

Unravelling species and pathotypes diversity of *Pyricularia* blast pathogens affecting wheat and grasses in Minas Gerais Cerrado, Brazil (Desvendando as espécies e a diversidade de patótipos de *Pyricularia* infectando trigo e gramíneas no Cerrado de Minas Gerais, Brasil)

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Pyricularia spp. cause blast diseases in cereals and grasses. Studies with a global collection of strains have disentangled the complex into separate species, as well as pathotypes (host-specialized groups) within P. oryzae. Previous taxonomic, biological and pathogenicity studies were conducted using a contemporary collection of Brazilian strains gathered from wheat and other hosts, all bordering wheat field. We hypothesized that new knowledge on genetic diversity and evolution is gained from the analyses of larger number of strains sampled from both wheat and non-wheat hosts, at increasing distances from wheat fields, including non-wheat regions. We conducted two surveys in the state of Minas Gerais during March (planting time) and May-June (during wheat maturation). Sites located at non-wheat and two wheat regions (South and Triângulo) were visited and blastsymptomatic leaves were collected and geotagged. The samples were grouped into three categories: a) wheat or grass within/bordering a wheat field, b) grass within wheat region and c) grass at non-wheat region. In total, 176 isolates were obtained from 440 plant samples: 14.7% were obtained from wheat fields, 63% from wheat region and 21.6% from non-wheat region. Represented hosts were: Digitaria spp. (30.1%), Eleusine sp. (22.1%), Urochloa spp. (13%), Cenchrus sp. (10.2%), Rhynchelytrum sp. (9%), Triticum sp. (8%), Panicum sp. (5.7%), Cynodon sp. (1.1%), Pennisetum sp. (0.5%). DNA was extracted from all strains and PCR assays were conducted using CH7-BAC9 (P. oryzae-specific) and MoT3 (Triticum pathotype-specific) loci. CH7 BAC9 was amplified from 60% of the strains, all isolates from Triticum sp., Rhynchelytrum sp., Panicum sp., 37/39 strains from Eleusine sp., 20/23 from Urochloa spp., 8/53 from Digitaria spp.. Of all P. oryzae, 35% (37/106) were represented by the Triticum pathotype (MoT) based on the amplification of the MoT3 marker. Among sampled hosts, MoT was most dominant in Triticum sp. (12/14 strains), Panicum sp. (8/10) and Urochloa spp. (15/20), less in Digitaria spp. (1/8), rare in Eleusine sp. (1/37). These preliminary results show a clear decline in the frequency of the Triticum pathotype when sampling at increasing distances from wheat fields. Ongoing studies include new isolations from plants of all sampling sites, but more from wheat, and identification of the non-P. oryzae species and the non-Triticum pathotypes.

Palavras-chave: Alternative hosts; Epidemiology; Wheat blast

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