

1 **Exploiting the Efficacy of Entomopathogenic Fungi Against Common**  
2 **Floricultural Pests: A Focus on Aphids (*Myzus persicae*), Whiteflies**  
3 **(*Bemisia tabaci B-biotype*) and Western Flower Thrips (*Frankliniella***  
4 ***occidentalis*)**

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7 **Abstract**

8 This study evaluated the efficacy of entomopathogenic fungi (EPF) as biocontrol agents  
9 against aphids, whiteflies and western flower thrips. The research employed a leaf disc bioassay  
10 with various conidia concentrations to determine lethal concentration (LC) and time (LT) for  
11 pest eradication. Additionally, the study assessed the activity of cuticle-degrading enzymes  
12 produced by EPF (Chitinase, Protease, and Lipase) to understand their pathogenic mechanisms.  
13 Molecular identification using ITS region of 18S rDNA identified virulent isolates. Results  
14 indicated that four isolates, ENPF-16, 24, 41 and 60, achieved significant mortality rates (95%  
15 to 100%) at a concentration of  $1 \times 10^8$  conidia/mL after nine days. *Akanthomyces* sp. (ENPF-  
16 41) exhibited the highest enzyme activity, followed by *Beauveria* sp. (ENPF-60). The virulent  
17 fungal isolates were identified as *Beauveria bassiana* and *Akanthomyces lecanii*. Among EPF,  
18 *Akanthomyces lecanii* (MT997935) displayed greater virulence against all three test insects  
19 with lower LC<sub>50</sub> and LT<sub>50</sub> values compared to other EPF. In summary, all fungal isolates  
20 induced mortality in the tested pests, but their effectiveness varied. *Akanthomyces lecanii*  
21 (MT997935) emerged as a promising biocontrol candidate due to its broad host range and  
22 strong virulence.

23 **Key words:** Biological control, Sucking pests, *Lecanicillium*, *Beauveria*, Chitinase, Cuticle  
24 degrading enzymes.

25 **Introduction**

26 Entomopathogenic fungi (EPF) are potentially the most diverse and versatile biological  
27 control agents due to their wide host range that often results in natural epizootics. An attractive  
28 feature of these fungi is that they infect by contact and act through penetration (Mondal et al.  
29 2016). They have certain advantages in pest control programs over other insect pathogens as  
30 they infect all stages of insects and directly infect through the cuticle, while other agents need  
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33 to be ingested. Mass production techniques for these fungi are simpler, easier and cheaper, and  
34 the persistent nature making them a potential candidate in pest control programs over other  
35 insect pathogens. The mode of entry and action of these fungi make them a promising option  
36 for combating sucking and piercing insects (Ramanujam et al. 2014). Approximately 750 to  
37 1,000 fungi are considered entomopathogens placed in over 100 genera many of these have  
38 great significance in insect pest management (Kachhawa, 2017; Ranadev et al. 2023). The most  
39 common and extensively studied entomopathogens are *Metarhizium* spp. *Beauveria* spp.  
40 *Nomuraea rileyi*, *Akanthomyces* spp., *Isaria* and *Hirsutella* spp. (Lacey et al. 2008; Kachhawa,  
41 2017). The *Akanthomyces* spp. and *Beauveria* spp. have been extensively used to combat  
42 different sucking pests under both greenhouse and field conditions.

43 Sucking pests, also known as sap-sucking insects, encompass aphids, thrips, whiteflies,  
44 mites, and leafhoppers. Their name stems from their piercing and sucking mouthparts,  
45 employed to extract sap from host plants. This sap removal leads to stunted growth, distorted  
46 appearance, chlorophyll loss, reduced strength, and premature leaf shedding in affected plants.  
47 Beyond transmitting diseases, some of these pests inject toxic substances into host plants  
48 during feeding, causing substantial agricultural losses (15-45%) and heightened management  
49 costs. Despite the use of approximately 2.5 million tonnes of pesticides annually, crop losses  
50 decreased from 41.1% during 1988-90 to 32.1% during 2001-03 (Dhaliwal et al. 2015). The  
51 use of pesticides has led to various problems, including the development of resistance and  
52 resurgence of sucking pests (Vandoorn and Vos, 2013) and residual toxic effects on humans,  
53 animals, insect parasites and predators. Additionally, the use of pesticides increases the cost of  
54 production. To address these challenges, it is essential to find a sustainable and environmentally  
55 friendly pest management solution.

56 Despite extensive research on biological control of insect pests, limited information exists on  
57 the biocontrol of sucking insect pests. Screening of local fungal isolates for their virulence  
58 characteristics is crucial for the success of biocontrol strategies (Sun et al. 2021). Conservation  
59 and periodic improvement of the efficacy of these biological control agents will significantly  
60 aid in crop protection and help to produce pesticide residue-free agricultural commodities,  
61 reducing the usage of pesticides in agriculture. The present study was aimed to isolate and  
62 identify the virulent native entomopathogenic fungal strains from soil and insect cadaver  
63 samples collected from various flower growing regions of India. Further the isolates were  
64 screened for biocontrol potential against the sap sucking insect pests viz. aphids, thrips and  
65 whiteflies in flower crops.

66 **Materials and Method**

67 **Sample collection and isolation of entomopathogenic fungi**

68 A systematic survey was conducted to collect soil and mummified insect samples from  
69 different locations of agro-climatic zones (eastern dry zone and southern dry zone) of  
70 Karnataka, India. The EPF isolates were isolated by insect bait method, serial dilution and  
71 plating on specific media, and directly placing the surface sterilized insect cadavers on potato  
72 dextrose agar (PDA). The spore suspension of isolates was prepared by adding 10mL 0.5 %  
73 sterile tween-80 to ten days old cultures and various concentration of conidial suspension was  
74 prepared by serial dilution. The conidial count was determined using an improved Neubauer  
75 Haemocytometer.

76

77 **Laboratory bioassay to assess the virulence of entomopathogenic fungal isolates**

78 The leaf disc bioassay, based on Nazir et al. (2018) protocol was used with slight  
79 modifications, involved using 8 cm diameter healthy gerbera leaf discs. These discs were  
80 surface sterilized with 70% alcohol and immersed in fungal spore suspensions at a  
81 concentration of  $1 \times 10^8$  conidia per mL for 10 seconds. A control group was treated with sterile  
82 distilled water. After air drying, the leaf discs were placed on sterile Petri plates with filter  
83 paper for humidity control during incubation. Each treatment had three replications. Twenty  
84 adult aphids (*Myzus persicae*), 2<sup>nd</sup> instar thrips larvae (*Frankliniella occidentalis*) and whitefly  
85 nymphs (*Bemisia tabaci B-biotype*) were placed on treated and control leaf discs using a sterile  
86 brush. The setup was incubated at  $25 \pm 1^\circ\text{C}$ . Observation for insect mortality occurred at two-  
87 day intervals from the 3<sup>rd</sup> to the 9<sup>th</sup> day after treatment (DAT). Dead insects were transferred  
88 to PDA plates and incubated at  $25 \pm 1^\circ\text{C}$  with 90% humidity to confirm fungal infection as the  
89 cause of death.

90

91 **Cuticle degrading enzyme activity of entomopathogenic fungal isolates**

92 The cuticle degrading enzyme index was determined by measuring the clear zone formed by  
93 degradation of specific organic compounds in the medium and the zone of clearance around  
94 the well was measured after five days after incubation. The enzymatic index was calculated by  
95 dividing the diameter of the halo zone by the diameter of the well, using the formula described  
96 by Bai et al. (2012). The chitinase enzyme index was determined using chitin yeast extract  
97 agar, Protease index was determined on skim milk agar and tributyrin agar was used for lipase  
98 enzyme index determination.

99 The chitinolytic activity of fungal isolates was determined by measuring the amount of  
100 reducing saccharides released from colloidal chitin using dinitro salicylic acid measuring the  
101 absorbance of the reaction mixture at 540 nm. The enzyme activity was expressed as unit per  
102 microliter. One unit of enzyme activity was defined as the amount of enzyme that released  
103 1 $\mu$ mol of N-acetyl D-glucosamine per minute under described conditions (Bai et al. 2012).  
104 Proteinase activity was estimated by measuring the absorbance at 280 nm. The number of  
105 amino acids released was calculated from a standard curve plotted against known  
106 concentrations of tyrosine (Hossain et al. 2006). Lipase activity was determined by a modified  
107 method of Pignede et al. (2018).

108

#### 109 **Determination of lethal concentration (LC<sub>50</sub>) and Lethal time (LT<sub>50</sub>)**

110 The selected fungal isolates that showed high virulence against the sucking pests of gerbera,  
111 including aphids, thrips, mites and whiteflies, were subjected to further testing to determine  
112 their lethal concentrations and lethal times. This was done at six different conidia  
113 concentrations (1 $\times$ 10<sup>4</sup>, 1 $\times$ 10<sup>5</sup>, 1 $\times$ 10<sup>6</sup>, 1 $\times$ 10<sup>7</sup>, 1 $\times$ 10<sup>8</sup> and 1 $\times$ 10<sup>9</sup> conidia per mL) using both the  
114 leaf disc method and the whole leaf method (Trinh et al. 2020).

115

#### 116 **Molecular identification of virulent entomopathogenic fungal isolates.**

117 The DNA was extracted from fungal mycelia by the cetyl trimethyl ammonium bromide  
118 (CTAB) extraction buffer (Doyle & Doyle, 1987) method, followed by purification through  
119 phenol/chloroform extraction and precipitation with isopropanol or ethanol (Ashktorab and  
120 Cohen, 1992). The ITS1 and ITS2 of regions of 18S rDNA were amplified using universal  
121 primers, ITS1 (5'-TCCGTAGGTGAACCTGCGG-3') and ITS4 (5'-  
122 TCCTCCGCTTATTGATATGC-3'), each at a concentration of 0.5  $\mu$ L (Hirata and Takamatsu,  
123 1996). The PCR products were sequenced through Sanger sequencing using outsourcing  
124 services Barcode BioSciences, an ISO-certified company based in Bangalore. The sequences  
125 were identified by using BLAST at NCBI website. Further, detailed phylogenetic analyses  
126 were conducted in MEGA 6.06 based on partial 18S rDNA gene sequences.

127

#### 128 **Data analysis**

129 The data obtained from the results was analyzed using ANOVA (Analysis of Variance) with  
130 the help of the software Web Agri Stat Package 2.0  
131 (<https://ccari.icar.gov.in/wasp2.0/index.php>, accessed on October 22, 2022), and the means  
132 were compared using post-hoc test (Duncan's multiple range test) at a 5% level. The LC<sub>50</sub> and

133 LT<sub>50</sub> values were calculated using the Probit analysis (Finney, 1971) function in the IBM SPSS  
134 Statistics v 20 software (IBM Corp., Armonk, NY, USA). The LC<sub>50</sub> values for  
135 entomopathogenic fungal isolates were considered significantly different if the 95 %  
136 confidence intervals (CIs) did not overlap with the CIs of other isolates.

137

## 138 **Results**

### 139 **Sample collection and isolation of entomopathogenic fungi**

140 In the present study, total of eighty-one fungal isolates were isolated from 26 insect cadavers  
141 and 55 soil samples and coded serially as ENPF. Most of the isolates were isolated from soil  
142 samples and few isolates were isolated from insect cadavers. The isolates were identified as  
143 *Aspergillus*, *Penicillium*, *Metarhizium*, *Beauveria*, *Trichoderma*, *Fusarium*, *Isaria* and  
144 *Hirsutella* spp. based on macro and microscopic observations (Ranadev et al. 2023).

145

### 146 **Virulence of entomopathogenic fungal isolates**

147 The results obtained from the virulence assay were diverse in terms of the mortality caused  
148 by different isolates over time, with an increase in the rate of mortality. All the  
149 entomopathogenic isolates successfully caused mycosis, starting from the third day after  
150 treatment (3 DAT) to the ninth day after treatment (9 DAT).

151 *Beauveria* isolate ENPF-60 and *Akanthomyces* isolates ENPF-24 and ENPF-41 demonstrated  
152 significantly highest aphid mortality, reaching 42.20% on the third day post-treatment. ENPF-  
153 16 recorded 33.33% mortality, while *Metarhizium* isolate ENPF-67 had the lowest at 4.44%.  
154 *Isaria*, *Aspergillus*, and *Hirsutella* isolates caused mortality ranging from 6% to 25%. Over  
155 subsequent days, mortality rates increased for all isolates, with *Akanthomyces* isolates showing  
156 pronounced mortality (75-88%) on the fifth to seventh days. *Beauveria* isolate ENPF-60  
157 exhibited significantly higher mortality (80.00% and 91.07%) on the fifth and seventh days  
158 compared to other *Beauveria* isolates. *Metarhizium* and *Aspergillus* isolates were less effective,  
159 causing 30% to 60% mortality between the fifth and ninth day. *Akanthomyces* (ENPF-24) and  
160 *Beauveria* (ENPF-60) isolate achieved the statistically highest mortality (100% and 92.00%,  
161 respectively) on the ninth day post-treatment (Table 1; Figure 3).

162 The results of the virulence assay on nymphs of whiteflies showed diverse outcomes in terms  
163 of the mortality caused by different isolates over time. Only the *Akanthomyces* isolates (ENPF-  
164 24 & ENPF-41) caused significantly higher mortality at all days after treatment (Table 1). The  
165 *Akanthomyces* isolate ENPF-41 showed the highest mortality, with 17.78, 73.33, 86.67, and  
166 100.00 per cent of the whitefly's mortality at 3, 5, 7, and 9 days after treatment, respectively.

167 *Beauveria* isolate ENPF-16 caused 22.22, 66.67, 80.00, and 91.11 per cent mortality at the  
168 same respective time points. The statistically lowest mortality of nymphs, at 33.33 percent, was  
169 reported for the *Hirsutella* isolate ENPF-58. In general, the mortality of whiteflies caused by  
170 different isolates ranged from 30 to 100 percent on the 9<sup>th</sup> day after treatment.

171 On the third day after treatment, only half of the isolates (*Beauveria*, *Akanthomyces* and  
172 *Isaria*) were successful in causing mortality in thrips, while the *Aspergillus*, *Metarhizium* and  
173 *Hirsutella* isolates failed to cause any mortality. The statistically highest percent mortality of  
174 thrips, 20% was caused by *Akanthomyces* isolate (ENPF-24), followed by *Beauveria* isolates  
175 (13.33%). The mortality rate increased as the days after treatment increased. On the 5th, 7th,  
176 and 9th day after treatment, both *Akanthomyces* isolates (ENPF-24 and ENPF-41) and  
177 *Beauveria* isolates (ENPF-16 and ENPF-60) showed significance in causing mycosis in thrips.  
178 The mortality caused by *Akanthomyces* isolates ranged from 66% to 93%, while it was around  
179 53% to 90% in *Beauveria* isolates. The *Aspergillus*, *Metarhizium* and *Hirsutella* isolates caused  
180 significantly lower mortality, ranging from 13% to 33.66% between the 5th to the 9th day after  
181 treatment. All the results related to the mortality of thrips caused by entomopathogenic fungal  
182 isolates are given in Table-1).

### 183 **Cuticle degrading enzyme activity of entomopathogenic fungal isolates**

184 Enzyme activity analysis revealed that among the sixteen EPF isolates, *Beauveria* and  
185 *Akanthomyces* isolates displayed the highest chitinase index (1.9 to 2.5), while *Hirsutella* and  
186 *Isaria* isolates showed lower chitinase indices of 1.8 and 1.4, respectively. For the proteolytic  
187 index, *Akanthomyces* spp. (ENPF-24) and *Beauveria* spp. (ENPF-16) exhibited significantly  
188 higher values of 3.4 and 3.3, followed by ENPF-60 and ENPF-41, with the lowest (1.6)  
189 recorded for *Aspergillus* spp. (ENPF-79). *Akanthomyces* spp. (ENPF-24) and *Beauveria* spp.  
190 (ENPF-16 & ENPF-16) showed similar lipolytic indices (3.3), followed by ENPF-41, while  
191 the lowest was observed in *Aspergillus* (ENPF-79) isolate. The enzyme activity index results  
192 highlighted variations in enzyme production among different isolates, even within the same  
193 genera (Figure 1).

194 The chitinase activity varied significantly among the sixteen isolates, with *Akanthomyces*  
195 (ENPF-24) showing the highest activity (0.53, 0.71, and 0.49 U/mL at 3rd, 6th, and 9th Days  
196 after incubation (DAI), respectively), followed by *Beauveria* (ENPF-60). The lowest  
197 chitinolytic activity was recorded in *Aspergillus* spp. (ENPF-79). In terms of protease activity,  
198 all sixteen isolates exhibited activities in the range of 0.55 to 1.21 U/mL. *Beauveria* (ENPF-

200 60) showed increased protease production (0.56, 1.57, and 1.49 U/mL at 3rd, 6th, and 9th DAI)  
201 followed by *Akanthomyces* (ENPF-24), while *Aspergillus* isolates had comparatively lower  
202 protease production (0.39-0.55 U/mL) than the other screened isolates. The same trend was  
203 observed for lipase activity, with *Beauveria* spp. (ENPF-16) producing the highest lipase  
204 activity (1.27 U/mL), followed by ENPF-24 (1.20 U/mL) (Figure 1).

### 205 **Lethal concentrations and lethal times for EPF isolates against sucking pests**

207 The results of the probit analysis of dose mortality of aphids 7 days after treatment showed  
208 that the LC<sub>50</sub> value for *Beauveria* isolate ENPF-16 was  $1.8 \times 10^6$  and for *Beauveria* isolate  
209 ENPF-60 was  $1.6 \times 10^6$ . Meanwhile, the LC<sub>50</sub> values for *Akanthomyces* isolates ENPF-24 and  
210 ENPF-41 were  $6.2 \times 10^6$  and  $9.4 \times 10^5$  respectively. Both *Beauveria* and *Akanthomyces* isolates  
211 showed high efficiency against aphids with low lethal concentrations. The lethal time to cause  
212 50 percent mortality in aphids was determined at  $1 \times 10^7$  conidia per mL, and the results  
213 uncovered that *Beauveria* isolate ENPF-60 had the shortest lethal time, causing 50 percent  
214 mortality in 4.21 days. *Akanthomyces lecanii* isolate ENPF-24 had a lethal time of 4.58 days.  
215 The highest lethal time was observed in *Beauveria* isolate ENPF-16, causing 50% mortality in  
216 5.64 days (Table 2). The results revealed that higher conidial spore concentrations ( $10^7$  to  $10^{10}$   
217 conidia/mL) were required to achieve 50 and 90 percent mortality in early days, compared to  
218 prolonged days of more than 10 days.

219 Probit analysis of whiteflies' lethal concentration at 7 days post-treatment showed no  
220 significant variation in LC<sub>50</sub> values for both *Akanthomyces* and *Beauveria* isolates, with all four  
221 isolates falling within the  $\times 10^5$  range. However, on the 9<sup>th</sup> day, differences emerged, notably  
222 with *Akanthomyces* isolate ENPF-41 displaying a lower LC<sub>50</sub> value ( $2.2 \times 10^4$ ). This suggests  
223 varying dose requirements (spore concentration) for mortality among entomopathogenic fungi  
224 of the same species. Survivorship analysis indicated that both *Akanthomyces* isolates achieved  
225 50 percent mortality in approximately 5.5 days, faster than *Beauveria* isolates at 6.2 days (Table  
226 2).

227 The probit analysis results on virulence of entomopathogenic fungi against 2<sup>nd</sup> instar larvae  
228 of thrips uncovered that *Akanthomyces* isolates were more effective in causing mortality of  
229 larvae of thrips at relatively lower conidial concentrations, with LC<sub>50</sub> value  $\times 10^5$  at 7 days after  
230 treatment. Meanwhile, the LC<sub>50</sub> values for *Beauveria* isolates were in the range of  $\times 10^6$  conidia  
231 per mL at 7 DAT. This trend was observed again at 9 days after treatment, with lower LC<sub>50</sub>.  
232 The results of Kaplan-Meier survival curves analysis of lethal time (50% mortality) revealed

233 that there was no significant difference between both *Akanthomyces lecanii* isolates at 7th day  
234 after treatment. The  $LT_{50}$  values for the *Akanthomyces* isolates were 5.51 and 5.95 days at a  
235 concentration of  $1 \times 10^7$  conidia per mL. Meanwhile, *Beauveria* isolates took around 6 days to  
236 cause 50% mortality in thrips (Table 2).

237

### 238 **Molecular identification of selected entomopathogenic fungal isolates**

239 The results of the phylogenetic tree analysis showed two broad distinct clusters for *Beauveria*  
240 *bassiana* isolates. The relationships obtained through pairwise sequence similarities were  
241 confirmed by the phylogenetic trees generated using different treeing methods. The partial 18S  
242 rDNA gene sequences of entomopathogenic fungal isolates ENPF-16 and ENPF-60 were  
243 identical to those of the species *Beauveria bassiana* with accession numbers MT997933 and  
244 MT997937, respectively. On the other hand, ENPF-24 and ENPF-41 were identified as  
245 *Akanthomyces lecanii* with accession numbers MT997934 and MT997935, respectively. The  
246 ENPF-16 and ENPF-60 isolates showed a similarity of nearly 93% and 98% to the *B. bassiana*  
247 strains Bb-9016 and IHBF14, respectively. Meanwhile, ENPF-24 and ENPF-41 isolates were  
248 found to be identical to *Akanthomyces lecanii* strains 6514 and 6543 with a similarity of over  
249 90 % (Figure 2).

250

### 251 **Discussion**

252 Although all the fungal isolates were able to cause mortality in *Myzus persicae*, *Bemisia*  
253 *tabaci* and *Frankliniella occidentalis* the percent mortality varied significantly among the  
254 isolates. This might be due to various factors such as the higher efficiency or virulence of  
255 strains isolated from closely related hosts against the same insects (Vu et al. 2008), variations  
256 in the production of extracellular enzymes such as protease, chitinase, lipase, endoprotease,  
257 esterase, and carboxypeptidase, and the role of mycotoxins like beauvericin, bassianolide,  
258 aphidiocolin (specific to aphids), and beauverolide (anti-immune activity) in causing mycosis  
259 (Kim et al. in 2013; Safavi, 2013; Sayed et al. in 2019). Other factors may include genetic  
260 virulence, conidia production and germination (Sani et al. 2020). The varying mortality rate  
261 among *Beauveria* and *Akanthomyces* isolates is intriguing. This variation could be due to the  
262 host specificity of the species, even though all four *Beauveria* isolates belong to the same genus  
263 there will be variation in the strain type. The host preference of the isolates might vary based  
264 on the species, or it could be attributed to differences in the production of extracellular enzymes  
265 is analyzed and published by several researchers like Senthil Kumar et al. (2015), Raheem and

266 Keridis, (2017), Sayed et al. (2019), Zhang et al. (2019), Trinh et al. (2020), Arthurs et al.,  
267 (2020) and Eski et al. (2022).

268 Insect cuticle is the foremost obstacle to pathogenesis. The entomopathogenic fungi must  
269 invade the cuticle barrier to successfully cause mycosis of insect pests (Mondal et al. 2016).  
270 Fascinatingly, results of exocellular enzymes activity by all six virulent entomopathogenic  
271 native isolates revealed that there is a variation in enzyme production of EPF isolates and even  
272 isolates belonging to the same species showed variation in enzyme production (Chitinase,  
273 Protease and lipase). The isolate ENPF-24 (*Akanthomyces lecanii*) has significantly high  
274 chitinase activity with lower protease and lipase activity whereas ENPF-60 (*B. bassiana*) has  
275 significantly highest protease activity. The variation of enzyme activity of an isolate reveals  
276 that enzyme activity varies based on their native host, virulence and gene expression (Nahar et  
277 al. 2004). The quantitative studies on enzyme production of sixteen isolates revealed that the  
278 production of the enzymes was more on the 6<sup>th</sup> day after inoculation compared with the results  
279 on the 3<sup>rd</sup> and 9<sup>th</sup> days after inoculation. This was noticed by several researchers like De Moraes  
280 et al. (2003) and Bai et al. (2012). This might be because of degradation and denaturation of  
281 enzymes produced during later stages of growth, lack of nutrients, and accumulation of other  
282 secondary metabolites (Bai et al. 2012; Demir et al. 2013).

283 Enzyme production on the 6th day was compared to the highest insect pest mortality during  
284 *in vitro* virulence studies on the 7th day. The parallel relationship between enzyme production  
285 and insect mortality indicates that the enzyme activity of isolates significantly contributes to  
286 insect mycosis, along with other factors. Cuticle-degrading enzymes from EPF, specifically  
287 chitinolytic enzymes, play a crucial role in the pathogenicity of isolates (Dhar and Kaur, 2009).  
288 This correlation is consistent with previous studies on various insect pests, such as *Plagioderia*  
289 *versicolora* (Demir et al. 2013), *Pieris brassicae* (Dhawan and Joshi, 2017) and *Myzus*  
290 *persicae* (Khan et al. 2012), confirming the importance of cuticle-degrading enzyme  
291 production in causing insect mortality.

292 The low lethal concentration and lethal time observed in *Beauveria* and *Akanthomyces*  
293 isolates may be due to their specificity, growth rate, enzyme production rate, and conidial  
294 germination (Nazir et al. 2018; Trinh et al. 2020). This is because as the spore concentration  
295 increases, the number of conidia per square area also increases, increasing the likelihood of  
296 causing mycosis in the insect pest.

297 The results of the phylogenetic analysis indicate the presence of two major distinct clusters  
298 of *B. bassiana* isolates. The molecular characterization of entomopathogenic fungal isolates

299 showed that ENPF-16 and ENPF-60 were identified as *Beauveria bassiana*, while ENPF-24  
300 and ENPF-41 were identified as *Akanthomyces lecanii*. The results are consistent with the  
301 findings of previous studies (Ozcelik and Guven, 2015), which have also identified *B. bassiana*  
302 and *Akanthomyces lecanii* as common entomopathogenic fungi in different regions. These  
303 findings provide valuable information on the distribution and identification of  
304 entomopathogenic fungi in different ecosystems, which is essential for the development of  
305 effective biocontrol strategies for insect pests.

306

### 307 **Conclusion**

308 The results imply that various entomopathogenic fungal isolates exhibit distinct levels of  
309 mortality in *Myzus persicae*, *Bemisia tabaci B-biotype* and *Frankliniella occidentalis*.  
310 *Akanthomyces* isolates prove most virulent, causing significant mortality across all three insect  
311 pests, particularly whiteflies. *Beauveria* isolates also demonstrate effectiveness against thrips  
312 and aphids. The correlation between enzyme production and insect mortality underscores the  
313 pivotal role of enzyme activity in fungal virulence. The concentration of conidial spores per  
314 milliliter significantly influences lethal concentration and time, with higher concentrations  
315 leading to increased mortality within a shorter timeframe. These findings align with prior  
316 research publications, underscoring the potential of *Akanthomyces* and *Beauveria* as biological  
317 control agents for whiteflies, thrips, and aphids. Nonetheless, further research is necessary to  
318 fully comprehend the factors contributing to the variation in lethal concentration and time  
319 among different fungal isolates.

320

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429

**Table 1.** Virulence of entomopathogenic fungal isolates on per cent cumulative and corrected mortality under *in vitro* conditions.

Insects	DAT	ENPF-3	ENPF-6	ENPF-8	ENPF-9	ENPF-16	ENPF-24	ENPF-26	ENPF-33	ENPF-41	ENPF-48	ENPF-53	ENPF-58	ENPF-60	ENPF-67	ENPF-68	ENPF-79
Aphids	3rd	26.7 (31.1) <sup>c</sup>	13.3 (21.4) <sup>e</sup>	20.0 (26.6) <sup>d</sup>	13.3 (21.4) <sup>e</sup>	33.3 (35.3) <sup>b</sup>	40.0 (39.2) <sup>a</sup>	13.3 (21.4) <sup>e</sup>	8.9 (17.3) <sup>f</sup>	40.0 (39.2) <sup>a</sup>	20.0 (26.6) <sup>d</sup>	13.3 (21.4) <sup>e</sup>	6.7 (15.0) <sup>g</sup>	42.2 (40.5) <sup>a</sup>	20.0 (26.6) <sup>d</sup>	13.3 (21.4) <sup>e</sup>	11.1 (19.4) <sup>ef</sup>
	5th	60.0 (50.8) <sup>d</sup>	33.3 (35.3) <sup>g</sup>	40.0 (39.2) <sup>f</sup>	33.3 (35.3) <sup>g</sup>	66.7 (54.7) <sup>c</sup>	73.3 (58.9) <sup>b</sup>	33.3 (35.3) <sup>g</sup>	31.1 (33.9) <sup>gh</sup>	73.3 (58.9) <sup>b</sup>	53.3 (46.9) <sup>e</sup>	31.1 (33.9) <sup>gh</sup>	26.7 (31.1) <sup>h</sup>	80.0 (63.4) <sup>a</sup>	34.0 (35.7) <sup>fg</sup>	33.3 (35.3) <sup>g</sup>	31.1 (33.9) <sup>gh</sup>
	7th	73.3 (58.9) <sup>c</sup>	44.4 (41.8) <sup>e</sup>	53.3 (46.9) <sup>d</sup>	48.9 (44.4) <sup>de</sup>	82.2 (65.0) <sup>b</sup>	86.7 (68.6) <sup>ab</sup>	46.7 (43.1) <sup>de</sup>	46.7 (43.1) <sup>de</sup>	88.9 (70.5) <sup>a</sup>	66.7 (54.7) <sup>c</sup>	53.3 (46.9) <sup>d</sup>	33.3 (35.3) <sup>f</sup>	91.1 (72.6) <sup>a</sup>	44.4 (41.8) <sup>e</sup>	53.3 (46.9) <sup>d</sup>	48.9 (44.4) <sup>de</sup>
	9th	80.0 (63.4) <sup>c</sup>	60.0 (50.8) <sup>ef</sup>	66.7 (54.7) <sup>e</sup>	57.8 (49.5) <sup>ef</sup>	95.6 (77.8) <sup>b</sup>	100.0 (90.0) <sup>a</sup>	53.3 (46.9) <sup>fgh</sup>	55.8 (48.3) <sup>fg</sup>	95.6 (77.8) <sup>b</sup>	73.3 (58.9) <sup>c</sup>	60.0 (50.8) <sup>ef</sup>	46.7 (42.8) <sup>h</sup>	100.0 (90.0) <sup>a</sup>	51.1 (45.6) <sup>gh</sup>	57.8 (49.5) <sup>ef</sup>	57.8 (49.5) <sup>ef</sup>
	<b>X</b>	<b>60.0</b>	<b>37.8</b>	<b>45.0</b>	<b>38.3</b>	<b>69.4</b>	<b>75.0</b>	<b>36.7</b>	<b>35.6</b>	<b>74.4</b>	<b>53.3</b>	<b>39.4</b>	<b>28.3</b>	<b>78.3</b>	<b>37.4</b>	<b>39.4</b>	<b>37.2</b>
Whitefly	3rd	17.8 (24.9) <sup>c</sup>	4.4 (12.2) <sup>g</sup>	15.6 (23.2) <sup>d</sup>	8.9 (17.3) <sup>e</sup>	22.2 (28.1) <sup>ab</sup>	24.4 (29.6) <sup>a</sup>	6.7 (15.0) <sup>f</sup>	6.7 (15.0) <sup>f</sup>	17.8 (24.9) <sup>c</sup>	20.0 (26.6) <sup>bc</sup>	6.7 (15.0) <sup>f</sup>	8.9 (17.3) <sup>e</sup>	24.4 (29.6) <sup>a</sup>	4.4 (12.2) <sup>g</sup>	6.7 (15.0) <sup>f</sup>	4.4 (12.2) <sup>g</sup>
	5th	46.7 (43.1) <sup>e</sup>	26.7 (31.1) <sup>g</sup>	53.3 (46.9) <sup>d</sup>	33.3 (35.3) <sup>f</sup>	66.7 (54.7) <sup>b</sup>	63.3 (52.7) <sup>bc</sup>	26.7 (31.1) <sup>g</sup>	20.0 (26.6) <sup>h</sup>	73.3 (58.9) <sup>a</sup>	53.3 (46.9) <sup>d</sup>	26.7 (31.1) <sup>g</sup>	13.3 (21.4) <sup>i</sup>	60.0 (50.8) <sup>c</sup>	22.2 (28.1) <sup>gh</sup>	33.3 (35.3) <sup>f</sup>	26.7 (31.1) <sup>g</sup>
	7th	60.0 (50.8) <sup>d</sup>	40.0 (39.2) <sup>ef</sup>	66.7 (54.7) <sup>d</sup>	46.7 (43.1) <sup>e</sup>	80.0 (63.4) <sup>b</sup>	80.0 (63.4) <sup>b</sup>	40.0 (39.2) <sup>ef</sup>	33.3 (35.3) <sup>f</sup>	86.7 (68.6) <sup>a</sup>	73.3 (58.9) <sup>c</sup>	40.0 (39.2) <sup>ef</sup>	26.7 (31.1) <sup>g</sup>	73.3 (58.9) <sup>c</sup>	40.0 (39.2) <sup>ef</sup>	40.0 (39.2) <sup>ef</sup>	33.3 (35.3) <sup>f</sup>
	9th	68.9 (56.1) <sup>d</sup>	48.9 (44.4) <sup>ef</sup>	73.3 (58.9) <sup>d</sup>	51.1 (45.6) <sup>ef</sup>	91.1 (72.7) <sup>b</sup>	93.3 (75.0) <sup>b</sup>	48.9 (44.4) <sup>ef</sup>	40.3 (39.4) <sup>gh</sup>	100.0 (90.0) <sup>a</sup>	82.2 (65.0) <sup>c</sup>	44.4 (41.8) <sup>fg</sup>	33.3 (35.3) <sup>h</sup>	84.4 (66.8) <sup>c</sup>	51.1 (45.6) <sup>ef</sup>	53.3 (46.9) <sup>e</sup>	46.7 (43.1) <sup>fg</sup>
	<b>X</b>	<b>48.3</b>	<b>30.0</b>	<b>52.2</b>	<b>35.0</b>	<b>65.0</b>	<b>65.3</b>	<b>30.6</b>	<b>25.1</b>	<b>69.4</b>	<b>57.2</b>	<b>29.4</b>	<b>20.6</b>	<b>60.6</b>	<b>29.4</b>	<b>33.3</b>	<b>27.8</b>
Thrips	3rd	13.3 (21.4) <sup>c</sup>	0.0 (0.28) <sup>f</sup>	11.1 (19.4) <sup>d</sup>	0.0 (0.28) <sup>f</sup>	11.1 (19.4) <sup>d</sup>	20.0 (26.6) <sup>a</sup>	0.0 (0.28) <sup>f</sup>	0.0 (0.28) <sup>f</sup>	15.6 (23.2) <sup>b</sup>	13.3 (21.4) <sup>c</sup>	0.0 (0.28) <sup>f</sup>	0.0 (0.28) <sup>f</sup>	13.3 (21.4) <sup>c</sup>	0.0 (0.28) <sup>f</sup>	2.2 (8.56) <sup>e</sup>	0.0 (0.28) <sup>f</sup>
	5th	53.3 (46.9) <sup>c</sup>	13.3 (21.4) <sup>fg</sup>	40.0 (39.2) <sup>d</sup>	15.6 (23.2) <sup>f</sup>	53.3 (46.9) <sup>c</sup>	66.7 (54.7) <sup>b</sup>	20.0 (26.6) <sup>e</sup>	13.3 (21.4) <sup>fg</sup>	66.7 (54.7) <sup>b</sup>	53.3 (46.9) <sup>c</sup>	20.0 (26.6) <sup>e</sup>	13.3 (21.4) <sup>fg</sup>	73.3 (58.9) <sup>a</sup>	20.0 (26.6) <sup>e</sup>	20.0 (26.6) <sup>e</sup>	11.1 (19.5) <sup>g</sup>
	7th	66.7 (54.7) <sup>b</sup>	26.7 (31.1) <sup>de</sup>	53.3 (46.9) <sup>c</sup>	28.9 (32.5) <sup>d</sup>	66.7 (54.7) <sup>b</sup>	80.0 (63.4) <sup>a</sup>	33.3 (35.3) <sup>de</sup>	26.7 (31.1) <sup>de</sup>	82.2 (65.0) <sup>a</sup>	66.7 (54.7) <sup>b</sup>	28.9 (32.5) <sup>d</sup>	20.0 (26.6) <sup>e</sup>	80.0 (63.4) <sup>a</sup>	26.7 (31.1) <sup>de</sup>	26.7 (31.1) <sup>de</sup>	20.0 (26.6) <sup>f</sup>
	9th	73.3 (58.9) <sup>d</sup>	36.7 (37.3) <sup>gd</sup>	60.0 (50.8) <sup>e</sup>	33.3 (35.3) <sup>hi</sup>	80.0 (63.4) <sup>c</sup>	93.3 (75.0) <sup>a</sup>	30.0 (33.2) <sup>ij</sup>	33.3 (35.3) <sup>hi</sup>	91.1 (72.6) <sup>b</sup>	73.3 (58.9) <sup>d</sup>	33.3 (35.3) <sup>hi</sup>	26.7 (31.1) <sup>j</sup>	90.0 (71.6) <sup>b</sup>	46.7 (43.1) <sup>f</sup>	40.0 (39.2) <sup>g</sup>	26.7 (31.1) <sup>j</sup>
	<b>X</b>	<b>51.7</b>	<b>19.2</b>	<b>41.1</b>	<b>19.4</b>	<b>52.8</b>	<b>65.0</b>	<b>20.8</b>	<b>18.3</b>	<b>63.9</b>	<b>51.7</b>	<b>20.6</b>	<b>15.0</b>	<b>64.2</b>	<b>23.3</b>	<b>22.2</b>	<b>14.4</b>

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**Note:** Values in the parentheses are ARCSINE transferred values. The values represented by same letters in each row are statistically on par with each other by DMRT mean of three replications at 95 % confidence interval (CI). **X**; Mean

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**Table 2.** Virulence (LC<sub>50</sub> & LT<sub>50</sub>) of entomopathogenic fungal isolates to *M. persicae*, *B. tabaci* and *Frankliniella occidentalis*.

Insect pest	Fungus	$\chi^2$	LC <sub>50</sub> <sup>1</sup> (Conidia/mL)	95% CI <sup>3</sup>		$\chi^2$	LT <sub>50</sub> (±SE) <sup>2</sup> (Days)	95% CI <sup>3</sup>	
				Lower	Upper			Lower	Upper
<b>Aphids</b> ( <i>Myzus persicae</i> )	<i>B. bassiana</i> (ENPF-16)	1.20	1.8×10 <sup>6</sup>	2.6×10 <sup>5</sup>	3.7×10 <sup>7</sup>	0.78	5.47±0.1	4.47	6.57
	<i>B. bassiana</i> (ENPF 60)	2.54	1.6×10 <sup>6</sup>	3.3×10 <sup>5</sup>	4.9×10 <sup>7</sup>	1.28	5.06±0.2	3.98	6.02
	<i>A. lecanii</i> (ENPF-24)	0.68	9.0×10 <sup>6</sup>	1.5×10 <sup>6</sup>	7.8×10 <sup>7</sup>	0.69	4.59±0.1	3.17	5.76
	<i>A. lecanii</i> (ENPF-41)	1.24	9.4×10 <sup>5</sup>	1.0×10 <sup>5</sup>	2.3×10 <sup>7</sup>	0.20	4.83±0.1	3.59	5.83
<b>Whitefly</b> ( <i>Bemisia tabaci</i> )	<i>B. bassiana</i> (ENPF-16)	3.23	6.5×10 <sup>5</sup>	4.8×10 <sup>4</sup>	7.3×10 <sup>6</sup>	2.26	6.46±0.2	5.12	7.56
	<i>B. bassiana</i> (ENPF 60)	2.46	8.4×10 <sup>5</sup>	6.2×10 <sup>4</sup>	1.2×10 <sup>7</sup>	1.55	6.18±0.1	5.06	7.38
	<i>A. lecanii</i> (ENPF-24)	1.67	6.3×10 <sup>5</sup>	4.4×10 <sup>4</sup>	7.4×10 <sup>6</sup>	0.87	5.78±0.1	4.57	6.69
	<i>A. lecanii</i> (ENPF-41)	2.14	1.5×10 <sup>5</sup>	9.3×10 <sup>3</sup>	6.3×10 <sup>6</sup>	0.95	5.89±0.1	4.71	6.83
<b>Thrips</b> ( <i>Frankliniella occidentalis</i> )	<i>B. bassiana</i> (ENPF-16)	0.32	4.9×10 <sup>6</sup>	3.5×10 <sup>5</sup>	1.0×10 <sup>8</sup>	1.03	6.76±0.2	5.68	7.71
	<i>B. bassiana</i> (ENPF 60)	0.09	3.8×10 <sup>6</sup>	3.0×10 <sup>5</sup>	6.0×10 <sup>7</sup>	1.63	6.17±0.2	4.87	7.22
	<i>A. lecanii</i> (ENPF-24)	1.76	2.3×10 <sup>5</sup>	1.4×10 <sup>4</sup>	6.2×10 <sup>6</sup>	2.60	5.51±0.1	4.48	6.46
	<i>A. lecanii</i> (ENPF-41)	1.49	1.5×10 <sup>5</sup>	8.9×10 <sup>3</sup>	7.9×10 <sup>6</sup>	0.93	5.94±0.1	4.75	6.89

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**Notes:** <sup>1</sup>The LC<sub>50</sub> values were calculated by Probit analysis using IBM SPSS v 20.0 from the mortality data collected from 7 various conidia concentration at 7 days after treatment.

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<sup>2</sup>LT<sub>50</sub> values for mortality were estimated by survivorship analysis (Kaplan-Meier survival curves) with censored data for insects surviving >8d incubation period in bioassay studies

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and survival curves were compared using the log-rank test  $\chi^2$  (chi-square) value at P ≤ 0.05. <sup>3</sup>95% Confidence intervals that did not overlap indicate differences between LC<sub>50</sub> and LT<sub>50</sub>

436

values.

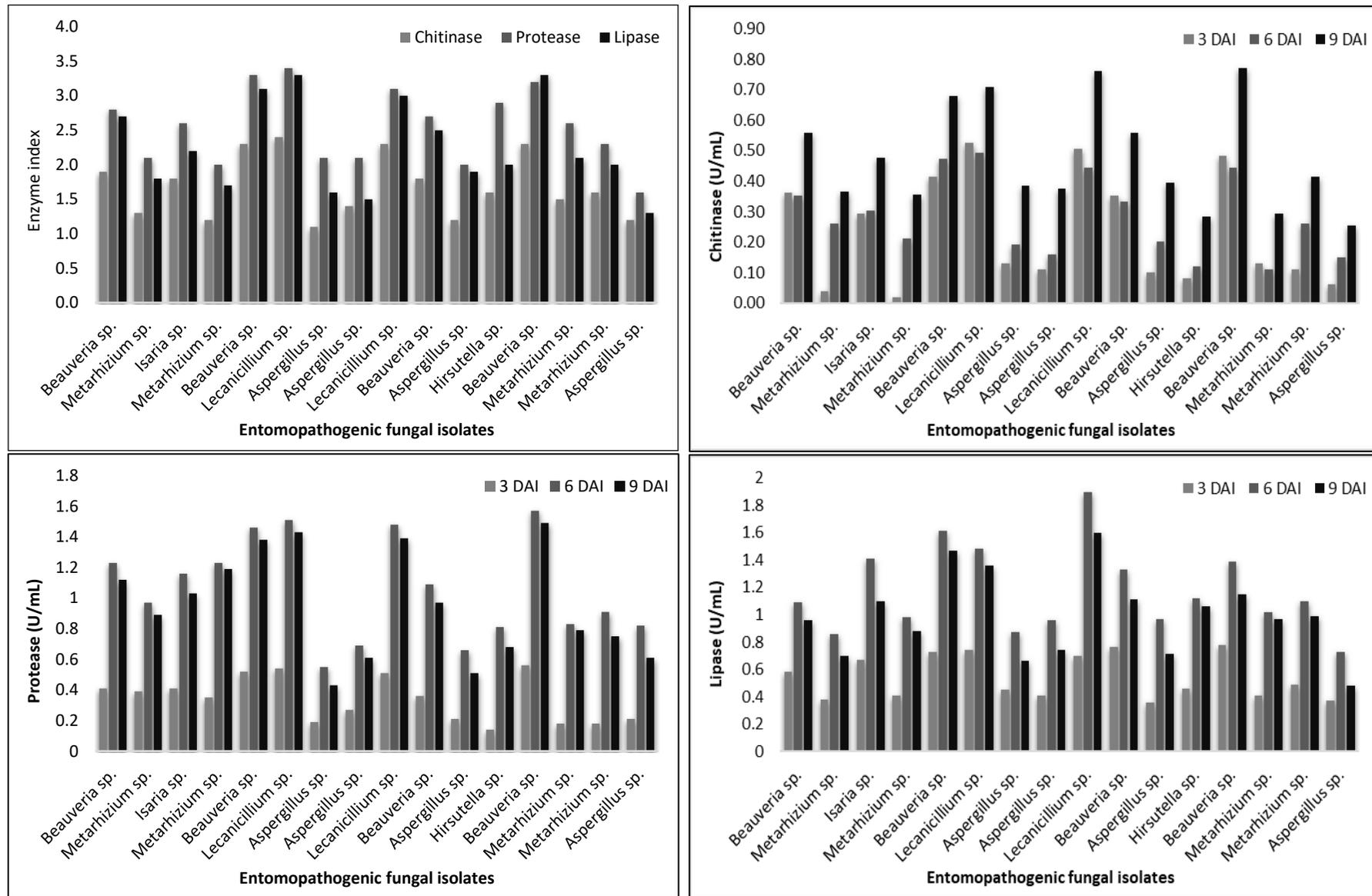
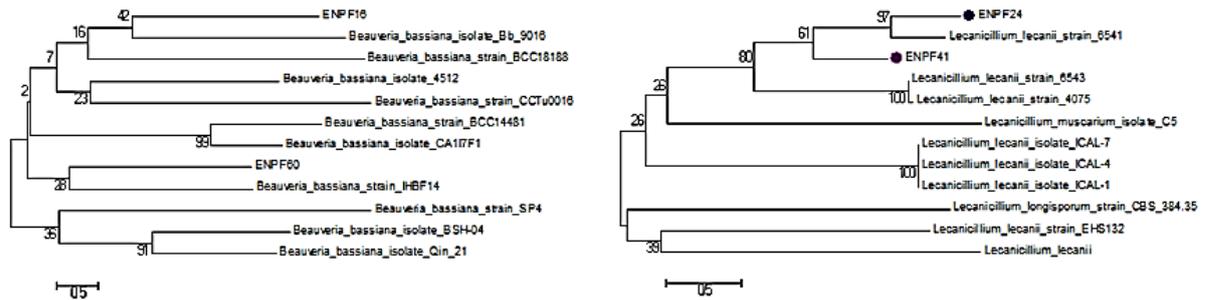
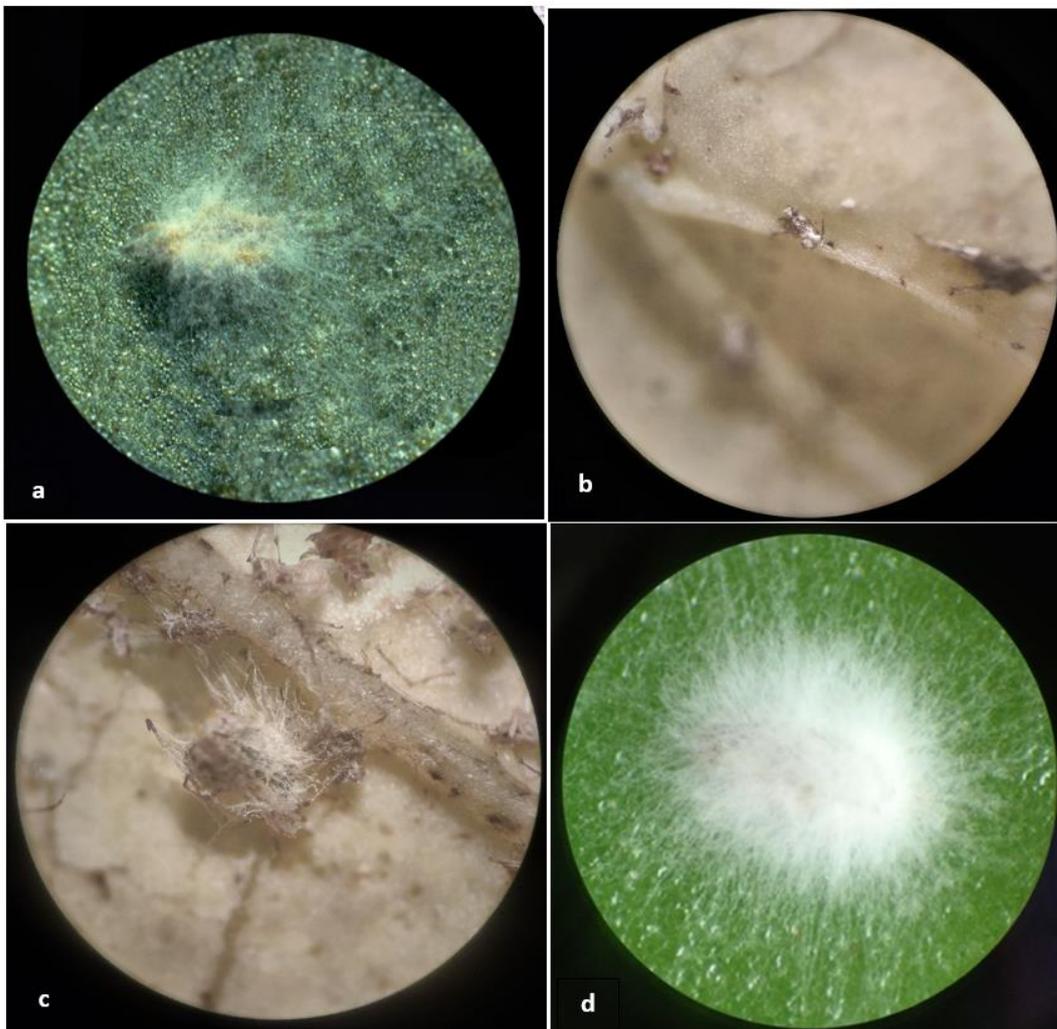


Figure 1: Cuticle degrading enzyme activity of entomopathogenic fungal isolates (Days after incubation).



**Figure 2.** A. Phylogenetic placement of *Beauveria* isolates B. Phylogenetic placement of *Lecanicillium* isolates, based on nearly full-length 18S rRNA gene sequences. The tree was calculated with the Neighbour-joining method. Bootstrap values are based on 1,000 replicates. Numbers shown above branches are bootstrap percentages for clades supported above the 70% level. The bar indicates 5% sequence divergence.



457 **Figure 3.** Mortality of various insect sucking insect pests caused by EPF isolates. *a*:  
 458 *Akanthomyces sp.* infected cadaver of aphid. *b*: *Beauveria sp.* infected cadaver of thrips. *c*:  
 459 *Akanthomyces sp.* infected cadaver of thrips. *d*: *Akanthomyces sp.* infected cadaver of whitefly.