**Rust Disease of Pea: A Review**

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**Abstract**

Pulses being important source of protein are essential adjunct to predominantly cereal based diet of large Indian population. Among all major pulses grown in India, pea (Pisum sativum L.) is considered as one of the important pulse crop. Pea diseases are major constraints to pea production in the developing countries. These diseases affects the crop both quantitatively (yield) as well as qualitatively (seed quality). Among these, the rust of pea caused by *Uromyces viciae–fabae* (Pers.) J. Schrot is considered as most important under warm and humid conditions. This review explains the geographical distribution, biology, epidemiology of pea rust pathogen and finally the different management aspects of rust disease of pea, such as the alteration in date of sowing, use of resistant cultivars, role of biotic and abiotic elicitors in induction of host plant resistance and lastly chemical control measures which cannot be avoided and must be taken into consideration up to environmentally safe level.

**Keywords**

Rust Disease, Pea, *Pisum sativum*

**Introduction**

A large proportion of Indian population is vegetarian and pulses are the main source of protein for them. The protein content in pulses is about 18-25 per cent which makes pulse one of the cheapest source of protein for human consumption. Pulses are the member of the family leguminosae, capable of utilizing *Rhizobium* bacterium in their root nodules, thus fixing atmospheric nitrogen and helps in improving soil fertility. Pulses leave behind reasonable quantity of nitrogen in the soil and add up to 30 kg N/ha to it. Pulses are also suitable for various crop rotations under rainfed conditions and they play vital role in sustainable agriculture in our country. In crop rotation, it helps in improvement of soil fertility and yield of succeeding crops (Rana and Sharma, 1993).

India is the largest producer, consumer and importer of pulses in the world. Pulses are grown about 24-26 million hectares of area producing 17-19 million tonnes of pulses annually in India which accounts for over one
third of the total world area and over 20 per cent of total world pulse production. Per capita production and availability of pulses in the country has observed quick decline. Per capita net pulse availability has declined from around 60 grams per day in the 1950s to 40 grams in the 1980s and further to around 35 grams per day in 2000s. However, in the last four years, there has been significant increase in consumption averaging around 50 grams due to higher production, because of National Food Security Mission (NFSM), with major emphasis on pulses and their imports, mostly of dry peas from Canada and Australia (IIPR, 2014).

Major pulses grown in India include chickpea or bengal gram (Cicer arietinum), pigeonpea or red gram (Cajanus cajan), lentil (Lens culinaris), urdbean or black gram (Vigna mungo), mungbean or green gram (Vigna radiata), lablab bean (Lablab purpureus), moth bean (Vigna aconitifolia), horse gram (Dolichos uniflorus), grass pea or khesari (Lathyrus sativus), cowpea (Vigna unguiculata), and broad bean or faba bean (Vicia faba).

Pea (Pisum sativum L.), the famous plant in which G.J. Mendel worked out Mendel Laws and Genetic Principles, is a noble and aristocratic vegetable. The crop is cultivated for its tender and immature pods for use as vegetable and mature dry pods for use as a pulse. In both cases, seeds are separated and used as vegetable or pulse. Tender seeds are also used in soups. Canned, frozen and dehydrated peas are very common for use during off-season. Like any other legume crop, pea is an integral component of sustainable agriculture due to its soil enriching and conditioning properties (Singh, 1984). Based on genetic diversity Vavilov (1926) listed different centre of origin for pea comprising Central Asia, the Near East, Abyssinia and the Mediterranean. Cultivated garden pea is not seen in wild state and it might have been originated from wild field pea or other related species.

During 2012-13, Pea (Pisum sativum L.) occupies an area of 0.76 million hectares with a production 0.84 million tonnes and productivity of 1100 kg/ha in our country (NCAER, 2014). Pea is a high quality protein rich pulse and vegetable crop. Dry pea generally contains 23 per cent protein, 48 per cent starch, eight per cent sugar, four per cent lipid, seven per cent crude fibre and three per cent ash (Duke and Ayensu, 1985).

Pea is affected by a number of fungal (rust, powdery mildew, downy mildew, root rot, alternaria blight, aschochyta blight, wilt, anthracnose, cercosphora leaf spot, damping off, seedling rot etc.), bacterial (bacterial blight and brown spot), nematode (cyst nematode, lesion nematode and root-knot nematode) and viral diseases (cucumber mosaic virus, pea early browning virus, pea enation mosaic, pea mosaic, pea seed borne mosaic, pea streak and pea stunt). These diseases, under the right conditions, can significantly decrease both yield and quality. Among these, the rust of pea caused by Uromyces viciae–fabae (Pers.) J. Schrot (syn. Uromyces fabae (Pers.) de Bary) is considered the most important under warm and humid conditions (Chand et al., 2004).

**Symptomatology**

The first symptoms appear with the development of aecia. The yellow aecia appear first on the undersurface of the leaves, stems and petioles. The formation of aecial stage is preceded by a slight yellowing which gradually turns brown. The uredopustules are powdery light brown in appearance. All the four stages develop on every green part of the host including the pods. The teleutopustules occur in the same sources as the uredia and
develop from the same mycelium (Singh, 1973). Thatcher (1939) studied the effect of *U. fabae* on pea. He pointed out that fungus increased the permeability of the host cell by secreting some metabolites, which ultimately prove fatal. Hahn *et. al.* (1977) also reported that a putative amino acid transporter was specifically expressed in haustoria of the rust fungus *Uromyces-fabae*, which may be the cause of increased permeability of the host cell. Staples (1968) and Haung and Staples (1982) proposed the synthesis of proteins during differentiation of the bean rust fungus. Staples and Stahmann (1964) have also reported the change in protein and several enzymes in susceptible bean leaves after rust infection.

**Biology of *Uromyces fabae***

**Pathogen description**

Two species of *Uromyces* have been reported to cause rust of pea. One of them *U. pisi* (Persoon) de Bary, has been reported from several European countries (Deutelmoser, 1926; Mayer, 1947; Palter and Stetbiner, 1957). It is a heteroecious species having its aecial stage in *Euphorbia cyparissias* and rarely occurs in India. In India, another species *U. fabae* (Pers.) de Bary has been found to cause pea rust (Butler, 1918; Prasada *et al.*, 1948; Kapooria *et al.*, 1966).

*Uromyces fabae* (*Uromyces viciae- fabae*) the rust of pea was first reported by Persoon in 1801. Later de Bary (1862) changed the genus and renamed it as *Uromyces fabae* (Pers.) de Bary. The pathogen *U. fabae* is described as autecious rust with aeciospores, urediospores and teliospores found on the same host plant (Arthur and Cummins, 1962; Gaumann, 1998). Gaumann proposed that the fungus be classified into nine formae speciales each with host range limited to two or three species. The isolates of *Uromyces viciae-fabae* share so many hosts in common that it is impossible to classify them into formae speciales (Conner and Bernier, 1982). Based on the distinctive shape and dimension of substomatal vesicle, *U. viciae-fabae* has been described as a species complex (Emeran *et al.*, 2005). The peridium of aecium in *U. fabae* is short, whitish and cup shaped. The aeciospores are round to angular or elliptical, yellow in colour with fine warts. They measure 14-22 microns in diameter. The urediospores are round to ovate light brown echinulate with 3-4 germ pores and measure 20-30 × 18-26 microns. The teliospores are subglobose to 2 0 ovate, thick walled, with flattened apex, smooth, single celled, pedecellate and measure 25-38 × 18-27 microns in size (Singh, 1973). Prasada and Verma (1948) working with *Uromyces fabae* from lentil found that infection with aeciospores at lower temperatures (17-26°C) results in the formation of secondary aecia, while at 25°C the infection causes development of uredia. No infection by aeciospores occurs at 30°C. Optimum temperature for germination of urediospores is 16-22°C, while urediospores germination does not occur at 28-29°C. The teleutospores of lentil rust can germinate at 12-22°C. The fungus completes its life cycle on peas and is further endowed with survival potential in the telial stage (Singh, 1973).

**Taxonomy and Nomenclature**

*Uromyces fabae* is an autococious and heterothallic fungus forming all the four type of spores *viz.*, pycniaospores/spermatiospores, aeciospores, urediospores and teliospores on pea only. Pycnia are small, flask shaped and produced as yellowish flecks on upper surface of leaves with a common nector drop at mouth. As the haploid pustules remained unfertilized the formation of pycnia, with separate scanty nectar drops on the lower surface of the leaves was observed (Prasada
and Singh, 1975). Rust of pea is caused by fungus *Uromyces viciae–fabae* (Pers.) J. Schrot. belongs to the phylum *Basidiomycota*, class *Urediniomycetes*, order *Uredinales* (rust fungi) and family *Pucciniaceae* (Alexopoulos et al., 1996). According to recent classification by Kirk et al., (2001, 2008) the systemic position of the *U. viciae–fabae* belongs to kingdom Fungi, phylum *Basidiomycota*, class *Pucciniomycetes*, order *Pucciniales* and family *Pucciniaceae*.

**Host range**

Prasada and Verma (1948) found that several species of *Vicia, Lathyrus, Pisum* and *Lentil* are susceptible to *U. fabae* in India and abroad. In India, species of *Vicia, Lathyrus*, and *Pisum* are described as host plant for *U. viciae–fabae* (Pers.) J. Schrot (Kapooria and Sinha, 1966). Bilgrami et al., (1979) reported the occurrence of this pathogen on various host species of pea, lentil and *lathyrus*. *Vicia faba* L., *V. biennes* L., *V. hirsuta* L., and *V. arborescis* L. were described as highly susceptible to *Uromyces fabae* and *Vicia sativa* and *Lathyrus aphaca* were found to be disease free. Conner and Bernier (1982) reported a total of 52 species of *Vicia faba* and 22 species of *Lathyrus* to be infected by *U. viciae–fabae* (Pers.) J. Schrot. Uppal (1993) has also reported that *U. fabae* infect several species of *Vicia, Lathyrus, Pisum* and *Lentil* in India and abroad.

**Geographical distribution**

Pea rust (*U. fabae*) is of worldwide occurrence and attacks number of host species belonging to different genera of the family *Leguminosae* in the Indo-Gangetic plains (Butler, 1918). There were reports of occurrence of *U. fabae* from most of the places of India including eastern India (Gupta, 1990; Chand et al., 1997), central India (Narsinghani et al., 1980), southern parts of India (Sokhi et al., 1974; Kumar et al., 1994) and from Himalayan region of Uttarakhand and Himachal Pradesh (Chauhan et al., 1991; Sharma, 1998). Survey of pea growing region of three districts of Bihar (Lal et al., 2007) and six district of Himachal Pradesh (Chauhan, 1988) state of India revealed that *U. fabae* was very serious in pea. Prasada and Verma, (1948) also reported the occurrence of *U. fabae* on lentil crop from Delhi. Roy (1949) in his list of fungi of Bengal recorded the prevalence of *U. fabae* on the leaves and stems of pea (*Pisum sativum*). Mitter and Tondon (1930); Pavgi and Upadhyay (1966) and Kapooria and Sinha (1966) reported the distribution of this pathogen in the regions of Uttar Pradesh, respectively. Baruah (1980) reported that rust infection on the pea plants is caused by both *U. fabae* and *U. pisi* of which *U. pisi* is of rare occurrence in India. Occurrence of *U. fabae* have been reported from Canada, Europe, Ethiopia, Australia and Iran in mild to severe forms on pea, lentil, alfalfa, broad bean and faba bean are also available (Conner and Bernier, 1982; Xue and Warkentin, 2002 and Sadravi et al., 2007). In the last few years, disease has been observed in almost epiphytotic form and could cause up to 20-100% losses in yield (Upadhyay et al.2015; Sharma, 1998).

**Life cycle**

*Uromyces fabae* is a macrocyclic rust fungus, it exhibits all five spore forms known for the *Uredinales*. It is autoecious, as all spores are produced by single host (Mendgen, 1997). After overwintering on residual plant material, diploid teliospores germinate in the spring with a metabasidium. After meiosis, the latter produces four haploid basidiospores with two different mating types. These spores after landing on a leaf of a host germinate and produce infection structures. Pycnia are produced which contain pycniospores. Pycniospore are exchanged between pycnia of
different mating types and after spermatization, dikaryotization occurs in aecial primordial. An aecium differentiates and dikaryotic aeciospores are produced. These aeciospores germinate and form infection structures from which uredia develops, which produce urediospores. Urediospore is the major asexual spore form of rust fungi produced in massive amount through repeated infection of host plants during the summer. Urediospores are dispersed aerially and can travel thousands of kilometers (Brown and Hovmoller, 2002).

Environmental factors affecting disease development

Decision to apply one or more fungicide spray will depend on the risk of rust epidemic in a particular year. Rust epidemic is determined by interaction of three important factors namely, susceptible host, virulent pathogen and most important i.e. favourable environment for a particular period of time. Therefore, it is necessary to know the correlation between different meteorological parameters and rust severity. Rust disease of pea caused by *Uromyces fabae* is very severe under warm and humid conditions in Tarai region. Prasada and Verma (1948) reported that relatively low temperatures, 17-22°C result in formation of secondary aecia while at 25°C development of uredia takes place. Infection and pustules formation was high at 20°C under greenhouse and laboratory conditions. It was observed that relationship between severity of pea rust and duration of leaf wetness at above 20°C temperature may be useful in predicting disease outbreak if initial inoculum is present (Chauhan and Singh 1995). Atmospheric temperature around 20°C maximum and 5°C minimum with high RH (60-70% mean weekly) and light shower or drizzle favour *Uromyces viciae-fabae* development and spread whereas temperature above 25°C and below 7-8°C along with rains disfavour rust spread (Mittal, 1997). Number of rainy days and rainfall during the crop season, play an important role in the spread of pea rust disease than any other weather parameters (Singh and Tripathi, 2004). Khare and Agrawal (1978) reported that high humidity, cloudy or drizzling weather with temperature of 20-22°C favours disease and those plants are more susceptible at flowering in lentil for *Uromyces viciae-fabae*. Hazarika *et al.*, (2000) demonstrated the effect of eight sowing dates on leaf spots and rust of groundnut in relation to weather factors during the crop season. They observed that, there was significant and positive correlation between the incidence of disease (leaf spot and rust disease) and weather factors i.e., rainfall, relative humidity and temperature. Negussie *et al.*, (2005) observed that at 20°C, dew period of at least three hours was required for minimum infection of lentil rust, whereas maximum infection occurred with a dew period of 24 hrs. Infection efficiency increased linearly as the duration of dew period increased from 0 to 24 hrs. The optimum germination of aeciospores, urediospores and teliospores was recorded at 20°C. Viability of aeciospores and urediospores of *U. viciae-fabae* (Pers.) de Bary decreased with increase in time, whereas, germination of teliospore after eight months of storage gave positive results (Joshi and Tripathi, 2012). They also found that age of plant had no direct relationship with rust appearance in lentil, while, 24 h leaf wetness after inoculation was found to be optimum for rust development. Singh *et al.*, (2012) found significant and positive correlation between rust severity and temperature. However, disease severity has a strong negative correlation with grain yield (kg/ha), rainfall and relative humidity. Similar observations were recorded by Bal and Kumar (2012). Upadhyay *et al.*, (2017) stated that rust disease was observed at a maximum temperature of 16.85 to 24.79°C, 8.09 to
12.27°C minimum temperature, 90.30 to 95.70 percent morning Relative Humidity (RH), 54.80 to 78.40 percent afternoons RH, 0.10 to 5.45 mm rainfall and wind velocity of 3.93 to 4.23 km/hr. Decision to spray fungicides will depend on the risk of rust epidemic. To help farmers in determining rust epidemic risk, there is need to work on developing forecast model for pea rust.

**Disease management strategies**

**Cultural practices viz., planting time, planting geometry, intercropping and row spacing**

Using principle of avoidance through alteration in date of sowing can be an effective way to disturb the interaction of three important factors namely host, pathogen and environment important for disease development and thus can be utilized as an effective cultural practice for the management of rust disease in field pea but the yield parameters should be taken into consideration. From past, many researchers have worked on these aspects which are mentioned here under:

Delayed in sowing *i.e.* after 15th October, increased the incidence of *Uromyces viciae-fabae* and decreased grain yield (Sangar and Singh, 1994). Similarly, Singh et al., (1996) reported that incidence of rust (*Uromyces viciae-fabae*) increased as sowing was delayed. In contrary to this, Bhardwaj and Sharma (1996) reported that plants from 15 October sowing were taller, produced the highest number of marketable pods and highest green pod yield (4.74 t/ha) with lowest percent disease index of rust (*Uromyces viciae-fabae*). Similar observation was observed by Rai & Gupta (2003) that rust intensity was found very high in late planted and closer spaced pea crop. In contrary to this, Singh Mittal (1997) observed that incidence of disease declined from the early to late sowing. Whereas, Tripathi and Rathi (2003) also studied on effects of different dates of sowing, inter-row spacing and intercropping on disease severity and grain yield of field pea. They reported that delayed sowing not only increased disease severity but also lower grain yield in plants having narrow spacing as compared to wider row spacing. They further emphasized that minimum disease severity was recorded in pea + mustard inter cropped plants followed by the pea + wheat, pea + linseed and pea + rajma. In oppose to Tripathi and Rathi (2003), Singh *et al.*, (2012) found least rust severity when pea was planted on October 15th during all the three crop seasons. The crop when sown lately *i.e.* sown on November 14, 29 and December 13th recorded highest severity of rust. Similarly, Singh *et al.*, (2014) studied the effects of cultural practices viz., planting time, planting geometry, intercropping and row direction on disease severity of field pea rust caused by *Uromyces viciae fabae* and grain yield. They found that late planting of pea has recorded the highest disease severity and minimum grain yield. They have also noticed that planting geometry *i.e.* row spacing has significant influence on disease severity *i.e.* wider row spacing showed less rust severity than close spacing. Similarly to other researchers, he found that minimum rust severity was recorded when field pea was intercropped with mustard. However, planting direction has not significantly influenced rust severity. Upadhyay *et al.*, (2018) studied the effect of alteration in date of sowing on rust severity and grain yield in field pea. Their investigation indicate that, early sown crop in 31st October, 7th November and 14th November face lower disease severity (8.67-17.50 percent) with low area under disease progress value (81-198.67) and produce good yield (690.90-775.39 kg/ha) and test weight (162.34-175.34 g) whereas crop sown in 21st November, 28th November, 5th December
and 12th December succumb to high disease severity (40-54.17 percent) showing high area under disease progress value (383.50-549.17) with low yield (429.06-581.95 kg/ha) and test weight (146.67-153.73 g).

Screening of germplasms for disease resistance

The use of host plant resistance is the best means of rust control (Bayaa and Erskine, 1998). Screening of field pea germplasms under field conditions for resistance to rust has been reported in India (Singh et al., 1995). Screening for rust severity indicated wide range of variations for rust resistance in the germplasm lines of pea and none of the genotypes tested were found to be free from infection (Narshinghani et al., 1980; Singh and Srivastava, 1985; Gupta, 1990; Kumar et al., 1994; Xue and Warkentin, 2002; Chand et al., 2004, Upadhyay et al., 2017). Rust severity is greatly influenced by the environment during infection initiation and disease development. This is the major bottleneck in screening and selection for rust resistance. Use of molecular markers would allow indirect selection for rust resistance independent of environmental effects (Rai et al., 2011). For the development of rust resistant varieties there is need for phenotypic screening as well as molecular screening of existing lines/germplasms/cultivars. Several researches that have been carried from past in these aspects are mentioned below:

Pal et al., (1980) screened a total of 292 accessions of pea (Pisum spp.) under field conditions for resistance to powdery mildew (Erysiphe polygoni) and rust (Uromyces fabae). Only three accessions--PJ207508, PJ222117, and EC109188 were resistant to rust. PJ207508 was resistant to both powdery mildew and rust disease. Likewise, Kumar et al., (1994) tested thirty tall genotypes of field pea against rust severity. Variety Pant P-8 had lowest pea rust cover under disease progress curve (AUDPC) value, growth rate (c) and apparent infection rate (r). However, in general KFP 106, DMR 11, HUP 8603, type 163 and KPMR 22 showed high level of slow resistance, being conditioned by a number of genes with small effects is more desirable. Similarly, total of 648 accessions of Vicia faba was screened for resistance to faba bean rust (Uromyces viciae-fabae) by Sillero et al., 2000. They identified two distinct types of resistance, both resulting in reduced disease severity (DS) and area under the disease progress curve (AUDPC), but differing in the expression of hypersensitivity i.e. one as incomplete non hypersensitive resistance and the other as incomplete resistance with late hypersensitivity. These two types of resistance were characterized by three macroscopic components of resistance: increased latent period (LP), decreased colony size (CS) and a relatively reduced infection frequency (IF), both on seedlings and on adult plants. Xue and Warkentin (2002) studied 93 field pea varieties to three isolates of U. viciae-fabae with symptoms (LAS) under control condition. Significant difference (P<0.5) was observed from pea varieties and rust isolates, and variety x isolate interaction. Similarly, three hundred and forty five accessions of pea of diverse origin, height, leaf types and disease reaction were screened for rust disease severity and area under disease progress curve (AUDPC) by Chand et al., 2006. Of the 345 accessions, forty-four genotypes were evaluated for disease intensity, which was converted into AUDPC, number of pustules/leaf and pustule size. They found fast rusting genotypes exhibiting lower AUDPC, accompanied with increased seed yield and seed weight when grown under the protected condition, as compared to those raised under the unprotected condition whereas the genotypes Pant P 11, FC 1, HUDP 16, JPBB 3 and HUP 14 appeared as slow rusting genotypes. Kushwaha et al.,
conducted field and polyhouse studies to determine the appropriate time for the assessment of slow rusting in pea to *Uromyces fabae* (Pers de Bary). The critical time occurred when disease severity on the susceptible (check) genotype HUVP 1 had crossed 90% but was <20% on the resistant (check) genotype FC 1. The disease assessment at critical time revealed precise differentiation between resistance and susceptible reactions in the F$_2$ generation of the cross HUVP 1/FC 1. Reduction in 100-seed weight of inoculated F3 progeny rows showed high correlation with rust severity at the critical time and AUDPC based on two assessments in the field. Significant reduction in 100-seed weight was observed only for susceptible lines whereas; reductions in moderately resistant and resistant lines were not significant. Mishra *et al.*, (2009) evaluated 107 genotypes of field pea against rust (*Uromyces viciae-fabae*), out of which genotypes P 9-77, P 2432; P2572 and P 2930 were found resistant, whereas 27 exhibited moderate reaction. Likewise, total of 2759 pea accessions was screened for resistance against *Uromyces pisi* (Pers.) Wint by Barilli *et al.*, (2009). All accessions displayed a compatible interaction (high infection type) both in adult plants under field conditions and in seedlings under growth chamber conditions, but with varying levels of disease reduction. The identified resistance was based on reduction of disease severity with no associated host cell necrosis, which fits the definition of Partial Resistance. No complete resistance or incomplete resistance based on hypersensitivity was observed. In present era, molecular markers associated with pea rust resistance would be useful in marker assisted selection (MAS). Utility of molecular markers associated with the pea rust resistance were evaluated in 30 diverse pea genotypes using four SSR markers (AA446 and AA505 flanking the major QTL Qruf; AD146 and AA416 flanking the minor QTL, Qruf1) by Singh *et al.*, (2015). QTL, Qruf flanking markers were able to identify all the resistant genotypes when used together, except Pant P 31. While, SSR markers AD146 and AA416 flanking the minor QTL, Qruf1 were able to identify all the pea resistant genotypes used for validation, except for HUDP-11 by AD146 and Pant P 31 by AA416. Similarly, SSR markers AA446 and AA505 were able to identify all the susceptible pea genotypes, except IPFD 99–13, HFP 9415 and S-143. SSR markers AD146 and AA416 were together able to identify all the pea susceptible genotypes used for validation, except KPMR 526, KPMR632 and IPFD 99–13. On the basis of marker allele analysis, they concluded that SSR markers (AA446, AA505, AD146 and AA416) can be used in MAS of pea rust resistance. Rai *et al.*, (2011) suggested that the Ruf gene proposed by Vijayalakshmi *et al.*, (2005) be now redesigned as Qruf to signify the quantitative nature of its action and detected another minor quantitative trait loci (QTL) (named Qruf1). Both QTLs were located on LGVII. Qruf was flanked by SSR markers, AA505 and AA446 (10.8 cM), explaining 22.2–42.4% and 23.5–58.8% of the total phenotypic variation for IF and AUDPC, respectively. Qruf was consistently identified across four environments. Therefore, the SSR markers flanking Qruf would be useful for marker-assisted selection for *U. viciae-fabae* resistance. The minor QTL was environment-specific, and it was detected only in the polyhouse (logarithm (base 10) of odds values 4.2 and 4.8). It was flanked by SSR markers, AD146 and AA416 (7.3 cM), and explained 11.2–12.4% of the total phenotypic variation. Similarly, Upadhyay *et al.*, (2017) screened 46 numbers of total germplasms, out of which two germplasms Pant P 244 and Pant P 42 showed moderate resistant, 13 germplasms were moderately susceptible, 29 germplasms were found susceptible and two germplasms HFP-4 and HUDP 1 were found highly
susceptible. Moderately resistant germplasm showed low AUDPC value (160.83-188.33) with slow infection rate (0.054-0.062). Pustule appeared on these genotypes were small (1.5-1.7mm) as compare to other susceptible genotypes whereas moderately susceptible genotypes scored AUDPC value from 175.83-437.50 with infection rate of 0.051-0.095. Size of the pustules showed high variation of 1.3-4.4mm. Genotypes with susceptible reaction showed AUDPC value of 292.50-797.50. Infection rate was ranged from 0.055-0.113 with pustule size of 2.9-4.6mm. Those genotypes which fall under highly susceptible reaction (HFP-4 and HUVP-1) scored highest AUDPC value of 1078.33-1223.33 with 0.064-0.075 infection rate. They showed largest pustule size of 4.2-4.6mm. Upadhyay and co-workers (2017) also did molecular screening of 32 number of phenotypically selected genotypes using four SSR markers - AA446 and AA505 flanking the major QTL Qruf; AD146 and AA416 flanking the minor QTL, Qruf1 associated with pea rust resistance. They have also concluded that SSR markers (AA446, AA505, AD146 and AA416) if used together, can be effective in marker assisted selection (MAS) of pea rust resistance.

Molecular markers linked to resistance genes could helps in assisting the selection of rust resistant segregants and thus improve efficiency of breeding. So far, works on molecular mapping of resistance against U. pisi are inadequate and more strong markers are required.

Breeding works for rust resistance is slow due to still inadequate genomic resources and because of the limited knowledge of the biology of various rust pathogens, their existence of races and their distribution. Therefore, to provide significant input in this area, it is important to improve the existing knowledge of biology of the causal agents as well as of the plant, resistance breeding will be efficiently accelerated.

Induction of host defence through biotic and abiotic elicitors

Plants can be induced for a more rapid or extra intense mobilization of defence responses leading to improved resistance to biotic or abiotic stresses (Beckers and Conrath, 2007). Many factors such as prior pathogen attack (biotic) and various chemical and environmental stimuli (abiotic) may act on plants to induce systemic acquired resistance (SAR) to subsequent pathogen attack (Kauss et al., 1992; Kessmann et al., 1994; Dann and Deverall, 1995; Barilli et al., 2010). SAR has been reported to be effective against a broad spectrum of pathogens including viruses, fungi, bacteria, nematodes and parasitic weeds (Beckers and Conrath, 2007). Induction of systemic resistance is associated with gene induction, the activation of a wide range of resistance mechanisms and the production of a wide range of defence compounds. It is race non-specific and is often effective against a broad spectrum of pathogenic agents (Kuc, 1995; Walters and Fountaine, 2009). Thus, study on induction of host defence through biotic and abiotic elicitors can be considered as one of the effective sustainable approaches in disease management.

Walters and Murray (1992) observed that inoculation of the lowest two leaves of broad bean (Viciae fabae) with urediospores of the rust fungus (U. viciae fabae), caused the upper leaves to become resistant to challenge inoculation with the same pathogen one, three, six and nine days later. The resistance was observed as diminished infected areas on the leaves and fewer uredia per standard area for up to 29 days from challenge inoculation. The resistance was very high when the difference between treatment and challenge
inoculation was one day but had disappeared when 12 days separated the two. In further experiments, Walters and Murray (1992) reported that treatment of the first two leaves with either 10mM tri-potassium phosphate or 5mM ethylene diamine tetraacetic acid (EDTA) also induced development of resistance in upper leaves to challenge inoculation and the induced resistance was observed for 21 days after challenge inoculation. Rust infection was reduced by 15.0 and 34.0 per cent, if the upper leaves were inoculated 24 hrs after potassium phosphate or EDTA treatment respectively while, there was 77.0 per cent reduction in infection if the interval between treatment and inoculation was increased to 12 days. Dann and Deverall (1995) reported that inoculation of unifoliate leaves of nine days old green bean (Phaseolus vulgaris) with spore suspension of Colletotrichum lindemuthianum (10^4 conidia/ml), causing local lesions, or spraying with 2-6-dichloroisonicotinic acid (20µg/ml) induces development of resistance in the upper leaves against challenge inoculation of U. appendiculatus afterwards. Rauscher et al., (1999) reported that treatment of broad bean leaves with salicylic acid or 2, 6, dichloroisonicotinic acid induces resistance against the rust fungus Uromyces viciae-fabae resulting in reduced rust pustules density. Inhibition of the rust infection hyphae in acquired resistance broad bean plants was found mainly due to antifungal activity of PR- 1 protein synthesized in plants in response to salicylic acid or dichloroisonicotinic acid application. Dann and Deverall (2000) observed that, when inoculation of first expanded leaves of pea seedlings with an avirulent strain of Pseudomonas syringae pv. pisi or treatment with sprays of benzothiadiazole (20 or 100 mg a.i/ml), decreased susceptibility of subsequent leaves 7 or 14 days later to challenge inoculation with Uromyces viciae-fabae causing pea rust was found. Effective treatment enhanced the activity of enzymes β-1,3glucanase and chitinases in untreated upper leaves. Similarly, Katoch et al., (2005) observed that when pea (Pisum sativum L.) plants treated with different concentrations of salicylic acid and 4-aminobutyric acid increased activities of phenol metabolizing enzymes implicated in the defense of plants. The enzymes peroxidase, polyphenol oxidase, phenylalanine ammonia-lyase and superoxide dismutase responded to treatment with variation in their activities. Phenolic content also varied following treatment with the inducers. Similarly, Systemic acquired resistance (SAR) induction on plant-pathogen interaction was developed using both biotic (U. pisi and U. appendiculatus) and abiotic (salicylic acid (SA), benzo-(1,2,3)-thiadiazole-7-carbothionic acid (BTH) and DL-β-aminobutyric acid (BABA)) inducers (Barilli et al., 2010b). Results obtained showed a significant reduction of infection levels locally and systemically with BTH and BABA foliar treatments, whereas neither biotic inducers nor SA had any significant effect hampering the rust development. Barilli et al., (2010a) found that Benzothiadiazole (BTH) and DL-β-aminobutyric acid (BABA) induced systemic resistance in susceptible and resistant pea genotypes against Uromyces pisi. Resistance was characterized by reduced infection frequency mainly due to decreases in appressorium formation, stomatal penetration, growth of infection hyphae and haustorium formation. Changes in β-1,3-glucanase, chitinase, phenylalanine ammonia-lyase and peroxidise activities and in total phenolics content, demonstrate that U. pisi resistance is induced by BTH and BABA treatments at early and late stages of the fungal infection process, but that the chemicals operate via different mechanisms.

Exogenous applications of salicylic acid (SA) and benzothiadiazole (BTH) solutions have been used in faba bean to induce systemic
acquired resistance (SAR) to rust (*Uromyces viciae-fabae*), ascochyta blight (*Ascochyta fabae*) and broomrape (*Orobanche crenata*). Both SA and BTH solutions were effective inducing SAR to *U. viciae-fabae* and *A. fabae* on susceptible accessions under controlled conditions, although SA was less effective than BTH for *A. fabae*. BTH treatments reduced the infection of all pathogens studied under field conditions in susceptible accessions, and rust infection was also reduced by SA applications. Moderately resistant accessions became immune to ascochyta blight with BTH treatment, and showed a lower degree of infection to rust after SA or BTH treatments. No effect was observed in the highly resistant accessions (Sillero et al., 2012). Barilli et al., (2012) studied systemic acquired resistance (SAR) to *Uromyces pisi* in pea by using a proteomic approach. Two-dimensional electrophoresis (2-DE) was used in order to compare the leaf proteome of two pea genotypes displaying different phenotypes (susceptible and partial resistance to the fungus), and in response to parasite infection under the effect of two inducers of SAR, BTH and BABA. Multivariate statistical analysis identified 126 differential protein spots under the experimental conditions (genotypes/treatments). All of these 126 protein spots were subjected to MALDI-TOF/TOF mass spectrometry to deduce their possible functions. A total of 50 proteins were identified using a combination of peptide mass finger printing (PMF) and MSMS fragmentation. Most of the identified proteins corresponded to enzymes belonging to photosynthesis, metabolism, biosynthesis, binding and defence response, whose behavior pattern was different in relation to susceptibility/ resistance of the genotypes studied and to the BTH/BABA induction to pathogen response. Results obtained in their work suggested that plants could reduce their photosynthesis and other energy metabolism and enhance the production of defence-related proteins to cope the stress. On the other side, they postulated that resistance induced by the chemicals operates via different mechanisms: BABA inducer could act via phenolic biosynthesis pathway, whereas resistance provided by BTH inducer seems to be mediated by defence and stress-related proteins. Recently, Upadhyay et al., (2016) studied total of fifteen elicitors tested alone/or in combination for induction of defense related enzymes in pea against *U. viciae-fabae* (Pers.) J. Schrot. They observed significant induction of total phenols, peroxidase, polyphenol oxidase and phenylalanine ammonia lyase in all the treatment as compare to control. Salicylic acid, *Pseudomonas fluorescens*, salicylic acid + *Pseudomonas fluorescens* were found most effective in induction of total phenols and peroxidase at 72 hrs after spray of elicitors. Polyphenol oxidase induction was found significantly high in oxalic acid, *Pseudomonas fluorescens* + *Trichoderma harzianum* and chitosan + *Pseudomonas fluorescens* at 72 hrs after spray of elicitors. Among all the treatments, maximum induction of Phenylalanine ammonia lyase activity was found in oxalic acid, *Trichoderma harzianum* + *Pseudomonas fluorescens* and isonicotinic acid + *Trichoderma harzianum* after 48hrs of spray of elicitors. Effect of different elicitors on percent disease index (PDI) 20 days after inoculation with uredospores of *U. Viciae-fabae* showed least PDI in salicylic acid, *Trichoderma harzianum* + *Pseudomonas fluorescens* and chitosan + *Pseudomonas fluorescens* treated plants.

**Chemical control**

Breeding for rust resistance is considered the most adequate control strategy, but only moderate levels of resistance are available in commercial cultivars. This reinforces the need
to integrate several control strategies including chemical control. Therefore, search for the best fungicides in management of pea rust under field condition should be carried out in order to find out its effectiveness in integrated disease management (IDM) strategy. Many experiments were previously conducted to find out the efficacy of fungicides which are mentioned here under:

Several inorganic sulphur preparations are reported to give effective control of *U. fabae* (El-Healy, 1939; Zaumeyer, 1946; Jacks, 1954; Accantino, 1964). Similarly, organic sulphur fungicides like ferbam, ziram, thiram and zineb have been reported to give good control of *U. fabae* (Jacks, 1954; Jacks and Webb, 1956; Accantino, 1964). Hiremath and Pavgi (1971) obtained complete inhibition of aeciospores germination of *U. fabae* with aureofungin to 20µg/ml and recommended early application of higher aureofungin concentrations to control rust disease. Sugha et al., in 1994 reported sensitivities of aeciospores and urediospores to benzimidazole and triazole fungicides. They concluded that benomyl, carbendazim, thiobendazol and thiophanate methyl have very good potential for suppressing the early establishment of pea due to aeciospores, whereas benomyl, flutriafol and myclobutanil should be effective in suppressing the late infection due to urediospores. They found hexaconazole (0.10%) and difenaconazole (0.01%) were best against rust and increase yield. Similarly, Singh and Tripathi (2004) tested ten different fungicides against rust of pea in a field experiment and they found that two to three sprays of Baycor 0.1 % at 15 days interval was most effective in reducing the disease severity and resulted in appreciable increase in grain yield. Likewise, Sugha et al., (2008) evaluate the efficacy of 22 fungicides against pea rust during crop (rabi) at farmer’s field. They observed that three fortnightly foliar sprays, starting with the appearance of disease, individually of bayleton, score, tebuconazole (folicur and tebuconazole) and hexaconazole (contaf and sitara) among systemic and at 10 days intervals of antracol and microsul share among non-systemic fungicides proved effective for combating the disease and in ameliorating the crop yield significantly. Khan et al., (2009) conducted research to find out the response of five different pea cultivars and efficacy of three different fungicides against *Uromyces pisi* (Pers) de Barry under field conditions for the control of pea rust. He found that all fungicides caused reduction in disease severity. The lowest disease attack was recorded in plants treated with Mancozeb followed by those with Bayleton showing a subsequent increase in yield. The effectiveness of eleven foliar-applied fungicides on faba bean rust (*Uromyces viciae-fabae* (Pers.) J. Schröt.) and on the
seed yield of faba bean (Vicia faba L.) were studied in growth chambers and in the field by (Emeran et al., 2011). Fungicides were tested at recommended and reduced rates. All the fungicides tested provided very effective preventive control in their growth chamber studies. Triazoles (difenoconazol, epoxiconazol, tebuconazol) and their mixtures with benzimidazoles (carbendazim-flutriafol and carbendazim-flusilazole) provided the most effective curative effect, even at 25% of recommended concentrations which were followed by dithiocarbamates, copper dithiocarbamate mixture, carboxamide and chlorothalonil. Triazoles, benzimidazole-triazole mixtures and carboxamide maintained their effect until 15 days after fungicide application. Under field conditions, rust infection caused 22-26% yield reduction.

All fungicides except mancozeb caused a significant decrease in disease severity under field conditions, but only treatments with triazoles and benzimidazole-triazole mixtures provided significant yield increases (22.7-15.6%) when applied twice. Three applications of oxycarbosin or copper-mancozeb were needed to provide a significant yield increase. Dithiocarbamates (thiram, maneb or mancozeb) or chlorothalonil reduced rust severity but did not provide a significant yield increase (Emeran et al., 2011).

Similarly, Singh (2012) conducted experiment to control the pea rust disease with foliar sprays of new strobilurin fungicides viz. Amistar and triazoles viz. Score (difenoconazole) and Tilt (propiconazole) in different combinations. He found a very significant disease control of 81.8% was obtained when two sprays of Score @0.1% were given at 20 days interval followed by Score @0.05% (66.5%). The minimum disease severity of 15.05% was observed with Score @0.1% over control plots. Basandrai et al., (2013) evaluated some commercially available fungicides for pea rust management. They found that three foliar sprays of tebuconazole 250 EFW resulted in the least mean rust disease severity (6.2%) followed by propiconazole 25 EC (23.7%) and hexaconazole 5 EC (26.1%) compared with 52.2% in no spray check. Similarly, Upadhyay et al., (2018) tested the efficacy of total of sixteen chemicals fungicides alone and/or in combination against rust disease of pea.

Their study revealed that all the sixteen fungicides were found effective for the management of disease as compare to control (water spray). However tebuconazole, carbendazim + tebuconazole, Mancozeb + tebuconazole, carbendazim + flusilazole, penflufen + trifloxystrobin were found equally and very effective among all. These fungicides showed considerable reduction in rust severity (12.50-16.67%) and area under disease progress curve (AUDPC) value (195.83-291.67) with high total yield (86.72-76.30 kg/ha) and test weight (160.94-180.93 g) as compare to control which showed highest rust severity (54.17%) and AUDPC value (1058.33) with lowest total yield (405.30 kg/ha) and test weight (144.0g). Correlation of AUDPC values with test weight and total yield were found significantly negatively correlated whereas no correlation was found with apparent rate of infection.

Out of various disease management approaches, each one of them has its own importance in managing the rust disease of pea. But, if each approach can be integrated with each other in best manner, they can perform more effectively. However, the compatibility of these combinations needs to be carried out in field condition before being adopted.
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