

Molecular identification of 16SrI-A, 16SrI-B, 16SrI-C, and 16SrI-L subgroups of phytoplasmas in gramineous plants in Lithuania

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Abstract

Diseased plants of family *Poaceae* (*Gramineae*) were observed in cereal crop and forage feed plant fields in the Vilnius, Kėdainiai and Raseiniai regions of Lithuania. Disease symptoms included yellowing of leaves and spikes, general stunting, sterility and deformation of spikes, dwarfed spikes, and twisted awns, indicating possible phytoplasmal infections. A phytoplasma-characteristic fragment of 16S rDNA was amplified from all the symptomatic plants that were tested in nested polymerase chain reaction (PCR), using phytoplasma-specific primers. RFLP analysis of the amplified 16S rDNA indicated that detected phytoplasmas infecting gramineous plants in Lithuania belong to several different subgroups in group 16SrI (aster yellows phytoplasma group): 16SrI-A (in *Avena sativa*); 16SrI-B (in *A. sativa*, *Hordeum vulgare*, *Triticosecale* and *Bromopsis inermis*); 16SrI-L (in *A. sativa* and *Lolium multiflorum*) and 16SrI-C (in *Poa pratensis* and *Festuca arundinaceae*).

Key words: gramineous plants, phytoplasma, 16S rRNR gene, PCR, RFLP.

Introduction

Considering that cereal crops and forage feeds are important in Lithuania, a great attention was paid to the presence of phytoplasmal diseases in these crops. The observation of gramineous plants in several locations in Lithuania during seven years indicated widespread presence of phytoplasmas belonging to group 16Sr-I (table 1). Comparatively little is known about phytoplasmal diseases of cereal crops in Europe and other places in the world. This work summarizes data of molecular evidence indicating that phytoplasmas could damage yields of gramineous plants, causing economic losses.

Materials and methods

Samples of diseased plants exhibiting yellowing of leaves and spikes, general stunting, sterility and deformation of spikes, dwarfed spikes, and twisted awns were collected in fields in Vilnius, Kėdainiai and Raseiniai districts in Lithuania. Template DNA was extracted from the tissues using Genomic DNA Purification Kit (MBI Fermentas, Vilnius, Lithuania) and used in polymerase chain reaction (PCR) for amplification of phytoplasmal 16S rDNA. In nested PCR, the first reaction was primed by phytoplasma-universal primer pair P1/P7 (Deng and Hiruki, 1991) and the second (nested) PCR primed by primer pair R16F2n/R16R2 (F2n/R2) (Gundersen and Lee, 1996). PCR products were analyzed by electrophoresis through 1% agarose gel, stained with ethidium bromide, and DNA bands visualized using a UV transilluminator.

Products (1,2 kbp) of the nested PCR, obtained with primer pair R16F2n/R16R2, were subjected to enzymatic restriction fragment polymorphism (RFLP) analysis using restriction endonucleases *AluI*, *HaeIII*, *HhaI*,

HinfI, *HpaII*, *MseI*, *RsaI*, *TagI*, *KpnI* and *Sau3AI* (MBI Fermentas) and electrophoresis through 5% acrylamide gel, staining with ethidium bromide, and visualization using a UV transilluminator. Phytoplasmas were classified in groups and subgroups, through comparisons of RFLP patterns previously published, in accordance with the classification scheme of Lee *et al.* (1998) and Marcone *et al.* (2000).

Results

Phytoplasma characteristic 1.2 kbp 16S rDNA PCR products were amplified from the total DNA of all symptomatic plants, but not from healthy plants, indicating that these plants were infected by phytoplasmas. The phytoplasmas strains were classified on the basis of RFLP analysis of 16S rDNA amplified in PCR, according to the classification scheme established by Lee *et al.* (1998) and Marcone *et al.* (2000). RFLP analysis of the amplified 16S rDNA indicated that the detected phytoplasmas infecting gramineous plants in Lithuania belong to several different ribosomal subgroups in group 16SrI (aster yellows phytoplasma group, AY): 16SrI-A (in *Avena sativa* L.); 16SrI-B [in *A. sativa*, *Hordeum vulgare* L., *Triticosecale* Wittm. ex A. Camus (*Triticum* L. × *Secale* L.), and *Bromopsis inermis* (Leyss.) Holub.]; 16SrI-L (in *A. sativa* and *Lolium multiflorum* Lam.) and 16SrI-C (in *Poa pratensis* L. and *Festuca arundinaceae* Schreb.) (table 1).

Discussion

The symptomatic gramineous plants were infected by phytoplasmas belonging to group 16SrI (AY phytoplasma group) (Lee *et al.*, 1998; Marcone *et al.*, 2000).

Table 1. Diseases of gramineous plants associated with infection by phytoplasma strains belonging to diverse subgroups of 16S rDNA ribosomal group 16SrI in Lithuania.

Disease (Phytoplasma strain)	Natural host plant	16S rDNA Group-subgroup	Reference
Oat proliferation (OatP)	<i>Avena sativa</i> L.	I-A	Jomantiene <i>et al.</i> , 2002
Oat stunt (OatSt)	<i>Avena sativa</i> L.	I-B	Urbanavičienė <i>et al.</i> , 2006
Oat yellows (Oat Y)	<i>Avena sativa</i> L.	I-L	Urbanavičienė <i>et al.</i> , 2004
Smooth brome grass stunt (BrS)	<i>Bromopsis inermis</i> (Leys.)	I-B	Urbanavičienė, 2005
Fescue yellows (FesY)	<i>Festuca arundinaceae</i> Schreb.	I-C	Valiūnas <i>et al.</i> , 2007
Barley deformation (BaDef)	<i>Hordeum vulgare</i> L.	I-B	Urbanavičienė <i>et al.</i> , 2004
Ryegrass yellows (RgY)	<i>Lolium multiflorum</i> Lam.	I-L	Urbanavičienė <i>et al.</i> , 2005
Poa stunt (PoaS)	<i>Poa pratensis</i> L.	I-C	Valiūnas <i>et al.</i> , 2007
<i>Triticosecale</i> stunt (TrSt)	<i>Triticosecale</i> Wittm. ex A. Camus (<i>Triticum</i> L. × <i>Secale</i> L.)	I-B	Urbanavičienė <i>et al.</i> , 2004

The strains of this phytoplasma group are spread worldwide, however little is known about phytoplasma infections in plants of family *Poaceae*. Some data show, for example, that AY phytoplasma disease may cause severe losses in barley and wheat (Chiykowski, 1963). At least six species of grass have been reported to belong to the host range of AY phytoplasma in the United States and Canada (Bantari, 1966). In Finland, AY phytoplasma was readily transmitted by *Macrosteles laevis* (Rib.) to and from graminaceous plants (Murtomaa, 1966). The literature contains only few reports of phytoplasmas possibly infecting oats. In some early reports, only phytoplasma-characteristic disease symptoms were reported. Interestingly, phytoplasma strains belonging to subgroup 16SrI-C were detected in common meadow-grass and tall fescues (Valiūnas *et al.*, 2007). Previously, *Poa annua* was identified as host of a subgroup 16SrXIV-A phytoplasma in Italy (Lee *et al.*, 1997). The detection of phytoplasmas infecting gramineous plants in Lithuania provides new insights on threatening diseases of grass species, including important grains, in Europe.

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