Incidence of viruses and viroids in Polish hop gardens

Marcin Przybyś

Department of Plant Breeding and Biotechnology – Institute of Soil Science and Plant Cultivation ul. Czartoryskich 8, 24-100 Puławy, POLAND

Abstract. Viruses and viroids cause losses in hop production and are very difficult to eliminate, they spread easily and infected plants become a source of infection for other plants in the hop garden. Hop infected with viruses or viroids usually do not show any symptoms of disease, which makes it difficult to identify infected plants and prevents their elimination from the plantation. Monitoring is of utmost importance to detect and eradicate at an early stage first sources of infection with pathogens that have potentially catastrophic impacts on hop production. The aim of the study was determine the incidence of Hop latent virus (HpLV), Arabis mosaic virus (ArMV), Hop stunt viroid (HSVd), Apple fruit crinkle viroid (AFCVd) and Citrus bark cracking viroid (CBCVd) in Polish hop gardens. Hop leaves from commercial hop gardens and a propagation facility were tested by RT-PCR. On the basis of the obtained results, HpLV was found to occur in a few hop gardens in all regions of hop cultivation in Poland. HSVd was found in one hop garden and ArMV, AFCVd and CBCVd were not found in any of the tested samples. The results provided knowledge about these pathogens and will be useful in effective management of the risk of these dangerous diseases.

Keywords: virus, viroid, hop, HpLV, HSVd

INTRODUCTION

The hop (*Humulus lupulus*) is one of the two species belonging to the genus *Humulus*. In Poland, hop is grown on an area of about 1420 ha, which ranks our country in the third position in Europe and the fifth in the world, in terms of cultivation area and production volume (IHGC, 2015). Hop is grown mainly for the needs of the brewing industry, as a raw material being the source of the characteristic flavour and aroma. Due to their anti-inflammatory and antioxidant properties, hop is also used in the cosmetics industry

Corresponding author:

Marcin Przybyś e-mail: mprzybys@iung.pulawy.pl phone: +48 81 4786 934 as an ingredient in shampoos and creams which delay the process of skin aging. In addition, extracts of hop cones are used in the pharmaceutical industry, for the production of sleep-inducing and sedative drugs, aiding in digestion and soothing ailments linked to menopause (Bohr et al., 2005; Corrêa et al., 2018; Forino et al., 2016).

Hop is a perennial species, cultivated for many years in the same position, without crop rotation. After a garden is established, over time, a gradual accumulation of pathogens in the plants and soil follows. Viruses and viroids are pathogens which accumulate particularly easily in plants because there are no effective chemical methods to combat them. Chemical agents can only reduce the population of vectors transmitting these pathogens. An additional problem promoting infections is the ease with which they are transmitted during agrotechnical procedures when mechanical damage to plants occurs (Skomra, 2015). Viral diseases rarely cause visible symptoms on hop, only in extreme cases can a viral or viroid infection lead to morphological changes of leaves, stunting or death of susceptible hop cultivars (Pethybridge et al., 2008). Despite the fact that infected plants do not show any symptoms, they often provide lower yield of cones with adversely modified chemical composition. The yield and content of alpha acids depend to a large extent on the variety and can be reduced by as much as 35% and 50%, respectively, in infected plants (Barbara et al., 1990; Patzak et al., 2001; Sano, 2013).

In Poland, so far, research has been conducted on the most common viruses: *Hop mosaic virus* (HpMV), *Prunus necrotic ring spot virus* (PNRSV), *Arabis mosaic virus* (ArMV), *American hop latent virus* (AHLV) and *hop latent viroid* (HLVd) (Skomra, 2001; Grudzińska, Solarska, 2005). In 2018, was started monitoring of hop gardens comprising all Polish hop growing regions to detect economically important hop viruses: *Hop latent virus* (HpLV), *Arabis mosaic virus* (ArMV), *Hop stunt viroid* (HSVd), *Apple fruit crinkle viroid* (AFCVd), *Citrus bark cracking viroid* (CBCVd). In the first year of monitoring, the pre-

sence of HpLV was found in 6, and HSVd in 2 out of 900 tested hop plants (Przybyś et al., 2019). HpLV belonging to the Carlavirus genus does not usually cause symptoms in hops, although sometimes it causes chlorotic spots on the leaves (Ziegler, et al. 2014). Its vector is hop aphid (Phorodon humuli) (Adams, Barbara, 1982). HpLV is not spread through seeds (Brunt, et al., 1996). Due to the latent nature of the infection, the exact extent of HpLV is unknown, but it has been found in hop gardens in Europe, the United States, New Zealand, Australia, China, South Africa and Japan (Pethybridge, et al., 2008, Seigner et al., 2014). ArMV belonging to the genus Nepovirus has a very large group of natural hosts including many weed species and crop species (Brunt, et al., 1996). Its spread depends on the occurrence of nematodes Xiphinema diversicaudatum, X. coxi, and X. bakeri, which are vectors of this virus, in hop gardens. Unlike HpLV, it can spread through seeds (Brunt, et al., 1996). Its occurrence in Europe has been established in England, France and Germany.

A separate group of the pathogens of hops is constituted by viroids, which are a bare, self-replicating form of single-stranded RNA, without a protein coat. HSVd has already been found in hop gardens in Japan, Korea, China, USA and Europe, which, in addition to hop, can infect grapevine, almond, plum, peach and apricot (Luigi, Faggioli, 2013; Sano, 2013). The infection with this pathogen causes stunting, yellowing and leaf curling in plants, but symptoms often appear several years later, in hop often causing a significant reduction in the content of alpha acids (Sano, 2013). HSVd is spread only mechanically through agrotechnical treatments damaging plants, and the manner of the long-distance spread of the disease is transmission along with infected seedlings (Pethybridge et al., 2008). Another viroid causing similar symptoms to HSVd, which has been found in hop gardens in Japan, is AFCVd (Sano, et al., 2008). Sometimes there occur asymptomatic infections with this pathogen, but invariably the effects of infection include a significant reduction in the content of alpha acids in hop cones. A very dangerous viroid that first appeared in Slovenian hop gardens in 2007 is CBCVd (Jakse et al., 2015). It has been shown to cause HSVd-like symptoms: stunting, yellowing and leaf curling, and plant death, but the incubation period is much shorter with CBCVd infection, and the disease progresses more aggressively. The disease spreads very quickly in hop gardens, usually along rows. Due to the recent epidemic in Slovenia and the level of threat to hop in Europe, a warning about CBCVd was published in June 2015 on the EPPO (European and Mediterranean Plant Protection Organization) alert list.

The aim of the research was to assess the harmfulness of viral diseases for Polish hop gardens by determining the occurrence of viruses (HpLV and ArMV) and viroids (CBCVd, HSVd, AFCVd) in Poland.

MATERIALS AND METHODS

Plant material and sampling

The research was conducted during the period of 2018–2019. Samples of hop leaves were collected from gardens located in all of the hop-growing regions in Poland: Lublin, Greater Poland and Lower Silesia (Fig. 1). Samples were collected three times during the growing season from both bitter and aromatic varieties. For this purpose, each plant from which a sample was taken for testing was labelled, so that it was possible to collect two additional samples at a later date. The first term covered the period of training hop plants onto the guiding strings, the second one covered the flowering phase and the third one covered the stage of maturity. The leaves from various storeys of the plant were collected into sterile 50 ml tubes and cooled to 4°C while still being in the garden.

RNA isolation

In order to perform isolation, 50 mg of material from several leaves was used and placed in 2 ml tubes, filled with 6 ceramic balls with a diameter of 2.8 mm. The samples thus prepared were cooled in liquid nitrogen and powdered in a Tissue Lyser homogeniser (Qiagen, Germany) at a shaking frequency of 30 Hz for 3 minutes. The homogenised material was subjected to total RNA isolation using RNA isolation kits - RNeasy PowerPlant Kit (Qiagen), in accordance with the procedure recommended by the manufacturer with a modification resulting from the high content of phenolic compounds in hop leaves. The modification consisted in reducing the volume of MBL/ β -ME lysis buffer from 600 µl to 550 µl and adding 50 µl of Phenolic Separation Solution in order to reduce formation of nucleic acid complexes with oxidized phenolic compounds, which would decrease the amount of isolated RNA. The isolated RNA was additionally purified to remove DNA residue using the DNase Max kit (Qiagen) in accordance to the procedure recommended by the manufacturer.

Reverse transcription

To obtain cDNA, RNA viruses and viroids were subjected to reverse transcription. 1 μ l of oligo d(T)₁₅ (50 μ M) - ArMV and HpLV or 1 μ l of inverted specific primer (2 μ M) – HSVd, CBCVd, AFCVd were added to 9 μ l of sterile RN-ases free water (Table1), 2 μ l RNA (1 μ g/ μ l) and 1 μ l of dNTP 10 mM mixture (Invitrogen, USA). The whole mixture was incubated at 65°C for 5 minutes and then cooled on ice for 1 minute. 4 μ l of 5x First-Strand buffer (Invitrogen), 1 μ l of 0.1M DTT (Invitrogen), 1 μ l of RNasin (40 u/ μ l) – RN-ases inhibitor (ThermoFisher Scientific) and 1 μ l of reverse transcriptase M-MLV Superscript III (200 u/ μ l) (Invitrogen) were added to the cooled mixture. The final volume of the reaction mixture was 20 μ l. The reaction of reverse transcription was carried out



Figure 1. Locations of hop garden from which samples for research were collected, according to hop-growing regions in Poland.

at 50°C for 60 minutes, after which it was heated to 70°C for 15 minutes in order to stop the reaction and denaturation of the reverse transcriptase.

DNA amplification

DNA amplification was carried out using 2 μ l of cDNA obtained through reverse transcription, 25 μ l of Platinum Green Hot Start PCR 2x Master Mix (Invitrogen), 0.2 μ M of each of the specific PCR reaction starters (Table 1). The reaction mixture was replenished with nuclease-free water to a final volume of 50 μ l. PCR parameters consisted of a denaturation step at 94°C for 2 minutes, followed by 35 cycles: 94°C for 30 seconds, 60°C (ArMV, CBCVd), 56°C (HpLV), 54°C (HSVd), 55°C (AFCVd) for 30 sec, 72°C for 30 or 90 seconds (HpLV), to finish with an extension step at 72°C for 5 minutes. The obtained amplicons were separated on a 2% agarose gel stained with ethidium bromide and visualized in UV light.

RESULTS AND DISCUSSION

In total, 1800 hop samples were collected over the period of 2018–2019 (Table 2). Samples were obtained from both symptomatic and asymptomatic plants. The research material was collected from 33 gardens with a total area of 55 ha. A larger number of samples collected in the Lublin region resulted from a greater number of gardens located in this region of hop cultivation.

In the research conducted, the occurrence of particular studied pathogens was determined using RT-PCR screening method (Fig. 2). In order to detect HpLV, PCR primers, which amplify the 5'-end of the viral genome, were used. The design of these primers was based on the gene encoding methyltransferase, which is highly evolutionarily conserved. In order to detect ArMV, a highly conservative region encoding coat protein (CP) was amplified, and all of the viroids: HSVd, CBCVd, AFCVd were detected with the use of the primers which amplified their full genomes (Table 1).

Different pathogens can be detectable in infected hops at different stages of plant development. In the case of ArMV detection, it is best to obtain leaf samples during the spring season because only then is it possible to detect infection by this virus, while HpLV is best detected at a later period (Wetzel et al., 2002). In the conducted research, the occurrence of hop latent virus (HpLV) was found in all regions of hop cultivation in Poland (Table 3). In the Lublin and Greater Poland regions, the presence of the virus

was found in two gardens of the cv. Marynka, one per each region. In the Lower Silesian region, infections caused by HpLV were found in two gardens where the cv. Hallertau Tradition and cv. Magnum were cultivated. HpLV was detected solely in the samples obtained during the second and third term. This was in agreement with the observations of Ziegler et al. (2014) and Tsai et al. (2012), where HpLV was also detected when the samples were composed of mature leaves from the lower part of the plant obtained in the second half of the growing season.

As a result of this research, infections with the hop stunt viroid (HSVd) were also established, after having been detected only in the Greater Poland hop cultivation region on the cv. Magnum in one hop garden. As in the case of HpLV, HSVd infestation was found in samples collected during the second and third term. HSVd was first discovered in Japan in the 1970s, but its occurrence was also later reported in South Korea and the USA (Yamamoto et al., 1973; Sano et al., 1989; Lee et al., 1990; Pethybridge et al., 2008). It was found for the first time in Europe in 2012 (Radisek et al., 2012). In the current research, the hop plant from which samples were obtained for testing showed signs of leaf yellowing (Fig. 3). Similar signs were observed by Eastwell and Nelson (2007). Furthermore, 3-5 years after infection they recorded inhibition of the plant growth. The inhibition of the growth of infected plants leads to a decrease in harvest yield and a decrease in the level of alpha acids (Sano, 2003). In the reserach conducted, no stunting of the infected plant was observed, which may suggest that its infection occurred no earlier than 3 years ago. Like other viroids, HSVd is transmitted mechanically. It has many hosts: plum, peach, citrus and vine (Sano et al., 1989, Diener et al., 1988, Matoušek et al., 2003), which may be a reservoir of the pathogen.

In total, as part of the research, HpLV was detected in 11 samples taken from 4 hops, which constitutes 0.6% of the tested samples. This is a low infection rate especially compared to a study by Pethybridge (2005), who found HpLV infections in 6-87% of hop plants located in Australia, depending on the variety. In Europe, in Germany, ba-

M 5 6 KN 3

M – Ladder, lanes 1-5 – hop latent virus (HpLV), 1116 bp, lane 6 - hop stunt viroid (HSVd), 297 bp, KN - negative control.

Figure 2. Electrophoretic analysis of RT-PCR products in 2% agarose gel. The gel was stained with ethidium bromide.

sed on the monitoring studies carried out by Seigner et al. (2014) it was found that HpLV is widely distributed

In this research HSVd was found in 2 out of 1800 samples obtained from one hop garden. The first hop infections in Europe were found in 2007 in Slovenia, where 1 to 30% of plants were infected in various gardens (Radisek et al., 2012). The presence of HSVd was also confirmed in monitoring studies carried out in Germany, where infections

Table 1. PCR primers used for the detection of viruses and hop viroids.

PathogenPrimerSequence (5'-3')Product (bp)ReferencesArMVArMV-FACCAGTGCCTACAAGAGTGTGTCC213Komínek et al. 2003ArMV-RTTGATTCCAGTTGTTAGTGACCCC213Komínek et al. 2003HpLVHpLV 5'MluCGCACGCGTGGATAAACAAACATACAA1116Ziegler et al. 2014HpLV 3'-1100GCTTAGCAATTGCGGATTGCAC1116Ziegler et al. 2014					
ArMVArMV-FACCAGTGCCTACAAGAGTGTGTCC213Komínek et al. 2003ArMV-RTTGATTCCAGTTGTTAGTGACCCC1116Ziegler et al. 2014HpLVHpLV 5'MluCGCACGCGTGGATAAACAAACATACAA1116Ziegler et al. 2014HpLV 3'-1100GCTTAGCAATTGCGGATTGCAC1116Ziegler et al. 2014	Pathogen	Primer	Sequence (5'-3')	Product (bp)	References
ArMV-RTTGATTCCAGTTGTTAGTGACCCCHpLV 5'MluCGCACGCGTGGATAAACAAACATACAA1116Ziegler et al. 2014HpLV 3'-1100GCTTAGCAATTGCGGATTGCAC1116Ziegler et al. 2014	ArMV	ArMV-F	ACCAGTGCCTACAAGAGTGTGTCC	213	Komínek et al. 2003
HpLVHpLV 5'MluCGCACGCGTGGATAAACAAACATACAA1116Ziegler et al. 2014HpLV 3'-1100GCTTAGCAATTGCGGATTGCAC1116Ziegler et al. 2014		ArMV-R	TTGATTCCAGTTGTTAGTGACCCC		
HpLV 3'-1100 GCTTAGCAATTGCGGATTGCAC	HpLV	HpLV 5'Mlu	CGCACGCGTGGATAAACAAACATACAA	1116	Ziegler et al. 2014
		HpLV 3'-1100	GCTTAGCAATTGCGGATTGCAC		
HSVd HpSVd3-160 GACGATCGATGGTGTTTCGAAG 297 Ziegler et al. 2014	HSVd	HpSVd3-160	GACGATCGATGGTGTTTCGAAG	297	Ziegler et al. 2014
HpSVd5-160 ATCGATCGTCCCTTCTTTAC		HpSVd5-160	ATCGATCGTCCCTTCTTCTTTAC		-
AFCVd AF-F TTGTCGACGAAGGGTCCTCA 382 Sano et al. 2004	AFCVd	AF-F	TTGTCGACGAAGGGTCCTCA	382	Sano et al. 2004
AF-R TTGTCGACGACGAGTCACCA		AF-R	TTGTCGACGACGAGTCACCA		
CBCVd CVd-IV-F1 GGGGAAATCTCTTCAGAC 284 Bernad, Duran-Vila, 2006	CBCVd	CVd-IV-F1	GGGGAAATCTCTTCAGAC	284	Bernad, Duran-Vila, 2006
CVd-IV-R1 GGGGATCCCTCTTCAGGT		CVd-IV-R1	GGGGATCCCTCTTCAGGT		



Dagian		Total				
Region	Aroma hop		Bittering hop		- Iotai	
Lublin	Lomik	72	Marynka	300	1080	
	Lubelski	192	Magnum	402		
	Perle	66				
	Sybilla	48				
Greater Poland	Lubelski	30	Marynka	60	402	
	Perle	30	Magnum	252		
	Hallertau Tradition	30				
Lower Silesia	Lubelski	60	Marynka	72	318	
	Hallertau Tradition	72	Magnum	114		
Total		600		1200	1800	

Table 2. Hop samples used in the tests.

Table 3. Incidence of viruses and viroids in hop garden.

	Pathogen						
Region	HpLV			HSVd			
	Cultivar	NHG	Samples	Cultivar	NHG	Samples	
Lublin	Marynka	1	3		0	0	
Greater Poland	Marynka	1	2	Magnum	1	2	
Lower Silesia	Hallertau Tradition	1	2		0	0	
	Magnum	1	2				
	Total	4	9		1	2	

NHG - number of hop gardens with pathogen detected





Figure 3. Hop leaf cv. Magnum a) no symptoms, b) with HSVd symptoms.

with this pathogen were detected in 9 out of 1444 tested plants (Seigner et al., 2014).

None of the tested samples showed the presence of *Arabis mosaic virus* (ArMV), *Apple fruit crinkle viroid* (AFCVd) and *Citrus bark cracking viroid* (CBCVd).

Monitoring the occurrence of viral diseases amongst plants is a very important activity, the purpose of which is to support early detection of threats and prevent the subsequent spread of diseases. The results indicate that HpLV and HSVd are the biggest threat to hop cultivations in Poland. However, it should be noted that in the case of HSVd only 2 samples from one infected hop plant was identified. Despite failure to find CBCVd, AFCVd and ArMV, the existence of this problem cannot be ruled out, especially that infections caused by these pathogens are known in Europe (Pethybridge et al., 2008; Radisek et al., 2012). Because genetic sources of resistance to HpLV and HSVd are unknown, it is crucial that the health of a hop garden be maintained at a high level by using high-quality seedlings as well as following phytosanitary recommendations, which will protect hop gardens against an infection and its subsequent spreading.

CONCLUSIONS

1. HpLV was found in only a few gardens in all of the hop cultivation regions in Poland.

2. HSVd was found in one hop garden in the Greater Poland region in Poland.

3. There were no ArMV, AFCVd and CBCVd in Polish hop gardens.

4. Due to the recent reports on the emergence of new pathogens which have been absent in Europe so far, such as HSVd or CBCVd, constant monitoring of the health of hop plants should be carried out in hop gardens.

REFERENCES

- Adams A.N., Barbara D.J., 1982. Host range, purification and some properties of two carlaviruses from hop (*Humulus lupulus*): hop latent and American hop latent. Annual Applied Biology, 101(3): 483-494, doi: 10.1111/j.1744-7348.1982. tb00849.x.
- Barbara D.J., Morton A., Adams A.N., Green C.P., 1990. Some effects of hop latent viroid on two cultivars of hop (*Hu-mulus lupulus*) in the UK. Annals of Applied Biology, 117: 359-366, doi: 10.1111/j.1744-7348.1990.tb04222.x.
- Bernad L., Duran-Vila N., 2006. A novel RT-PCR approach for detection and characterization of citrus viroids. Molecular Cellular Probes, 20(2):105-113, doi: 10.1016/j.mcp.2005.11.001.
- Bohr G., Gerhäuser C., Knauft J., Zapp J., Becker H., 2005. Anti-inflammatory Acylphloroglucinol Derivatives from Hops (*Humulus lupulus*). Journal of Natural Products, 68: 1545-1548, doi: 10.1021/np050164z.
- Brunt A.A., Crabtree K., Dallwitz M.J., Gibbs A.J., Watson L., 1996. Viruses of plants: descriptions and lists from the VIDE database. CAB International, Wallingford, UK.

- Corrêa R.C.G., Peralta R.M., Haminiuk C.W.I., Maciel G.M., Bracht A., Ferreira I.C.F.R., 2018. New phytochemicals as potential human anti-aging compounds: Reality, promise, and challenges. Critical Reviews in Food Science and Nutrition, 58: 942–957, doi: 10.1080/10408398.2016.1233860.
- Diener T.O., Smith D.R., Hammond R.H., Albanese G., Larosa R., Davino M., 1988. Citrus-B viroid identified as a strain of hop stunt viroid. Plant Disease, 72: 691-693, doi: 10.1094/ pd-72-0691.
- Eastwell K.C., Nelson M.E., 2007. Occurrence of viroids in commercial hop (*Humulus lupulus* L.) production areas of Washington State. Online. Plant Health Progress, doi: 10.109/ PHP2007-1127-01-RS.
- Forino M., Pace S., Chianese G., Santagostini L., Werner M., Weinigel C., 2016. Humudifucol and bioactive prenylated polyphenols from hops (Humulus lupulus cv."Cascade"). Journal of Natural Products, 79(3): 590-597. doi: 10.1021/ acs.jnatprod.5b01052
- Grudzińska M., Solarska E., 2005. The elimination of viruses and hop latent viroid from hop (*Humulus lupulus* L.) in Poland. Acta Horticulturae, 668: 149-152, DOI: 10.17660/Acta-Hortic.2005.668.19.
- IHGC International Hop Grovers Convention. 2015. Economic Commision – Summary reports. http://www.hmelj-giz.si/ ihgc/doc/2015%20JUL%20IHGC%20EC%20report.pdf
- Jakse J., Radisek S., Pokorn T., Matousek J., Javornik B., 2015. Deep-sequencing revealed *Citrus bark cracking viroid* (CBCVd) as a highly aggressive pathogen on hop. Plant Pathology, 64: 831-842, doi: 10.1111/ppa.12325.
- Komínek P., Svoboda P., Abou Ghanem-Sabanadzovic N., 2003. Improved detection of Arabis mosaic virus in grapevine and hop plants. Acta Virologica, 47: 199-200.
- Lee J.Y., Lee S.H., Sänger H.L., 1990. Viroid diseases occurring on Korean hop plants. Korean Journal of Plant Pathology, 6: 256-260.
- Luigi M., Faggioli F., 2013. Development of a quantitative real-time RT-PCR (qRT-PCR) for the detection of hop stunt viroid. European Journal of Plant Pathology, 137: 231-235, doi: 10.1007/s10658-013-0243-2.
- Matoušek J., Orctová L., Patzak J., Svoboda P., Ludvíková I., 2003. Molecular sampling of hop stunt viroid (HSVd) from grapevines in hop production areas in the Czech Republic and hop protection. Plant, Soil and Environment, 49(4): 168-175.
- Patzak J., Matoušek J., Krofta K., Svoboda P., 2001. Hop latent viroid (HLVd)-caused pathogenesis: effects of HLVd infection on lupulin composition of meristem culture-derived *Humulus lupulus*. Biologia Plantarum, 44: 579-585.
- Pethybridge S.J., Hay F.S., Barbara D.J., Eastwell K.C., Wilson C.R., 2008. Viruses and Viroids Infecting Hop: Significance, Epidemiology, and Management. Plant Disease, 92(3): 324-338, doi: 10.1094/PDIS-92-3-0324.
- Pethybridge S.J., 2005. Epidemiology and management of viruses of hop in Australia. Acta Horticulturae, 668: 131-142, doi: 10.17660/actahortic.2005.668.17.
- Przybyś M., Skomra U., Korbecka-Glinka G., 2019. The occurrence of previously unmonitored viruses (HpLV, ArMV) and viroids (HpSVd, AFCVd, CBCVd) in hop gardens in Poland. Biuletyn Instytutu Hodowli i Aklimatyzacji Roślin, 286: 371-374.
- Radisek S., Majer A., Jakse J., Javornik B., Matoušek J., 2012. First Report of Hop stunt viroid Infecting Hop in Slovenia, Disease Notes, 96(4):592, doi: 10.1094/PDIS-08-11-0640-PDN.

- Sano T., 2013. History, origin, and diversity of hop stunt disease and *Hop stunt viroid*. pp. 87-96, In: III International Humulus Symposium; editors J. Patzak, A. Koutoulis, doi: 10.17660/ ActaHortic.2013.1010.9.
- Sano T., Isono S., Matsuki K., Kawaguchi-Ito Y., Tanaka K., Kondo K.-i., et al., 2008. Vegetative propagation and its possible role as a genetic bottleneck in the shaping of the apple fruit crinkle viroid populations in apple and hop plants. Virus Genes, 37: 298-303, doi: 10.1007/s11262-008-0270-9.
- Sano T., Yoshida H., Goshono M., Monma T., Kawasaki H., Ishizaki K., 2004. Characterization of a new viroid strain from hops: evidence for viroid speciation by isolation in different host species. Journal of General Plant Pathology, 70(3): 181-187.
- Sano T., 2003. Hop stunt viroid. pp. 207-212. In: Viroids; (eds.): Hadidi A, Flores R., Randles J.W., Semancik J.S.; Collingwood, Australia, CSIRO Press.
- Sano T., Hataya T., Terai Y., Shikata E., 1989. Hop stunt viroid strains from dapple fruit disease of plum and peach in Japan. Journal of General Virology, 70: 1311-1319, doi: 10.1099/0022-1317-70-6-1311.
- Seigner L., Lutz A., Seigner E., 2014. Monitoring of important virus and viroid infections in German hop (*Humulus lupulus* L.) yards. Brewing Science, 67: 81-87.

- Skomra U., 2015. Metodyka integorwanej ochrony chmielu. Instytut Uprawy Nawożenia i Glebozawstwa - Państwowy Instytut Badawczy, Puławy.
- Skomra U., 2001. Virus occurrence in hop plants (*Humulus lupulus* L.) in the region of Lublin. Pamiętnik Puławski, 126: 107-124. (in Polish + summary in English)
- Tsai C.W., Daugherty M.P., Almeida R.P.P., 2012. Seasonal dynamics and virus translocation of *Grapevine leafroll-associated virus 3* in grapevine cultivars. Plant Pathology, 61: 977-985, doi: 10.1111/j.1365-3059.2011.02571.x.
- Wetzel T., Jardak R., Meunier L., Ghorbel A., Reustle G.M., Krczal G., 2002. Simultaneous RT/PCR detection and differentiation of arabis mosaic and grapevine fanleaf nepoviruses in grapevines with a single pair of primers. Journal of Virological Methods, 101(1-2): 63-69, doi: 10.1016/S0166-0934(01)00422-0.
- Yamamoto H., Kagami Y., Kurokawa M., Nishimura S., Ukawa S., Kubo S., 1973. Studies on hop stunt disease in Japan. Report of the Research Laboratories of Kirin Brewery Co., Ltd., 16: 49-62.
- Ziegler A., Kawka M., Przybys M., Doroszewska T., Skomra U., Kastirr U., Matoušek J., Schubert J., 2014. Detection and molecular analysis of Hop latent virus and Hop latent viroid in hop samples from Poland. Journal fur Kulturpflanzen, 66(7): 248-254, doi: 10.5073/JFK.2014.07.04.

The research was carried out as part of basic research for biological progress in plant production on the basis of the decision of the Minister of Agriculture and Rural Development HOR.hn.802.11.2018, Task 107

Author ORCID Marcin Przybyś 0000-0002-6567-2954

received – 26 November 2019 revised – 12 March 2020 accepted – 6 November 2020



This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution-ShareAlike (CC BY-SA) license (http://creativecommons.org/licenses/by/4.0/).