

To define biological properties of the pathogens, we conducted a host assay using indicator plants. Indicator plants were inoculated with sap obtained from cactus plants demonstrating virus-like symptoms. Necrotic local lesions observed on *Chenopodium murale* and *Gomphrena globosa* were typical for Cactus virus X (CVX) [5,6]. In indirect ELISA we observed positive reactions with sap *Mamillaria microhelix*, *Sulcorebutia* sp., *Bolivocereus samupatanus*, *Opuntia* sp., *Ferocactus* sp., *Mamillaria magnimamma*, *Opuntia* sp., *Cereus* sp., *Astrophytum capricorne*, *Astrophytum myriostigma* and *Ritterocereus pruinosus* with antiserum to Potato virus X (PVX, which is serologically related to Cactus virus X).

Comparing the results of bioassay and ELISA tests we further focused on samples of *Mamillaria microhelix*, *Sulcorebutia* sp., *Bolivocereus samupatanus*, *Opuntia* sp., *Ferocactus* sp., *Mamillaria magnimamma*, *Opuntia* sp., *Cereus* sp., *Astrophytum capricorne*, *Astrophytum myriostigma* from collection of Nikita Botanical Garden and *Ritterocereus pruinosus* from the collection of Karazin' Botanic Garden of Kharkiv National University. These plants were probably infected with CVX. To confirm our assumption about CVX infection and to study the morphology of the pathogen we carried out transmission electron microscopy. In sap of all plants we registered filamentous virions with size $580 \times 13 \pm 2$ nm (Fig.3), which is typical for *Cactus virus X*.

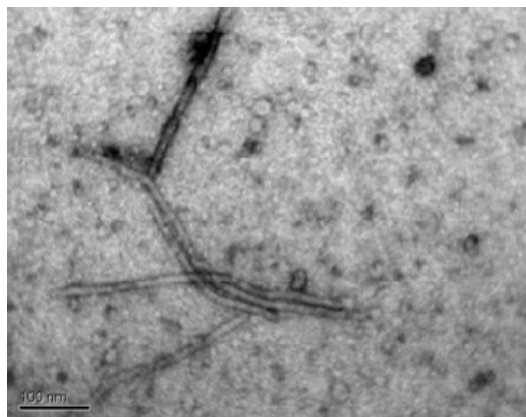


Fig.3. Electron micrograph of flexible particles from plant material from *Ferocactus* sp.

Summarizing the obtained results it is possible to assert that collections of Nikitsky Botanical Garden and Botanical garden of Karazin's Kharkiv National University were contaminated by Cactus virus X. Exchange collection material without testing on virus infection, can increase chances of uncontrolled distribution of viral infections. Some infections are symptomless. In this case the asymptomatic virus infection in cactus makes a danger because these plants can become the source of distribution of pathogens in collections. In addition, cactuses could support reproduction of viruses of other types of plants and, thus, be the reservoirs of plant virus infections. A timely exposure and permanent control of the state of population of these cultural plants is the obligatory link of the system of their protecting from this group of pathogens.

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Надійшла до редколегії 24.04.12

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ELECTRON MICROSCOPY AND BIOLOGICAL PROPERTIES OF THE PATHOGEN AFFECTING *PLATANThERA BIFOLIA*

Проведено обстеження хворих рослин *Platanthera bifolia*, отриманих в Канівському природному заповіднику. Була доведена інфекційність соку, а також можливість механічної передачі вірусу. Методом трансмісійної електронної мікроскопії в зразках уражених рослин були виявлені сферичні частинки, діаметром около 50 нм. Згідно морфологічних особливостей, вірус, виділений з рослин *Platanthera bifolia*, може належати до родини *Caulimoviridae*.

The plants of *Platanthera bifolia* collected in Kaniv National Reserve were observed. The infectivity of plant sap and viral capability to transmit in mechanical manner were confirmed. The spherical particles with diameter about 50 nm were observed in preparations from symptomatic leaf tissue in transmission electron microscope. According to viral morphology the pathogen of *Platanthera bifolia* may be a member of *Caulimoviridae*.

Introduction. *Orchidaceae* Juss. is one of the biggest families of flowering plants which includes about 35 000 species. They can be found all over the world excluding deserts and Polar Region [2]. Viral diseases of orchids are known from the middle of XX century [4]. Until now more over 30 viruses of orchids have been described [8, 10]. The

majority of these viruses were detected as pathogens of tropical orchids which are cultivated *in situ*. On the other hand the viruses of terrestrial orchids of temperate zone are studied not enough. The infection of *Cypripedium* sp., *Orchis* sp., *Ophrys* sp. by Tobacco rattle (TRV) and Turnip mosaic virus (TuMV) were described [5]. Also the antigens

to *Arabis mosaic virus* (ArMV), *Bean yellow mosaic virus* (BYMV), *Tobacco mosaic virus* (TMV), *Tomato aspermy virus* (TAV), TRV and TuMV were detected in orchids of natural Ukrainian flora [1].

Materials and methods. Plant material was collected in Kaniv Natural Reserve (Snakes' Islands). For further investigation we used plants of *Platanthera bifolia* with typical viral symptoms (fig. 1).



Figure 1. Necrotic lesions on the leaves of *Platanthera bifolia*

Virus identification was carried out using standard indirect ELISA and DAS-ELISA with polyclonal antisera to *Cucumber mosaic virus* (CMV), *Tomato spotted wilt virus* (TSWV), *Potato virus Y* (PVY), BYMV, TAV, TMV, TRV, TuMV and *Cauliflower mosaic virus* (CaMV).

Biological properties of viruses were studied using following test-plants: *Chenopodium murale*, *Gomphrena globosa*, *Lycopersicon esculentum*, *Nicotiana glauca*, *N. benthamiana*, *N. rustica*, *N. tabacum*, *Petunia hybrida* and *Zinnia elegans*. The test-plants were inoculated in early stages of growth by mechanical sap transmission, applying carborundum as an abrasive. The inoculum was prepared by homogenizing plant tissue with 0.1 M phosphate buffer (pH 7.4) and followed by centrifugation (5000 rpm for 20 min).

The morphology of virions were studied in leaf dip preparations negatively stained with 2% uranyl acetate EM using a JEM-1400 electron microscope, magnification 60 000.

Results and discussion. The ELISA testing revealed no antigens of BYMV, CMV, TAV, TMV, TRV, TSWV, TuMV and PVY in samples of infected *P. bifolia*.

The sap of necrotic leaves from *P. bifolia* were experimentally inoculated to the range of test-plants, common for many polyhostal viruses [3, 7]. All inoculated plants show no local reaction but *L. esculentum*, *N. benthamiana*, *N. rustica* and *N. tabacum* reacted systemically. The systemic symptoms of viral infection appeared two months after inoculation. No visual symptoms were observed on *C. murale*, *G. globosa*, *P. hybrida* and *Z. elegans* (table 1).

Table 1. Reaction of test plants inoculated with sap from *Platanthera bifolia*

Test-plant	Symptoms
<i>Chenopodium murale</i>	-/-*
<i>Gomphrena globosa</i>	-/-
<i>Lycopersicon esculentum</i>	Chlorotic lesions, leaf deformation
<i>Nicotiana glauca</i>	-/-
<i>Nicotiana benthamiana</i>	Mosaic
<i>Nicotiana rustica</i>	Mosaic
<i>Nicotiana tabacum</i>	Chlorosis
<i>Petunia hybrida</i>	-/-
<i>Zinnia elegans</i>	-/-

*"-/-" – no visual symptoms

EM examination of infected plants tissue revealed the presence of spherical virus-like particles about 50 nm in diameter (fig. 2). Such morphology are typical for represen-

tatives of *Caulimoviridae*: *Caulimovirus*, *Cavemovirus*, *Petuvirus*, *Soymovirus* [9].

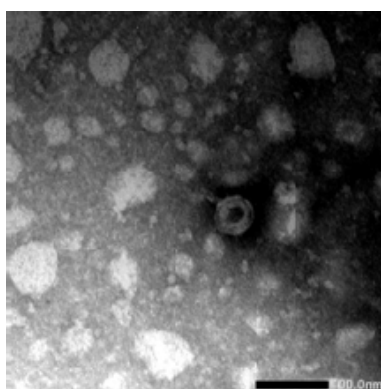


Figure 2. Electron micrograph of virus-like particles in the sap of infected plants

The majority of viruses belonging to the listed genera can be transmitted by mechanical inoculation. The reaction of test-plants eliminates such viruses as *Blueberry red ringspot virus*, *Cestrum leaf curling virus*, *Petunia vein clearing virus* and *Tobacco vein clearing virus* which are not transmitted in mechanical manner [3, 6].

For further identification of virus DAS-ELISA test were used. No positive reaction with antiserum to CaMV was obtained. According to results of DAS-ELISA we can exclude infection of *P. bifolia* by CaMV and serologically related viruses: *Carnation etched ring virus*, *Dahlia mosaic virus*, *Horseradish latent virus* and *Strawberry vein banding virus* [3].

In conclusion, host assay of sap samples extracted from infected plants of *P. bifolia* approved its infectivity. EM studies confirmed the presence of virus-like particles in infected plant tissues. The isometric morphology and size range allows to suggest that the viral pathogen of *P. bifolia* can be a representative of family *Caulimoviridae*. The cases of similar virus infection of orchids are not described in literature data. However, there are not much investigation about spreading

viruses in natural ecosystems. Further work will be focused on identification of the virus infecting *P. bifolia*.

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Надійшла до редколегії 24.04.12

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BIOTECHNOLOGICAL (GM) CROPS IN THE WORLD AND EUROPE, PLUM HONEYSWEET RESISTANT TO PPV, RESEARCH IN THE CR: THE PAST, THE PRESENT AND THE FUTURE

Быстрое увеличение площадей посевов ГМ-культур, делает их наиболее высокоадаптивными посевными культурами в истории сельского хозяйства. Здесь представлены результаты девяти лет тестирования слив клона C5 (трансформирован капсидным белком вируса шарки сливы) при высокой и постоянной инфекционной нагрузке PPV-Rec, как самостоятельного агента, так и в сочетании с вирусами карликовости чернослива и кольцевой хлоротичности яблони.

Fast increasing hecтарage of GM crops makes them the highly adopted crop technology in the history of agriculture. Results of nine years testing of plum clone C5 (transformed with the Plum pox virus coat protein) under the high and permanent infection pressure of PPV-Rec alone and in combinations with Prune dwarf virus, and Apple chlorotic leafspot virus both from graft inoculation and natural aphid vectors are presented.

Introduction. Research in genetic engineering started in late seventies (Colwell et al., 1985). The first genetically modified (GM), syn. biotechnological (Biotech) crops were commercialized in 1995 (cotton, company Monsanto; potato, company Syngenta). 1.7 million hectares of Biotech crops (cotton, corn, potato) were planted in 1996 already (USDA-APHIS Biotechnology Permits Database, available at <http://www.nbiap.vt.edu/> (March 2001). Increase in hecтарage to 160 million hectares in 2011 (Fig. 1) makes Biotech crops the fastest adopted crop technology in the his-

tory of agriculture. The International Service for the Acquisition of Agri-biotech Applications (ISAAA) is every year publishing "Global status of commercialized Biotech/GM crops". The last one was published as ISAAA Brief No. 43 in 2011 (James C., 2011). The objective of this Brief is to provide information and knowledge to the scientific community and society on Biotech/GM crops and its contribution to global food, feed, fiber and fuel security, to sustainable agriculture.

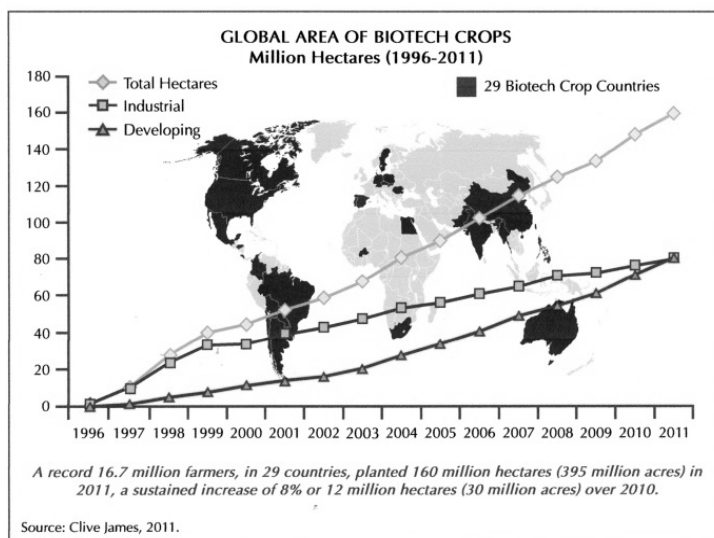


Fig. 1. Global area of biotech crops