand Naine [21]. SOD
267 constitutes the front-line of defense against ROS and oxidative stress in plant cells and also
268 is one of the important scavenging enzymes. It is also reported that the induction of
269 antioxidant enzymes, including SOD, is vital for the development of plant stress tolerance.
270 Based on the present result, it can be concluded that higher SOD activity in Rasthali
271 compared to Grand Naine might contribute to increased level of tolerance to BBTV infection
272 in Rasthali and Grand Naine.

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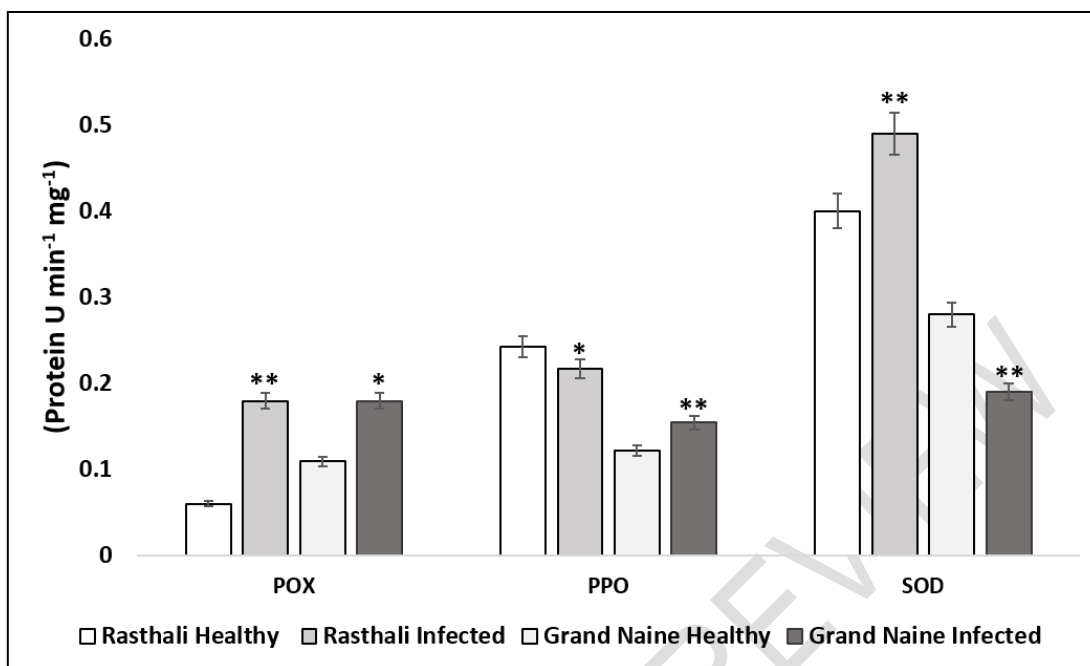
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Fig. 5. Changes in enzyme activities of CAT, APX and GPX in healthy and BBTV infected Rasthali and Grand naine banana plants.
Data represents the mean \pm standard error of mean of four independent replications. Significant differences in healthy and infected from each cultivar analysed by Student's t test ($P < 0.05$, ** $P < 0.01$) are shown.*



285

286 **Fig.6. Changes in enzyme activities of POX, PPO and SOD in healthy and BBTV**
 287 **infected Rasthali and Grand Naine banana plants.**

288 *Data represents the mean \pm standard error of mean of four independent replications. Significant*
 289 *differences in healthy and infected from each cultivar analysed by Student's t test (* P <0.05, ** P <0.01)*
 290 *are shown.*

291 **4. CONCLUSION**

292 It is well known that plant defense mechanism is complex, and the evolution of new strains
 293 of pathogens makes it a very difficult task to study. Various physiological and biochemical
 294 parameters were analyzed in BBTV infected and healthy banana cultivars Grand Naine and
 295 Rasthali. Our results indicated significant increase in defense enzyme activities in the BBTV
 296 infected cultivars compared to the healthy. There was a significant increase in amount of
 297 phenol and polyphenols in Rasthali in comparison to Grand Naine. The level of difference of
 298 biochemical constituents between the genotypes reverberates the variation of genotypes in
 299 defense against the BBTV. The findings of this study will help in better understanding of
 300 various physiological changes that occur in banana species against the BBTV and will
 301 contribute to plant resistance mechanisms which in turn will provide new tools for crop
 302 improvement.

303

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310 REFERENCES

- 311 1. Tornero P, Chao RA, Luthin WN, Goff SA, Dangl JL. Large-scale structure–
312 function analysis of the Arabidopsis RPM1 disease resistance protein. The Plant
313 Cell. 2002;14(2):435-50.
- 314 2. Jones JD, Dangl JL. The plant immune system. Nature. 2006;444(7117):323.
- 315 3. Vanitha SC, Niranjana SR, Umesha S. Role of phenylalanine ammonia lyase and
316 polyphenol oxidase in host resistance to bacterial wilt of tomato. J. Phytopathology.
317 2009;157(9):552-7.
- 318 4. Zhao CJ, Wang AR, Shi YJ, Wang LQ, Liu WD, Wang ZH, Lu GD. Identification of
319 defense-related genes in rice responding to challenge by *Rhizoctonia solani*. Theor.
320 Appl. Genet. 2008;116(4):501-16.
- 321 5. Doyle JJ, Doyle JL. 1987. A rapid DNA isolation procedure from small quantity of
322 fresh leaf material. Phytochemical Bulletin. 1987;119: 11–15.
- 323 6. Hiscox JD, Israelstam GF. A method for the extraction of chlorophyll from leaf
324 tissue without maceration. Can J Bot. 1979 Jun 15;57(12):1332-1334.
- 325 7. Dubois M, Gilles KA, Hamilton JK, Rebers PT, Smith F. Colorimetric method for
326 determination of sugars and related substances. Anal Chem. 1956;28(3):350-6.
- 327 8. McCready RM, Guggolz J, Silveira V, Owens HS. Determination of starch and
328 amylose in vegetables. Anal Chem. 1950;22(9):1156-8.
- 329 9. Folin O, Ciocalteu V. On Tyrosine and Tryptophan determinations in proteins. J
330 Bio chem. 1927;73(2):627-50.
- 331 10. Bradford MM. A rapid and sensitive method for the quantitation of microgram
332 quantities of protein utilizing the principle of protein-dye binding. Ana biochem.
333 1976;72(1-2):248-54.
- 334 11. Malick CP, Singh MB. Plants enzymology. New Delhi: Kalyani Publishers.
335 Anubis, with *Schistosoma mansoni* scadinarian. Journal of laboratory animal.
336 1980;34:119-26.
- 337 12. Ngadze E, Icishahayo D, Coutinho TA, Van der Waals JE. Role of polyphenol
338 oxidase, peroxidase, phenylalanine ammonia lyase, chlorogenic acid, and total
339 soluble phenols in resistance of potatoes to soft rot. Plant Dis. 2012;96(2):186-92.

- 340 13. Maechly AC, Chance B. The assay of catalase and peroxidase. Methods of
341 Biochemical Analysis. Interscience Inc., New York. 1954:357-424.
- 342 14. Chen GX, Asada K. Ascorbate peroxidase in tea leaves: occurrence of two
343 isozymes and the differences in their enzymatic and molecular properties.
344 Plant Cell Physiol. 1989;30(7):987-98.
- 345 15. Upadhyaya A, Sankhla D, Davis TD, Sankhla N, Smith BN. Effect of
346 paclobutrazol on the activities of some enzymes of activated oxygen metabolism and
347 lipid peroxidation in senescing soybean leaves. J. pl. physiol. 1985;121(5):453-61.
- 348 16. Dhindsa RS, Plumb-Dhindsa P, Thorpe TA. Leaf senescence: correlated with
349 increased levels of membrane permeability and lipid peroxidation, and decreased
350 levels of superoxide dismutase and catalase. J Exp Bot. 1981;32(1):93-101.
- 351 17. Goodman RN, Király Z, Zaitlin M. The biochemistry and physiology of infectious
352 plant disease. The biochemistry and physiology of infectious plant disease. 1967.
- 353 18. Balachandran S, Hurry VM, Kelley SE, Osmond CB, Robinson SA, Rohozinski J,
354 Seaton GG, Sims DA. Concepts of plant biotic stress. Some insights into the stress
355 physiology of virus-infected plants, from the perspective of photosynthesis. Physiol
356 Plant. 1997;100(2):203-13.
- 357 19. Sinha A, Srivastava M. Biochemical changes in mungbean plants infected by
358 Mungbean yellow mosaic virus. Int J Virol. 2010;6(3):150-7.
- 359 20. Zhuang, Jun, Wenwu L, Christopher JC, Pengxiang S, Taiyun W, Zujjian W,
360 Laihui X. Cleavage of the Babuvirus movement protein B4 into functional peptides
361 capable of host factor conjugation is required for virulence. Virologica sinica. 2019. 1-
362 11.
- 363 21. Anuradha C, Selvarajan R, Vasantha S, Suresha GS. Biochemical
364 Characterization of Compatible Plant Virus Interaction: A Case Study with Bunchy
365 Top Virus-Banana Host-Pathosystem. 2015;14 (4): 212-222.
- 366 22. Gaddam SA, Kotakadi VS, Reddy MN, Saigopal DV. Antigenic relationships of
367 citrus yellow mosaic virus by immunological methods. Asian J. Plant Sci. Res.
368 2012;566-9.
- 369 23. Jain AK, Yadava HS. Biochemical Constituents of Finger Millet Genotypes
370 Associated with Resistance to Blast Caused by *Pyricularia grisea* Sacc. Ann Plant
371 Prot Sci. 2003;11(1):70-4.

- 372 24. Kushwaha KP, Narain U. Biochemical changes in pigeon-pea leaves infested
373 with *Alternaria tenuissima*. Ann Plant Prot Sci. 2005:13(2):415-7.
- 374 25. Parashar A, Lodha P. Phenolic estimation in *Foeniculum vulgare* infected with
375 ramularia blight. Annals of Plant Protection Sciences. 2007:15(2):396-8.
- 376 26. Niyongere C, Ateka E, Losenge T, Blomme G, Lepoint P. Screening Musa
377 genotypes for Banana Bunchy top disease resistance in Burundi. Acta horta.2011:
378 897.
- 379 27. Manohar Jebakumar R, Selvarajan R. Biopriming of micropropagated banana
380 plants at pre-or post-BBTV inoculation stage with rhizosphere and endophytic
381 bacteria determines their ability to induce systemic resistance against BBTV in
382 cultivar Grand Naine. Biocontrol Sci Technol. 2018:28(11):1074-90.
- 383 28. Carvalho DD, Ferreira RA, Oliveira LM, Oliveira AF, Gemaque RC. Proteins and
384 isozymes electrophoresis in seeds of *Copaifera Langsdorffii* Desf. (*Leguminosae*
385 *caesalpinioideae*) artificially aged. Revista Árvore. 2006:30(1):19-24.
- 386 29. Király Z, Barna B, Kecskés A, Fodor J. Down-regulation of antioxidative capacity
387 in a transgenic tobacco which fails to develop acquired resistance to necrotization
388 caused by TMV. Free Radic Res. 2002:36(9):981-91.
- 389 30. Dieng H, Satho T, Hassan AA, Aziz AT, Morales RE, Hamid SA, Miake F,
390 Abubakar S. Peroxidase activity after viral infection and whitefly infestation in juvenile
391 and mature leaves of *Solanum lycopersicum*. J Phytopathol. 2011:159(11-12):707-
392 12.
- 393 31. Devanathan M, Ramaiah M, Sundar AR, Murugan M. Changes of peroxidase and
394 polyphenol oxidase in bunchy top nana virus infected and healthy cultivars of
395 banana. AoB Plants. 2005:19(1):114.
- 396 32. Meena RK, Patni V, Arora DK. Study on phenolics and their oxidative enzyme in
397 *Capsicum annum* L. infected with Geminivirus. Asian J. Exp. Sci. 2008:22(3):307-
398 10.
- 399 33. Siddique Z, Akhtar KP, Hameed A, Sarwar N, Imran-UI-Haq, Khan SA.
400 Biochemical alterations in leaves of resistant and susceptible cotton genotypes
401 infected systemically by cotton leaf curl Burewala virus. J Plant Interact.
402 2014:9(1):702-11.

- 403 34. Rai VP, Jaiswal N, Kumar S, Singh SP, Kumar R, Rai AB. Response of total
404 phenols and peroxidase activity in Chilli exposed to pepper leaf curl virus disease.
405 *Vegetable Science*. 2010;37(1):78-80.
- 406 35. Kobeasy MI, El-Beltagi HS, El-Shazly MA, Khattab EA. Induction of resistance in
407 *Arachis hypogaea* L. against Peanut mottle virus by nitric oxide and salicylic acid.
408 *Physiological and Mol Plant Pathol*. 2011;76(2):112-8.
- 409 36. Hakmaoui A, Pérez-Bueno ML, García-Fontana B, Camejo D, Jiménez A, Sevilla
410 F, Barón M. Analysis of the antioxidant response of *Nicotiana benthamiana* to
411 infection with two strains of Pepper mild mottle virus. *J Exp Bot*. 2012;63(15):5487-
412 96.
- 413 37. Rodríguez M, Taleisnik E, Lenardon S, Lascano R. Are Sunflower chlorotic
414 mottle virus infection symptoms modulated by early increases in leaf sugar
415 concentration. *J Plant Physiol*. 2010;167(14):1137-44.
- 416 38. Chatterjee A, Ghosh SK. Alterations in biochemical components in mesta plants
417 infected with yellow vein mosaic disease. *Brazilian journal of plant physiology*.
418 2008;20(4):267-75.

419 **ABBREVIATIONS**

- 420 APX Ascorbate Peroxidase
421 CAT Catalase
422 GPX Guaiacol peroxidase
423 PPO Poly Phenol Oxidase
424 POX Peroxidase
425 SOD Superoxide dismutase