

Virus Diseases of Faba Bean (*Vicia faba* L.) in Asia and Africa

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ABSTRACT

Faba bean (*Vicia faba* L.) is the fourth most important pulse crop in the world. Consumed as dry seeds, green vegetable, or as processed food, its products are a cheap source of high-quality protein in the human diet, while its dry seeds, green haulm and dry straw are used as animal feed. This crop is naturally infected by around 50 viruses worldwide, and the number continues to increase. Fortunately only few are of major economic importance in Asian and African countries. This paper will not review the literature on all of these viruses, as there are already a number of comprehensive reviews. Rather will it deal with those of major importance in Asian and African countries and focus on the research progress made over the last two decades. Surveys conducted over the last two decades by ICARDA scientists have shown that the viruses of major economic importance on faba bean in Asia and Africa are: *Faba bean necrotic yellows*, *Bean leafroll*, *Bean yellow mosaic*, *Broad bean mottle* and *Pea seed-borne mosaic viruses*. Other viruses such as *Alfalfa mosaic*, *Beet western yellows*, *Broad bean wilt*, *Broad bean true mosaic*, *Broad bean stain*, *Chickpea chlorotic dwarf*, *Cucumber mosaic*, *Milk vetch dwarf*, *Pea early browning*, *Pea enation mosaic* and *Soybean dwarf viruses* are important in specific locations in specific countries. Significant progress has been made at ICARDA in virus characterization and diagnosis over the last 15 years. The availability of highly sensitive serological methods and specific diagnostic reagents currently permit more accurate detection of viruses. These accomplishments have a positive impact on screening for virus diseases resistance.

Keywords: Algeria, China, control, detection, Egypt, Ethiopia, Iran, Iraq, Japan, Jordan, Lebanon, Libya, Morocco, Pakistan, Sudan, Syria, transmission, Tunisia, Turkey, Yemen

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INTRODUCTION

Faba bean (*Vicia faba* L.) also known as broad bean, horse bean, field bean or tick bean, and is one of the earliest domesticated crops. It is one of the major winter-sown legume crops in the world and has considerable importance as a low-cost food, rich in proteins (20-25% in seeds) and carbohydrates. It is the principal protein source for poor people

in some Asian and African countries.

Faba bean's high adaptability is reflected in the diverse ecological regions implying that it is grown from the equator to almost the Arctic Circle, and from sea level to very high altitudes. The different names reflect the variation within the species, with broad bean meaning the large-seeded cultivars grown for human food, while horse bean and field bean refer to cultivars with smaller, harder seeds

Table 1 Faba bean (dry), area harvested, production and yield in different major production regions in the world during 2002, 2003 and 2004.

Regions	Area harvested (1000 ha)			Yield per hectare (tonnes/ha)			Production (1000 tonnes)		
	2002	2003	2004	2002	2003	2004	2002	2003	2004
WORLD	2659.14	2632.47	2584.29	1.59	1.66	1.65	4226.27	4359.74	4268.10
AFRICA	802.87	860.36	840.63	1.43	1.20	1.30	1150.15	1036.68	1089.40
Algeria	33.61	45.00	45.10	0.68	0.68	0.67	22.93	30.7	30.00
Egypt	127.19	106.07	101.03	3.15	3.18	3.27	400.91	336.84	330.49
Ethiopia	371.67	389.99	380.31	1.22	0.93	1.12	453.13	361.68	426.89
Libya	9.90	9.90	9.96	1.31	1.31	1.30	13.00	13.00	13.00
Morocco	154.10	151.40	150.00	0.58	0.68	0.67	88.78	103.06	100.00
Sudan	58.00	58.00	58.00	2.52	2.41	2.41	146.00	140.00	140.00
Tunisia	43.60	95.00	91.24	0.51	0.51	0.50	22.30	48.00	45.60
ASIA	1298.84	1172.6	1092.02	1.68	1.89	1.88	2179.09	2221.62	2056.53
China	1254.00	1130.00	1050.00	1.67	1.90	1.88	2100.00	2142.00	1976.00
Iraq	*	*	*	*	*	*	*	*	*
Japan	0.10	0.10	0.10	1.00	1.00	1.00	0.10	0.10	0.10
Jordan	*	*	0.34	*	*	2.12	*	*	0.71
Lebanon	*	0.37	0.28		2.16	1.07	*	0.8	0.30
Syria	15.50	15.50	15.49	2.02	2.02	2.02	31.3	31.3	31.30
Turkey	18.00	17.00	17.00	1.78	1.94	1.94	32.00	33	33.00
Yemen	3.28	3.24	2.23	1.35	1.34	1.92	4.44	4.35	4.28
EUROPE	247.35	277.98	282.43	2.60	2.42	2.82	644.21	671.74	796.68
SOUTH AMERICA	112.13	125.93	129.42	0.93	0.90	0.86	104.77	113.05	111.44
N&C AMERICA	40.95	40.60	41.79	0.98	0.98	1.11	40.05	39.65	46.55
AUSTRALIA	157.00	155.00	198.00	0.69	1.78	0.85	108.00	277.00	167.50

Source: FAOSTAT (2007)

* No data available

used for animal feed. Large-seeded cultivars are used as a vegetable, either green or dried, fresh or canned. It is a common breakfast food in the Middle East, Mediterranean region, China and Ethiopia (Bond *et al.* 1985). Roasted seeds are eaten like peanuts in India (Duke 1981). Straw from faba bean harvest fetches a premium in Egypt and Sudan and is considered as a cash crop (Bond *et al.* 1985). The straw is also used for brick making and as a fuel in parts of Sudan and Ethiopia. Faba bean also serves as a break crop in continuous cereal rotations, to improve soil productivity.

Faba bean originated in South Western Asia, probably in the "Fertile Crescent" around the rivers of Tigris and Euphrates. While common beans did not reach Europe before the Spaniards brought them back from America in the early 16th century, the faba bean has been a part of eastern Mediterranean diet for some 8000 years. From this region the faba bean has spread around the world and is now, according to FAO, grown in 50 countries. The major geographical regions that contribute to faba bean production include (a) Far-East, (b) West Asia, (c) North Africa, (d) Nile Valley and Ethiopia, (e) Europe, and (f) Central and South America. The largest producer of faba bean is China (50% of total world production comes from China), followed by Ethiopia and Egypt (FAOSTAT, 2007). The Asian-African countries account for nearly 75% of total global production of faba bean (Table 1), and the average yield in this region is nearly the same as the world average.

The world's largest collections of faba bean germplasm are found at the International Centre for Agricultural Research in the Dry Areas (ICARDA) in Syria, and at the Vavilov Institute in Russia. According to the FAO, there are about 26,000 accessions of faba bean held in genebanks around the world.

Abiotic and biotic factors constitute major constraints that limit the realization of the full yield potential of faba bean and cause yield instability. Faba bean is known to be susceptible to over 100 different pathogens (Schmidt *et al.* 1980; Cockbain 1983; Bos *et al.* 1988; Nene *et al.* 1988; van Emden *et al.* 1988). Many of the diseases which affect faba bean, especially those induced by viruses, can also infect other food and forage legumes. Their relative importance, however, varies depending on the geographical location and the agroecological conditions of the crop production system. Worldwide, around 50 viruses have been reported to infect faba bean (Schmidt *et al.* 1980; Cockbain 1983; Bos *et al.* 1988; Makkouk *et al.* 2003c), 16 of which

were reported to infect this crop in Asia and Africa. The surveys and inventory of virus diseases affecting faba bean in Asia and Africa are incomplete and every year one or more viruses are added to the list of viruses prevalent in these regions.

In this review, we have attempted to review the work done on the most economically important viruses that attack faba bean in the major production countries in Asia and Africa, their ecology and epidemiology, transmission, sensitive assays available for detection, and appropriate measures for their control.

VIRUSES REPORTED TO INFECT FABA BEAN IN AFRICA AND ASIA

Large-scale surveys carried out during the last two decades in Africa and Asia have identified 16 viruses that are the most economically important: 6 causing yellowing/stunting/necrosis and 10 causing mosaic/mottling symptoms (Table 2). These viruses are widespread in specific regions within specific countries.

Viruses causing yellowing, stunting and necrosis

Viruses causing yellowing/stunting are the most important virus diseases in many regions of the world, and were considered for many years (1960-1985) to be caused mainly by infection with *Bean leafroll virus* (BLRV, family *Luteoviridae*). Work conducted between 1985 and 2000 has clearly shown that there are a number of distinct viruses, in addition to BLRV, each of which can produce yellowing/stunting symptoms in faba bean in Asia and Africa. These viruses are *Beet western yellows virus* (BWYV, genus *Polerovirus*, family *Luteoviridae*), *Faba bean necrotic yellows virus* (FBNYV), *Milk vetch dwarf virus* (MDV) (genus *Nanovirus*, family *Nanoviridae*), *Chickpea chlorotic dwarf virus* (CpCDV, genus *Mastrevirus*, family *Geminiviridae*) and *Soybean dwarf virus* (SbDV, family *Luteoviridae*). The distribution of these viruses in Asia and Africa is presented in Table 2. FBNYV was found to be the most prevalent virus, followed by BLRV, BWYV and CpCDV. MDV, a virus related to FBNYV, has so far been found only in China and Japan.

The symptoms induced by the above six viruses are mostly leaf-rolling, yellowing and stunting of infected plants (Fig. 1). Old leaves of infected plants tend to be leathery. All these viruses are phloem-limited and transmit-

Table 2 Viruses reported to naturally infect faba bean in Africa and Asia.

Continent/ Country	Virus reported on faba bean** (reference number)															
	Viruses causing yellowing/stunting/necrosis symptoms						Viruses causing mosaic/ mottling/ enation symptoms									
	FBNYV	MDV	BLRV	BWYV	SbDV	CpCDV	PEMV	BYMV	PSbMV	BBWV	AIMV	CMV	BBMV	BBSV	BBTMV	PEBV
AFRICA																
Algeria	+ ^{C,P}		+ ^C	+ ^C				+ ^C	+ ^L				+++			+++
	(32)		(5)	(4)				(4)	(42)				(71)			(38)
Egypt	+++		+++			+++	+++	+++	+++	+++	+++	+++	+++	+++	+++	
	(32, 43)		(40, 43)			(49)	(40)	(24, 40)	(7, 42)	(24, 43)	(16, 43)	(40, 51)	(40)	(7, 40)	(7, 51)	
Ethiopia	+++		+++	+++	+++	+++	+ ^L	+++	+++	+ ^C	+ ^P	+ ^P	+ ^C	+++	+++	+ ^P
	(13, 32)		(13)	(13)	(13)	(2)	(63)	(2, 13)	(2, 42)	(63)	(1)	(1)	(13)	(2)	(2)	(1)
Libya	+++		+++					+++	+++		+++	+++				+++
	(19)							(68)	(42)		(28)	(28)				(14)
Morocco	+++		+++	+ ^C			+++	+++	+++	+++	+++	+ ^C	+++	+++	+++	+++
	(15)		(22, 40)	(23)			(22)	(22, 40)	(22, 42)	(41)	(22)	(17)	(22, 40)	(22, 42)	(22)	(22, 37)
Sudan	+++		+++			+++	+++	+++	+++	+++	+++	+++	+++	+++	+++	
	(48)		(40, 44)			(44)	(40)	(40, 44)	(40, 44)	(41)	(44, 58)	(40, 52)	(40, 54)	(40)	(40)	
Tunisia	+++		+++	+++	+++		+++	+++	+++	+++	+++	+++	+++	+++	+++	
	(55)		(40, 55)	(55)	(56)		(40)	(40)	(40, 42)	(40, 41)		(40, 55)	(40, 55)	(40, 55)	(40)	
ASIA																
China		+++	+++	+++			+++	+++	+++	+++		+++	+++	+++	+++	
		(34)	(67)	(62)			(69)	(62, 67)	(62)	(66, 67)		(62, 70)	(21)	(67)	(21)	
Iran	+++		+++	+ ^{C,L}	+ ^{C,L}	+ ^{C,L}	+++	+++	+++	+++	+++	+++	+++	+++	+ ^L	
	(50)		(29, 50)	(50)	(50)	(50)	(20)	(30, 50)	(29)	(59)	(29)	(29)	(49)	(50)		
Iraq	+++		+++	+++	+++	+++		+++	+ ^C	+++	+++	+++	+++	+++		
	(18)		(18)	(18)	(18)	(18)		(18)	(31)	(18)	(18, 60)	(18)	(6)			
Japan		+++			+ ^S			+++	+++	+++		+++				
		(27, 61)			(64)			(26)	(65)	(26)		(65)				
Jordan	+++		+++	+++			+++	+++	+ ^L	+++	+++				+ ^L	
	(11, 32)		(35)	(35)				(10)	(8)	(9)	(57)			(8)		
Lebanon	+++		+++	+ ^C				+++	+++		+++			+++	+++	
	(32)		(40)	(25)				(39, 57)	(40)		(57)			(40)	(57)	
Pakistan	+ ^{C,L}			+ ^{C,L}		+ ^{C,L}		+++	+ ^{C,L}		+ ^C	+ ^{C,L}				
	(47)			(47)		(47)		(3)	(47)		(47)	(47)				
Syria	+++		+++	+++	+ ^C	+ ^C	+++	+++	+++	+++	+++	+++	+++	+++	+++	+++
	(32)		(40)	(53)	(45)	(33)	(40)	(40)	(40)	(40, 41)	(53)	(40)	(40)	(40)	(40)	
Turkey	+++		+ ^C	+ ^C		+ ^C		+++	+ ^L						+ ^L	
	(32)		(25)	(25)		(35)		(36)	(42)					(12)		
Yemen	+++		+++			+++		+++	+++		+++					
	(46)		(46)			(46)		(46)	(46)		(46)					
Manner of transmission in faba bean*	Aph-P	Aph-P	Aph-P	Aph-P	Aph-P	Leaf-hoppers	Sa, Aph-P	Sa, Se, Aph-NP	Sa, Se, Aph-NP	Sa, Se, Aph-NP	Sa, Aph-NP	Sa, Se, Aph-NP	Sa, Se, Beetles	Sa, Se, Beetles	Sa, Se, Beetles	Sa, Se, Nematodes

* Sa= Sap, Se= seeds, Aph-NP= Aphids in non-persistent manner; Aph-P= Aphids in persistent manner.

** +++: reported on faba bean, +: reported in this country but not on faba bean, +^S reported on chickpea, +^L: reported on lentil, +*^P: reported on pea, +*^S: reported on soybean, FBNYV= *Faba bean necrotic yellows virus*, MDV= *Milk vetch dwarf virus*, BLRV= *Bean leaf roll virus*, BWYV= *Beet western yellows virus*, SbDV= *Soybean dwarf virus*, CpCDV= *Chickpea chlorotic dwarf virus*, PEMV= *Pea enation mosaic virus*, BYMV= *Bean yellow mosaic virus*, PSbMV= *Pea seed-borne mosaic virus*, BBWV= *Broad bean wilt virus*, AIMV= *Alfalfa mosaic virus*, CMV= *Cucumber mosaic virus*, BBMV= *Broad bean mottle virus*, BBSV= *Broad bean stain virus*, BBTMV= *Broad bean true mosaic virus*, PEBV= *Pea early browning virus*, the numbers between brackets are reference numbers as follows: (1) Abraham and Makkouk 2002; (2) Abraham et al. 2000; (3) Aftab et al. 1989; (4) Ait Yahia et al. 1997; (5) Ait Yahia et al. 1999; (6) Al-Ani and Al-Azzawi, 1987; (7) Allam et al. 1979; (8) Al-Mabrouk and Mansour 2000; (9) Al-Musa et al. 1986; (10) Al-Musa et al. 1987; (11) Al-Nsour et al. 1998; (12) Bayaa et al. 1998; (13) Bekele et al. 2005; (14) Bos et al. 1993; (15) El-Amri 1999a; (16) El-Attar et al. 1971; (17) El-Maataoui and El-Hassani 1984; (18) El-Muadhidi et al. 2001; (19) Fadel et al. 2005; (20) Farzadfar and Izadpanah 2001; (21) Ford et al. 1981; (22) Fortass and Bos 1991; (23) Fortass et al. 1997; (24) Gamal-Eldin et al. 1982; (25) Horn et al. 1995; (26) Inouye 1969; (27) Inouye et al. 1968; (28) Ismail and Hassan 1995; (29) Kaiser et al. 1968; (30) Kaiser 1973; (31) Kassim 1997; (32) Katul et al. 1993; (33) Kumari et al. 2004; (34) Kumari et al. 2007; (35) Kumari 2002; (36) Kurçman 1977; (37) Lockhart and Fischer 1976; (38) Mahir et al. 1992; (39) Makkouk et al. 1982; (40) Makkouk et al. 1988b; (41) Makkouk et al. 1990; (42) Makkouk et al. 1993b; (43) Makkouk et al. 1994; (44) Makkouk et al. 1995; (45) Makkouk et al. 1997; (46) Makkouk et al. 1998a; (47) Makkouk et al. 2001; (48) Makkouk et al. 2003a; (49) Makkouk et al. 2003b; (50) Makkouk et al. 2003d; (51) Mazyad et al. 1975; (52) Milles and Ahmed 1984; (53) Mouhanna et al. 1994; (54) Murant et al. 1974; (55) Najjar et al. 2000; (56) Najjar et al. 2003; (57) Nienhanus and Saad 1967; (58) Nour and Nour 1962; (59) Parvin and Izadpanah 1978; (60) Salama and El-Behadli 1979; (61) Sano et al. 1998; (62) Shiyang et al. 2007; (63) Tadesse et al. 1999; (64) Tamada 1975; (65) Tanaka et al. 1973; (66) Xu et al. 1988; (67) Xu Zhiang et al. 1985; (68) Younis et al. 1992; (69) Yu 1979; (70) Yu et al. 2005; (70) Zagh and Ferault 1980.

ted by aphids in a persistent manner except CpCDV that is persistently transmitted by a leafhopper. They are neither transmitted mechanically nor by seeds. In some countries (e.g. Egypt, Syria and Tunisia) these viruses have, in some years, caused almost complete failure of the faba bean crop (Makkouk et al. 1994, 1998b; Najjar et al. 2000).

Viruses causing mosaic/mottling

Faba bean viruses causing mosaic/mottling (Fig. 1) have been reported in 17 countries in Africa and Asia (Table 2). These viruses are: *Alfalfa mosaic virus* (AIMV, genus *Alfalmovirus*, family *Bromoviridae*), *Bean yellow mosaic virus* (BYMV, genus *Potyvirus*, family *Potyviridae*), *Broad bean wilt virus* (BBWV, genus *Fabavirus*, family *Comoviridae*), *Broad bean mottle virus* (BBMV, genus *Bromovirus*, family *Bromoviridae*), *Broad bean stain virus* (BBSV), *Broad bean true mosaic virus* (BBTMV) (genus *Comovirus*, family *Comoviridae*), *Cucumber mosaic virus* (CMV, genus *Cucumovirus*, family *Bromoviridae*), *Pea enation mosaic virus-1* (PEMV-1, genus *Enamovirus*, family *Luteoviridae*) and *Pea early browning virus* (PEBV, genus *Tobravirus*).

High incidence of BYMV was detected in most African and Asian countries, followed by PSbMV, CMV and AIMV. PEBV so far has been reported only from North African countries (Lockhart and Fischer 1976; Mahir et al. 1992; Bos et al. 1993). This virus is transmitted by free-living Trichodorida nematodes, which tend to remain localized in soil but can subsist on many host species including weeds. Disease distribution in crops therefore is usually localized, and such infection spots may only gradually enlarge even in the presence of sensitive crops.

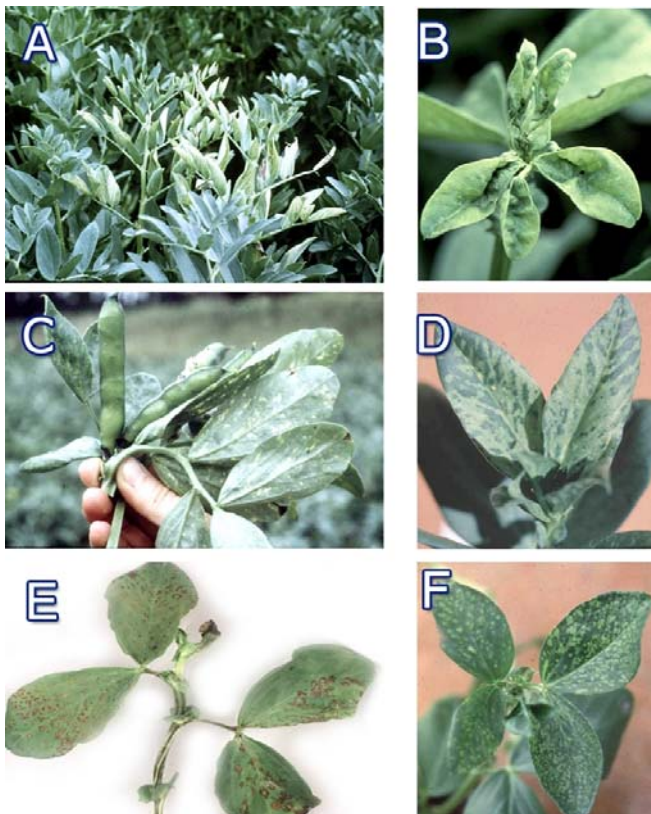


Fig. 1 (A) Chlorosis and leaf rolling symptoms in faba bean caused by *Bean leafroll virus*, (B) leaf cupping and chlorosis of growing point of faba bean plants caused by *Faba bean necrotic yellows virus*, (C) irregular chlorotic spots of pods and leaves of faba bean plant caused by *Pea enation mosaic virus*, (D) mosaic symptoms on faba bean leaves caused by *Bean yellow mosaic virus*, (E) necrotic symptoms on stem and leaves of faba bean plant caused by *Alfalfa mosaic virus*, (F) mottling with chlorotic spots on faba bean leaves caused by *Broad bean mottle virus*.

Other viruses

In addition to the viruses mentioned above, a few other viruses were reported to cause damage to faba bean in specific countries and in limited areas, such as *Tomato spotted wilt virus* (TSWV, genus *Tospovirus*, family *Bunyaviridae*) in China (Yu 1947); *Broad bean necrosis virus* (BBNV, genus *Pomovirus*) in Japan (Inouye and Nakasone 1980); and *Chickpea chlorotic stunt virus* (CCSV, genus *Polerovirus*, family *Luteoviridae*) in Ethiopia (Abraham *et al.* 2006). In addition, broad bean phyllody has been reported in Sudan (Jones *et al.* 1984), caused by a phytoplasma. Phyllody is characterized by the replacement of flower parts by green leaf-like structures.

ECONOMIC IMPORTANCE

Viral disease epidemics and associated crop losses not only depend on the incidence of infection, but also on symptom severity and growth stage at which infection occurs. The growth stage when plants become virus-infected and the proportion of plants infected in the crop are critical factors in determining the extent of yield losses. Losses are generally greatest when plants become infected at vulnerable early growth stages and incidence reaches high levels.

Viruses causing yellowing/stunting can have a marked effect on yield. During the growing season of 1991/1992, a severe FBNYV epidemic affected faba bean in Middle Egypt leading to yield losses of over 90% (Makkouk *et al.* 1994). In Iran, faba bean plants inoculated with BLRV during or after flowering yielded, respectively, 89 and 59% less than uninoculated plants; plants inoculated before flowering failed to produce any seed (Kaiser 1973).

In a field yield experiment in Syria, using 4 x 1.8 m plots and a plant density of 12 seeds/m², artificial infection with BYMV, BBMV and BBSV 11 weeks after sowing (pre-flowering) led to 81, 54 and 84% yield loss, respectively. Inoculation with the same viruses 15 weeks after sowing (flowering) and 20 weeks after sowing (pod setting) led to 56, 84 and 18%, and 39, 37 and 18% yield loss, respectively. The mixed infection of BBMV and BYMV caused almost complete failure of the crop when inoculation was made before or during flowering (Makkouk *et al.* 1988b). In Iran, faba bean plants inoculated with BYMV before, during or after flowering yielded, respectively, 44, 42 and 23% less than uninoculated plants (Kaiser 1973). Inoculation of faba bean variety "Syrian Local" with a Syrian isolate of BBWV 14 weeks (pre-flowering) and 16 weeks after sowing (flowering) led to 25.8 and 1.8% yield loss, respectively (Makkouk *et al.* 1990). Seed yield loss of faba bean inoculated with PSbMV at the flowering stage was 40.5% (Makkouk *et al.* 1993b). In Egypt, the number of pods/plant was 55.5, 11.25 and 20.33% less than that of healthy plants when faba bean plants were inoculated with BYMV, BBTMV and BBSV, respectively (Allam *et al.* 1979).

BYMV is worldwide in distribution, infects many wild and cultivated legumes and is the most common cause of mosaic symptoms in faba bean. Field incidence with BYMV can vary greatly among locations. High incidence, up to 100%, was observed in some regions of Egypt, Sudan and the coastal areas of Syria. These locations are known for their relatively warm winters which favor increased aphid population and movement. Studies conducted in the region suggested that potential yield loss due to BYMV infection could vary from 15 to 45% depending on the time of infection (Kaiser 1973; Makkouk *et al.* 1988b).

Several viruses impair the quality of faba bean seeds thereby rendering them less attractive to consumers. For example, PSbMV induces necrotic rings and line patterns and malformation in faba bean seed coats (Makkouk *et al.* 1993b). Likewise, BBSV infection leads to undesirable staining of faba bean seed coat, which renders the seeds useless for canning (Bos *et al.* 1988) (Fig. 2). Kaiser (1973) found that pods of faba beans, infected with BYMV, occasionally developed necrotic ring spotting and discolored seeds.

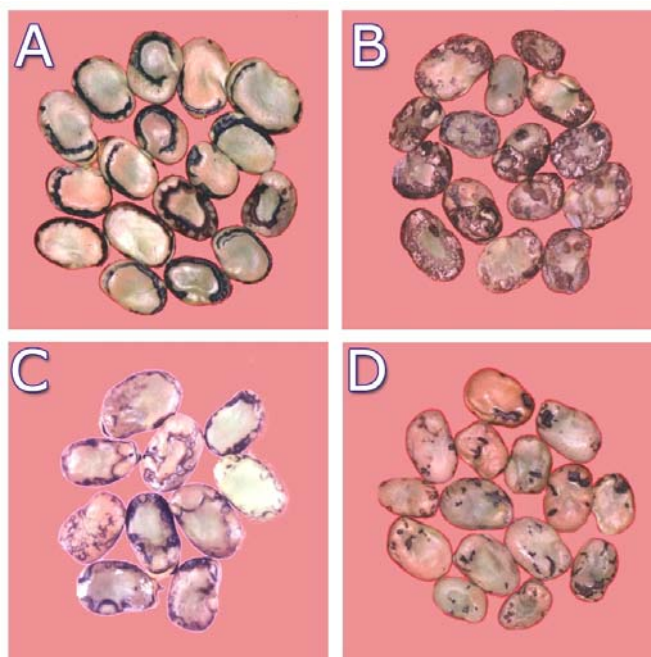


Fig. 2 Seed-coat symptoms of faba bean seeds from plants infected with *Broad bean stain virus* (A), *Broad bean mottle virus* (B), *Pea seed-borne mosaic virus* (C) and *Bean yellow mosaic virus* (D).

VIRUS TRANSMISSION

Virus spread by vectors

Insects are particularly important vectors because they are usually highly mobile and may travel long distances. All faba bean viruses reported in Asia and Africa are known to be spread by vectors, with the majority (15 viruses) being transmitted by insects. One of the insect-borne viruses is transmitted by leafhoppers (CpCDV), three by beetles (BBSV, BBMV and BBTMV) and 11 by aphids. Six viruses are transmitted by aphids in a persistent manner and five viruses in the non-persistent manner. Only PEBV is transmitted by nematodes (Table 2). Spread of the persistent viruses (e.g. FBNYV, BLRV) can be over a long distance, with the possibility that an individual insect can transmit the virus to many plants. The beetle-transmitted viruses can persist in the vector for days or weeks. Table 3 summarizes the most important vectors reported to transmit faba bean viruses worldwide.

As is the case of many diseases, the progress curve for virus-infected plants is usually sigmoid. The extent and rate of increase, however, depend on the distance from the inoculum source and the wind direction. A typical distance of spread from the source for non-persistent viruses, such as BYMV, is 100 m upwind and 250 m downwind, and the virus movement from the source to the crop is according to a gradient. On the other hand, persistent transmission by aphids, such as BLRV, is over a long distance. Consequently, the source of infection is difficult to trace, and virus movement to the crop is usually at random and not according to a gradient. In areas where alfalfa is grown, infection with BLRV is usually high because most alfalfa varieties are susceptible to BLRV and often do not show symptoms.

Our observations in faba bean fields in Egypt, Syria and Tunisia indicated that aphid vectors, transmit FBNYV to a few plants in the field early in the season (September-October), often close to the field border. When weather conditions favor aphid population build-up and activity, these single-plant foci increase to patches of 50-100 infected plants within 2-3 weeks. This type of spread suggests that faba bean is a favorable host for both the virus and its vectors and that the spread of FBNYV mainly depends on colonizing aphid species within the crop in a fashion typical of persistently transmitted viruses. In addition, environmental conditions play an important role in the spread of FBNYV. High incidence has been observed in regions characterized by mild winters such as Middle Egypt (Beni Suef and Minia governorates), the Jordan Valley and the coastal areas of Syria and Turkey. In all these regions winter temperatures rarely fall below 5°C permitting the aphid vector to overwinter parthenogenetically. When temperatures rise the aphids become active, multiply and spread the virus.

Seeds infected with seed-borne viruses are dispersed at random during planting. The spread of virus by insect vectors from these foci can be rapid. Kaiser *et al.* (1968) reported that in Iran, the incidence of BYMV in field trials of faba bean increased from 0.2% seed-borne infections to 85% disease incidence in three months. Seed certification for seed-borne viruses is mainly to limit economic loss rather than to eradicate the pathogen. Tolerance levels, i.e., levels of seed-borne virus that do not lead to economic losses, may be high in regions where viruses have no efficient vector or the vector's population density is low. For example, when faba bean seeds infected with BBSV were multiplied for several years in Scotland, where the beetle vectors *Apion vorax* and *Sitona lineatus* were either absent or rare, virus infection of the seeds declined from year to year (Jones 1980).

Seed transmission

Virus transmission via seed is of dual importance. Virus-infected seeds act both as sources of inoculum and as vehicles of virus dissemination. Out of the 16 viruses reported to in-

fect faba bean in Asia and Africa, eight are seed-transmitted (Table 4). Rates of seed transmission greatly depend on the virus, virus strain, host species and host cultivar. Viruses that infect the embryo may also be transmitted in pollen to seeds on virus-free mother plants. Infection of faba bean after flowering rarely leads to infection of the seed. Commonly, infected seeds appear to be normal-looking except in a few cases where visible symptoms are observed, as in the case of BBSV. Thus, visual inspection of seeds cannot be used to eliminate seed-borne viruses. BBSV was thought to have been imported into England with seed from Morocco (Gibbs *et al.* 1968) and, indeed, was later confirmed to be widespread in Morocco (Fischer and Lockhart 1976).

Surveys conducted in the region, have revealed the occurrence of a few viruses with wide distribution and often high incidence in faba bean crops (Table 2). Most are potentially of great significance for crop improvement programs because of their transmission in seed. Serious consideration must be given to those viruses in international breeding programs and in systems of commercial seed production. For example, BBMV is known to occur in Syria, Egypt, Sudan, Tunisia, Morocco (Makkouk *et al.* 1988b) and Algeria (Zagh and Ferault 1980). A detailed survey in Morocco revealed its occurrence in 56% of the fields inspected, with a maximum incidence of 33% recorded in one field (Fortass and Bos 1991). The virus is beetle- and seed-transmitted and may cause serious losses.

VIRUS DETECTION

Accurate diagnosis combined with sensitive, rapid and early detection of viral diseases is critical for effective management of faba bean-based cropping systems. Appropriate control procedures can only be applied effectively if the disease is correctly identified and distribution in an area is known. The last three decades have witnessed significant developments in the sensitivity of the methods used to detect faba bean viruses.

Immunological (protein-based) methods

The development of enzyme-linked immunosorbent assay (ELISA) for plant virus detection (Clark and Adams 1977) was a major step forward and replaced earlier serological methods such as gel diffusion, especially for large-scale testing. This was enhanced further with the development of monoclonal antibody technology and its application to a large number of faba bean viruses. Specific reagents (monoclonal and/or polyclonal antibodies) have been developed for most viruses affecting faba bean (Table 5), and their utilization in a variety of ELISA tests has made faba bean virus diagnosis simple and fast. Since then a number of ELISA variants were developed which further improved the sensitivity of faba bean virus testing. Sensitivity of detection was increased further by using chemiluminescent substrates (e.g. BYMV in faba bean) (Makkouk *et al.* 1993a). In many laboratories in developing countries, facilities for sophisticated tests are lacking. Accordingly, the simpler procedure the more widely it can be used. For this reason Tissue-blot immunoassay (TBIA) was developed to identify most faba bean viruses (Makkouk and Kumari 1996). TBIA does not require expensive equipment, is simple to conduct, inexpensive, sensitive and can be completed within 3-4 hours. In addition, it is fairly easy to differentiate infected from healthy plants. TBIA is a useful test to detect all faba bean viruses reported and it is especially recommended for virus surveys (Najar *et al.* 2000; El-Muadhidi *et al.* 2001; Makkouk *et al.* 2001, 2003d) and for evaluating virus-host interactions (Makkouk *et al.* 2002; Kumari and Makkouk 2003) (Fig. 3).

Molecular (nucleic acid-based) methods

Nucleic acid hybridization has been used successfully for the detection of many viruses (Hammond and Hammond

Table 3 Major insect and nematode vectors reported to transmit faba bean viruses.

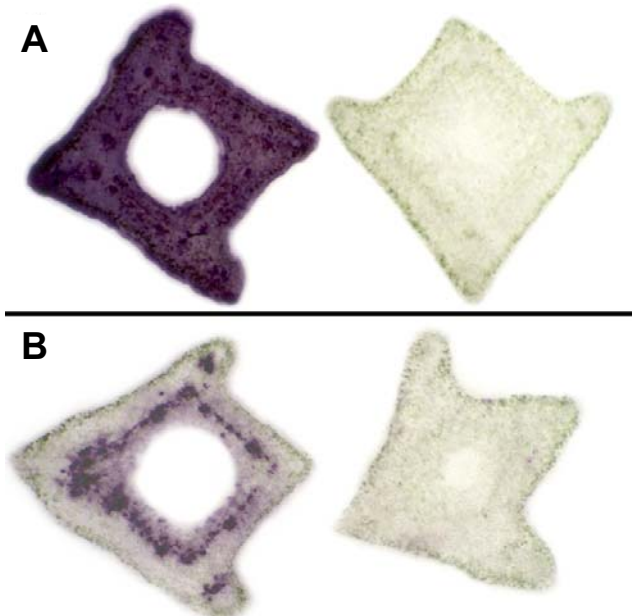
Vector	Virus transmitted	Reference	
Aphids			
<i>Acyrtosiphon pisum</i> Harris	AIMV, BWYV, MDV	Cockbain 1983	
	BBWV	Makkouk <i>et al.</i> 1990	
	BLRV	Cockbain 1983; Johnstone <i>et al.</i> 1984; Skaf and Makkouk, 1988; Ait Yahia <i>et al.</i> 1999	
	BYMV	Cockbain 1983; Skaf and Makkouk, 1988	
	CMV	Edwardson and Christie 1986	
	FBNYV	Katul <i>et al.</i> 1993; Franz <i>et al.</i> 1998; Al-Nsour <i>et al.</i> 1998; Al-Amri 1999b	
	PEMV	Cockbain 1983; Cockbain <i>et al.</i> 1986; Hagedorn 1996; Kumari <i>et al.</i> 2001b	
	PSbMV	Makkouk <i>et al.</i> 1993b	
	SbDV	Makkouk <i>et al.</i> 1997; Damsteegt <i>et al.</i> 1999; Terauchi <i>et al.</i> 2003	
	<i>Aphis fabae</i> Scopoli	AIMV	Cockbain 1983
BBWV		Makkouk <i>et al.</i> 1990	
BLRV		Kaiser 1973; Johnstone <i>et al.</i> 1984; Skaf and Makkouk, 1988; Ait Yahia <i>et al.</i> 1999	
BYMV		Cockbain 1983; Skaf and Makkouk, 1988	
CMV		Edwardson and Christie 1986	
FBNYV		Katul <i>et al.</i> 1993; Al-Amri 1999b	
PSbMV		Makkouk <i>et al.</i> 1993b	
<i>Aphis craccivora</i> Koch.	AIMV, PEMV	Cockbain 1983	
	BBWV	Makkouk <i>et al.</i> 1990	
	BLRV	Cockbain 1983; Johnstone <i>et al.</i> 1984; Skaf and Makkouk, 1988; Ait Yahia <i>et al.</i> 1999	
	BWYV	Boswell and Gibbs 1983	
	BYMV	Cockbain 1983; Skaf and Makkouk, 1988	
	CMV	Edwardson and Christie 1986	
	FBNYV	Katul <i>et al.</i> 1993; Franz <i>et al.</i> 1998; ; Al-Nsour <i>et al.</i> 1998; Al-Amri 1999b	
	MDV	Cockbain 1983; Sano <i>et al.</i> 1998	
	PSbMV	Makkouk <i>et al.</i> 1993b	
	<i>Myzus persicae</i> (Sulz.)	AIMV	Edwardson and Christie 1986; Wang <i>et al.</i> 2006
BBWV		Makkouk <i>et al.</i> 1990	
BLRV		Cockbain 1983; Johnstone <i>et al.</i> 1984	
BYMV, BWYV, MDV		Cockbain 1983	
CMV		Edwardson and Christie 1986	
PEMV		Cockbain <i>et al.</i> 1986; Edwardson and Christie 1986; Hagedorn 1996; Kumari <i>et al.</i> 2001b	
PSbMV		Makkouk <i>et al.</i> 1993b	
<i>Macrosiphum euphorbiae</i> (Thomas)		AIMV, BYMV, CMV	Edwardson and Christie 1986
		BBWV	Stubbs 1960
		BLRV	Cockbain 1983; Johnstone <i>et al.</i> 1984
	PEMV	Edwardson and Christie 1986; Hagedorn 1996	
<i>Rhopalosiphum padi</i> L.	CMV, PEMV	Edwardson and Christie 1986	
	PSbMV	Makkouk <i>et al.</i> 1993b	
<i>Aulacorthum solani</i> (Kaltenbach.)	AIMV, CMV, PEMV	Edwardson and Christie 1986	
	BWYV	Cockbain 1983	
	SbDV	Tamada 1975; Cockbain 1983; Honda 2001; Terauchi <i>et al.</i> 2003	
Beetles			
<i>Acalymma trivittata</i> Mannerheim	BBMV	Walters and Surin 1973	
<i>Apion aestivum</i> Germ.	BBSV	Cockbain <i>et al.</i> 1975	
<i>Apion aethiops</i> Hbst.	BBSV, BBTMV	Cockbain <i>et al.</i> 1975	
<i>Apion arrogans</i> Wencher	BBMV, BBSV	Makkouk and Kumari 1989, 1995a	
<i>Apion radiolus</i> Kirby	BBMV	Fortass and Diallo 1993	
<i>Apion vorax</i> Hbst.	BBMV	Cockbain 1983	
	BBSV, BBTMV	Cockbain <i>et al.</i> 1975	
<i>Colaspis flavida</i> Say	BBMV	Walters and Surin 1973	
<i>Diabrotica undecimpunctata</i> Mannerheim	BBMV	Walters and Surin 1973	
<i>Hypera variabilis</i> Herbst	BBMV	Fortass and Diallo 1993	
<i>Pachytychius strumarius</i> Gyll	BBMV	Fortass and Diallo 1993	
<i>Sitona lineatus</i> var. <i>viridifrons</i> Motsch	BBMV	Borges and Louro 1974	
<i>Sitona crinita</i> Herbst	BBSV	Makkouk and Kumari 1995a	
<i>Sitona limosa</i> Rossi	BBMV, BBSV	Makkouk and Kumari 1995a	
<i>Sitona lineatus</i> L.	BBMV, BBSV	Makkouk and Kumari 1995a	
	BBMV	Fortass and Diallo 1993	
	BBSV, BBTMV	Cockbain <i>et al.</i> 1975	
<i>Spodoptera exigua</i> Hübner	BBMV	Ahmed and Eisa 1999	
<i>Smicronyx cyaneus</i> Gyll	BBMV	Fortass and Diallo 1993	
Leafhoppers			
<i>Orosius orientalis</i> (Matsumura)	CpCDV	Horn <i>et al.</i> 1993	
<i>Orosius albicinctus</i> Distant	CpCDV	Kumari <i>et al.</i> 2004	
Nematodes			
<i>Trichodorus primitivus</i> , <i>T. viruliferous</i> , <i>Paratrichodorus anemones</i> , <i>P. pachydermus</i> , <i>P. teres</i>	PEBV	Boulton 1996	

Table 4 Viruses that can be transmitted via faba bean seeds.

Virus	Reported seed transmission rate (%)	Reference
BYMV	0.1-15	Kaiser 1973; Murrant <i>et al.</i> 1974; Sasaya <i>et al.</i> 1993; Latham and Jones 2001a
BBMV	1-2	Makkouk <i>et al.</i> 1988a; Murrant <i>et al.</i> 1974; Fortass and Bos 1992
BBSV	1-10	Gibbs and Smith 1970; Jones 1978; Makkouk <i>et al.</i> 1987a; Allam <i>et al.</i> 1979
BBTMV	1-17	Blaszczak 1970; Cockbain <i>et al.</i> 1976; Jones 1978; Allam <i>et al.</i> 1979
BBWV	0.4-0.6	Putz and Kuszala, 1973; Makkouk <i>et al.</i> 1990
CMV	0.1	Latham and Jones 2001a
PEBV	1-10	Fiedorow 1983
PSbMV	0.2-2	Makkouk <i>et al.</i> 1993b; Musil 1980; Latham and Jones 2001b

Table 5 Antibodies available for the detection of faba bean viruses

Virus	Antibodies type	Reference
AIMV	Polyclonal	Makkouk <i>et al.</i> 1987b
	Monoclonal	Hajimorad <i>et al.</i> 1990; Gallo and Matisova 1993
BBMV	Polyclonal	Makkouk <i>et al.</i> 1987b, 1988a; Fortass and Bos 1992
BBSV	Polyclonal	Jones and Barker 1976; Makkouk <i>et al.</i> 1987a
	Monoclonal	Subr <i>et al.</i> 1994
BBTMV	Polyclonal	Jones and Barker 1976; Kumari and Makkouk (ICARDA, unpublished data)
BBWV	Polyclonal	Uyemoto and Provvidenti 1974; Xu <i>et al.</i> 1988; Makkouk <i>et al.</i> 1990; Qi <i>et al.</i> 2002
	Monoclonal	Qing <i>et al.</i> 2000
BLRV	Polyclonal	D'Arcy <i>et al.</i> 1989; Ashby and Huttinga 1979; Ait Yahia <i>et al.</i> 1999; Kumari and Makkouk (ICARDA, unpublished data)
	Monoclonal	Katul 1992
BWYV	Polyclonal	Govier 1985; Kumari and Makkouk (ICARDA, unpublished data)
	Monoclonal	Rabenstein <i>et al.</i> 1984; D'Arcy <i>et al.</i> 1989; Herrbach <i>et al.</i> 1991; Ellis and Wiczorek 1992
BYMV	Polyclonal	Makkouk <i>et al.</i> 1988c; Subr and Matisova 1999
	Monoclonal	Scott <i>et al.</i> 1989; Jordan and Hammond 1991; Werkmeister and Shukla 1991; Subr and Matisova 1999
CMV	Polyclonal	Kumari and Makkouk (ICARDA, unpublished data)
	Monoclonal	Haase <i>et al.</i> 1989; Porta <i>et al.</i> 1989; Wahyuni <i>et al.</i> 1992; Hayes <i>et al.</i> 1994; Hsu <i>et al.</i> 2000; Yu <i>et al.</i> 2005
CpCDV	Polyclonal	Horn <i>et al.</i> 1993; Kumari <i>et al.</i> 2006
FBNYV	Polyclonal	Katul <i>et al.</i> 1993; Kumari <i>et al.</i> 2001a
	Monoclonal	Franz <i>et al.</i> 1996
PEBV	Polyclonal	Makkouk and Kumari 1998
PEMV	Polyclonal	Gibbs <i>et al.</i> 1966; Izadpanah and Shepherd 1966; Mahmood and Peters 1973; Kumari <i>et al.</i> 2001b
PSbMV	Polyclonal	Makkouk <i>et al.</i> 1993b
	Monoclonal	Jordan and Hammond 1991
SbDV	Polyclonal	Kojima and Tamada 1976; Makkouk <i>et al.</i> 1997; Damsteegt <i>et al.</i> 1999
	Monoclonal	D'Arcy <i>et al.</i> 1989; Mikoshiba <i>et al.</i> 1996; Damsteegt <i>et al.</i> 1999

**Fig. 3** Detection of *Bean yellow mosaic virus* (A) and *Faba bean necrotic yellows virus* (B) by tissue blot immunoassay (TBIA) in infected faba bean stem blot (left) as compared to a healthy plant (right).

1985; Pesic and Hiruki 1988; Martin and D'Arcy 1990; Herrbach *et al.* 1991). Cloning of plant viral nucleic acids and the development of nonradioactive detection methods have increased the utility of nucleic acid hybridization for virus detection. The development of polymerase chain reac-

tion (PCR) has greatly improved the sensitivity and utility of hybridization and other nucleic acid based assays. Immunocapture PCR is a very sensitive detection method that combines the advantages of serology and PCR (Phan *et al.* 1997; Shamloul *et al.* 1999; Yu *et al.* 2005). Faba bean viruses for which PCR detection assays have been reported are listed in **Table 6**. The technique has been adapted to the detection of both DNA and RNA viruses (with either single- or double-stranded genomes (**Table 6**). In addition to faba bean virus detection in infected plants, PCR has also been adapted for the detection of viruses in their vectors (van der Wilk *et al.* 1994; Shamloul *et al.* 1999).

VIRUS DISEASE CONTROL

As there is no direct practical way of curing crops from virus infection, all current control strategies emphasize measures that prevent or reduce infection. Control measures can be classified into (i) those that control the virus, (ii) those that are directed towards avoidance of vectors or reducing their incidence, and (iii) integrated approaches, which combine all possible control components in such a way that they complement each other and can be applied at farm level as one control package.

Methods directed to control the virus

Targeting sources of infection

Since around 50% of viruses affecting faba bean (Bos *et al.* 1988) are seed-borne (**Table 4**), it is always recommended to use virus-free seed for planting, especially when the virus is also transmitted by active vectors. The removal of symp-

Table 6 Primers available for detection of faba bean viruses by PCR.

Virus	Genome	Sequence of primer pairs 5' – 3'	Amplified fragment (bp)	Reference
AIMV	ssRNA	CGTCAGCTTTCGTCGAACA GCCGTCGCGCATGGTAAT	288	Braiana <i>et al.</i> 1994
BBWV-II	ssRNA	AACTGAAACTCGCCATCTCC ATAGTTTCCGTGGCTGATTC	444	Uga 2005
BLRV	ssRNA	TCCAGCAATCTTGGCATCTC GAAGATCAAGCCAGGTTCA AAAGAGGTTCTACAGGCCAC GATCAAGTTCCTCGCAGAAC	391 440	Ortiz <i>et al.</i> 2005 Kumari <i>et al.</i> 2006
BWYV	ssRNA	ATGAATACGGTCTGGGTAC GATAGTTGAGGAAAGGGAGTTG GTCTACCTATTTGG ATGGTCGCTAGAGG ATGCAATTTCTCGCTCACGCAAACA TCATACAAACATTTCCGGTGTGAC	429 950 750	Kumari <i>et al.</i> 2006 Fortass <i>et al.</i> 1997 Hauser <i>et al.</i> 2000
BYMV	ssRNA	GGTTTGGCYAGRATGCTTTTG GAGAATTTAAAGACGGATA CAGTTTATTATGCAGCGG GTTATCATCAATCTTCCTGC	240 644	Braiana <i>et al.</i> 1994 Uga 2005
CMV	ssRNA	CGAGTCATGGACAAATCTGAATCAA AGYCCTTCCGAAGAAAYCTAGGAGA GCCGTAAGCTGGATGGACAA TATGATAAGAAGCTTGTTTCGCG GGCGAATTCGAGCTCGCCGTAAGCTGGATGGAC CTCGAATTCGGATCCGCTTCTCCGCGAG TATGATAAGAAGCTTGTTTCGCGCA TTTTAGCCGTAAGCTGGATGGACAACCC GTTTATTTACAAGAGCGTACGG GGTTCGAAAGTATAACCGGG AGTGACTTCAGGCAGT GCTTGTTTCGCGCATTCA	879-881 482-501 920 500 650 436	Uga 2005 Wylie <i>et al.</i> 1993 Abdullahi <i>et al.</i> 2001 Braiana <i>et al.</i> 1994 Sclavounos <i>et al.</i> 2006 Davino <i>et al.</i> 2005
FBNYV	ssDNA	TACAGCTGTCTTTGCTTCCT CGCGGAGTAATTAATCAAAT ACATCGAAGAGCAGTATCTGG ACGTTGTCGTTTTACCTTGG TTTCCCCTTCGCTAAGTTAA ACACCTCCTTGAACTGGTATAA CATTTCGGATGAACATCTGGG ATGAACTATCAAGCGATGGAG	666 487 931 1002	Kumari <i>et al.</i> , 2007 Shamloul <i>et al.</i> 1999 Shamloul <i>et al.</i> 1999 Shamloul <i>et al.</i> 1999
MDV	ssDNA	TCTCTATAAAAAGCTGTTA AAATGATTGTTGATTTTCATT TAATGTAATGAAGAACA CAGTCAATATACTACTAT CATAGATGGACCTTGGGAG GCGGTTTCTTTCTTCTGGC	608 997 1002	Kumari <i>et al.</i> 2007 Sano <i>et al.</i> 1998 Sano <i>et al.</i> 1998
PEBV	ssRNA	GGACCCTAATAGGAGGTGCC CATTACAAACAGTTAAATGAACACCC	886	Vellios <i>et al.</i> 2002
PEMV	ssRNA	GAGGGTGCCACCACGACTAC TGAAAATTAGATAAGGAAAACCCAAG	114	Skaf <i>et al.</i> 2000
PSbMV	ssRNA	GATTTCTTCGTTGTTTGT CTTGAGTGTGGCGTGGTT GCTCTAGACTCGAGGGGAARTCRAAAGCTAAAAC GTCCTAGAGCTTGCACATWGGATTGTA TACATCTAGATTACATGGCTCTCATTCCGAGAAG CAAACGCGTGACGAAACCAAGGATGATGAAAG TACATCTAGATTACATGGCTCTCATTCCGAGAAG GGTTGCTCGAGGGTGTGAGACCAAGATGAAAG	494 654 888 958	Phan <i>et al.</i> 1997 Phan <i>et al.</i> 1997 Roberts <i>et al.</i> 2003 Roberts <i>et al.</i> 2003
SbDV	ssRNA	CTGCTTCTGGTGATTACACTGCCG CGCTTTCATTTAACGYCATCAAAGGG AGGCCAAGGCGGCTAAGAG AAGTTGCCTGGCTGCAGGAG GGAATATCACTTTCCGGCCGCTCT GGCATGATACCAGTGAAGACC GCGGTTAGCAATGTCGCAATAC CATAAGCGATGGAACCTGACGA	110 440 281 372	Phibbs <i>et al.</i> 2004 Kumari <i>et al.</i> 2006 Harrison <i>et al.</i> 2005 Wang <i>et al.</i> 2006

tomatic plants, known as roguing, is a phytosanitary control measure that is widely used to remove sources of virus infection from affected fields. When practiced 2-3 times early during the growing season, roguing of FBNYV-infected crops was effective in minimizing the incidence of primary infection foci inside small faba bean fields in Egypt (Mak-

kouk *et al.* 1998b). Overwintering or oversummering crops which could play the role of sources of infection should be avoided through spatial isolation. Such methods are more effective with non-persistent viruses than with persistent viruses. With non-persistent viruses, such as BYMV, a few hundred meters may suffice, or better 1000 m. Whereas,

Table 7 Faba bean genotypes reported to be resistant to virus diseases

Genotypes/Cultivars	Resistant to	Reference
2N138	BYMV	Ghad and Bernier 1984
BPL 5247 through to BPL 5255	BYMV	Makkouk and Kumari 1995b
BPL 5271 through to BPL 5285	BLRV	Makkouk <i>et al.</i> 2002
G-2/78	BYMV and PEMV	Schmidt <i>et al.</i> 1989
B-1/5	BYMV	Schmidt <i>et al.</i> 1989
B-1/33	PEMV	Schmidt <i>et al.</i> 1989
cvs. Fiord, Barkool, Icarus, Ascot	BYMV	McKirdy <i>et al.</i> 2000
cvs. Ascot, Fiord, Icarus	CMV	Latham <i>et al.</i> 2001
cvs. Fiord, Barkool, Icarus, Ascot	AIMV	Latham and Jones 2001b
Several genotypes	BBTMV	Mazyad <i>et al.</i> 1975

persistently transmitted viruses such as BLRV can be carried from lucerne fields over a long distance, making it difficult to avoid the source of infection.

Selection and breeding for virus resistance

Host resistance, when available is the most acceptable component of virus control because it is environment-friendly, practical and economically feasible for resource-poor farmers. Several workers have identified faba bean lines resistant to viruses (**Table 7**). Resistance to BYMV in faba bean was found to be controlled by two recessive complementary genes, *bym-1* and *bym-2* (Rohloff and Stupnagel 1984; Schmidt *et al.* 1985). Combined resistance to BYMV, *Clover yellow vein virus* (genus *Potyvirus*, family *Potyviridae*) and *Aphis fabae* (Schmidt *et al.* 1986) and to BYMV and PEMV (Schmidt *et al.* 1989) was identified in faba bean. However, host resistance for a number of economically important viral diseases of faba bean persistently transmitted by insects (e.g. FBNYV) has not yet been identified, and control of such diseases is dependent on the availability of other control options. With advances in the development of regeneration systems of faba bean, there is good potential for producing transgenic faba bean through genetic engineering to reduce losses due to luteoviruses, FBNYV or CpCDV. At present, a number of legumes are being transformed with viral genes (coat protein, replicase, etc.) by different groups to produce virus-resistant legumes. There are no cultivars available in the market for immediate use by farmers, but genetic transformation could be a useful alternative, especially where resistance genes are not found in the existing genotypes of legume crops, including faba bean.

Methods directed towards avoidance of vectors or reducing their incidence

Cultural practices

Cultural practices, such as planting date, high seeding rate and narrow row spacing, use of early maturing cultivars, and planting of with borders non hosts have proved to be effective in reducing virus incidence in faba bean crops. Manipulation of planting date to avoid exposing young plants to peak vector populations at the most vulnerable early growth stage is a standard virus control measure that is widely recommended for use with crop legumes (Thresh 2003). For example, in Syria and Egypt, faba bean crops planted early in September are often severely attacked, leading to 100% FBNYV infection (Makkouk *et al.* 1998b). In such circumstances, farmers plough the crop under and re-plant with another crop. Delaying sowing until October or November often resulted in lower virus infection, and consequently reduced crop losses due to fewer viruliferous aphid vectors arriving from neighboring virus sources. The exact date of sowing in any region, however, should be based on a forecasting system, which is dependent on monitoring the population of viruliferous aphids migrating from sources of infection to the faba bean crop.

Virus vector control (chemical control)

Application of insecticides helps to decrease the spread of some faba bean viruses vectored by insects. However, it is often ineffective because success depends on the mechanism of transmission of the virus and the mode of action of the pesticide selected. In general, success in reducing virus spread by chemical control of vectors is considerably greater with persistently than with non-persistently transmitted viruses. This is mostly because incoming viruliferous vectors carrying non-persistently transmitted viruses tend not to be killed fast enough to prevent probing and subsequent virus inoculation to sprayed plants. In a three-year trial covering large areas of faba bean in East Germany (DDR), insecticide application reduced the average infection by BYMV, PEMV and BLRV by 63, 72 and 71%, respectively (Schmidt *et al.* 1977). Oil sprays can be used instead, but are rarely cost effective because of the repeated applications required. The most effective types of insecticides for control of non-persistently transmitted aphid-borne viruses are the newer generation of synthetic pyrethroids (Loebenstein and Raccach 1980), because of their rapid knockdown and greater anti-feedant activity. Field experiments at ICARDA showed that the use of a systemic seed treatment insecticide Imidacloprid (Gaucho[®]) at a rate of 0.5-2.8 g a.i./kg seed gave significant protection of faba bean plots (2 x 1.5 m plots and a plant density of 25 seeds/m²), against FBNYV and BLRV infection, which lasted for two months after sowing. Incidence of FBNYV was reduced from 28% in untreated plots to 2% and 1% in plots treated with 1.4 and 2.8 g a.i./kg of seed, respectively, and the yield loss from 37% in untreated plots to 0% in plots treated either with 1.4 or 2.8 g a.i./kg of seed (Makkouk and Kumari 2001).

Breeding for vector resistance

Genetic resistance to the vector was advocated some decades ago, but not much progress has been made in this area. The availability of cheap and effective chemicals in the 1950s and 1960s reduced interest in investigations of vector resistance in plants. However, the development of insect resistance to pesticides and the public awareness of environmental hazards resulting from their heavy use renewed the interest in breeding for insect resistance in plants (Jones 1987; Ajirilo *et al.* 2006). Resistance of faba bean to the aphids *A. craccivora*, *A. fabae* and *Acyrtosiphon pisum* has been reported (Clement *et al.* 1994). In Egypt, over 1000 lines were screened for aphid resistance, and 36 were classified as resistant. The faba bean line BPL 23 was resistant to both *A. craccivora* and *A. fabae* (Bond *et al.* 1994). Whether or not the use of such cultivars could lead to reduced spread of faba bean viruses in the field awaits further evaluation.

Integrated approach

Each of the control measures mentioned provides only partial control, but combining genetic resistance, cultural practices, and chemical sprays is expected to lead to improved disease control. The use of host resistance, whether obtained by classical breeding or genetic engineering, and one or two well-timed sprays coupled with optimal planting date and early roguing of virus-infected plants could offer reasonable and economic control, and stabilize faba bean production. However, selecting the best mix of measures for each virus-crop combination and production system requires knowledge of the epidemiology of the causal virus, and the mode of action and effectiveness of each individual control measure. Each strategy needs to be affordable by farmers and fulfill the requirements of being environmentally and socially responsible. It must also be compatible with control measures already in use against other pests and pathogens. An example of an integrated approach being widely used for practical virus control in a faba bean crop is the one for FBNYV in Egypt. This strategy combines planting late in

the growing season, use of high seeding rate, application of one or two systemic insecticide sprays (well timed during the early stages of the crop development), and rouging of infected plants early in the growing season. Its use has led to significant reduction of FBNYV infection and more profitable faba bean production in Egypt (Makkouk *et al.* 2003c).

CONCLUDING REMARKS

The most serious viruses which affect faba bean production are FBNYV and luteoviruses (e.g. BLRV). More work is needed on the epidemiology of these viruses in faba bean to provide information which will assist with the development of improved control strategies. There is also need to determine which aphids species are important in spreading faba bean viruses and understand the conditions which favor colonization of faba bean by aphids. In addition, other aphid species may be important in virus transmission without actually colonizing faba bean, which many occur during probing activity as they migrate over the crop (e.g. cereal aphids such as *Rhopalosiphum padi*). Similarly, aphids colonizing weeds within faba bean crops could also act as vectors.

Many countries in Africa and Asia do not have well-equipped virus research laboratories which permit workers to effectively implement plant quarantine controls. Therefore provision of diagnostic tools and training is essential to expedite progress in many countries.

Controlling FBNYV and luteoviruses which are well adapted to a faba bean crop is not an easy task. These viruses are disseminated by aphids and most currently grown cultivars are susceptible. Application of insecticides helps to reduce spread of these viruses, but this practice is only effective when applied at regular intervals, a process which increases the cost of production. In addition, environmental concerns impose limitations on pesticide use. Under these circumstances genetic resistance seems to be the most appropriate approach even though complete immunity has not been achieved. In the long term, emphasis should focus on the use of virus-resistant faba bean cultivars. It will be necessary for plant virologists to cooperate closely with faba bean breeders to achieve this goal. In addition, faba bean lines, with useful levels of resistance to virus infection by aphids or with decreased rate of seed transmission (e.g. BBSV, BYMV) can be used in breeding programs. In the future, transgenic faba bean expressing viral genes which can inhibit virus replication may form an integral part of an environmentally safe viral disease management package for faba bean.

Many scientists in developed and developing nations continue to work on the molecular characterization of virus strains, although the outcome of this research has not significantly influenced genetic improvement strategies. The development of molecular markers for virus resistance genes in faba bean, on the other hand, is expected to expedite the selection of virus-resistant faba bean genotypes.

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