## **RESEARCH ARTICLE**

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# Nuclear and structural dynamics during the establishment of a specialized effector-secreting cell by *Magnaporthe oryzae* in living rice cells

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### Abstract

**Background:** To cause an economically important blast disease on rice, the filamentous fungus *Magnaporthe oryzae* forms a specialized infection structure, called an appressorium, to penetrate host cells. Once inside host cells, the fungus produces a filamentous primary hypha that differentiates into multicellular bulbous invasive hyphae (IH), which are surrounded by a host-derived membrane. These hyphae secrete cytoplasmic effectors that enter host cells presumably via the biotrophic interfacial complex (BIC). The first IH cell, also known as the side BIC-associated cell, is a specialized effector-secreting cell essential for a successful infection. This study aims to determine cellular processes that lead to the development of this effector-secreting first IH cell inside susceptible rice cells.

**Results:** Using live-cell confocal imaging, we determined a series of cellular events by which the appressorium gives rise to the first IH cell in live rice cells. The filamentous primary hypha extended from the appressorium and underwent asymmetric swelling at its apex. The single nucleus in the appressorium divided, and then one nucleus migrated into the swollen apex. Septation occurred in the filamentous region of the primary hypha, establishing the first IH cell. The tip BIC that was initially associated with the primary hypha became the side BIC on the swollen apex prior to nuclear division in the appressorium. The average distance between the early side BIC and the nearest nucleus in the appressorium was estimated to be more than 32 µm. These results suggest an unknown mechanism by which effectors that are expressed in the appressorium are transported through the primary hypha for their secretion into the distantly located BIC. When *M. oryzae* was inoculated on heat-killed rice cells, penetration proceeded as normal, but there was no differentiation of a bulbous IH cell, suggesting its specialization for establishment of biotrophic infection.

**Conclusions:** Our studies reveal cellular dynamics associated with the development of the effector-secreting first IH cell. Our data raise new mechanistic questions concerning hyphal differentiation in response to host environmental cues and effector trafficking from the appressorium to the BIC.

**Keywords:** Appressorium, Biotrophic interfacial complex, Biotrophy, Effector, Live-cell imaging, Mitosis, Rice blast fungus, Hyphal differentiation

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#### Background

The filamentous ascomycete fungus Magnaporthe oryzae causes the economically important blast disease on rice and other crops [1]. Several key cellular events that occur during rice infection have been documented (Fig. 1a). On the rice leaf surface, the fungus forms a specialized infection structure, called an appressorium, which produces a narrow penetration peg to breach the plant surface [2]. Upon penetration, the peg expands to form a filamentous primary hypha that grows into the rice cell and subsequently differentiates into bulbous invasive hyphae (IH) with a pseudohyphal appearance due to regular constrictions at presumed septa [3, 4]. Cell cycle regulation is important for the appressorium development and IH proliferation [5–7]. Semi-closed mitosis has been proposed to occur in M. oryzae, during which the nuclear envelope remains intact, but nuclear pore complexes partially dissociate in the appressorium and IH [8, 9]. The primary hypha

22-26 hpi

and IH invade living host cells (biotrophy) while contained within a host-derived extra-invasive hyphal membrane (EIHM) [4, 10]. The IH grow throughout the first invaded cell before invading adjacent cells [4] (Fig. 1a). The IH colonize each new cell biotrophically, but eventually kill host cells (necrotrophy); the invaded cells lose viability by the time the IH move into adjacent cells [4, 11].

During the biotrophic invasion, the primary hypha and IH secrete both cytoplasmic effector proteins (effectors; i.e., Pwl2, Bas1, Bas107, and AvrPiz-t) that enter the host cytoplasm and apoplastic effectors (i.e., Bas4, Bas113, and Slp1) that are retained in the EIHM compartment [10, 12–17]. These effectors are known or presumed to suppress host defense and facilitate infection [10, 14, 16]. Cytoplasmic effectors are preferentially localized in the biotrophic interfacial complex (BIC), which has been hypothesized as the site of effector translocation into the host cytoplasm [12, 18]. Two stages of BICs have

36-40 hpi



26-28 hpi

been previously described [12] (Fig. 1a). The tip BIC is associated with the apex of the primary hypha. As the hypha develops, the BIC is positioned on the side of the first IH cell, and thus it is called the side BIC. The first IH cell, also known as the side BIC-associated cell, is a specialized effector-secreting cell essential for a successful infection. Giraldo et al. (2013) have shown that this IH cell uses two distinct pathways for effector secretion. One is the conventional secretory pathway for secretion of apoplastic effectors into the EIHM compartment; the other is the exocyst components Exo70 and Sec5-involved pathway for secretion of cytoplasmic effectors into the BIC. In addition, this first IH cell is critical in its role as the mother cell of the subsequent biotrophic IH cells.

Although our knowledge of early biotrophic invasion has increased, questions remain to be answered, such as how the first IH cell develops inside live host cells, what mechanisms control hyphal differentiation, and what cellular processes are involved in BIC development and effector trafficking. In this study, we provide insights into some of these questions by using live-cell imaging of *M. oryzae* invasion of susceptible rice cells. Our studies reveal a series of cellular events associated with the first IH cell development. These studies lead to new mechanistic questions concerning *in planta* hyphal differentiation and effector trafficking.

#### **Results and discussion**

#### Apical swelling of the primary hypha precedes mitosis

We generated *M. oryzae* transformants constitutively expressing EYFP (labeling the cytoplasm with green fluorescence) and histone H1-tdTomato fusion protein (H1-tdTomato; labeling nuclei with red fluorescence). These transformants exhibited wild-type morphology and pathogenicity (Additional file 1). To characterize the development of the first IH cell upon host penetration (transition between the first two stages in Fig. 1a), we used one of these transformants, CKF2138, and imaged 124 random infection sites from 20 separate inoculations and microscopy sessions at 22-28 h post inoculation (hpi). Quantitative analysis of these images revealed three sequential growth stages (Fig. 1b): (1) the filamentous primary hypha with one nucleus in the appressorium, (2) the apically bulbous primary hypha with one nucleus in the appressorium, and (3) the apically bulbous primary hypha that contains a nucleus with another nucleus in the appressorium. We quantified these observations by measuring the diameter of the primary hypha close to the appressorium and the diameter at the apical region of the primary hypha before or after swelling (Fig. 1c). At all growth stages, the diameter of the primary hypha close to the appressorium was conserved with a mean of 2.3  $\mu$ m (standard deviation = 0.4  $\mu$ m, n = 124). In infection sites with the anucleate primary hypha and a single nucleus present in the appressorium, the diameter of the primary hypha tip had a mean of 3.1  $\mu$ m (standard deviation = 1.0  $\mu$ m, *n* = 55). In infection sites displaying one nucleus in the appressorium and another one in the bulbous portion of the primary hypha, the diameter of the bulbous portion had a mean of 5.6  $\mu$ m (standard deviation = 0.9  $\mu$ m, *n* = 69), frequently ranging from 4.0 to 6.9  $\mu$ m. Taken together, these data indicate that nuclear division and migration typically occur after the apical tip of the primary hypha has swollen to over 4  $\mu$ m in diameter.

## Mitosis in the appressorium and subsequent migration of a single nucleus into the first bulbous IH cell

To determine nuclear dynamics during development of the first IH cell, we conducted time-lapse microscopy of rice cells invaded by M. oryzae mitotic reporter strain CKF1962 expressing H1-tdTomato and GFPnuclear localization signal (NLS) [8]. Consistent with the previous study [9], after plant penetration, nuclear division occurred in the appressorium, and subsequently one mitotic nucleus migrated through the primary hypha to arrive in the swollen portion of the primary hypha, while the other nucleus remained in the appressorium (Fig. 2a; n = 14). The timing of nuclear division was consistent with our prediction that nuclear division occurs in infections that had a swollen primary hypha tip lacking a nucleus (Fig. 1c). Intriguingly, the nucleus that remained in the appressorium appeared to associate with the penetration pore as the other nucleus migrated through the primary hypha to the swollen hypha tip (Fig. 2a; n = 19). Our observation of nuclear division within the appressorium contrasts a previous report, in which the appressorial nucleus migrates to the primary hypha and then undergoes mitosis [5]. We considered the possibility that the discrepancy might have arisen because different strains of M. oryzae and rice were used, specifically M. oryzae O-137-derived strain (H1-tdTomato) infecting rice strain YT16 in our study and M. oryzae Guy11-derived strain (H1-tdTomato) infecting rice strain CO-39 in the other study [5]. However, we ruled out this possibility because we observed the consistent pattern of nucleus division within the appressorium followed by migration of a single nucleus into the primary hypha when M. oryzae Guy11-derived strain (H1-tdTomato) infects YT16 (Fig. 2b; n = 4) or CO-39 (Fig. 2c; n = 3). The reason for this discrepancy remains to be determined. It is worth noting, however, that our findings are consistent with Colletotrichum gloeosporioides, in which nuclear division occurs in the appressorium, and then one nucleus migrates into the penetration hypha, whereas the other nucleus remains in the appressorium [19].



## Formation of the first septum after nuclear division and migration

To define the location and timing of septation during development of the first IH cell, we generated *M. oryzae* transformants expressing fluorescently labeled septa together with H1-tdTomato. The septa were visualized by constitutively expressing the plasma membrane-targeted PLC delta:PH as a translational fusion to GFP (GFP-PLC delta:PH) in *M. oryzae*. PLC delta:PH binds to PI(4,5)P2 lipids [20] and has been shown to localize to the plasma membrane and septa in *Neurospora crassa* [21]. *M. oryzae* transformants expressing GFP-PLC delta:PH showed green fluorescence localized at the plasma membrane of the appressorium and primary hypha (Fig. 3 top). After nuclear division, one nucleus moved to the bulbous tip of the primary hypha, and the other nucleus remained in the appressorium (Fig. 3 middle). Subsequently, the green fluorescence localized near the midpoint of the filamentous



**Fig. 3** The first post-penetration septum forms in the primary hypha. Shown are maximum intensity projections of confocal optical sections of a rice cell invaded by *M. oryzae* strain CKF2686 at 27 hpi. Merged nuclear tdTomato (*red*) and GFP fused to PLCdelta:PH (*green*) are shown. *Top:* One nucleus (*arrowhead*) was located in the appressorium (outlined by *dotted line*) and GFP-PLCdelta:PH was localized to the plasma membrane. *Middle:* The same infection site imaged 50 min later showing one appressorial nucleus and one nucleus in the bulbous hypha. *Bottom:* The same infection site after an additional 25 min showing GFP fluorescence concentrating at the location of the first septum (*arrow*). *Bar* = 5 µm

portion of the primary hypha, indicating the site of the first septum (Fig. 3 bottom). The primary hypha was previously characterized by Heath et al. (1990) and Kankanala et al. (2007): it is a filamentous structure that differentiates from the penetration peg, an extremely narrow structure that originates at the appressorium and forces through the plant cell wall, and that subsequently differentiates into bulbous IH. Our septation result further reveals that the primary hypha is a filamentous part of two distinct cell types, constituting the invasive extension of the single-celled appressorium and the first IH cell.

#### Development of BICs during hyphal differentiation

Two-stage development of BICs has been previously documented [12] (Fig. 1a): the tip BIC is associated with the apex of the primary hypha; as the primary hypha switches to bulbous IH growth, the BIC maintains its position on what is now the side of the bulbous IH. To determine how the tip BIC becomes the side BIC, we used our earlier observations about the time of nuclear division relative to hyphal swelling (Figs. 1c and 2) to design an experiment. We imaged M. oryzae transformant CKF1651 expressing cytoplasmic EYFP and nuclear mRFP as well as mRFP fused to the BIC-localized cytoplasmic effector Pwl2 [12] at different growth stages analogous to the three panels in Fig. 1b. In cases of no bulbous differentiation, the BIC was at the tip of the primary hypha (Fig. 4a left). When bulbous differentiation had occurred, but nuclear division had not, the BIC was already at the side, indicating that the BIC is displaced from its tip position as the primary hypha swells (Fig. 4a middle). In infections where nuclear division and migration had taken place, we observed the BIC still at the side, where it remained as a new hypha began to grow at the tip (Fig. 4a right). These results indicate that the bulbous swelling of the primary hypha occurs prior to nuclear division and is responsible for the BIC being "left" on the side of the cell. This displacement of the BIC indicates that there is depolarization at the apex of the primary hypha during bulbous differentiation. The position of the BIC on one side of the BIC-associated cell suggests that the opposite side of the primary hypha



**Fig. 4** BIC development. **a** Tip to side BIC differentiation and first nuclear division event at 26 hpi. Shown are maximum intensity projections of confocal optical sections (18.9–28.5  $\mu$ m in total depth). Images are of *M. oryzae* CKF1651, expressing cytoplasmic EYFP (*green*), nuclear mRFP (*red*), and BIC-localized mRFP (*red*; cytoplasmic effector Pwl2 fused to mRFP). *Arrows* = BICs. *Arrowheads* = nuclei. *Asterisk* = appressorium. *Left*: Tip BIC stage. *Middle*: Side BIC stage. *Right*: After nuclear division, the side BIC remained at the same position. *Bars* = 5  $\mu$ m. **b** Representative measurement from the penetration point to the side BIC in *M. oryzae* CKF1616 at 29 hpi. Images show BIC-localized mCherry (*red*; cytoplasmic effector Pwl2 fused to mCherry:NLS), fungal hypha outlined by EGFP (*green*; apoplastic effector Bas4 fused to EGFP), and mCherry and EGFP overlapping in the side BIC (*yellow*). *Top*: A maximum intensity projection of confocal optical sections (23  $\mu$ m in total depth) and 2-dimensional length (*gray line*; length = 31.4  $\mu$ m) measured with the open Bezier tool in the Zen Black software. *Bottom*: A 3-dimensional reconstruction of the same infection site shows 3D length (*gray line*; distance = 33.2  $\mu$ m) measured with the Imaris dendrite creation algorithm (still image of the last frame of Additional file 2: Movie S1). *Bars* = 5  $\mu$ m. **c** Comparative graphical summary of measurements taken with 3D Imaris and with 2D Zen Black software of the distance between the penetration point to the side BIC. Individual infection sites (*black dots*) have both a 3D measurement (x-axis) and a corresponding 2D measurement (y-axis). The vertical distance between a data point and the 1-to-1 line (*blue*) on the graph corresponds to the difference in length between the 2D and 3D measurements associated with that data point

grows more than the BIC side. This asymmetric swelling produces an oval cell offset from the center of the hypha (Figs. 1 and 4) rather than a round cell, which would result from equal swelling in all directions. This growth is clearly distinct from the polarized growth observed during filamentous primary hypha development, of which the tip BIC is characteristic. Polarity in filamentous fungi is established and maintained by multiple proteins, especially those associated with control of cytoskeletal positioning [22]. The Spitzenkörper (Spk) and polarisome have been established as instrumental in determining the direction of hyphal tip growth in filamentous fungi [23-25]. In M. oryzae, Spk-associated motor protein Mlc1 is localized to the growing tip of the primary hypha before differentiation of the BIC-associated cell [18]. After differentiation, Mlc1 is not present at the growing IH tip, but can be visualized near septa and in the subapical BIC-associated cell [18]. In contrast, the polarisome component Spa2 is localized to the growing tip of the primary hypha and IH [18]. Because an IH cell develops from the growing end of the BIC-associated cell, there must be a repolarization event after asymmetric swelling occurs. The Spa2 and other polarisome proteins are likely involved in repolarization of the IH growing tip.

## Regular growth from the appressorium penetration to the side BIC

The filamentous primary hypha appeared to extend from the appressorium to a consistent distance before widening at its apical region. We reasoned that the side BIC would be positioned at a defined distance from the appressorium. To test this, we measured the distance from the appressorium penetration point to the side BIC using *M. oryzae* transformant CKF1616, expressing the BIC-localized cytoplasmic effector Pwl2 fused to mCherry:NLS and also IH-outlining apoplastic effector Bas4 fused to EGFP [12]. The penetration point was identified by examining the EGFP fluorescence outlining the primary hypha in conjunction with the bright-field, and the BIC was identified based on the strongly localized mCherry fluorescence. We conducted two-dimensional measurements of these images

using the Zen Black open Bezier tool (Version 8.1, Zeiss). The distance along the primary hypha from the penetration point to the side BIC has an average of 36  $\mu$ m, ranging from 30 to 45  $\mu$ m (*n* = 22) (Fig. 4b top; Fig. 4c). To obtain more accurate measurements of the same infection sites, we also used the three-dimensional dendrite measurement algorithm in Imaris (Version 7.6, Bitplane). An additional movie file shows the threedimensional measurement in more detail (see Additional file 2: Movie S1). The average distance was 43  $\mu$ m, ranging from 32 to 54  $\mu$ m (*n* = 22) (Fig. 4b bottom; Fig. 4c; Additional file 2: Movie S1). Our finding of this considerable distance raises an intriguing question concerning effector secretion. Cytoplasmic effectors such as Pwl2 are secreted into the tip BIC and the side BIC, which are located more than 32  $\mu m$ from the appressorial nucleus (Fig. 4a left and middle; Fig. 4b-c) in which the PWL2 gene is expressed (Zhu and Khang, unpublished). There must be a mechanism to transport effectors from the appressorium through the primary hypha for their secretion to the distantly located BIC, but the mechanism remains to be determined.

#### Hyphal growth inside heat-killed rice cells

We hypothesized that the development we observed during the early biotrophic invasion is specific to live host cells. To test this, we used a heat-killed cell assay, inoculating nonviable rice tissue with conidia. When inoculated on this tissue, M. oryzae produced the normal appressorium (Fig. 5), indicating that the surface cues were still sufficient for appressorium differentiation. After penetration into heat-killed host cells, the fungus did not show distinct differentiation from filamentous to bulbous growth as observed during invasion of live host cells, but elongated, filamentous-like growth throughout the rice cell and its neighbors (Fig. 5). Hyphae growing in heat-killed cells spread rapidly into adjacent cells rather than completely filling the first cell, which the IH in living cells do over the course of 12 h [4]. In living host cells, BICs are easily recognized in bright-field images due to their predictable locations on the tip of the primary hypha, the side of the first bulbous IH cell, and the tips and sides of hyphae moving into adjacent host cells [12] (Fig. 1a). We did not observe any indications of BICs in the bright-field images of infections in heat-killed cells. The absence of BICs in dead host cells is consistent with Giraldo et al. (2013) showing that BICs are derived from host membrane components such as the plasma membrane and endoplasmic reticulum; therefore, BICs are unlikely to form in host cells, which lack intact membranes.



(green) and nuclear tdTomato (red, but shown in yellow due to overlap of green and red), growth in heat-killed rice cells at 25 hpi. Shown are merged fluorescence confocal images of a single focal plane (*top* and *middle*), or a projection (29  $\mu$ m in total depth) (*bottom*) of the same infection site. Rice cell walls are shown as *white lines. Top:* A single focal plane shows the appressorium (*asterisk*) with a nucleus, germ tube (*star*), and hypha (*x*) that grew on the rice surface. *Middle:* A focal plane 12  $\mu$ m beneath the appressorium shows 2 filamentous-like IH branches (*plus signs*) and locations where IH had crossed the rice cell wall into neighboring cells (*arrows*). *Bottom:* A maximum intensity projection shows all features of both focal planes (*top* and *middle*). *Bar* = 10  $\mu$ m

#### Conclusions

We provide cellular details of early post-penetration development of *M. oryzae* during invasion of susceptible rice cells. This study defines developmental stages, in which the single-celled appressorium gives rise to the first IH cell, and the tip BIC becomes the side BIC when the tip BIC-associated hyphal apex undergoes asymmetric swelling prior to mitosis in the appressorium (Fig. 6). Development of BICs and BIC-associated cells is induced during invasion of live rice cells. Our study provides a descriptive framework for early biotrophic invasion of rice leaf sheath tissue. It remains to be determined whether the cellular processes reported in this study are conserved during infection of other rice tissues and of other plant species. This study also leads to new mechanistic questions concerning effector trafficking from the appressorium to the BIC and hyphal differentiation in response to host environmental cues.



**Fig. 6** Schematic diagram of early post-penetration growth in susceptible rice cells. Pre-penetration: *M. oryzae* conidium (*white*; three-celled) germinates and forms an appressorium (*circle*) containing one nucleus (*blue circle*). Post-penetration: (Stage 1) The appressorium penetrates the host cell wall, forming an anucleate filamentous primary hypha (*gray*) with the tip BIC (*red*). (Stage 2) The tip of the filamentous primary hypha swells and the tip BIC becomes the side BIC (*red*). The primary hypha remains anucleate. (Stage 3) Nuclear division occurs in the appressorium. (Stage 4) One mitotic nucleus migrates into the bulbous portion of the primary hyphal tip, while the other nucleus remains in the appressorium. (Stage 5) The first septum forms in the filamentous portion of the effector-secreting BIC-associated cell. (Stage 6) Polarized growth resumes from the BIC-associated cell

#### Methods

#### Plasmids, strains, and fungal transformation

For nuclear-localized tandem-dimer Tomato (tdTomato), the binary plasmid pCK1287 was constructed by cloning the histone H1 gene from N. crassa (H1) at the 5' end of the tdTomato gene under control of the M. oryzae ribosomal protein 27 (P27) promoter as follows: the 2.2 kb EcoRI-BamHI fragment (P27:H1) isolated from pBV229 and the 1.7 kb BamHI-HindIII fragment (tdTomato:Nos terminator) isolated from pBV359 were ligated between EcoRI and HindIII sites of the binary vector pBV1 (pBHt2) [26]. For the membrane-localized GFP, the binary plasmid pCK1417 was constructed by cloning the following three fragments between EcoRI-HindIII sites of pBV141 (pBGt) [27]: (a) the 0.5 kb EcoRI-NheI fragment (P27) isolated from pCK1374, (b) the 1.4 kb NheI-HpaI fragment (GFP-PLCdelta) isolated from pGFP-C1-PLCdelta:PH [20] (Addgene plasmid # 21179), and (c) the 0.5 kb PvuII-HindIII fragment of pBV210 (N. crassa β-tubulin terminator). To construct pCK1374, the 0.5 kb PCR product containing the P27 promoter was amplified from pBV167 using the (5'-GAATTCGAATTGGGTACTprimers CKP23 CAAATTGG-3') and CKP355 (5'- GCTAGCTTTGAA-GATTGGGTTCCTAC-3') and subsequently cloned in pJET1.2 (Thermo Scientific). The underlined sequences in CKP23 and CKP355 correspond to EcoRI and NheI sites, respectively. pBV plasmids (pBV1, pBV141, pBV167, pBV210, pBV229, and pBV359 as well as pBV377 described below) were obtained from Dr. Barbara Valent (Kansas State University).

Plasmids were transformed into M. oryzae O-137 or O-137-derived transformants using Agrobacterium tumefaciens-mediated transformation [28]. O-137 is a highly aggressive isolate collected from rice (Oryza sativa) in Hangzhou, Zhejiang, China [29]. CKF2138 (used in Figs. 1b, c and 5) was made by transforming KV1 with pCK1287 (P27:H1-tdTomato). KV1 is an O-137-derived strain that constitutively expresses cytoplasmic EYFP (P27:EYFP) [4]. CKF1962 (used in Fig. 2a) was made by sequentially transforming O-137 with pCK1288 (P27:3xGFP:nuclear localization signal or NLS) and pCK1287 [8]. M. oryzae Guy11 (H1-RFP; used in Fig. 2b and c) has been described in Fernandez et al. (2014). CKF2686 (used in Fig. 3) was made by sequentially transforming O-137 with pCK1287 and pCK1417 (P27:GFP-PLCdelta:PH). CKF1651 (used in Fig. 4a) was made by transforming the O-137-derived CKF110 (P27:EYFP and P27:H1-mRFP) with pBV377 (expressing cytoplasmic effector protein Pwl2 fused to mRFP with the native PWL2 promoter) [12]. CKF1616 (alias KV121; used in Fig. 4b) is an O-137-derived strain that expresses cytoplasmic effector Pwl2 fused to mCherry:NLS with the native PWL2 promoter and apoplastic effector Bas4

fused to EGFP with the native *BAS4* promoter [12]. Fungal transformants were purified by the isolation of single germinating spores. Wild-type strain O-137 and transformants were stored dehydrated and frozen at -20 °C to maintain full pathogenicity and cultured on oatmeal agar plates at 24 °C under continuous light [30].

#### Infection assays

Rice sheath inoculations were performed as previously described [11, 31]. Briefly, excised leaf sheaths (5-9 cm long) from 17- to 21-day old plants were inoculated with a spore suspension  $(5 \times 10^4 \text{ spores/mL in sterile water})$ . The inoculated sheaths were hand-trimmed at 22-28 hpi and immediately used for confocal microscopy. For heat-killed sheath inoculation assays, we first optimized heat-treatment conditions as follows: Trimmed sheaths were incubated in 70 °C water for 10 or 25 min, immersed in 0.75 M sucrose, and then visually examined under a 20× objective lens for indications of plasmolysis. Sheath cells treated for 10 min underwent some plasmolysis, but those treated for 25 min did not, indicating the 25 min incubation was sufficient to kill cells. In subsequent heat-killed inoculations, pre-trimmed sheaths were incubated in 70 °C water for 25 min, allowed to cool to room temperature, and then inoculated with a spore suspension.

#### Microscopy and image analysis

Confocal microscopy was performed on a Zeiss Axio Imager M1 microscope equipped with a Zeiss LSM 510 META system using Plan-Apochromat 20×/0.8 NA and Plan-Neofluor 40×/1.3 NA (oil) objectives. Excitation/ emission wavelengths were 488 nm/505 to 530 nm (EGFP and YFP), and 543 nm/560 to 615 nm (mRFP, mCherry, and tdTomato). Images were acquired and processed using LSM 510 software (Version 3.2). Confocal images of CKF1616 used for measuring the distance from the appressorial penetration point to the side BIC were obtained from Dr. Barbara Valent. For 2-dimensional measurements, confocal z-stack images were processed using the Zen Black software (Version 8.1, Zeiss). Each image was converted to a maximum intensity projection. The appressorial penetration point was identified based on brightfield visualization of the appressorium and on where the green fluorescence (Bas4-EGFP) outline of the primary hypha stopped. The BIC was identified from the overlap of red (Pwl2-mCherry:NLS) and green fluorescence. The distance between these two points was measured along the center of the primary hypha using the open Bezier tool. For 3-dimensional measurements, confocal z-stack images were imported into Imaris software (Version 7.6, Bitplane). The penetration point and BIC were identified as described for 2D measurements. The dendrite creation algorithm was used to produce a filament between the two points based on green fluorescence outline of the primary hypha for determining the path, and the length of the created dendrite was measured.

#### **Additional files**

Additional file 1: Morphology and pathogenicity of *M. oryzae* transformant CKF2138 is consistent with wild-type strain. Single plane merged bright-field and fluorescence confocal images of the *M. oryzae* wild-type strain O-137 (A) and the transformant CKF2138 (B) invading sheath cells of the rice strain YT16 at 48 hpi. *M. oryzae* CKF2138 constitutively expresses cytoplasmic EYFP (green) and histone H1-tdTomato (red). By this time, both strains have fully colonized initially invaded host cells and have invaded one to two neighboring cells away. The transformant remains consistent in morphology and growth stage with the wild-type strain. Note the advantage of the transformant, whose fluorescent hyphae are more readily distinguished compared to the wild-type hyphae. Bar = 20 µm. (PPTX 1654 kb)

Additional file 2: Movie S1. Three-dimensional measurement from the appressorium penetration point to the BIC of *M. oryzae* strain CKF1616 growing in a rice cell at 29 hpi. Apoplastic effector Bas4 with a translational fusion to EGFP (green) outlined the invasive hypha. The BIC and the rice nucleus were visualized by BIC-accumulating cytoplasmic effector Pwl2 with a translational fusion to mCherry:NLS (red). Overlap of green and red signals is shown in yellow. Source image is a series of 1 µm confocal optical sections taken over a depth of 23 µm. Movie was created using the key frame animation feature in Imaris software (Version 7.6, Bitplane). At the start of the animation, the penetration point is on the left, and the BIC is on the right. As the animation rotates, a penetration point-to-BIC measurement (gray line) is overlaid on the image. The measurement was performed with the dendrite creation algorithm in the Imaris software (described in the Methods section) and has a length of 33.2  $\mu m.$  A maximum intensity projection of the same infection site in this movie is shown in Fig. 4b (top), and a mirrored still of the last frame in this movie is shown in Fig. 4b (bottom). Bar = 5 µm. (MOV 4288 kb)

#### Abbreviations

BIC: Biotrophic interfacial complex; EIHM: Extra-invasive hyphal membrane; hpi: Hours post inoculation; IH: Invasive hyphae

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#### Availability of data and materials

The data sets supporting the results of this article are included within the article and two additional files.

#### Authors' contributions

CHK conceived and designed the experiments. ENS, KJ, CBJ, DWK, JZ, and CHK performed the experiments. ENS, KJ, CBJ, CHK analyzed the data and wrote the paper. All authors have read and approved the final manuscript.

#### **Competing interests**

The authors declare that they have no competing interests.

#### Consent for publication

Not applicable.

#### Ethics approval and consent to participate

Not applicable.

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#### References

- Khang CH, Valent B. *Magnaporthe oryzae* and rice blast disease. In: Borkovich KA, Ebbole DJ, editors. Cellular and molecular biology of filamentous fungi. Washington DC: ASM Press; 2010. p. 593–606.
- Howard RJ, Valent B. Breaking and entering: host penetration by the fungal rice blast pathogen *Magnaporthe grisea*. Annu Rev Microbiol. 1996;50:491–512.
- Heath MC, Valent B, Howard RJ, Chumley FG. Interactions of two strains of Magnaporthe grisea with rice, goosegrass, and weeping lovegrass. Can J Bot. 1990;68:1627–37.
- Kankanala P, Czymmek K, Valent B. Roles for rice membrane dynamics and plasmodesmata during biotrophic invasion by the blast fungus. Plant Cell. 2007;19:706–24.
- Veneault-Fourrey C, Barooah M, Egan M, Wakley G, Talbot NJ. Autophagic fungal cell death is necessary for infection by the rice blast fungus. Science. 2006;312:580–3.
- Saunders DGO, Dagdas YF, Talbot NJ. Spatial uncoupling of mitosis and cytokinesis during appressorium-mediated plant infection by the rice blast fungus *Magnaporthe oryzae*. Plant Cell. 2010;22:2417–28.
- Fernandez J, Marroquin-Guzman M, Wilson RA. Evidence for a transketolasemediated metabolic checkpoint governing biotrophic growth in rice cells by the blast fungus *Magnaporthe oryzae*. PLoS Pathog. 2014;10(9):e1004354.
- Jones K, Jenkinson CB, Borges Araújo M, Zhu J, Kim RY, Kim DW, Khang CH. Mitotic stopwatch for the blast fungus *Magnaporthe oryzae* during invasion of rice cells. Fungal Genet Biol. 2016;93:46–9.
- Jenkinson CB, Jones K, Zhu J, Dorhmi S, Khang CH. The appressorium of the rice blast fungus *Magnaporthe oryzae* remains mitotically active during postpenetration hyphal growth. Fungal Genet Biol. 2017;98:35–8.
- Mentlak TA, Kombrink A, Shinya T, Ryder LS, Otomo I, Saitoh H, Terauchi R, Nishizawa Y, Shibuya N, Thomma BPHJ, Talbot NJ. Effector-mediated suppression of chitin-triggered immunity by *Magnaporthe oryzae* is necessary for rice blast disease. Plant Cell. 2012;24:322–35.
- Jones K, Kim DW, Park JS, Khang CH. Live-cell fluorescence imaging to investigate the dynamics of plant cell death during infection by the rice blast fungus *Magnaporthe oryzae*. BMC Plant Biol. 2016;16:1–8.
- Khang CH, Berruyer R, Giraldo MC, Kankanala P, Park S-Y, Czymmek K, Kang S, Valent B. Translocation of *Magnaporthe oryzae* effectors into rice cells and their subsequent cell-to-cell movement. Plant Cell. 2010;22:1388–403.
- Valent B, Khang CH. Recent advances in rice blast effector research. Curr Opin Plant Biol. 2010;13:434–41.
- Park C-H, Chen S, Shirsekar G, Zhou B, Khang CH, Songkumarn P, Afzal A, Ning Y, Wang R, Bellizi M, Valent B, Wang G-L. The *Magnaporthe oryzae* effector AvrPiz-t targets the RING E3 ubiquitin ligase APIP6 to suppress pathogen-associated molecular pattern–triggered immunity in rice. Plant Cell. 2012;24:4748–62.
- Giraldo MC, Valent B. Filamentous plant pathogen effectors in action. Nat Rev Micro. 2013;11:800–14.
- 16. Yi M, Valent B. Communication between filamentous pathogens and plants at the biotrophic interface. Annu Rev Phytopathol. 2013;51:587–611.
- Zhang S, Xu J-R. Effectors and effector delivery in *Magnaporthe oryzae*. PLoS Pathog. 2014;10:e1003826.
- Giraldo MC, Dagdas YF, Gupta YK, Mentlak TA, Yi M, Martinez-Rocha AL, Saitoh H, Terauchi R, Talbot NJ, Valent B. Two distinct secretion systems facilitate tissue invasion by the rice blast fungus *Magnaporthe oryzae*. Nat Commun. 2013;4:1996.
- Nesher I, Barhoom S, Sharon A. Cell cycle and cell death are not necessary for appressorium formation and plant infection in the fungal plant pathogen *Colletotrichum gloeosporioides*. BMC Biol. 2008;6:9.
- Stauffer TP, Ahn S, Meyer T. Receptor-induced transient reduction in plasma membrane PtdIns(4,5)P2 concentration monitored in living cells. Curr Biol. 1998;8:343–6.
- 21. Mähs A, Ischebeck T, Heilig Y, Stenzel I, Hempel F, Seiler S, Heilmann I. The essential phosphoinositide kinase MSS-4 is required for polar hyphal morphogenesis, localizing to sites of growth and cell fusion in *Neurospora crassa*. PLoS One. 2012;7:e51454.

- 22. Momany M. Polarity in filamentous fungi: establishment, maintenance and new axes. Curr Opin Microbiol. 2002;5:580–5.
- Crampin H, Finley K, Gerami-Nejad M, Court H, Gale C, Berman J, Sudbery P. Candida albicans hyphae have a Spitzenkörper that is distinct from the polarisome found in yeast and pseudohyphae. J Cell Sci. 2005;118:2935–47.
- 24. Berepiki A, Lichius A, Shoji J-Y, Tilsner J, Read N. F-actin dynamics in *Neurospora crassa*. Eukaryotic Cell. 2010;9:547–57.
- 25. Riquelme M. Tip growth in filamentous fungi: a road trip to the apex. Annu Rev Microbiol. 2013;67:587–609.
- Mullins E, Romaine CP, Chen X, Geiser D, Raina R, Kang S. Agrobacterium tumefaciens-mediated transformation of *Fusarium oxysporum*: an efficient tool for insertional mutagenesis and gene transfer. Phytopathology. 2001;91:173–80.
- Kim H-S, Park S-Y, Lee S, Adams EL, Czymmek K, Kang S. Loss of cAMPdependent protein kinase A affects multiple traits important for root pathogenesis by *Fusarium oxysporum*. Mol Plant-Microbe Interact. 2011;24:719–32.
- Khang CH, Park S-Y, Rho H-S, Lee Y-H, Kang S. Filamentous fungi (*Magnaporthe grisea* and *Fusarium oxysporum*). Agrobacterium Protocols. 2006;2:403–20.
- Orbach JM, Farrell L, Sweigard AJ, Chumley GF, Valent B. A telomeric avirulence gene determines efficacy for the rice blast resistance gene *Pi-ta*. Plant Cell. 2000;12:2019–32.
- Valent B, Farrall L, Chumley GF. Magnaporthe grisea genes for pathogenicity and virulence identified through a series of backcrosses. Genetics. 1991;127:87–101.
- Koga H, Dohi K, Nakayachi O, Mori M. A novel inoculation method of Magnaporthe grisea for cytological observation of the infection process using intact leaf sheaths of rice plants. Physiol Mol Plant Pathol. 2004;64:67–72.

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