

# Over 10 years' experience in potato virus proficiency tests as external proof of diagnostic laboratory competence



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Landwirtschaftskammer  
**Niedersachsen**

**MV** 

Mecklenburg-Vorpommern

Landesamt für Landwirtschaft,  
Lebensmittelsicherheit und  
Fischerei



Leibniz-Institut • DSMZ-Deutsche Sammlung von Mikroorganismen und Zellkulturen GmbH

# History of the proficiency test for potato viruses

- In accordance with the German Seed Potato Regulation (PflKartV), seed potatoes are usually tested for PVX, PVY, PLRV, PVM, PVA and PVS
- The requirement to participate in a PT stems from the ISO 17025 accreditation
- Initiated in 2010 by three national laboratories: Lower Saxony, Mecklenburg, Bavaria
- In the early years, the sample material consisted of fresh leaves
- Results were not reproducible and PT not accepted by auditors
- Since 2014 DSMZ as co-organizer for sample production and shipment



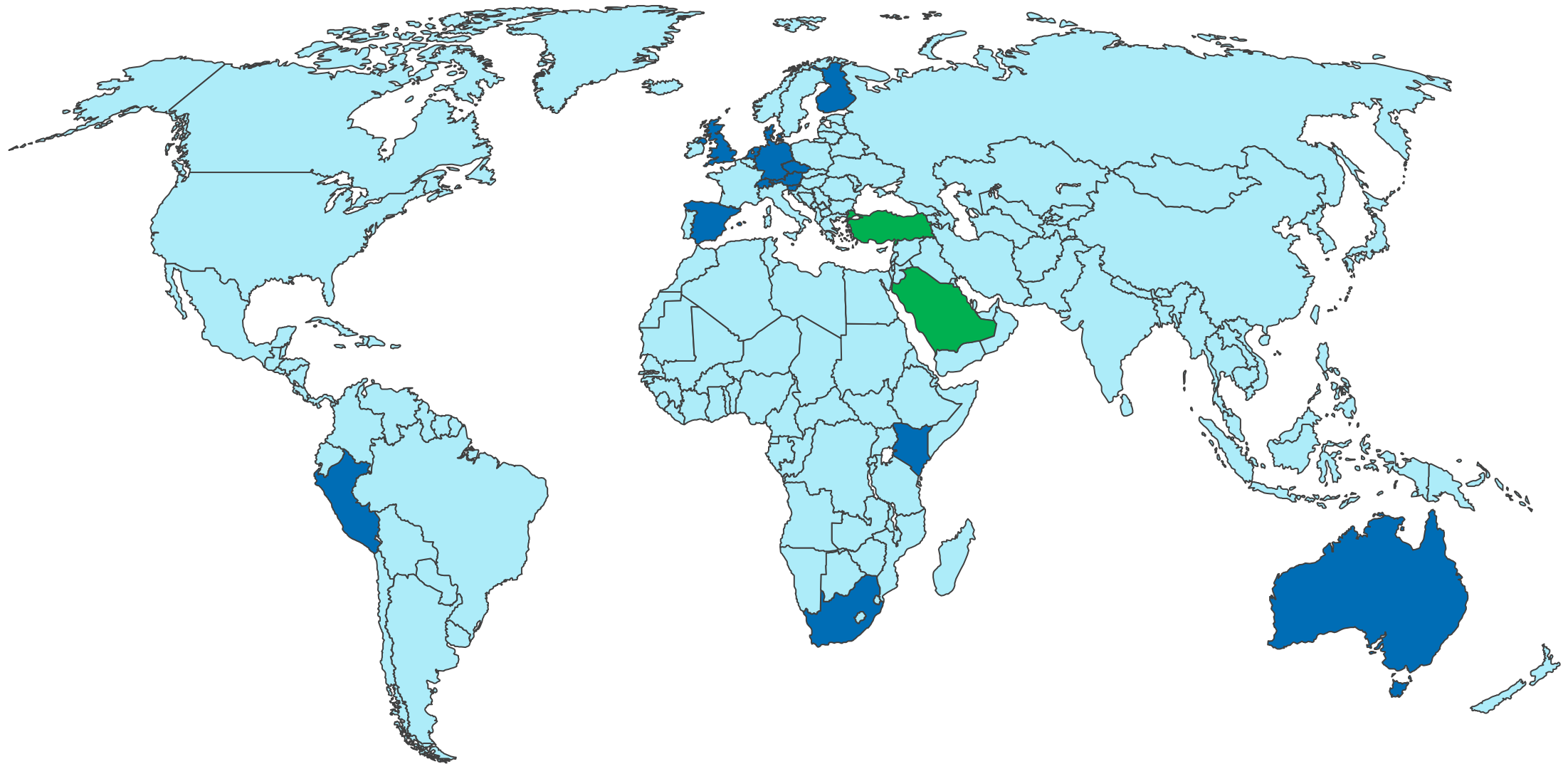
# Samples/virus isolates used in 2025

<u>Sample 2025</u>	<u>Virus</u>	<u>Acc. No. DSMZ (GenBank)</u>	<u>Origin</u>	<u>Host</u>
1	PVA	PV-0535 (MW197135)	Germany, BBA	<i>Nicotiana occidentalis</i> P1
2	PLRV	PV-0842 (MZ202339)	Germany, BBA	<i>Physalis floridana</i>
3	PVS	PV-1327 (OM471986)	Germany, Bioland	<i>Chenopodium quinoa</i>
4	PVM, PVY, PVX	W24-092	Germany, LALLF	potato
5	PVM	PV-0273 (MW582794)	Germany, BBA	<i>Nicotiana occidentalis</i> P1
6	PVX	W24-091	Germany, LALLF	potato
7	healthy	-	Germany, DSMZ	potato 'Amado'
8	healthy	-	Germany, DSMZ	potato 'Amado'

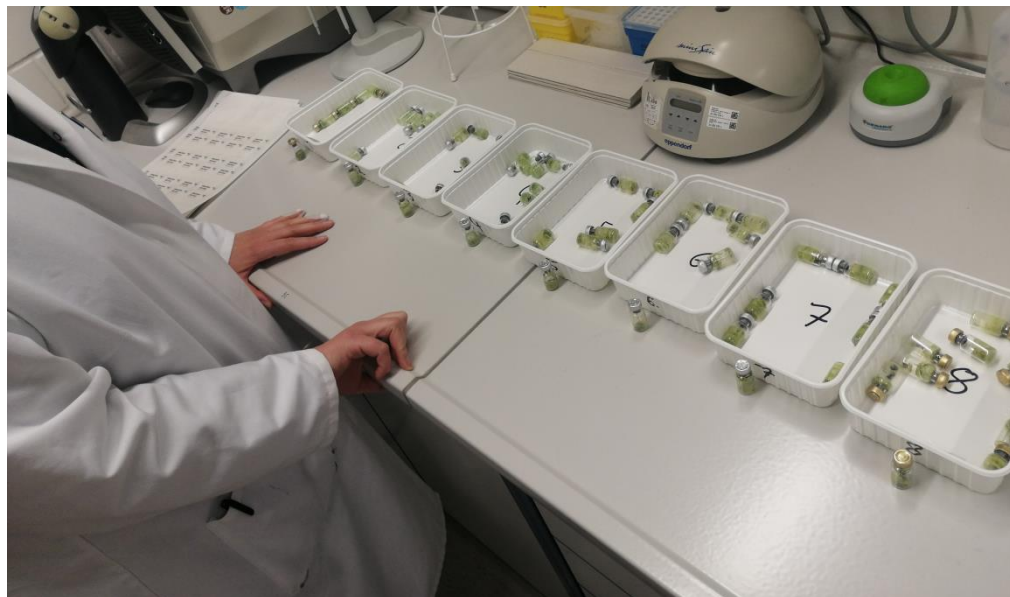
- W-Numbers are well characterized field samples, temporarily available
- All PV-numbers are permanent DSMZ reference isolates
- Sequences available at NCBI GenBank
- Prestesting done by ELISA (DSMZ) and RT-qPCR (LALLF)



# 2025: 39 participants from 18 countries with 59 sets of samples



# Anonymization and participants



Country 2025	Laboratories	Sample sets
Germany	16	24
Luxembourg	1	2
Slovenia	1	2
Denmark	1	1
Austria	1	1
Finland	2	2
Netherlands	1	2
Czech Republic	1	1
Switzerland	2	3
Scotland	2	4
Saudi Arabia	3	3
Turkey	1	1
England	2	4
South Afrika	1	2
Australia	1	1
Peru	1	2
Kenya	1	2
Spain	1	2
<b>Σ Countries = 18</b>	<b>39</b>	<b>59</b>

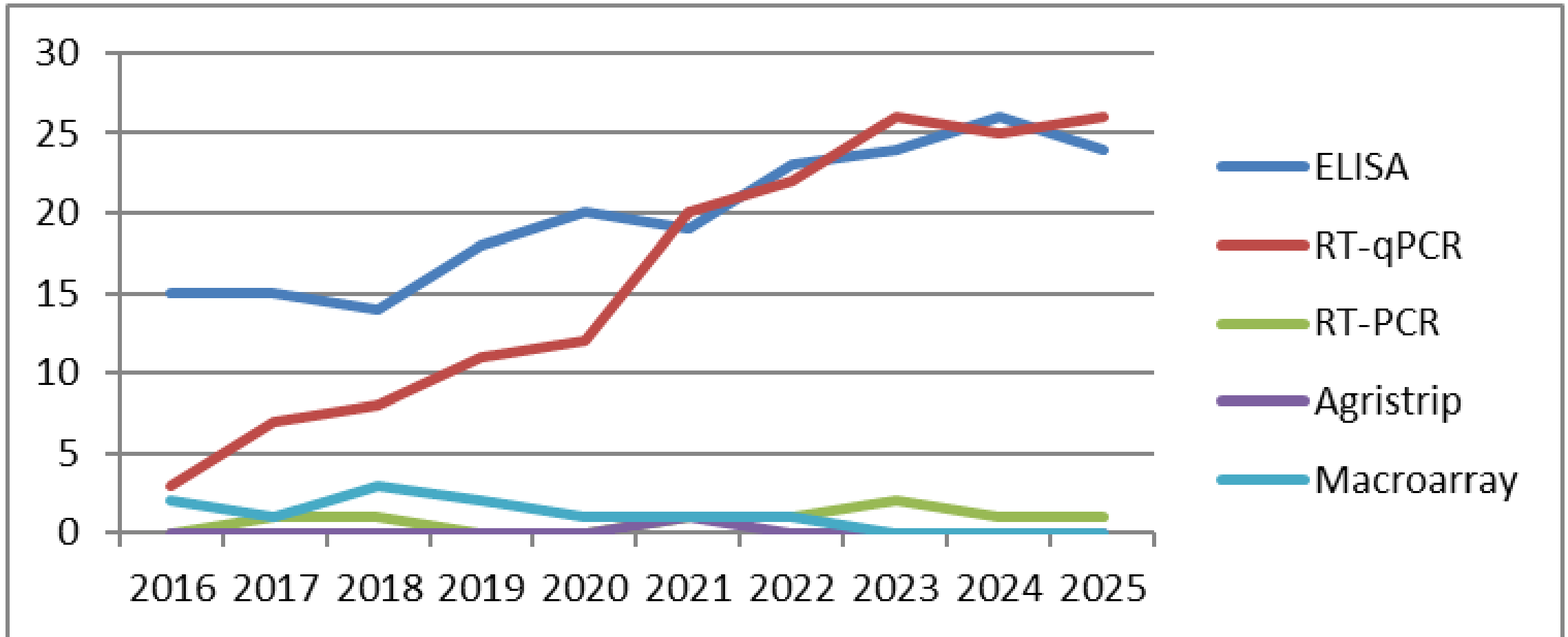


# Methods applied

Method	Sample sets analysed									
	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025
ELISA	15	15	14	18	20	19	23	24	26	24
RT-qPCR	3	7	8	11	12	20	22	26	25	26
RT-PCR	0	1	1	0	0	1	1	2	1	1
Agristrip	0	0	0	0	0	1	0	0	0	0
Macroarray	2	1	3	2	1	1	1	0	0	0
<b>Σ</b>	<b>20</b>	<b>24</b>	<b>26</b>	<b>31</b>	<b>33</b>	<b>42</b>	<b>47</b>	<b>52</b>	<b>52</b>	<b>51</b>

- **2025: results for 8 sets are pending**
- **RT-qPCR: Only 15 out of 26 laboratories test for all viruses (PVS, PVX, and PVM are often omitted)**
- **ELISA: Only one lab does not test for all viruses**

## Methods used



## Origin of kits/methods

<b>Supplier ELISA</b>	<b>Bioreba</b>	<b>Agdia</b>	<b>DSMZ</b>	<b>Prime</b>	<b>SASA</b>	<b>Loewe</b>
<b>Sets analysed</b>	<b>16</b>	<b>1</b>	<b>4</b>	<b>2</b>	<b>1</b>	<b>1</b>

<b>RT-qPCR</b>	<b>Bioreba kit</b>	<b>published /in house</b>
<b>Sets analysed</b>	<b>4</b>	<b>22</b>

# Sample results

Ergebniseingang	23.02.2026	11.02.2026	11.02.2026	17.02.2026	24.02.2026	13.02.2026	21.01.2026	21.01.2026	26.01.2026	26.01.2026	23.02.2026
Methode	ELISA	RT-qPCR	RT-qPCR	RT-qPCR	ELISA	ELISA	ELISA	ELISA	ELISA	ELISA	ELISA
Quelle	Bioreba	several	several	Pastrik et al.	Bioreba/DSMZ	DSMZ	Bioreba	Bioreba	Bioreba	Bioreba	Bioreba/Prime
Probe/Teilnehmer	<b>A</b>	<b>B</b>	<b>C</b>	<b>D</b>	<b>E</b>	<b>F</b>	<b>G</b>	<b>H</b>	<b>I</b>	<b>J</b>	<b>K</b>
1 PVA	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	PVY f-pos
2 PLRV	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺
3 PVS	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺
4 PVM, PVY, PVX	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺
5 PVM	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺
6 PVX	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺
7 Kartoffel gesund	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺
8 Kartoffel gesund	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺

Ergebniseingang	19.02.2026	23.02.2026	23.02.2026	01.01.2026	23.02.2026	23.02.2026	17.02.2026	17.02.2026	23.02.2026	20.02.2026	20.02.2026
Methode	RT-qPCR	RT-qPCR	ELISA	RT-qPCR	ELISA	RT-qPCR	RT-qPCR	ELISA	RT-PCR	ELISA	RT-qPCR
Quelle	several	Schumpp 202	Loewe	several	Bioreba	Bioreba	unpublished	Bioreba	several	Agdia	unknown
Probe/Teilnehmer	<b>AS</b>	<b>AT</b>	<b>AU</b>	<b>AV</b>	<b>AW</b>	<b>AX</b>	<b>AY</b>	<b>AZ</b>	<b>BA</b>	<b>BB</b>	<b>BC</b>
1 PVA	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺
2 PLRV	☺	☺	☺	PVY f-pos	☺	☺	☺	☺	☺	☺	☺
3 PVS	☺	☺	☺	☺	☺	☺	PVY f-pos	☺	☺	☺	☺
4 PVM, PVY, PVX	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	PVY f-neg
5 PVM	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺
6 PVX	☺	☺	☺	☺	☺	☺	☺	☺	PVX f-neg	☺	☺
7 Kartoffel gesund	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺
8 Kartoffel gesund	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺	☺

## In 2025: Only 7 deviations in 51 sample sets analysed (in total 408 samples, corresponding to ca. 2,400 individual tests)

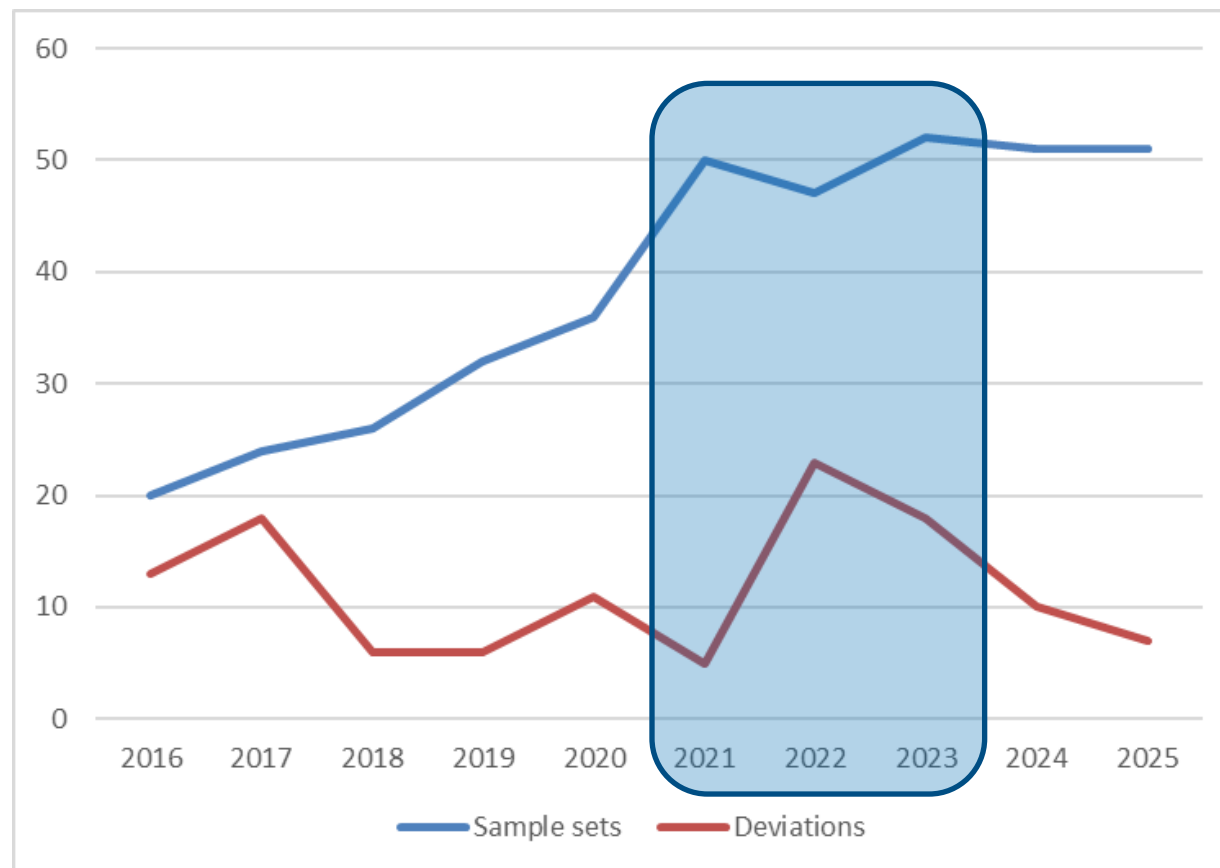
Virus	Deviations by method			
	ELISA	ELISA	RT-qPCR	RT-qPCR
	<b>false negative</b>	<b>false positive</b>	<b>false negative</b>	<b>false positive</b>
$\Sigma$	<b>0</b>	<b>1</b>	<b>2</b>	<b>4</b>
PLRV				<b>1</b>
PVY		<b>1</b>	<b>1</b>	<b>3*</b>
PVM				
PVA				
PVX			<b>1*</b>	
PVS				

\*RT-PCR

\*3 different batches



# Deviations over the years



	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025
Sample sets	20	24	26	32	36	50	47	52	51	51
Deviations	13	18	6	6	11	5	23	18	10	7



# First-time participants

<u>Ergebniseingang</u>	17.02.2023	09.02.2023	26.01.2023	17.01.2023	17.01.2023	17.01.2023	17.01.2023
<u>Methode</u>	RT-qPCR	RT-qPCR	ELISA	ELISA	Macroarray	RT-qPCR	RT-qPCR
<u>Quelle</u>	Bioreba/Kit	unpublished	Sediag	Bioreba	Bioreba	Bioreba/Kit	Bioreba/Kit
<u>Probe/Teilnehmer</u>	A	B	C	D	E	F	G
1 Kartoffel gesund	PVX f-pos	☺	☺	☺	☺	☺	☺
2 PVX	siehe 1)	☺	☺	☺	☺	☺	☺
3 PLRV, PVM, PVY, PVS	PVX f-pos	☺	☺	☺	☺	☺	☺
4 PVS	☺	☺	☺	☺	☺	☺	☺
5 PLRV, PVM	PVA f-pos	☺	☺	☺	☺	☺	☺
6 PVA	☺	☺	☺	☺	☺	☺	☺
7 PLRV, PVM, PVS	PVS f-neg	☺	☺	☺	☺	☺	☺
8 PVX	☺	☺	☺	☺	☺	☺	☺
	1) PLRV f-pos, PVY f-pos, PVS f-pos, PVM f-pos						

**Using a kit does not guarantee accurate results**

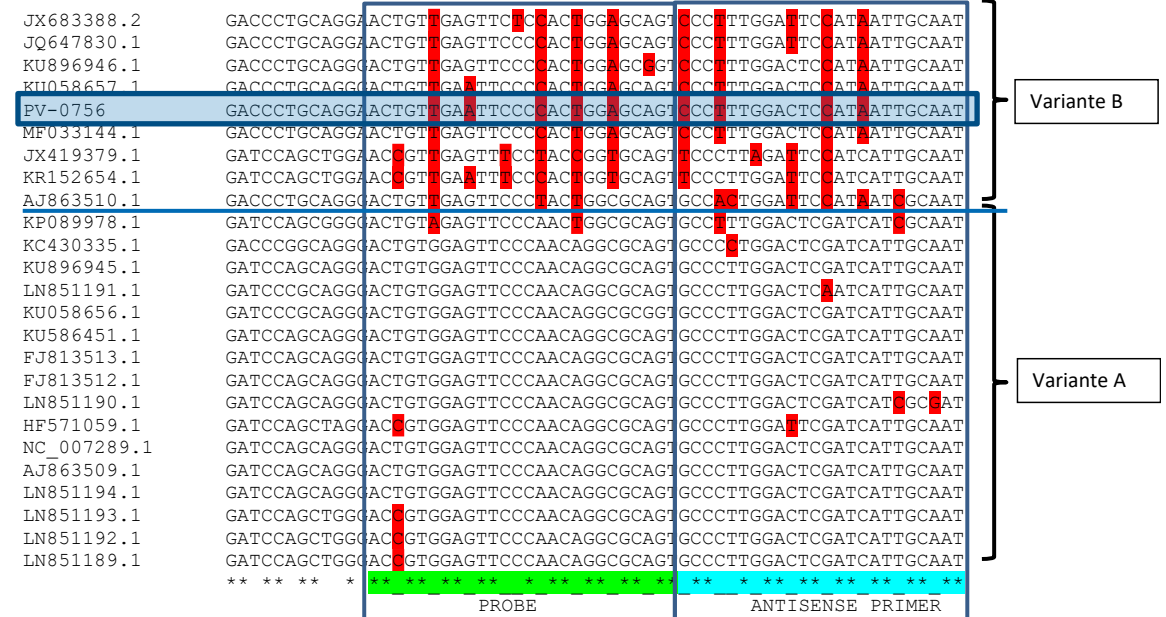
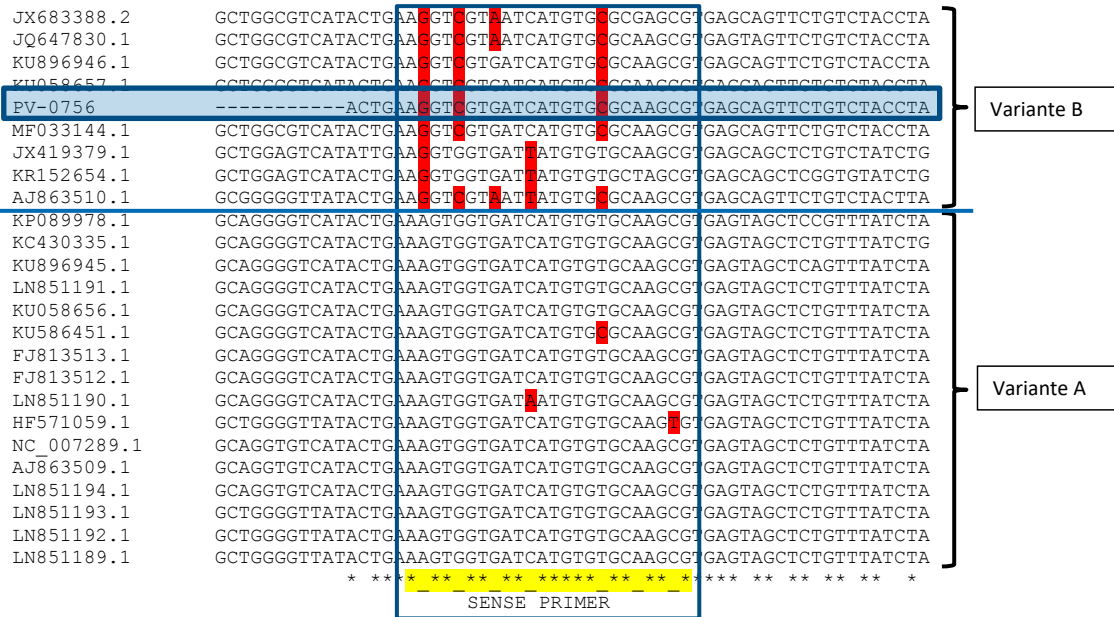
# Differences in ct Values for the Bioreba RT-qPCR Kit

Bioreba RT-qPCR (ct)	PVY	PVM	PLRV	PVX	PVS	PVA
Anwender 1	17	14, 15	18	17, 22	16	16
Anwender 2	18	13, 14	19	14, 17	14	14
Anwender 3	25	25, 26	24	24, 25	21	23
Anwender 4	29	24, 26	21	22, 23	26	23

# PVS variability and mismatches with primers/probe

- In the PT in 2017 most labs using RT-qPCR missed PVS (PV-0756, belonging to the “Andean strain”)
- Those labs used the primer/probe combination published by Mortimer-Jones et al.( 2009)
- These show numerous mismatches to Andean strain isolates including PV-0756

## Variant B: “Andean strain”



## Variant A: “ordinary strain”



# Summary and Conclusions

- **What started as a personal initiative has grown into a successful, self-organized PT**
- **High international demand highlights the global shortage of PTs**
- **The number of labs testing tubers directly using RT-qPCR has increased, but ELISA use remains stable**
- **Deviations stable at a low level**
- **“Newcomers” from the early 2020s have gotten their problems under control**
- **Successful application and sensitivity in RT-qPCR varies greatly depending on the user/laboratory**
- **PTs help identify weaknesses in published methods (PVS RT-qPCR, Mortimer-Jones et al. 2009)**



# Thank you for your attention!



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# Prestesting by ELISA (DSMZ) and RT-qPCR (LALLF)

A/B: 2 vials/batch tested

<u>Sample 2025</u>	<u>Virus</u>	<u>ELISA (OD 2h A/B)</u>	<u>RT-qPCR (ct A/B)</u>
1	PVA	1,5/1,6	11,8/11,7
2	PLRV	3,8/3,8	14,0/13,5
3	PVS	3,7/3,8	10,3/10,3
4	PVM, PVY, PVX	PVM: 1,5/1,5 PVY: 0,4/0,5 PVX: 2,3/2,5	PVM: 12,7/12,4 PVY: 13,4/12,4 PVX: 6,5/6,6
5	PVM	1,8/1,5	12,0/11,8
6	PVX	2,9/3,3	14,2/13,6
7	gesund	-	-
8	gesund	-	-

## Detailed information on participants' false-positive results

- **ELISA PVY false positive: OD 0,29**      **PVY true positive: OD 1,46**
- **RT-qPCR PLRV false positive: ct 28**      **PLRV true positive: ct 15**
- **RT-qPCR PVY false positive: ct 27**      **PVY true positive: ct 14**
- **RT-qPCR PVY false positive: ct 18**      **PVY true positive: ct 16**
- **RT-qPCR PVY false positive: ct 35**      **PVY true positive: ct 22**

<u>Sample 2025</u>	<u>Virus</u>	<u>RT-qPCR pretesting (ct A/B)</u>
2	PLRV	14,0/13,5
4	PVY	13,4/12,4