BOTRYOSPHAERIACEAE FUNGAL SPECIES AS POTENTIAL PATHOGENS OF MELIACEAE IN THE ARID AND SEMI-ARID LANDS OF KENYA

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ABSTRACT

Botryosphaeriaceae are significant and cosmopolitan pathogens worldwide. In Kenya, the fungal family has been reported on *Grevillea robusta* on-farm. Two *Meliaceae* tree species, exotic *Azadirachta indica* (A. Juss.) and indigenous *Melia volkensii* (Geurke) were sampled across Kenya. Symptoms found in the field included shoot die-back (45%), canker 14%, both canker and dieback (25%) while 16% of the trees sampled were healthy. *Botryosphaeriaceae* was most prevalent in *Melia volkensii* mortality rate of 28%. DNA results of the ITS and Tef1-alpha gene region for 86 isolates classified into 6 species of the *Botryosphaeriaceae*. 3 of these species belonged to the genus *Lasiodiplodia* namely *L. pseudotheobromae*, *L. theobromae* and *L. parva*. This is the first report of species *Spencermartinsia viticola* and *Macrophoma theicola* in Kenya. Pathogenicity tests done under glass house conditions showed that the *L. pseudotheobromae* species was most virulent to both *M. volkensii* and *A. indica* while *L. theobromae* was least virulent to both tree species. Wilting and necrosis was recorded within 7 days of inoculation but wound healing occurred on both species after 12 weeks. This study gave insight into disease resistance and tolerance of these dryland species for plantation establishment. It also revealed wider host diversity for *Botryosphaeriaceae* in the drylands.

Key words: Botryosphaeriaceae, Meliaceae, Disease tolerance.

Introduction

Over 60% of Kenya's land areas has been classified as arid or semi-arid under agro-ecological zoning. These areas receive low rainfall 250-500 mm annually, have high minimum and maximum temperatures 23-24°C and low to poor fertility of soils. The vastness of these areas in Kenya has led to afforestation efforts in a bid to create a green wall and prevent further desertification and as adaptation strategies to the effects of climate change. Meliaceae is a family of leguminous trees that consists of trees, shrubs and sometimes shrublets. It has about 50 genera and 650 species in tropical, subtropical and sometimes warm temperate regions of both hemispheres. The family has unique characteristics in the chemicals present in its plants i.e. limonoids, alkaloids and phenols that have been seen to contribute to the trees resistance to insect damage (Agrios, 2005). Most commonly planted species of Meliaceae in Kenya include native Melia volkensii and exotic Azadirachta indica (unpublished data) which have been widely planted in the ASALs as drought resistant and termite resistant tree species. They also have multiple uses such as timber production, fodder for livestock, medicinal use in treating several ailments. Botryosphaeria

species have been a common isolate as a causative agent of tree diseases in Kenya for the past 7 years. Botryosphaeria is a genera of fungi that has many members and has been found to be endophytic to many tree species (Slippers et al., 2004a). Its first report was on Grevillea robusta (Njuguna et al., 2011) common used as a shade tree for tea plantations and alley cropping. It has also been isolated from Eucalyptus tree species. The family contains 17 genera and over 150 species of fungus with a wide host range across the globe (Phillips et al., 2013). As environmental conditions worsen due to changing weather patterns this family of Fungi is crucial as it becomes a virulent pathogen at the onset of physiological stress. Presence of the pathogen on some existing plantations in the country also poses the risk of cross infection of native tree species from the exotic tree plantations. The present study was a survey on the presence of Botryosphaeriaceae on the existing plantations of representative species of Meliaceae and recommend management and control measures to be taken in controlling the disease and preventing an epidemic.

Material and Methods

248 trees in total of both natural stands and

Representatives of *Meliaceae* in Kenya showed susceptibility to species of *Botryosphaeriaceae* with symptoms of canker and die-back but showed resistance/tolerance during the pathogenicity test and have been recommended for climate change mitigation.

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woodlots were sampled. Disease incidence and severity was scored by two people standing 5 m away from the tree and assessing the percentage of the tree's above ground biomass covered by symptoms such as stem and branch cankers, shoot and tip dieback, bark cracking as well as gum exudation. Disease incidence was scored as a percentage derived from the portion of trees that exhibited disease symptoms out of the total number of trees in the plantation or cluster used. Samples of twigs and bark were then collected using sterilized secateurs and blades and taken to the lab for analysis. Tools used to collect samples were sterilized with 70% ethanol to avoid infecting other trees. Trees whose trunks or stems were scraped were sprayed with fungicide to prevent reinfection by other pathogens.

Laboratory analysis

Both moist chambering and surface sterilization were used to obtain cultures from the samples collected. For moist chamberingthe cut sections were placed in surface sterilized glass petri dishes. The petri dishes were then placed in sterile tins containing wet blotting papers and the tin covered to form a damp chamber to encourage microbe growth. The tins were left at room temperature (21°C) for at least 5 days until the fungi grows on the sampled material. In surface sterilization Sections of both symptomatic and asymptomatic tissue were cut into 6 small pieces washed with 33% Hydrogen peroxide for 2 minutes before rinsing them three times with distilled water. The pieces were then dried by blotting on sterile filter papers under a sterile laminar flow bench before being placed onto 2% Malt Extract Agar (Sigma-Aldrich) which was amended with 0.05% Streptomycin (Sigma-Aldrich) to inhibit bacterial growth. The petridishes were then sealed with Parafilm and incubated at 25°C in 12 hour alternating light and dark cycles. Mycelial growth was monitored and colonies developed counted after 3 days. Conidial Morphology was studied under a microscope and used to classify the isolates into their genus level. These characteristics were studied under a compound microscope to define the shape, form, size, pigmentation (colour), elevation, opacity and the colony morphology to identify the pathogen.

DNA based identification

Three day old mycelia was grinded using a mechanical grinder and steel balls to make a fine powder. Tris was then added to the mixture and incubated for 50 minutes in a warm bath. CTAB was then added to the mixture and incubated at 40°C for 30 minutes before adding 70% Sodium Chloride. PCR amplification using specific primers for ITS and Tef1-alpha were used. The

sequences were then analyzed using Gene Sequencer 5000 and the sequences edited to match them to the nearest possible species. Instead that sequences were edited using MEGA 4.0 software and a cladogram produced to show the same. A series of reactions with specific chemicals and reagents is used to extract rDNA from cells of the fungi which is then analyzed using a sequencer machine into bands made of base pairs of DNA. The downloaded sequences are then compared with existing sequences from Genbank and other culture collections to find the closest relative at 100%.

Results and Discussion

50.3% of the isolated fungal species were made up of Botryosphaeriaceae followed by Nectriaceae (32.3%) and then Pestalotiopsis (3.6%) Botryosphaeria species was also isolated from healthy trees characterizing it as an endophyte that would become pathogenic in the event of physiological stress on the tree. Well silviculturally managed trees had less disease symptoms than those in the wild.

Grand Total	247
Botryosphaeria sp.	125
Fusarium sp.	63
Pestalotia sp.	22
Nectria sp.	18
Phomopsis sp.	4
Alternaria sp.	3
Phoma sp.	3

Disease severity was higher in Melia volkensii than in Azadirachta indica however, there was no mortality recorded for A. indica and a few recorded for M. volkensii.

DNA results

Melia Volkensii isolates	Azadirachta indica isolates
Alternaria alternata	Cladosporium flabelliforme
Aspergillus flavus	Cytosporasp.
Diaporthe isoberliniae	Fusarium oxysporum
Diaporthe macintoshii	Lasiodiplodia crassispora
Lasiodiplodia crassispora	Lasiodiplodia parva
Lasiodiplodia parva	Lasiodiplodia pseudotheobromae
Lasiodiplodia parva /	Lasiodiplodia pseudotheobromae
<i>Botryosphaeria</i> sp	
Lasiodiplodia pseudotheobromae	Lasiodiplodia sp. KR1
Lasiodiplodiasp. KR1	Lasiodiplodia theobromae
Lasiodiplodia theobromae	Lasiodiplodia theobromae
Lasiodiplodia theobromae	Macrophoma theicola
Neofusicoccum parvum	Pestalotiopsissp.
Nigrosporasp.	
Penicillium chrysogenum	Phomasp.
Pestalotiopsissp.	
Spencermartinsia viticola	
Talaromyces purpureogenus	

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Aspergillus_niger_isolate_13E06[1
Penicillium_sp_J41221
Penicillium_chrysogenum
Aspengillus_flavus_voucher_RIFA_29-R1
Penicillium_sp_27_BRO-2013
 Penicillium_sp_F02
Phaeoacremonium_scolyti_voucher_DUCC406
Phaeoacremonium_tu:canum_strain_1Pal
 Penicillium_griscofalvum_strain_F22
  Sigmoidea_sp_C-4
 Trichodarma_gamsii_clone_2
   Microsphaeropsis_anundinis_isolate_P21-169
    Microphaeropos, amindosis, isotate, Jr.2
Phoma, pp., P107
Epicocum, sp., 22MS01
Curvolaria, sp., RFA, 73B
Phoma, sp., P48E6
A hemaria, alternata _strain_9C
A hemaria, solani_strain_UPT-10
are inn_decorollulars_toolpe_48.0V
   Fusarium, decemceflulare, isolate, 48.IV
   Facarium_solani_isolate_9_EL1-4
 Phomopus_sp_F122
   Diaporthe_macintoshii_strain_BRIP_55064a
   Diaporthe_isoberliniae_strain_CPC_22549
   Phomopsis_sp_CML_1937
    Pestalotiopsis_sp_PG52
    Pestalotiopsis_sp_TR061
  Postalotiopsis_sp_C6b2b
Postalotiopsis_sp_SD6tES
   Fusarium_oxysponum_strain_LCF32
Fusarium_sp_4_BRO-2013
    Fusarium_sp_G338
    Fusarium_oxysporum_strain_CHS-2
   Fusarium_sp_IBL_03157
   Cytospora_sp_T39_JC-2014
Cytospora_sp_T_AE-2013_strain_F4839
    Nigrospora, sp., TA26-42
Cladosporium_flabelliforme_inolate_P83-452
Cladosporium_cladosporioides_strain_F207
     Diescorea_alata_cultivar_GuangxiYulin_shunyao_ckme_1
     asiodiplodia_crassispora_strain_UCD27Co
   Las iodiplodia_crassispom_stmin_UCD24Co
 Las iodiplodia_crassispora_strain_CMM_0283
 Fusarium_sp_Vega(89
Fusarium_solari_strain_fse4
  Nectria_balansae_stmin_AR4446
 Tiarosporella_tritici_strain_CBS_118719
   Pseudofunicoccum_ofivaceum_strain_WTS_58
   Neofunicocoum_sp_M109
Neofunicocoum_sp_M28
    Neotusicoccum_parvum_isolate_A564
   Spencermartinsia_viticola_strain_UCP134
Dothiordla_iberica_strain_UASWS0893
   Dothiorella_longicollis_strain_CBS_122068
Dothiorella_sp_FIIVW-2008a_strain_CBS_121765
   Alanphillipsia_aloeigena_strain_CPC_21286
    Lasiodiplodia_theobromae_isolate_MBAi39CL
   Lasiodiplodia sp.
   Las iodiplodia_crassispora_strain_xsd08010
   Lasiodiplodia_sp._P34E1
Lasiodiplodia_theobromae_isolate_81.I
   Laciodiplodia_pseudotheobromae_isolate_7_DL5-2
   Laciodiplodia_pseudotheobromae_strain_B0361
   Lasiodiplodia_theobromae_strain_xsd08007
Macrophoma_sp_VPM-12
   Macrophoma_theicola_stmin_UPA-62
   Lasiodiplodia_theobromae_voucher_MAC_Karu-4
   Lusiodiplodia purva strain CBS 456.78
   Lasiodiplodia_perva_strain_CMW28295
   Lasiodiplodia theobromae strain Gr428
 Lasiodiplodia_puva_strain_CMW28333

    Diplodia_corticola_strais_CBS678_b

Talaromy ces_purpureo genus_strain_ZB076
Aspergillus flavus
                                                              - Diplodia_corticola_strain_CBS678_a
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Botryosphaeriaceae Theiss & P. Syd. (Fungi, Acsomycota, Pezizomycotina, Dothideomycetes, Botryosphaeriales) is a family of fungi with a cosmopolitan distribution across the globe except in the Polar Regions known to cause severe die-back and canker on many

woody plants. The characteristic of some of its species as endophytes makes a crucial species of interest especially as more and more wood lots are being established in lands that already experience moisture stress.

Lasiodiplodia, Dothiorella, Macrophoma, Sphaeropsis, Fusicoccum, Diplodia and Botryodiplodia include some of the most common genera linked to Botryosphaeria (Denman et al., 2000).

The family has 17 identifiable genera and over 103 species known from culture (Phillips et al., 2013). Species from this family have been isolated from native, exotic and naturalized tree species in most tropical countries from both plantation and agro-forestry systems. Many genera of plants including angiosperms and conifers are susceptible to Botryosphaeria attack. In Kenya Botryosphaeriaceae has been isolated from many exotic tree species commonly established in plantations across the country. A major predisposing factor has been poor species site matching by planting tree species outside their optimal ecological zone limit. These species include Eucalyptus camaldulensis, Eucalyptus clones, Grevillea robusta, Azadirachta indica, Senna siamea. An attack and epidemic of Diplodia pini (renamed to Sphaeropsis sapinea) on Pine plantations in the country led to the ban of planting of *Pinus radiata* in eastern Africa. Known to have some chemical properties that make them disease resistant *Meliaceae* continued to fight the symptoms of the disease. The high disease severity but no mortality in some of the trees sampled was seen as perceived disease resistance of the target tree species.

Conclusion

Control measures against *Botryosphaeriaceae* species has been proper species-site matching during tree establishment and management of trees planted to ensure optimum growth conditions with little or no physiological stress on the plants. This reduces the susceptibility of the plants to infection by *Botryosphaeriaceae*. Treatment of pruning wounds with fungicides has also proved to be helpful in control of new infections in plantations. Once affected the mechanical removal and burning of infected stumps is advised to remove the source of inoculums which can lead to an epidemic through wide spread of the fungal infection. Further research should be done on these tree species with field trials to ascertain the perceived versus the real threats for seedlings once planted in the field.

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केन्या के शुष्क एवं अर्ध-शुष्क भूमियों में मीलिएसीया के एक सक्षम रोग जनक के रूप में बॉट्रीओस्फेरिएसीया कवकी प्रजाति एंजीला एम. मुथेमा, जेनी डब्ल्यू जूगूना एवं फ्रान्सिस के. सांग

बॉट्रीओस्फेरिएसीया विश्वभर में महत्वपूर्ण एवं सर्वदेशीय रोगजनक हैं। केन्या में कवक कुल फार्म में ग्रीविलीया रॉबुस्टा पर सूचित किया गया है। दो *मीलिएसीया* वृक्ष प्रजातियों, विदेशज *ऐजैडिरैक्टा इंडिका* (ए. जस) और देशज *मीलिया वोल्कीन्सी* (गूर्की) का सम्पूर्ण केन्या में सैम्पल किया गया। क्षेत्र में पाए गए रोग लक्षण में प्ररोह पश्च क्षय (45%), कैंकर (14%), कैंकर और पश्च क्षय दोनों (25%), शामिल थे जबिक सैम्पल किए गए 16 प्रतिशत वृक्ष स्वस्थ थे। बॉट्टीओस्फेरिएसीया 28 प्रतिशत की मर्त्यता दर के साथ *मीलिया वोल्कीन्सी* में सबसे ज्यादा प्रचलित था। 86 आइसोलेटों के लिए इसके जीन क्षेत्र के डी.एन.ए. परिणामों को *बॉट्टीओस्फेरिएसीया* की 6 प्रजातियों में वर्गीकृत किया गया। इन प्रजातियों में से 3 जीनस *लेसिओडिप्लोडिया*, यथा- एल. स्युडोथीओब्रोमेई, एल. थीओब्रोमेई और एल. पर्वा से संबंधित हैं। केन्या में प्रजाति स्पेन्सरमार्टिन्सिए विटिकोला और मैक्रोफोमे थीकोला की यह पहली रिपोर्ट है। ग्लास हाउस अवस्थाओं के तहत रोग जनकता जाँच ने दर्शाया कि स्यूडोथीओब्रोमेई प्रजाति *मीलिया वोल्कीन्सी* एवं *ऐजैडिरैक्टा इंडिका* दोनों के लिए सबसे विषाक्त थी जबकि एल. *थीओब्रोमेई* दोनों वृक्ष प्रजातियों के लिए न्युनतम विषाक्त थी। संरोपण के 7 दिन की भीतर म्लानी और ऊतक क्षय अभिलिखित किया गया किन्त दोनों प्रजातियों में 12 हफ्ते के उपरान्त घाव भरने शरू हए। इस अध्ययन में रोपण स्थापना के लिए इन शुष्क भूमि प्रजातियों के रोग प्रतिरोध एवं सहनशीलता में अन्तर्दृष्टि दिया। इसने शुष्क भूमियों में *बॉट्रीओस्फेरिएसीया* के लिए व्यापक परपोषी विविधता को भी उद्घाटित किया।

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