

Occurrence of strawberry viruses in *Fragaria* germplasm and evaluation of cryotherapy as an eradication method for strawberry viruses

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Abstract

Strawberry plants are highly susceptible to viral infections, which pose significant threats to global strawberry production. This study aims to explore the efficacy of *in vitro* initiation and cryopreservation of shoot tips as a potential strategy for eradicating strawberry viruses. We tested plants for four important strawberry viruses namely: SMoV, SCV, SMEY and SVBV. The plants, which tested positive were either cultivated as *in vitro* cultures then returned to a green house or field collection cultivation, or treated by cryopreservation. After cryopreservation, the plants were cultivated again *in vitro* and then in the green house or field. The viruses were detected within each propagation step. Significant eradication effects were found for SMoV and SCV when plants were treated by *in vitro* initiation or with cryotherapy, but not for SMEY or SVBV. The results of this study show that cryotherapy or *in vitro* initiation can lead to the elimination of strawberry viruses, but the kind of therapy appears to depend on the type of virus.

Keywords: *Fragaria*, virus eradication, cryotherapy

Text

Strawberries are one of the most economically important temperate fruit crops, with an annual production of 9.157,127.5 t on an area of 389,665 ha worldwide in 2021 (FAO stat, <https://www.fao.org/faostat/en>). The main producing countries are USA, Netherlands, Morocco, Spain and Albania. With a percentage of 3.1% of the German

fruit production, strawberry cultivation was the third-largest in Europe with a yield of 130,630 tonnes on an area of 12,500 ha in 2021 (FAO stat, <https://www.fao.org/faostat/en>). For the successful cultivation of strawberries, it is necessary to provide virus-free plant material. Virus infections are one main reason for the degeneration of propagation material in strawberries. Once infected, vegetative propagation transmits the viruses from one propagation phase to the next. An infested plant weakens the plant in the long term, leading to increased pathogen susceptibility. However, the virus infection itself also leads to economic losses due to bad fruit quality, deformation of leaves and other symptoms (Martin and Tzanetakis 2006). More than 25 viruses have been described for strawberries to date (Fránová et al. 2019, Koloniuk et al 2022a), which were transmitted via insects, nematodes or other vectors (Bragard et al. 2019, Martin and Tzanetakis et al. 2006, Franova et al. 2019, Koloniuk et al 2022b). Martin and Tzanetakis (2006) reported aphid transmitted viruses, mainly, the *strawberry mottle virus* (SMoV), *strawberry mild yellow edge virus* (SMYEV), *strawberry crinkle virus* (SCV) and *strawberry vein banding virus* (SVBV) as the most economically important ones in strawberry cultivation areas of the world.. Although control of field infestation of the vector *Chaetosiphon fragaefolii* (strawberry aphid) is possible (reviewed in CABI 2022), once a plant is infected, the only way to stop virus dissemination is an eradication of infested plants (Greber 1979, Boxus 1989, Nazarov et al. 2020, Rubio et al. 2020). The generation of virus free plants is an important task for the provision of plants for vegetative propagation, cultivation and preservation of genetic resources. Methods for virus elimination are described for several cultivated plant species and are mainly chemotherapy (Faccioli 2001, Modarresi Chahardehi et al. 2016, AlMaarri et al. 2012), thermotherapy (Faccioli 2001, Wang et al. 2006, AlMaarri et al. 2012, Waswa et al. 2017, Zhao et a. 2018), electrotherapy (AlMaarri et al. 2012), cryotherapy (Zhao et al. 2018) or meristem culture (Faccioli 2001, Quazi and Martin, 1978, Wang et al. 2006, Zhang et al. 2019). For strawberries cryotherapy, thermotherapy and *in vitro* culture techniques were described for single virus eradication (Boxus 1976, McGrew 1965). However, cryotherapy has not been investigated for the eradication of different strawberry viruses. This study investigated the occurrence of strawberry viruses in the

germplasm repository in the Fruit Genebank of the Julius Kühn-Institute (JKI) Dresden-Pillnitz and used the well-established method of cryopreservation (Höfer et al. 2016) as a possible method for the eradication of different strawberry viruses.

The plant material was obtained from the *Fragaria* collection of the Fruit Genebank of the Julius Kuehn Institut (JKI). Seventy-seven cultivars and seven unassigned accessions of *Fragaria ×ananassa* as well as 168 accession of *Fragaria* wild species and hybrids were tested for four strawberry viruses in the field (see list of the tested cultivars and wild species accession in supplemental material table S1). PCR was used to test and detect four strawberry viruses namely: (SMoV - strawberry mottle virus, SCV – strawberry crinkle virus, SMYEV – strawberry mild yellow edge virus, SVBV – strawberry vein banding virus). A mix of different leaves of up to eight plant samples per accession (n=1-8) were collected (see table S1) for virus detection in the cultivar collection. For initial virus detection in the wild species collection, a mix of different leaves from up to three plants per accession was collected and tested as one sample (n=1). Between three to 10 plant samples (n=3-10) per cultivar were collected for the detection of viruses in the set of 19 cultivars for evaluation of virus eradication efficiency in the field, after in vitro initiation, after cryo-conservation and finally after transfer into the greenhouse again. RNA was isolated from 40 mg leaf material, and the invitrap Spin RNA Mini Kit (Invitac Molecular GmbH, Berlin, Germany) was used for extraction according to the manufacturer's protocol. The RNA obtained was diluted in 50 µl dd H₂O. The quantification of the isolated nucleic acid was performed on the NanoDrop 2000c device. Synthesis of cDNA was performed using the Revert Aid First Strand cDNA Synthesis Kit (Thermo Fisher) according to the manufacturer's protocol. Random hexamer oligos and oligo_dT18-nucleotides were used for the synthesis. A total of 1 µg RNA was the input amount for cDNA synthesis per sample. Successful cDNA synthesis was evaluated using a standard PCR method using elongation factor EF specific primers (EF_F und EF_R, Flachowsky et al. 2007). The PCR conditions were: 13,4 µl ddH₂O, 2.5 µl 10x DreamTaq Puffer with 20 mM MgCl₂, 2.5 µl dNTPs with 2 mM, 1.25 µl of 10 µM EF1αF and EF1αR, 1 µl 20x red buffer, 1 µl of 0,125% BSA (Zhang et al. 2014), 1 µl of 25% PVP (Koonjul et al. 1999) and 0,1 µl of 5 U µl⁻¹ DreamTaq polymerase. A total of 1µl cDNA was used for each

90 PCR reaction. PCR was performed with 1 x initial denaturation: 94 °C 3', 35 x
91 denaturation/annealing/elongation: 94 °C 30''/56 °C 1'/72 °C 1', 1 x final elongation:
92 72 °C 3' and 1 x cooling: 10 °C ∞. The primer sequences to proof strawberry leaf
93 material on the occurrence of strawberry viruses was obtained from the publication
94 listed in table S2 and PCR was performed according to the mastermix and conditions
95 in table S3. Amplificates of investigated samples, positive and negative control
96 samples will be separated by agarose gel electrophoresis. For each sample 10 µl PCR
97 product is loaded into a 1,5 % agarose gel and separated at 90 Volt. A 50 bp size
98 standard (Thermo-Fisher Scientific) is used. Positive samples amplify the specific
99 fragment, whereas negative samples obtained no fragment. The evaluation of virus
100 eradication effect by cryotherapy compared to *in vitro* initiation was tested on 19
101 cultivars (Coral, Dukat, Florika, Fraginetta, Gloria, Mieke Nova, Mrak, Pantagruella,
102 Papa Lange, Pegasus, Pervagata, Polka, Rosa Perle, Rubia, Senga Dulcita, Senga
103 Gigana, Symphony, Talisman, Triscana). For the evaluation of virus eradication effect
104 by cryotherapy, samples of the cultivars were obtained from the field (test phase – A).
105 Stolons of positive tested plants were obtained and shoot tips were isolated in the
106 laboratory according to the experimental procedures described in Höfer (2011). Up to
107 three single shoot tips of virus positive plants (n= up to 3) were dissected to establish
108 *in vitro* cultures before cryotherapy (test phase – B). Negatively tested plants obtained
109 from *in vitro* cultures were used for re-transmission from the laboratory into the
110 greenhouse (test phase – C) for virus retesting to study the effect of shoot tip
111 dissection on virus elimination. Up to three individual plants (n= up to 3) were used for
112 virus testing. *In vitro* apical shoot tips from positive tested cultivars were dissected
113 from up to 4-week-old *in vitro* plants and the method described in Höfer et al. 2016
114 was performed for cryopreservation and recovery of plant shoot tips. Up to 10 plants
115 per cultivar were (n= up to 10) were tested on the occurrence of viruses (in-vitro culture
116 after cryo, test phase – D). After transmission of recovered plants into the greenhouse
117 (test phase – E, plants were tested on the occurrence of viruses as described for initial
118 virus testing (n= up to 9). The frequency of positive tested plant samples per virus, the
119 percentage of positive and negative tested cultivars/species was calculated. For the

19 cultivars mentioned above, the frequency of positive tested samples for each virus was calculated per cultivar and test phase.

A total of 84 *Fragaria xananassa* accessions and 164 accessions of 22 *Fragaria* wild species and hybrids were tested on the occurrence of four strawberry viruses. An example of the detection results obtained by PCR for each single virus is shown in figure 1. Table 1 shows the percentage of positive tested samples. The highest virus frequency in *Fragaria xananassa* was observed for the SCV (73.2%) and SMYEV (72.1%). A lower frequency was obtained for SMoV (57.5%) and SVBV (4.3%). Single virus frequencies determined for each species are shown in table 1. To determine the frequency of each virus over all tested samples and species, a mean frequency of each virus was calculated. The most frequent virus was SMYEV (74.5%), whereas SCV (35.9%), SMoV (30.9%) and SVBV (11.8%) showed a lower mean frequency. Between 6.8 % (SVBV) and 82.9 % (SMYEV) of samples collected from 19 cultivars tested positive for all four viruses in field. The effect of virus elimination when shoot tips were isolated from stolons of infected strawberry plants to establish *in vitro* cultures resulted in 26.3 % (SMYEV) to 98.2 % (SVBV) virus free plants. After re-transmission into the green house between 2.6 % (SCV) to 76.3 % (SMYEV) of the tested plants were re-infected with viruses (table 2). The effect of cryotherapy was also investigated and 14.9 % (SMYEV) to 100 % (SCV) negative tested plants were obtained. After re-transmission of cryotherapy threatened plants into the green house between 22.6 % (SMYEV) and 100 % (SCV) of plants tested negative on the strawberry viruses. The results are shown in table 3.

Strawberries are highly susceptible to strawberry viruses, and sources of resistance to the viruses or vectors are not investigated so far (Shanks and Barrit 1974, Barrit and Shanks 1980). Chemical controls against the vectors are also possible, but only with very high application rates, which is contrary to current socio-economic developments. Once a plant is infected, it can only be eradicated and new virus-free plant material has to be made available. Providing virus-free plant material for new plantings is therefore the best strategy so far (Bettoni et al. 2022). In this study, we therefore investigated the effect of *in vitro* initiation and cryopreservation on virus

elimination on strawberry (Figure 1A). Significant eradication effects were found for all viruses by *in vitro* initiation and further by cryopreservation (Table 2 and 3). In potatoes, Bettonie et al. 2022 and Kushnarenko et al. 2017 showed a high elimination rate against three viruses by chemotherapy and cryotherapy. In other species such as raspberry (35%), sweet potato (100%), banana (34 to 90%), grapevine (96% to 100%), quince (33 to 37%), apple (35 to 100%) and *Prunus spec.* (50%), cryotherapy was also successfully performed to eliminate viruses (Harding et al. 2004, Helliott et al. 2002, Feng et al. 2013, Cui et al. 2015; Pathirana et al. 2015, 2019, Farhadi-Toolii et al. 2022, Wang et al. 2022a). In addition to cryotherapy, this study confirms that *in vitro* initiation (Table 2) already leads to a reduction on strawberry viruses, which was previously shown by Boxus (1976). However, the experiments also show that no effect could be detected for the eradication of SMYEV. Binhua et al. (2008) especially reports the successful elimination of SMYEV by freezing, which is contradictory to the results of that study. This virus showed the highest frequency in the tested plant assortment. Whether this virus can be successfully eliminated in combination with heat or chemotherapy (Bettonie et al. 2022) remains to be answered in future research projects.

Acknowledgements

We would like to thank Sabine Bartsch for her technical assistance with sampling, DNA isolation and virus testing, and Ute Sonntag and Katrin Winkler for her assistance with inculturing and cryotherapy of strawberries.

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Figures

Figure 1 (A) Schematic phases of virus elimination by in vitro initiation and cultivation
 or cryopreservation (A-E). (B) Detection of four strawberry viruses with RT-PCR. E1
 SMYEV –positive control of strawberry mild yellow edge virus with a band at 271 bp,
 E2 SCV –positive control of strawberry crinkle virus with a band at 345 bp, E3 SVBV
 –positive control of strawberry vein banding virus with a band at 435 bp, E4 –positive
 control of strawberry mottle virus with a band at 219 bp, E1-E4 –negative control using
 the AtropaNad2 band at 188 bp, 50bp – size marker.

Tables

Table 1 Frequency of four strawberry viruses in *Fragaria* germplasm.

Table 2 Results from the evaluation of strawberry virus eradication via stolon
 meristem explant isolation and re-transmission into the green house

320 **Table 3** Results from the evaluation of strawberry virus eradication via stolon
 321 meristem explant isolation, cryopreservation treatment and re-transmission into the
 322 green house

323

324 **Supplemental tables**

325 **Table S1** Tested accessions used in this study.

326 **Table S2** Primer sequences to proof strawberry leaf material on the occurrence of 4
 327 strawberry viruses.

328 **Table S3** Mastermix and PCR conditions for strawberry virus detection.

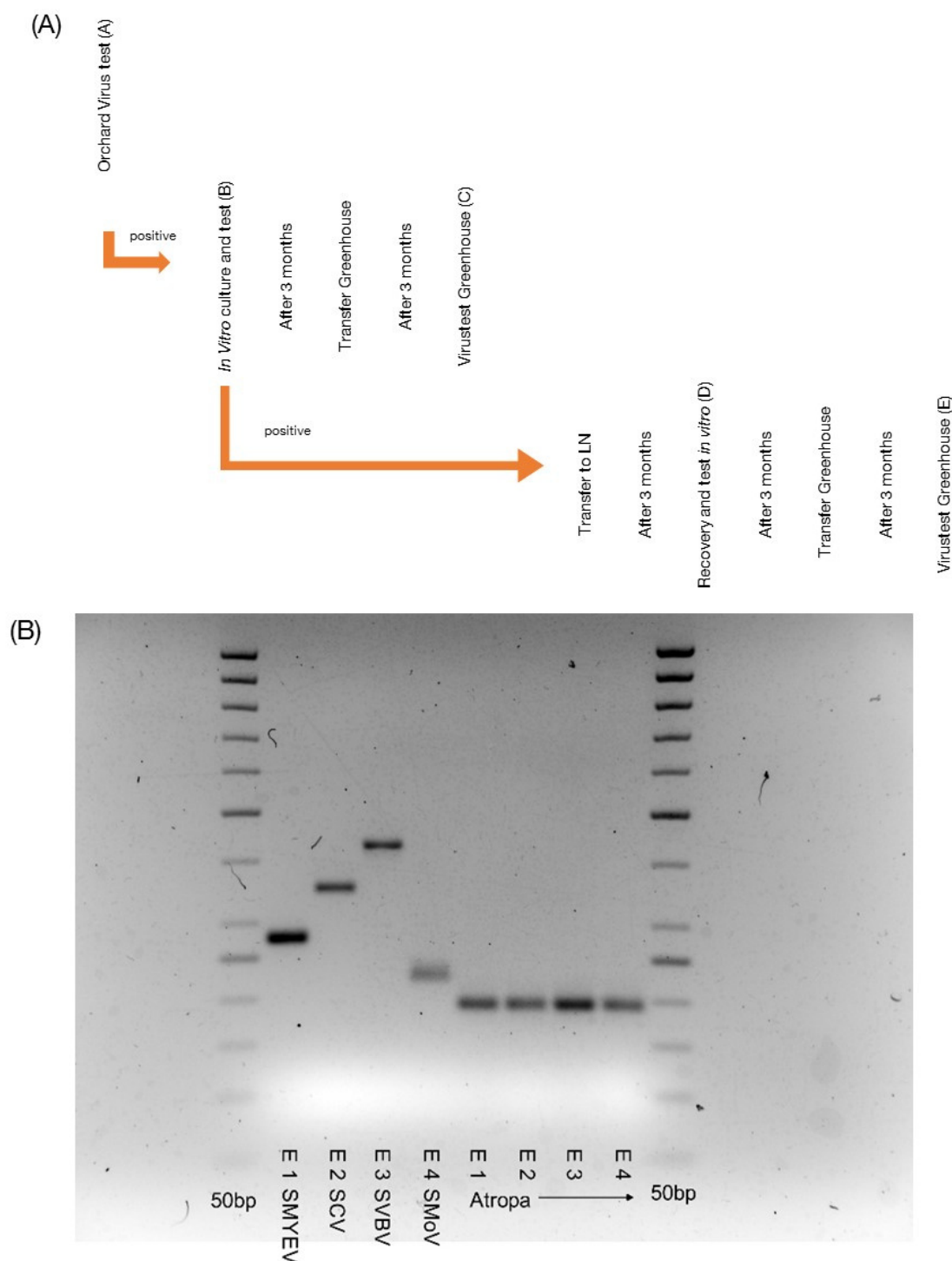


Figure 1A Schematic phases of virus elimination by in vitro initiation and cultivation or cryopreservation (A-E), LN – liquid nitrogen. 1B Detection of four strawberry viruses with RT-PCR. E1 SMYEV –positive control of strawberry mild yellow edge virus with a band at 271 bp, E2 SCV –positive control of strawberry crinkle virus with a band at 345 bp, E3 SVBV – positive control of strawberry vein banding virus with a band at 435 bp, E4 –positive control of strawberry mottle virus with a band at 219

bp, E1-E4 –negative control using the AtropaNad2 band at 188 bp, 50bp – size marker.

Table 1 Frequency of four strawberry viruses in *Fragaria* germplasm.

Species	No. of samples	% positive tested samples* / accessions			
		SMoV	SMYEV	SCV	SVBV
<i>Fragaria xananassa</i>	280	57.5	72.1	73.2	4.3
<i>Fragaria x bifer</i>	3	100.0	100.0	100.0	66.7
<i>Fragaria x bringhurstii</i>	1	0.0	100.0	100.0	0.0
<i>Fragaria bucharica</i>	8	12.5	75.0	37.5	0.0
<i>Fragaria chiloensis</i>	19	36.8	100.0	78.9	15.8
<i>Fragaria corymbosa</i>	9	22.2	44.4	0.0	33.3
<i>Fragaria gracilis</i>	4	50.0	0.0	0.0	0.0
<i>Fragaria hybr.</i>	2	50.0	100.0	50.0	0.0
<i>Fragaria iinumae</i>	1	0.0	100.0	0.0	100.0
<i>Fragaria mandshurica</i>	8	25.0	75.0	25.0	0.0
<i>Fragaria moschata</i>	10	30.0	10.0	40.0	10.0
<i>Fragaria moupinensis</i>	1	0.0	0.0	0.0	0.0
<i>Fragaria nilgerrensis</i>	5	0.0	100.0	0.0	20.0
<i>Fragaria nipponica</i>	7	14.3	100.0	42.9	0.0
<i>Fragaria nubicola</i>	6	0.0	100.0	33.3	0.0
<i>Fragaria orientalis</i>	7	28.6	100.0	28.6	0.0
<i>Fragaria pentaphylla</i>	3	66.7	0.0	0.0	0.0
<i>Fragaria sp.</i>	19	5.3	42.1	5.3	0.0
<i>Fragaria tibetica</i>	4	0.0	100.0	0.0	0.0
<i>Fragaria vesca</i>	18	38.9	100.0	83.3	5.6
<i>Fragaria virginiana</i>	16	43.8	93.8	50.0	0.0
<i>Fragaria viridis</i>	6	100.0	100.0	50.0	16.7
<i>Fragaria yezoensis</i>	7	28.6	100.0	28.6	0.0

* in case of *Fragaria xananassa* more then one sample per accession were tested

Table 2 Results from the evaluation of strawberry virus eradication via stolon meristem explant isolation and re-transmission into the green house

No.	test phase	plant material	virus	samples tested	no. positive	no. negative	%- pos.	%-difference from 100 % pos. tested plant material
A	from field collection	leaves from flower box	SMoV	76	51	25	69.8	-
			SMYEV		58	18	82.9	-
			SCV		58	18	73.3	-
			SVBV		5	71	6.8	-
↓ only positive plants from phase (I) were used in phase (II)								
B	in-vitro culture before cryo	explants from stolon meristem	SMoV	47	15	32	31.6	68.4
			SMYEV		35	12	73.7	26.3
			SCV		3	44	6.1	93.9
			SVBV		1	46	1.8	98.2
↓ only negative plants from phase (II) were used in phase (III)								
C	re-transmission from the lab into green house	leaves from flower box	SMoV	49	8	41	17.5	82.5
			SMYEV		38	11	76.3	23.7
			SCV		2	47	2.6	97.4
			SVBV		2	47	4.4	95.6

Table 3 Results from the evaluation of strawberry virus eradication via stolon meristem explant isolation, cryopreservation treatment and re-transmission into the green house

No.	test phase	plant material	virus	samples tested	no. positive	no. negative	%-pos.	%-difference from 100 % pos. tested plant material
↓ only positive plants from phase (I) were used in phase (II)								
B	in-vitro culture before cryo	explants from stolon meristem	SMoV	47	15	32	31.6	68.4
			SMYEV		35	12	73.7	26.7
			SCV		3	44	6.1	93.9
			SVBV		1	46	1.8	98.2
↓ only positive plants from phase (II) were used in phase (III)								
D	in-vitro culture after cryo	explants from stolon meristem	SMoV	111	12	99	9.0	91.0
			SMYEV		99	12	85.1	14.9
			SCV		0	111	0	100.0
			SVBV		2	109	1.1	98.9
↓ only negative plants from phase (III) were used in phase (IV)								
E	re-transmission from the lab after cryo	leaves from flower box	SMoV	108	10	98	9.1	90.9
			SMYEV		84	24	77.4	22.6
			SCV		0	108	0	100.0
			SVBV		1	107	0.4	99.6

Table S1 Tested accessions used in this study.

Accession no.	Species	cultivar name
ERB0018	Fragaria ×ananassa	Asinigra
ERB0048		Calea
ERB0065		Coral
ERB0077		Demerland
ERB0089		Dukat
ERB0094		Elsanta
ERB0115		Fraginetta
ERB0120		Fraroma
ERB0136		Gento
ERB0142		Gloria
ERB0144		Gorella
ERB0171		Imtraga
ERB0180		Joghana
ERB0186		Jurica
ERB0195		Korbinskaya rannyaya
ERB0201		Lihama
ERB0209		Machern
ERB0239		Optima
ERB0240		Orion
ERB0245		Papa Lange
ERB0251		Pervagata
ERB0253		Pink Panda
ERB0255		Polka
ERB0257		Senga Precosa
ERB0258		Senga Precosana
ERB0262		Prinz Julius Ernst
ERB0272		Redgauntlet
ERB0277		Rigensa
ERB0281		Rosella
ERB0283		Roter Regen
ERB0285		Rubia
ERB0288		Rupine
ERB0295		Sara
ERB0297		Schloß Horneburg
ERB0300		Seligra
ERB0301		Senga Dulcita
ERB0302		Senga Gigana
ERB0313		Silvia
ERB0322		Spadeka
ERB0331		Sturms Zuckersüße
ERB0336		Surprise des Halles
ERB0338		Sweetheart
ERB0339		Symphony
ERB0342		Talisman
ERB0346		Tenira
ERB0348		Thielesa
ERB0349		Thuriga
ERB0350		Tina

Accession no.	Species	cultivar name
ERB0355		Tribute
ERB0356		Triscana
ERB0362		Unermüdliche
ERB0390		Mrak
ERB0391		Pantagruella
ERB0392		Tioga
ERB0393		Royal Sovereign
ERB0398		Paula
ERB0401		Frabella
ERB0403		Tago
ERB0407		Pegasus
ERB0409		Profumata di Tortona
ERB0419		Mieze Nova
ERB0422		Multiplex
ERB0423		Rosa Perle
ERB0424		Quarantaine de Prin
ERB0425		Blanc Amélioré
ERB0426		Little Scarlet
ERB0427		Muricata
ERB0429		Sannié
ERB0430		Gartenfreude
ERB0432		Marie Charlotte
ERB0433		Ronja
ERB0434		Weiß Hagmann
ERB0435		Florika
ERB0436		Linné
ERB0437		Lucida Perfecta
ERB0438		Illa Martin
ERB0440		Ulrichsberg
FRA0001	<i>Fragaria bucharica</i>	-
FRA0002	<i>Fragaria bucharica</i>	-
FRA0003	<i>Fragaria bucharica</i>	-
FRA0004	<i>Fragaria bucharica</i>	-
FRA0005	<i>Fragaria bucharica</i>	-
FRA0006	<i>Fragaria bucharica</i>	-
FRA0007	<i>Fragaria bucharica</i>	-
FRA0011	<i>Fragaria chiloensis</i>	-
FRA0012	<i>Fragaria chiloensis</i>	-
FRA0013	<i>Fragaria chiloensis</i>	-
FRA0015	<i>Fragaria chiloensis</i>	-
FRA0022	<i>Fragaria chiloensis</i>	-
FRA0023	<i>Fragaria corymbosa</i>	-
FRA0024	<i>Fragaria corymbosa</i>	-
FRA0025	<i>Fragaria corymbosa</i>	-
FRA0026	<i>Fragaria corymbosa</i>	-
FRA0027	<i>Fragaria corymbosa</i>	-
FRA0028	<i>Fragaria corymbosa</i>	-
FRA0029	<i>Fragaria corymbosa</i>	-
FRA0030	<i>Fragaria corymbosa</i>	-

Accession no.	Species	cultivar name
FRA0031	<i>Fragaria corymbosa</i>	-
FRA0033	<i>Fragaria gracilis</i>	-
FRA0034	<i>Fragaria gracilis</i>	-
FRA0035	<i>Fragaria gracilis</i>	-
FRA0036	<i>Fragaria gracilis</i>	-
FRA0037	<i>Fragaria</i> sp.	-
FRA0038	<i>Fragaria</i> hybr.	-
FRA0039	<i>Fragaria iinumae</i>	-
FRA0041	<i>Fragaria mandshurica</i>	-
FRA0042	<i>Fragaria mandshurica</i>	-
FRA0045	<i>Fragaria mandshurica</i>	-
FRA0046	<i>Fragaria mandshurica</i>	-
FRA0048	<i>Fragaria moschata</i>	-
FRA0050	<i>Fragaria moschata</i>	-
FRA0052	<i>Fragaria moschata</i>	-
FRA0054	<i>Fragaria moschata</i>	-
FRA0057	<i>Fragaria moschata</i>	-
FRA0058	<i>Fragaria moschata</i>	-
FRA0061	<i>Fragaria moschata</i>	-
FRA0066	<i>Fragaria moschata</i>	-
FRA0068	<i>Fragaria moschata</i>	-
FRA0073	<i>Fragaria moschata</i>	-
FRA0075	<i>Fragaria</i> hybr.	-
FRA0076	<i>Fragaria moupinensis</i>	-
FRA0077	<i>Fragaria nilgerrensis</i>	-
FRA0078	<i>Fragaria nilgerrensis</i>	-
FRA0079	<i>Fragaria nilgerrensis</i>	-
FRA0080	<i>Fragaria nilgerrensis</i>	-
FRA0081	<i>Fragaria nilgerrensis</i>	-
FRA0084	<i>Fragaria nipponica</i>	-
FRA0085	<i>Fragaria nipponica</i>	-
FRA0087	<i>Fragaria nubicola</i>	-
FRA0088	<i>Fragaria nubicola</i>	-
FRA0089	<i>Fragaria nubicola</i>	-
FRA0090	<i>Fragaria orientalis</i>	-
FRA0091	<i>Fragaria orientalis</i>	-
FRA0092	<i>Fragaria orientalis</i>	-
FRA0093	<i>Fragaria orientalis</i>	-
FRA0095	<i>Fragaria orientalis</i>	-
FRA0096	<i>Fragaria pentaphylla</i>	-
FRA0097	<i>Fragaria pentaphylla</i>	-
FRA0098	<i>Fragaria pentaphylla</i>	-
FRA0099	<i>Fragaria</i> sp.	-
FRA0100	<i>Fragaria mandshurica</i>	-
FRA0101	<i>Fragaria mandshurica</i>	-
FRA0102	<i>Fragaria mandshurica</i>	-
FRA0103	<i>Fragaria</i> sp.	-
FRA0104	<i>Fragaria</i> sp.	-
FRA0105	<i>Fragaria</i> sp.	-

Accession no.	Species	cultivar name
FRA0106	<i>Fragaria</i> sp.	-
FRA0107	<i>Fragaria</i> sp.	-
FRA0108	<i>Fragaria</i> sp.	-
FRA0110	<i>Fragaria</i> sp.	-
FRA0111	<i>Fragaria</i> sp.	-
FRA0112	<i>Fragaria</i> sp.	-
FRA0113	<i>Fragaria</i> sp.	-
FRA0114	<i>Fragaria</i> sp.	-
FRA0115	<i>Fragaria</i> sp.	-
FRA0118	<i>Fragaria mandshurica</i>	-
FRA0119	<i>Fragaria</i> sp.	-
FRA0120	<i>Fragaria</i> sp.	-
FRA0121	<i>Fragaria</i> sp.	-
FRA0122	<i>Fragaria</i> sp.	-
FRA0123	<i>Fragaria</i> sp.	-
FRA0125	<i>Fragaria tibetica</i>	-
FRA0127	<i>Fragaria tibetica</i>	-
FRA0128	<i>Fragaria tibetica</i>	-
FRA0135	<i>Fragaria vesca</i>	-
FRA0140	<i>Fragaria vesca</i>	-
FRA0142	<i>Fragaria vesca</i>	-
FRA0150	<i>Fragaria vesca</i>	-
FRA0164	<i>Fragaria vesca</i>	-
FRA0172	<i>Fragaria vesca</i>	-
FRA0175	<i>Fragaria vesca</i>	-
FRA0178	<i>Fragaria vesca</i>	-
FRA0182	<i>Fragaria vesca</i>	-
FRA0185	<i>Fragaria vesca</i>	-
FRA0186	<i>Fragaria vesca</i>	-
FRA0195	<i>Fragaria vesca</i>	-
FRA0201	<i>Fragaria vesca</i>	-
FRA0205	<i>Fragaria vesca</i>	-
FRA0207	<i>Fragaria virginiana</i>	-
FRA0208	<i>Fragaria virginiana</i>	-
FRA0209	<i>Fragaria virginiana</i>	-
FRA0218	<i>Fragaria virginiana</i>	-
FRA0220	<i>Fragaria virginiana</i>	-
FRA0222	<i>Fragaria virginiana</i>	-
FRA0227	<i>Fragaria virginiana</i>	-
FRA0230	<i>Fragaria virginiana</i>	-
FRA0231	<i>Fragaria virginiana</i>	-
FRA0233	<i>Fragaria virginiana</i>	-
FRA0234	<i>Fragaria virginiana</i>	-
FRA0237	<i>Fragaria virginiana</i>	-
FRA0240	<i>Fragaria virginiana</i>	-
FRA0244	<i>Fragaria virginiana</i>	-
FRA0246	<i>Fragaria virginiana</i>	-
FRA0249	<i>Fragaria virginiana</i>	-
FRA0254	<i>Fragaria viridis</i>	-

Accession no.	Species	cultivar name
FRA0262	<i>Fragaria viridis</i>	-
FRA0263	<i>Fragaria viridis</i>	-
FRA0272	<i>Fragaria viridis</i>	-
FRA0280	<i>Fragaria viridis</i>	-
FRA0282	<i>Fragaria viridis</i>	-
FRA0283	<i>Fragaria ×ananassa</i>	-
FRA0284	<i>Fragaria ×ananassa</i>	-
FRA0286	<i>Fragaria ×ananassa</i>	-
FRA0287	<i>Fragaria ×ananassa</i>	-
FRA0288	<i>Fragaria ×ananassa</i>	-
FRA0289	<i>Fragaria ×ananassa</i>	-
FRA0290	<i>Fragaria ×ananassa</i>	-
FRA0292	<i>Fragaria ×bifera</i>	-
FRA0295	<i>Fragaria ×bifera</i>	-
FRA0296	<i>Fragaria ×bifera</i>	-
FRA0298	<i>Fragaria ×bringhurstii</i>	-
FRA0299	<i>Fragaria yezoensis</i>	-
FRA0301	<i>Fragaria yezoensis</i>	-
FRA0303	<i>Fragaria yezoensis</i>	-
FRA0305	<i>Fragaria yezoensis</i>	-
FRA0306	<i>Fragaria yezoensis</i>	-
FRA0308	<i>Fragaria yezoensis</i>	-
FRA0311	<i>Fragaria bucharica</i>	-
FRA0312	<i>Fragaria tibetica</i>	-
FRA0313	<i>Fragaria orientalis</i>	-
FRA0314	<i>Fragaria nubicola</i>	-
FRA0315	<i>Fragaria nubicola</i>	-
FRA0316	<i>Fragaria nubicola</i>	-
FRA0317	<i>Fragaria vesca</i>	-
FRA0319	<i>Fragaria yezoensis</i>	-
FRA0320	<i>Fragaria vesca</i>	-
FRA0322	<i>Fragaria nipponica</i>	-
FRA0323	<i>Fragaria nipponica</i>	-
FRA0324	<i>Fragaria nipponica</i>	-
FRA0325	<i>Fragaria nipponica</i>	-
FRA0326	<i>Fragaria nipponica</i>	-
FRA0327	<i>Fragaria chiloensis</i>	-
FRA0333	<i>Fragaria orientalis</i>	-
FRA0334	<i>Fragaria vesca</i>	-
FRA0335	<i>Fragaria chiloensis</i>	-
FRA0337	<i>Fragaria chiloensis</i>	-
FRA0340	<i>Fragaria chiloensis</i>	-
FRA0341	<i>Fragaria chiloensis</i>	-
FRA0344	<i>Fragaria chiloensis</i>	-
FRA0345	<i>Fragaria chiloensis</i>	-
FRA0346	<i>Fragaria chiloensis</i>	-
FRA0349	<i>Fragaria chiloensis</i>	-
FRA0350	<i>Fragaria chiloensis</i>	-
FRA0351	<i>Fragaria chiloensis</i>	-

Accession no.	Species	cultivar name
FRA0353	Fragaria chiloensis	-
FRA0355	Fragaria chiloensis	-
FRA0356	Fragaria chiloensis	-
FRA0372	Fragaria vesca	-

Table S2: Primer sequences to proof strawberry leaf material on the occurrence of 4 strawberry viruses.

Primer	Type	Sequence	Expected fragment size (bp)	Reference
SVBVdetaf	F	AGT AAG ACT GTT GGT AAT GCC A	435	Thompson et al. 2003
SVBVdetb	R	TTT CTC CAT GTA GGC TTT GA		
SCVdeta	F	CAT TGG TGG CAG ACC CAT CA	345	
SCVdetb	R	TTC AGG ACC TAT TTG ATG ACA		
SMYEVdeta	F	GTG TGC TCA ATC CAG CCA G	271	
SMYEVdetb	R	CAT GGC ACT CAT TGG AGC TGG G		
SMoVdeta	F	TAA GCG ACC ACG ACT GTG ACA AAG	219	
SMoVdetb	R	TCT TGG GCT TGG ATC GTC ACC TG		
AtropaNad2.1a	F	GGA CTC CTG ACG TAT ACG AAG GATC	188	
AtropaNad2.2b	R	AGC AAT GAG ATT CCC CAA TAT CAT		

Table S3 Mastermix and PCR conditions for strawberry virus detection.

Reagent (initial concentration)	µl per sample	Final concentration	PCR conditions
dd H ₂ O	13,4		Cycler: room 215 Programm: SVBV 1 x initial denaturation: 94 °C 3' 38 x denaturation/annealing/elongation: 94 °C 1'/55 °C 40''/72 °C 40'' 1 x final elongation: 72 °C 5' 1 x cooling: 10 °C ∞
10 x DreamTaq Puffer (20 mM MgCl ₂)	2,5	1 x	
2 mM dNTP's	2,5	0,2 mM	
SVBVdetaf (10 µM)	1,25	0,5 µM	
SVBVdetb (10 µM)	1,25	0,5µM	
20 x rot Puffer	1	0,8 x	
BSA ^a (0,125 %)	1	0,005 %	
PVP ^b (25 %)	1	1 %	
DreamTaq Polymerase (5 U/µl)	0,1	0,5 U	
DNA-Probe ^c	1		
Total	25		

^a 0,005 % BSA (Bovine serum albumin)/µl was added to the mastermix to prevent PCR-inhibitory substrates (Zhang et al. 2014).

^b 1 % PVP (Polyvinylpyrrolidone)/µl was added to the mastermix to prevent PCR-inhibitory substrates (Koonjul et al. 1999).

^c In general 10 ng/µl DNA per standard-PCR each was used, using cDNA no concentration was determined.