



Genome sequence of the necrotrophic plant pathogen *Alternaria brassicicola* Abra43

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Auteur	Belmas, Elodie [1], Briand, Martial [2], Kwasiborski, Anthony [3], Colou, Justine [4], NGuyen, Guillaume [5], Iacomi, B. [6], Grappin, Philippe [7], Campion, Claire [8], Simoneau, Philippe [9], Barret, Matthieu [10], Guillemette, Thomas [11]
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Résumé en anglais	<i>Alternaria brassicicola</i> causes dark spot (or black spot) disease, which is one of the most common and destructive fungal diseases of Brassicaceae spp. worldwide. Here, we report the draft genome sequence of strain Abra43. The assembly comprises 29 scaffolds, with an N50 value of 2.1 Mb. The assembled genome was 31,036,461 bp in length, with a G+C content of 50.85%.
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Lien vers le document	Genome sequence of the necrotrophic plant pathogen <i>Alternaria brassicicola</i> Abra43

Liens

- [1] <http://okina.univ-angers.fr/publications?f%5Bauthor%5D=24900>
- [2] <http://okina.univ-angers.fr/publications?f%5Bauthor%5D=12672>
- [3] <http://okina.univ-angers.fr/publications?f%5Bauthor%5D=25786>
- [4] <http://okina.univ-angers.fr/publications?f%5Bauthor%5D=26861>
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