ERRATUM

Erratum for Abdolrasouli et al., Genomic Context of Azole Resistance Mutations in *Aspergillus fumigatus* Determined Using Whole-Genome Sequencing

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Volume 6, issue 3, doi: 10.1128/mBio.00536-15. Figure 1 incorrectly displayed the location of the TR_{34}/L98H mutation in gene *cyp51a*. The revised Fig. 1 (below) shows the correct location.

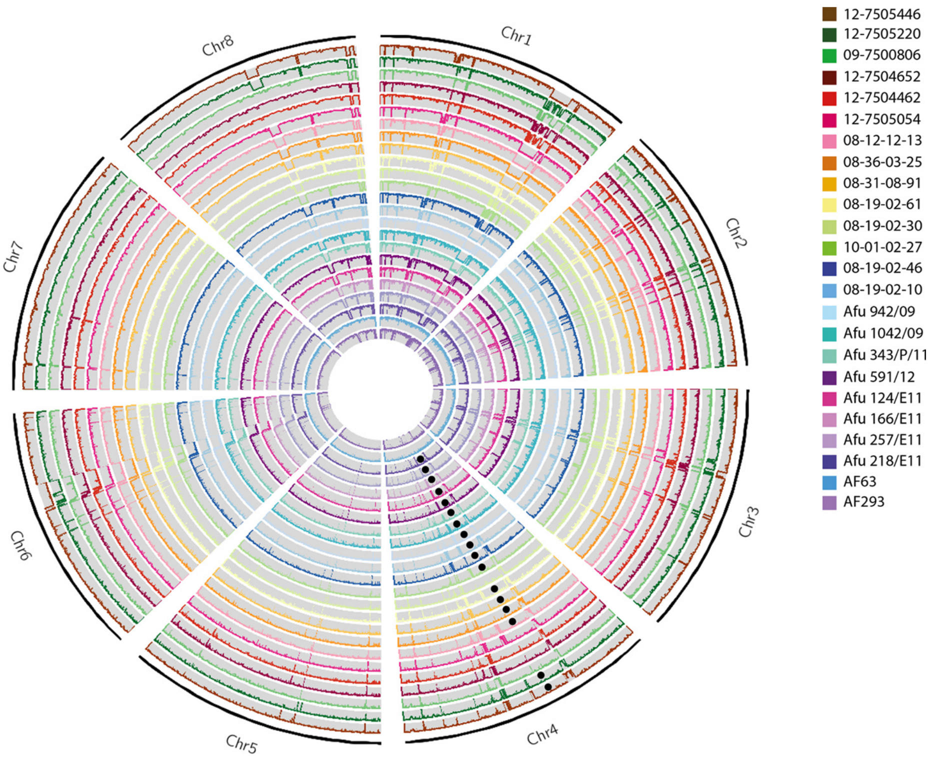

FIG 1 Circos (72) image of normalized whole-genome depth of coverage of all 24 *A. fumigatus* isolates (plotted as listed in the key), averaged over 10,000-bp bins. Black circles mark the presence of the TR_{34}/L98H mutation. Chromosomes 1 and 6 show large deletions spanning >300 kbp in most isolates, except AF65 and AF293, while chromosome 8 displays a 60-kbp deletion in all isolates except AF65, 09-7500806, 12-7504652, and 12-7504462.

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