

*Full Length Research Paper*

# Phenotypic and molecular screening of some tomato germplasm for resistance to tomato yellow leaf curl virus disease in Ghana

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Tomato yellow leaf curl virus (TYLCV) is a major tomato virus in Ghana and Africa as a whole. In this study, 30 accessions of *Solanum lycopersicum* L. with reported TYLCV-resistance from AVRDC were assessed for resistance to TYLCV in Ghana. Plants were grown in a field, a hot spot of the disease and the reactions of plants were evaluated based on the disease symptoms when they were 30, 45 and 60 days after transplanting. Molecular screening was also done to re-confirm the phenotypic evaluation. All the tomato accessions demonstrated various degrees of disease symptoms. Phenotypic evaluation was confirmed by amplification of TYLCV DNA fragment in all tested accessions. Based on the phenotypic and molecular evaluations, no accession provided complete resistance to TYLCV in Ghana. However, accessions with milder symptoms of TYLCV in the field were considered as tolerant. The high level of susceptibility to viral infection noted in the field was not observed in the molecular screening. The viral DNA was detected using six different primers and the primers indicated polymorphism. TYLCV was detected in 23 accessions using primer pair GhF and GhR. The results suggested that accessions that indicated symptoms of the disease on the field but had no TYLCV DNA amplification could be due to other viruses or virus strain. Accessions with reported resistance in other countries but collapsed in Ghana could be attributed to genotype – environment interactions and or the emergence of new mutants of the TYLCV in Ghana.

**Key words:** Molecular screening, tomato, tomato yellow leaf curl virus, virus resistance, DNA, primer.

## INTRODUCTION

In Ghana, tomato is a very popular and important vegetable crop which is consumed on nearly a daily basis by every household (Horna et al., 2006). It is used in preparing soups and stews. The highest quality fruits and greatest yields are obtained in the dry season with supplementary water (Tweneboah, 1998). Currently, more money is spent on tomato cultivation than on any other vegetable (Wolff, 1999).

Despite tomato's importance in Ghana, local production is not able to meet the domestic demand and tomatoes are often imported, mainly from Burkina Faso. This is a

drain on the country's economy. This situation is attributed to a number of constraints. Two such constraints are the pests and diseases that affect tomato production in Ghana. Among them, tomato yellow leaf curl virus (TYLCV) is of economic importance (Bhyan et al., 2007; Valizadeh et al., 2011).

TYLCV, a *Begomovirus* of the family Geminiviridae, is the most devastating virus of the tomato plant in tropical and subtropical regions including Ghana. The family Geminiviridae comprises plant viruses that have a circular, single-stranded DNA genome and geminate particles consisting of two incomplete icosahedra (Hull, 2002). Geminiviruses are classified into four genera based on the type of insect vector, host range, and genome organization (Rybicki et al., 2000; El-Din et al., 2005). The genus *Begomovirus* includes species with

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**Table 1.** Tomato accessions used for screening against tomato yellow leaf curl virus.

Entries	Code	Resistance source	Origin
FLA 505	A1	LA 1969 ( <i>L. chilense</i> )	J. Scott, Univ. Florida
FLA 456-4	A2	Tyking, LA2779 ( <i>L. chilense</i> )	J. Scott, Univ. Florida
FLA 478-6-3-0	A3	LA1938, Tyking, Fiona	J. Scott, Univ. Florida
FLA 653-3-1-0	A4	LA2779 ( <i>L. chilense</i> ), Tyking	J. Scott, Univ. Florida
FLA 496-11-6-1-0	A5	LA1932 ( <i>L. chilense</i> ), Tyking	J. Scott, Univ. Florida
TLB 111	A6	H24	AVRDC
TY52	A7	LA 1969 ( <i>L. chilense</i> )	D. Zamir, Hebrew Univ.
99S-C-39-20-11-24-17-0	A8	Unknown	Namdhari Seeds, India
H24	A9	<i>L. hirsutum</i> f.sp glabratum	G. Kallo, India
CLN2026D	A10	Susceptible check	AVRDC
Pimpinellifolium	G11	Unknown	CSIR-CRI
WSP2F1pt.3	G12	Unknown	CSIR-CRI
WS273.3 Large	G13	Unknown	CSIR-CRI
WSP2F7 (3) pt.3	G14	Unknown	CSIR-CRI
2641A	B16	Unknown	AVRDC
Tomato Money Maker	B17	Unknown	USA
Tomato Roma-Jam Vf	B18	Unknown	Burkina Faso
Parona	B19	Unknown	Local
Local Roma	B20	Unknown	Local
Rando	B21	Unknown	Local
Tomato Slumac	B22	Unknown	Holland
Tomato Tima	B23	Unknown	France
Tomato Red Cloud	B24	Unknown	Holland
Tomato Rio Grande	B25	Unknown	Holland
Petomech (Ghana/France)	B26	Unknown	France
Tomato Roma VF	B27	Unknown	USA
Petomech (Ghana/Burkina)	B28	Unknown	Burkina Faso
Petomech (Ghana)	B29	Unknown	Ghana
Tomato Ventura F	B30	Unknown	USA

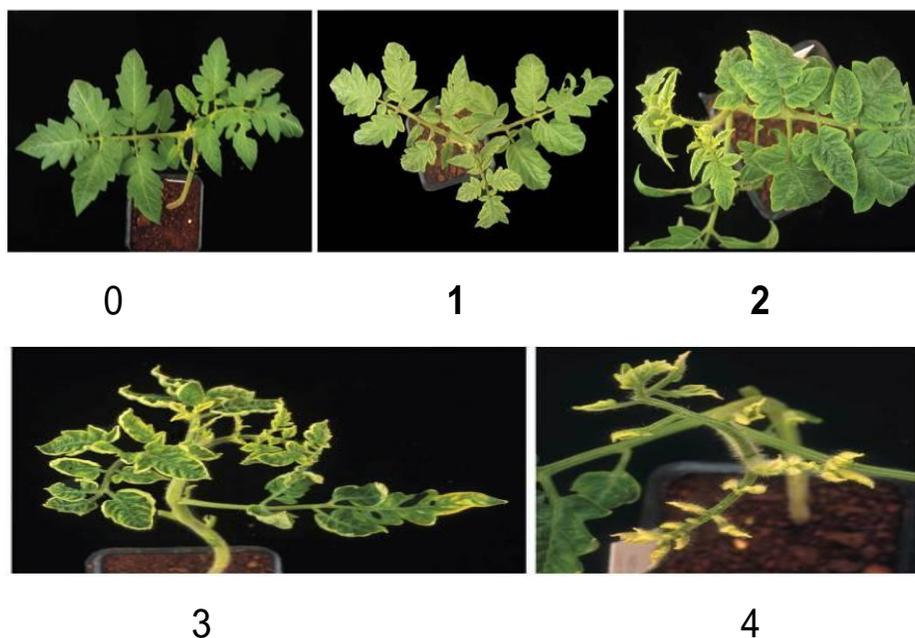
monopartite or bipartite genomes such as TYLCV that are transmitted by the whitefly *Bemisia tabaci* Gennadius (Hemiptera: Aleyrodidae) (Moriones and Navas Castillo, 2000; Cohen and Nitzany, 1966; Mazyad et al., 2007). *B. tabaci* has developed resistance against insecticides in recent years (Dittrich and Ernst, 1990) and therefore, a few viruliferous whiteflies may be enough for transmitting the virus to a large number of plants (Mazyad et al., 2007). Chemical control methods as well as integrated pest management (IPM) strategies employed for controlling the vector have not been successful in decreasing the incidence of TYLCV on tomato crop (Bhyan et al., 2007; Reynaud et al., 2003). Under these circumstances, breeding for resistance to TYLCV appears to be a promising and environmentally friendly approach for controlling the disease (Chague et al., 1997). Host plant resistance will therefore be an important component of an overall whitefly-transmitted geminivirus control strategy. Several techniques such as enzyme linked immunosorbent assay (ELISA), tissue-blotting immuno-binding assay (TBIA) and dot blotting

immuno-binding assay (DBIA) as well as polymerase chain reaction (PCR) provide a sensitive and specific means for the detection and identification of whitefly transmitted geminiviruses (WTGV) in the infected plants and their vector whitefly *B. tabaci* (Cohen et al., 1989; Mehta et al., 1994; Fargette et al., 1996; Tsai et al., 2006; El-Din et al., 2005; Bhyan et al., 2007). The objective of this work was to screen available tomato germplasm through convectional and molecular approaches to identify stable sources for breeding against the TYLCV disease.

## MATERIALS AND METHODS

### Screening of germplasm for TYLCV resistance

Thirty tomato germplasm collected from Burkina Faso, USA, Holland, France and Green seeds (Taiwan), Ghana and Asian Vegetable Research Development Centre, Taiwan (Table 1) were screened at Afari (a hot spot of the disease) for resistance to TYLCV disease.



**Figure 1.** TYLCV Symptom Severity Scale. Severity scores were based on 0 - 4 scale developed by AVRDC where; 0- No symptoms, 1- Slight yellowing (mild symptom); 2- Leaf curling and yellowing (moderate symptom); 3- Yellowing, Curling and Cupping (severe symptom); 4- Severe stunting, curling and cupping; plant stops growth (very severe symptom). Source: Lapidot and Friedman (2002).

### Experimental design

The tomato germplasm were planted on a plot size of 1863 m<sup>2</sup> (81 m × 23 m). A spacing of 90 cm × 70 cm in a randomized complete block design (RCBD) with three replications was used. They were planted in triple rows and each accession had 10 plants per row giving a total of 30 plants per accession. Standard agronomic practices such as weed control, fertilizer application (15-15-15) at 250 kg/ha and spraying of fungicides (Shavit F at a rate of 50 g/15 L) were used.

### Data collection

Scoring for incidence and severity of TYLCV disease on the plants was done at 30, 45 and 60 days after transplanting using a symptom severity scale (Figure 1). Other characteristics on the fruits were taken. Data were taken on the number of fruits per plant and plot. A refractometer was used in taking the brix after harvesting.

### Viral DNA detection: Plant DNA extraction

For molecular analysis, leaf samples were collected from 30 tomato accessions to confirm the phenotypic screening. DNA extraction was carried out using the DNA isolation method described by Egnin et al. (1998) with some modifications.

### Primer for PCR Amplification

PCR amplification was done using six primer pairs. These included PAR1c496/PAL1v1978, PTYc1121/PTYv787, AC1048/AV494, GHF/GHR, KR/KF and MR/MF. The first three were degenerated

primers; the last three were developed during this work (Table 2). The primer 3 program was used to develop the primers (Rozen and Skaletsky, 2000). The sequences in the last three are available in the NCBI. The primer pairs GHF/GHR, KR/KF and MR/MF have GenBank accession numbers EU350585, EU847739 and EU847740, respectively.

### PCR amplification of viral DNA

PCR was performed in 10 µl volume using six primer pairs. The reaction mixture composition was 1.5 mM of 5x Buffer A with MgCl<sub>2</sub>, 10 mM of dNTPs, 0.12 µM each of forward and reverse primer and 5 µl robust taq. Water was added to make a final volume. The same reaction mixtures were used. PCR amplification was carried out in a BIO-RAD Mycycler™ Thermal cycler. For primer pairs, PAL1v1978/PAR1c496, GHF/GHR, KF/KR and MF/MR, DNA amplification parameters were 30 cycles of denaturation for 1 min at 94°C, 50°C for 1 min, 72°C for 3 min and final extension at 72°C for 3 min. Amplification products were maintained at 4°C prior to electrophoresis. The DNA amplification parameters for primer pair AC1048 and AV496 also had an initial denaturation of 92°C for 1 min followed by 35 cycles at 92°C for 1 min, annealing at 60°C for 20 s and extension at 72°C for 30 s. The primer pair PTYv/PTYc profile amplified at 94°C for 1 min, 58°C for 1 min, 72°C for 2 min and a final extension at 72°C for 10 min.

### Gel electrophoresis

The PCR products were electrophoresed in a 1% agarose gel (7.5 µl ethidium bromide, 200 ml 1X TAE, 2.0 g agarose). 2 µl of loading dye was mixed with 10 µl of the sample and the mixture was loaded in a gel of 1 × TAE buffer. 4 kb of DNA ladder was used as a marker and water was used as a negative control. The samples

**Table 2.** Primer pairs used for PCR detection of TYLCV.

Primer sequence	Source	Reference
PARc1496/PAL1v1978 5'GCA/TCT/GCA/GGC/CCA/CAT/YGT/CTT/YCC/NGT (30 mer) 5'AAT/ACT/GCA/GGG/CTT/ CT/RTA/CAT/RGG (27 mer)	Metabion International Company	Rojas et al. (1993)
AV494/AC1048 GCCCATGTATAGAAAGCCAAG GGATTAGAGGCATGTGTACATG	Metabion International Company	Wyatt and Brown (1996)
PTYv787/PTYc1121 5-GTTTCGATAATGAGCCCAG-3 5-ATGTAACAGAAACTCATG-3	Metabion International Company	Zhou et al. (2008)
GHF/GHR F: GCCCGAAAGCTTCGTTGTT TTCCCGCT R: ACGGATGGCCGCTTTGGGT ATTCG	Metabion International Company	Osei et al. (2008)
KF/KR F: GGACCCGGCGCACTATTTAT GTTGGC R: ACCCCATTACCCCAATACCA	Metabion International Company	Osei et al. (2008)
MF/MR F: TGGCCGCGCCCTTCCTTTTGT R: ACCAATGGCTCCCCAAAGCGT	Metabion International Company	Osei et al. (2008)

were run in TAE buffer for 40 min at 120 V in a 150 ml BIO RAD electrophoretic apparatus. The gels were then observed by alpha imager software on a computer. Banding patterns were then observed and compared between individuals showing viral presence and absence.

## RESULTS

### Screening for TYLCV- resistance under field conditions

#### *Incidence (number of plants infected)*

The results of the screening for TYLCV-resistance on the field are given in Table 3. Of the 30 accessions studied, there were highly significant differences in terms of percentage of infected plants at 30, 45 and 60 days after transplanting (DAT). At 30 days after transplanting, accessions A3, A5, G14, B16, B1, B20, B22, B24 and B26 had fewer numbers of infected plants. B23 however, recorded the highest percentage of plants that were infected at 30 DAT. More than half (>50%) of tomato accession were affected at 45 days after transplanting except accessions A3, A5, G14, B18, B20 and B22. 100% infection was recorded in accessions A1, A7, A10, B21, B23 and B29 at 60 days after transplanting.

#### *Symptom severity of the disease*

At 30 days after transplanting, B20, B22 and B26 were

the only accessions with no symptoms. There were however, some accessions (A1, A6, A9, A10, G12, G14, G15 and B16) which showed mild symptoms. This changed to moderate and severe symptoms at 45 and 60 days after transplanting, respectively. Apart from tomato accessions B20, B26 which showed mild symptoms and G12, B24 which showed moderate symptoms of the disease respectively, the rest exhibited severe symptoms of the disease after 60 days of transplanting (Table 3).

#### **Fruit characteristics of tomato accessions studied**

Accession B23 gave the highest average fruit weight (61.85 g). This was followed by B21 (46.54 g) and A7 (43.42 g). Accession G12 however recorded the lowest average fruit weight per fruit (8.9 g). In terms of the average number of fruits per plant, tomato accession G12 gave the highest number of fruits (37). This was followed by G14 (35) and G11 (33). Accession A1 however gave the lowest number of fruits per plant. The fruit shapes of the accessions studied were round, high round, plumshaped, heartshaped, flattened, slightly flattened and lengthened cylindrical. 23% of the tomato accessions had round and high round fruit shape with 20% plum shaped and heart shaped. 33% of the tomato accessions had flattened and lengthened cylindrical fruit shape and 6.7% have slightly flattened fruit. 63.3% of the tomato accessions had red fruit colour at maturity and the rest representing 36.7% had orange fruit colour at maturity. The highest brix (soluble solids) among the accessions was four and the least was two (Table 4).

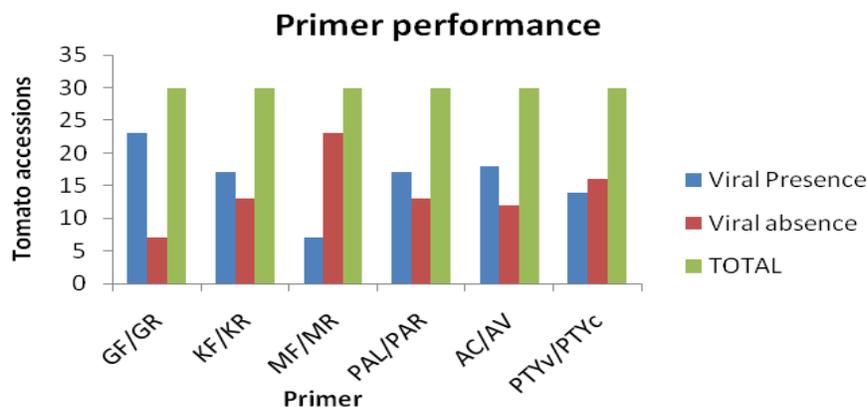
**Table 3.** Reaction of different tomato accessions to TYLCV on field assays.

Field code	Accession	% of TYLCV infected plants			Level of severity		
		30 DAT*	45 DAT	60 DAT	30 DAT	45 DAT	60 DAT
A1	FLA 505	60.00±00 <sup>BCD</sup>	90.00±00 <sup>AB</sup>	100.00±00 <sup>A</sup>	1.00±00	3.00±00	3.00±00
A2	FLA 456-4	50.00±00 <sup>CDE</sup>	76.67±00 <sup>BCD</sup>	76.67±00 <sup>FGH</sup>	2.00±00	3.00±00	3.00±00
A3	FLA478-6-3-0	20.00±5.7 <sup>FG</sup>	50.00±5.7 <sup>EF</sup>	56.67±3.3 <sup>IJ</sup>	2.00±00	3.00±00	3.00±00
A4	FLA 653-3-1-0	60.00±00 <sup>BCD</sup>	70.00±00 <sup>CD</sup>	70.00±00 <sup>EFG</sup>	3.00±00	3.00±00	3.00±00
A5	FLA496-11-6-1-0	20.00±00 <sup>FG</sup>	40.00±00 <sup>F</sup>	73.33±3.3 <sup>GH</sup>	3.00±00	3.00±00	3.00±00
A6	TLB111	60.00±11.5 <sup>BCD</sup>	80.00±11.5 <sup>BC</sup>	60.00±00 <sup>I</sup>	1.00±00	3.00±00	3.00±00
A7	TY52	60.00±5.7 <sup>BCD</sup>	80.00±5.7 <sup>BC</sup>	100.00±00 <sup>A</sup>	2.00±00	3.00±00	3.00±00
A8	99S-C-39-20-11-24	50.00±00 <sup>CDE</sup>	60.00±5.7 <sup>DE</sup>	86.67±3.3 <sup>CDE</sup>	2.00±00	3.00±00	3.00±00
A9	H24	50.00±00 <sup>CDE</sup>	70.00±5.7 <sup>CD</sup>	90.00±00 <sup>I</sup>	1.00±00	3.00±00	3.00±00
A10	CLN 2026D	60.00±00 <sup>BCD</sup>	83.33±3.3 <sup>A</sup>	100.00±00 <sup>A</sup>	1.00±00	3.00±00	3.00±00
G11	Pimpinellifolium	50.00±00 <sup>CDE</sup>	70.00±00 <sup>CD</sup>	93.33±3.3 <sup>ABC</sup>	2.00±00	3.00±00	3.00±00
G12	WSP2F1pt.3	50.00±00 <sup>FG</sup>	60.00±00 <sup>F</sup>	70.00±00 <sup>EFG</sup>	1.00±00	2.00±00	2.00±00
G13	WS273.3 Large	40.00±5.7 <sup>DEF</sup>	70.00±5.7 <sup>CD</sup>	80.00±00 <sup>EFG</sup>	2.00±00	3.00±00	3.00±00
G14	WSP2F7(3)pt.3	20±00.5.7 <sup>FG</sup>	40.00±00 <sup>F</sup>	80.00±00 <sup>EFG</sup>	1.00±00	2.00±00	2.00±00
G15	WSP27F7(3)pt.3	60.00±00 <sup>BCD</sup>	70.00±00 <sup>BC</sup>	83.33±6.7 <sup>DEF</sup>	1.00±00	2.00±00	3.00±00
B16	2641A	30.00±5.7 <sup>EF</sup>	90.00±00 <sup>AB</sup>	90.00±00 <sup>BCD</sup>	1.00±00	3.00±00	3.00±00
B17	Tomato Money M.	40.00±5.7 <sup>DEF</sup>	70.00±00 <sup>CD</sup>	76.67±3.3 <sup>FGH</sup>	3.00±00	3.00±00	3.00±00
B18	Tomato Roma-Jam	30.00±00 <sup>EF</sup>	50.00±00 <sup>EF</sup>	60.00±00 <sup>I</sup>	1.00±00	2.00±00	3.00±00
B19	Parona	80.00±00 <sup>AB</sup>	80.00±00 <sup>BC</sup>	96.67±00 <sup>AB</sup>	2.00±00	3.00±00	3.00±00
B20	Local Roma	20.00±00 <sup>FG</sup>	50.00±00 <sup>AB</sup>	50.00±00 <sup>J</sup>	0.00±00	1.00±00	1.00±00
B21	Rando	80.00±00 <sup>AB</sup>	80.00±00 <sup>BC</sup>	100.00±00 <sup>A</sup>	2.00±00	3.00±00	3.00±00
B22	Tomato Slumac	33.3±3.3 <sup>G</sup>	50.00±00 <sup>EF</sup>	50.00±00 <sup>J</sup>	0.00±00	3.00±00	3.00±00
B23	Tomato Tima	90.00±00 <sup>A</sup>	100.00±00 <sup>A</sup>	100.00±00 <sup>A</sup>	3.00 ±00	3.00±00	3.00±00
B24	Tomato Red Cloud	30.00±00 <sup>EF</sup>	70.00±00 <sup>CD</sup>	80.00±00 <sup>EFG</sup>	1.00±00	2.00±00	2.00±00
B25	Tomato Rio Grand	60.00±00 <sup>BCD</sup>	70.00±00 <sup>CD</sup>	80.00±00 <sup>EFG</sup>	1.00±00	2.00±00	3.00±00
B26	Petomech-gh/Fr.	40.00±00 <sup>DEF</sup>	70.00±00 <sup>CD</sup>	80.00±00 <sup>EFG</sup>	0.00±00	1.00±00	1.00±00
B27	Tomato Roma	80.00±00 <sup>AB</sup>	90.00±00 <sup>AB</sup>	100.00±00 <sup>A</sup>	2.00±00	3.00±00	3.00±00
B28	Petomech-Bk.	70.00±00 <sup>AB</sup>	70.00±00 <sup>CD</sup>	70.00±00 <sup>H</sup>	2.00±00	3.00±00	3.00±00
B29	Petomech-gh	80.00±00 <sup>AB</sup>	100.00±00 <sup>A</sup>	100.00±00 <sup>A</sup>	2.00±00	3.00±00	3.00±00
B30	Tomato Ventura F	60.00±00 <sup>BCD</sup>	70.00±00 <sup>CD</sup>	80.00±00 <sup>EFG</sup>	1.00±00	3.00±00	3.00±00
df	29.86	29.86	29.86	29.86	29.86	29.86	29.86
P value	<0.001	<0.001	<0.001	<0.001	<0.001	<0.001	<0.001

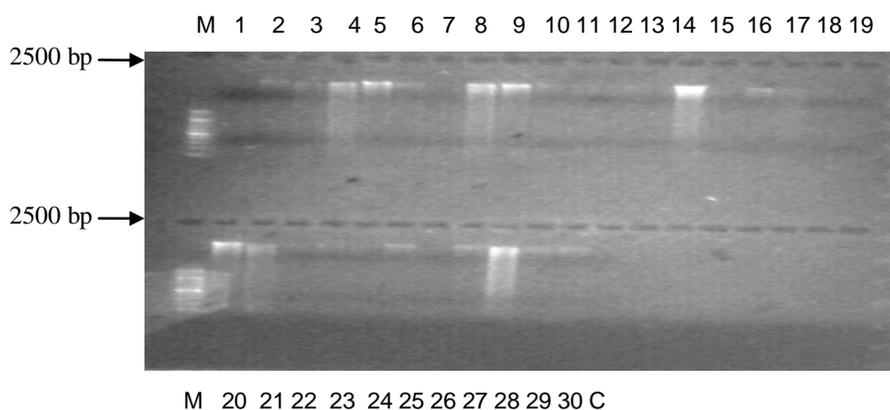
0 = No symptom (NS); 1 = mild symptom (MS); 2 = moderate symptom (MoS); 3 = severe symptom (SS); 4 = very severe symptom (VSS) \*DAT-Days after transplanting.

**Table 4.** Fruit and yield characteristics of tomato accessions.

Field code	Accession	Mean fruit weight (g)	Average yield/plant No. of fruits per plant	Brix	Fruit shape	Fruit colour
A1	FLA 505	19.44±0.13	3.00±0.58	3.00 ± 00	Round	Red
A2	FLA 456-4	26.98±0.59	16.00±1.53	4.00 ± 00	Round	Red
A3	FLA478-6-3-0	20.69±0.02	9.00±00	4.00 ± 00	High round	Red
A4	FLA 653-3-1-0	33.23±2.19	10.00±1.15	3.00 ± 00	Round	Red
A5	FLA496-11-6-1-0	28.18±0.12	20.00±1.73	2.00 ± 00	High round	Orange
A6	TLB111	22.47±0.20	15.00±0.58	3.00 ± 00	Slightly flattened	Red
A7	TY52	43.42±0.77	6.00±00	3.00 ± 00	Plum shaped	Orange
A8	99S-C-39-20-11-24-17-0	29.43±0.01	10.00±2.31	2.00 ± 00	Plum shaped	Orange
A9	H24	12.62±0.03	5.67±0.33	3.00 ± 00	Heart shaped	Red
A10	CLN 2026D	9.80±0.06	8.00±0.67	3.00 ± 00	High round	Orange
G11	Pimpinellifolium	15.10±00	33.00±2.59	4.00 ± 00	Round	Red
G12	WSP2F1pt.3	8.9±0.06	37.00±1.15	4.00 ± 00	Round	Red
G13	WS273.3 Large	33.1±2.08	20.00±0.58	4.00 ± 00	High round	Red
G14	WSP2F7(3)pt.3	25.9±0.58	35.00±1.15	4.00 ± 00	Round	Red
G15	WSP27F7(3)pt.3	19.00±1.20	14.00±00	4.00 ± 00	Round	Red
B16	2641A	19.24±00	7.00±1.15	4.00 ± 00	Plum shaped	Orange
B17	Tomato Money M	36.47±0.20	22.00±1.15	4.00 ± 00	Heart shaped	Red
B18	Tomato Roma-Jam Vf	10.00±1.15	11.00±0.58	2.00 ± 00	Plum shaped	Red
B19	Parona	26.92±00	8.00±00	3.00 ± 00	Slightly flattened	Red
B20	Local Roma	11.07±0.60	17.00±0.58	3.00 ± 00	High round	Red
B21	Rando	46.54±00	4.00±00	3.00 ± 00	Flattened	Orange
B22	Tomato Slumac	33.84±00	9.00±1.15	3.00 ± 00	Lengthened cylindrical	Red
B23	Tomato Tima	61.85±0.49	18.00±00	2.00 ± 00	Heart shaped	Orange
B24	Tomato red cloud	28.97±00	7.00±1.15	3.00 ± 00	Heart shaped	Red
B25	Tomato Rio grand	12.54±0.06	4.00±0.58	3.00 ± 00	Heart shaped	Red
B26	Petomech (gh/France)	24.68±0.33	13.00±00	3.00 ± 00	High round	Orange
B27	Tomato Roma	17.73±00	4.00±00	3.00 ± 00	Plum shaped	Red
B28	Petomech (gh/Burkina)	24.83±2.5	14.00±0.58	4.00 ± 00	Heart shaped	Orange
B29	Petomech (gh/France)	24.68±0.33	13.00±00	3.00 ± 00	High round	Orange
B30	Tomato Ventura F	16.48±0.56	4.00±0.58	3.00 ± 00	Plum shaped	Orange
df	29.86	29.86	29.86	29.86		
P value	<0.001	<0.001	<0.001	<0.001		



**Figure 2.** Primer performances in viral DNA amplification.



**Figure 3.** TYLCV amplified product on agarose gel using primer AC1048 and AV494.

Molecular analysis of tomato accessions for TYLCV resistance

### **Primer efficiency for PCR amplification**

The primers gave a high degree of polymorphism among the 30 tomato accessions. Figure 2 shows the efficiency of each primer in the amplification of the viral DNA between the accessions. The most consistent amplification of a viral DNA fragment was obtained with the primer pair GF and GR (Figure 3). It detected high number of viral DNA (23/30) than all the other primers used. However, primer pair MF and MR amplified low number of viral DNA (7/30).

### **PCR detection and amplification of viral DNA**

None of the six primer pairs used detected viral DNA in accession B24 (Table 5). All the primers amplified viral DNA in accessions A2, A5, A9 and G14. Five out of the six (5/6) primer pairs detected viral DNA in accessions

B16, B20 and B27. Four out of the six (4/6) primer pairs detected viral DNA amplification for accessions A7, A8, B17, B21 and B28. DNA fragment of expected sizes 75, 2200 and 2500 bp were amplified using TYLCV specific primer pairs GHF/GHR, KF/KR and MF/MR, respectively. The other degenerate primers PAL1v1978/PAR1c496, AC1048/AV494 and PTYv787/PTYc1121 also gave the expected amplified viral DNA of 2500, 2500 and 2700 bp, respectively (Figures 3 and 4).

## **DISCUSSION**

### **Screening for TYLCV- resistant plant under field conditions**

Significant differences that emerged from tomato accessions on the incidence of the disease could possibly be attributed to the fact that the whiteflies had affinity for some particular accessions than others and resulted in some accessions being more susceptible to the virus than others. This probably made them feed and

**Table 5.** Scores of viral detection using six primer pairs.

Tomato lines	Primers					
	GhF/GhR	KR/KF	MF/MR	PAL/PAR	AC/AV	PTYv/PTYc
A1	-	+	+	-	-	+
A2	+	+	+	+	+	+
A3	+	-	-	-	+	-
A4	-	+	-	-	+	+
A5	+	+	+	+	+	+
A6	+	-	-	-	+	-
A7	+	+	+	+	-	-
A8	+	+	+	-	+	-
A9	+	+	+	+	+	+
A10	+	+	-	+	-	-
G11	+	-	-	+	-	-
G12	-	-	-	+	-	-
G13	-	-	-	+	-	-
G14	+	+	+	+	+	+
G15	+	-	-	+	-	-
B16	+	+	-	+	+	+
B17	+	+	-	-	+	+
B18	+	-	-	+	-	+
B19	+	-	-	-	-	-
B20	+	+	-	+	+	+
B21	+	+	-	-	+	+
B22	+	-	-	-	-	-
B23	+	-	-	-	+	-
B24	-	-	-	-	-	-
B25	+	+	-	-	+	-
B26	+	+	-	-	-	-
B27	+	+	-	+	+	+
B28	+	-	-	+	+	+
B29	-	+	-	+	+	-
B30	-	-	-	+	+	+
C	-	-	-	-	-	-

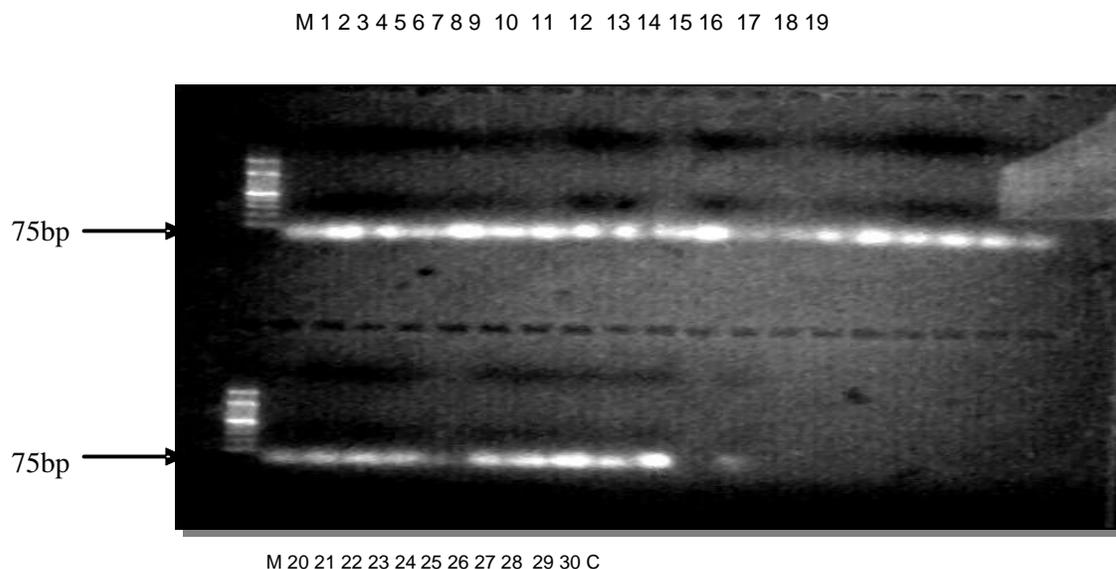
+ = Presence of viral DNA; - = Absence of viral DNA.

transmit the virus following their longer stay on those plants hence the high levels of viral DNA into the plants. The accessions that had fewer number of plants infected could be explained by the late occurrence of TYLCV infection related to whitefly population variation. Moreover, this difference in reaction could be due to the virus strain, vector genotype or altered feeding conditions of the vector (Delatte et al., 2006; Navas-Castillo et al., 1999). The accessions that exhibited varying range of the disease symptoms lacked resistance. Most of them were found to be highly susceptible to TYLCV and only few of them showed very mild symptoms and considered as tolerant. Accessions that had in earlier studies been reported to be TYLCV tolerant (AVRDC, 2001) were found to be susceptible in this study. Accessions A1, A4, A6, A7, A10, G15, B19, B21, B23, B27, B28, and B29 showed a much faster development of disease symptoms

at 30 days after transplanting than the rest and it could be due to the virus strain, vector genotype or altered feeding conditions of the vector (Delatte et al., 2006; Navas-Castillo et al., 1999). The tolerant accessions have been reported to be associated with the presence of exudates from trichome glands on the leaf surface, in which whiteflies become entrapped (Channarayappa and Shivashankar, 1992).

#### Viral DNA detection in tomato accessions

After plants were assessed on the field for the development of disease symptoms, amplification of viral DNA by PCR was done. This was made to confirm the phenotypic screening. The primer pair GhF and GhR detected quite high number of viral DNA. This suggested



**Figure 4.** TYLCV amplified product on agarose gel using GhF and GhR Gh. Primers.

that the primer set was efficient and useful as a reliable marker for detecting or screening susceptible plants to TYLCV. Viral DNA was not detected in tomato accession B24 using the six primer pairs in this study. Meanwhile this accession B24 showed characteristic symptoms of the TYLCV disease in the field and could be putative. Mutations at the primer binding sites in the accession B24 could also account for that. It could also be as a result of other plant viruses other than the TYLCV attacking the plant on the field and indicating similar characteristic symptoms of TYLCV.

In general, based on both phenotypic and molecular evaluations, four categories of accessions were identified, accessions with severe symptoms and presence of viral DNA, accessions with moderate symptoms but relatively low levels of viral DNA, accessions with mild symptoms but high levels of viral DNA and accessions with mild symptoms and no viral DNA. There was actually no accession that showed no symptoms on the field as well as TYLCV DNA amplification. As such, no accession was considered as resistant in this study. Tolerance and resistance are relative terms, largely related to the rate of virus replication (Pilowsky and Cohen, 1990). The results obtained in this study also agreed to this.

## Conclusion

Complete resistance to the tomato yellow leaf curl virus disease in Ghana following field screening and molecular screening was not identified in any of the tomato accessions studied. The tomato accession, B24, showed the highest level of tolerance since it produced weak or

mild symptoms and no viral DNA was detected with the six primer pairs used.

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