Mating Type and Aggressiveness of *Phytophthora infestans* (Mont.) de Bary in Potato-Growing Areas of Punjab, Pakistan, 2017–2018 and Identification of Genotype 13_A2 in 2019–2020

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Abstract

Samples of blighted potato leaves were collected from fields in six potato-growing districts of the Punjab Province of Pakistan in 2017–2018. A total of 149 isolates of *Phytophthora infestans* were obtained from six potato cultivars (Asterix, Barsenna, Burana, Caroda, Mazika, Sante). Of these isolates, 73% were A1 mating type, 23% were A2 mating type and 4% were self-fertile. Both mating types of *P. infestans* occurred in all six districts sampled, but in every case, the A1 mating type predominated. The foliar aggressiveness of 104 of these isolates (weakly pathogenic isolates were excluded) was tested on detached leaflets of potato cv. Caroda, and a composite aggressiveness index (CAI) calculated from lesion area (measured after 10 days), latent period and infection frequency was used to compare isolates. There was no difference in CAI between isolates obtained from different districts or cultivars. The A2 mating type isolates had significantly greater CAIs than A1 isolates but this does not imply a genetic linkage nor a general association between mating type and aggressiveness. It may be that the A2 isolates belonged to an aggressive clonal lineage such as 13_A2 that has been reported in other countries in the region. While it was not possible to test the isolates collected in 2017–2018, genotyping of samples collected in 2019–2020 showed the widespread occurrence of the EU_13_A2 clonal lineage in the same districts of Pakistan and supported this hypothesis. This is the first report of EU_13_A2 from Pakistan. The implications for potato late blight management in the Punjab are discussed.

Keywords Aggressiveness · Late blight · Mating type · *Phytophthora infestans* population
Introduction

Potato (Solanum tuberosum L.) is the most important vegetable crop worldwide and the third most important food crop after wheat and rice (Jansky et al. 2019); its tubers are a globally important dietary source of starch, protein, antioxidants and vitamins (Burlingame et al. 2009). Potatoes grow in a wide range of habitats from sea level to 4000 m altitude and from 47° S to 65° N latitude (Hijmans 2001). While potato consumption has been declining in Europe and North America, it has been steadily increasing in the developing world; now, more potato production takes place in developing countries than in the developed world and potatoes have an important role in food security of developing countries (Wijesinha-Bettoni and Mouillé 2019). Late blight of potato caused by Phytophthora infestans (Mont.) de Bary is the most devastating disease affecting the crop worldwide in terms of both quality and quantity of yield (Nowicki et al. 2012). It is one of the main limiting factors for potato production in the world, and if the disease is not controlled, losses can reach 100% (Ghorbani et al. 2004), and even with low infection levels, the crop may be unsuitable for storage (Fernández-Northcote et al. 2000).

*P. infestans* is a heterothallic oomycete with two mating types, A1 and A2. Until the 1980s, the A2 mating type had been found only in Mexico where Gallegly and Galindo (1958) showed that both mating types were present; elsewhere, only the A1 mating type had been identified. The A2 mating type was first reported outside Mexico by Hohl and Iselin (1984), and now, both mating types occur in most potato-producing regions worldwide (Fry et al. 2015). Where both mating types are present in the same field, sexual recombination may occur resulting in formation of sexual oospores which can survive in soil for several years (Turkensteen et al. 2000), cause early infections and affect the population structure (Fry 2008). Sexual recombination can lead to greater variability, rendering the population more adaptable and generating more aggressive genotypes, which may be fungicide-resistant (Drenth et al. 1993; Cooke et al. 2012). However, the presence of both mating types does not necessarily result in the pathogen population reproducing sexually, although it is obviously a prerequisite. *P. infestans* may reproduce predominantly clonally even if both mating types are present: two or more separate clonal lineages may exist in parallel, surviving on potato tubers during the winter, as when there was only a single mating type (Yuen and Andersson 2013).

Potatoes are a significant crop in Pakistan; in 2018, the annual production was about 4.58 million tonnes on about 0.20 million hectares (Pakistan Economic Survey 2018–19). The first report of late blight in Pakistan was from the Swat District (Khan et al. 1985), and it has since been found in the plains of the Punjab, the North-West Frontier Province (NWFP), Baluchistan and northern Pakistan. Ahmad and Mirza (1995) first identified the A2 mating type of *P. infestans* in Pakistan when they tested 45 isolates collected in 1994 from potato-growing areas of the Punjab and the NWFP and found that 49% were A2, while the remainder were A1. Subsequently, 281 isolates collected over 3 seasons between 1997 and 2000 were tested and both mating types were found in all areas except the lower valleys of the NWFP where only A1 was detected; the A1 mating type predominated in all years, but self-fertile isolates were also found in 1999–2000 (Ahmad et al. 2002). A further 178 isolates collected from the potato-growing areas of Pakistan in the Punjab, the Swat valley and the Upper Swat valley were tested for metalaxyl resistance and mating type (Ahmad et al. 2008); metalaxyl resistance occurred
among both A1 and A2 isolates except in the Upper Swat valley where resistance occurred only in the A1 population. In the last 10 years, there has been little published information on the population structure of *P. infestans* in Pakistan and it is unclear if the pathogen is sexually reproducing, although Ahmad et al. (2002) noted the presence of both mating types within a single field. As late blight of potato is an important threat to potato production in Pakistan and the occurrence together of the two mating types of *P. infestans* potentially impacts on disease management, an extensive study of their incidence in the Punjab, where over 95% of Pakistan’s potatoes are produced, was initiated. The foliar aggressiveness of selected isolates of *P. infestans* was investigated. Genotyping of the 2017–2018 isolates using the SSR markers which have been applied to *P. infestans* populations worldwide (Li et al. 2013) to elucidate further the structure of the pathogen population in Pakistan was unfortunately not possible, but samples of *P. infestans* DNA were collected in the same region in 2019–2020 and genotyped.

**Materials and Methods**

The research was conducted in the laboratory of Plant Pathology, College of Agriculture, University of Sargodha, Sargodha, Punjab, Pakistan during 2017–2020.

**Sample Collection and Isolation of Phytophthora infestans**

Leaves with the characteristic symptoms of late blight were collected from infected potato crops during field visits to potato-producing areas of the Punjab, Pakistan, in 2017 and 2018 (Table 1). Two methods were used to isolate *P. infestans* into agar culture. In the first, blighted leaves were incubated overnight in humid boxes at 15–20 °C to encourage sporulation. When sporulation had developed, *P. infestans* was isolated by picking off sporangia using small cubes of rye agar A (Caten and Jinks 1968) containing 200 mg sodium ampicillin L⁻¹ and 50 mg rifampicin L⁻¹ and placing the cubes onto antibiotic rye agar plates. The plates were incubated at 18 °C in darkness and checked regularly for growth of *P. infestans* hyphae under a low-power microscope. When colonies had developed, they were sub-cultured onto fresh antibiotic rye agar plates. Alternatively, sections of leaves with late blight lesions were placed under potato tuber slices (cut from surface-sterilised tubers) in Petri dishes sealed with parafilm and incubated at 18 °C for up to 7 days; sporulation usually developed on the upper surface of the slices in 3–4 days. Sporangia were then picked off using antibiotic rye agar cubes and isolated as described above. Sampling of late blight lesion DNA onto FTA cards according to the methods of Li et al. (2013) was conducted from potato crops in the same potato-producing areas in December 2019 and January 2020 (Table 2). The abaxial surfaces of four actively spreading foliar blight lesions per crop were pressed onto each FTA card, allowed to air dry and then each card stored individually in plastic bags for subsequent processing.

**Maintenance of Isolates**

Isolate cultures were maintained for periods of up to 6 months on 9-cm Petri plates using either rye agar A (Caten and Jinks 1968) or pea agar (Hollomon 1965). Cultures
were maintained in long-term storage on slopes of rye A agar (Caten and Jinks 1968) in 25-ml universal glass bottles (c. 16 ml agar per bottle) inclined approximately at 45°. Isolates were sub-cultured when required for experiments by removing a section of

Table 1  Potato-growing areas of the Punjab, Pakistan, from which late blight samples were collected in 2017 and 2018

| District | Location name               | Location                  | Year | Cultivar | No. of isolates obtained |
|----------|-----------------------------|---------------------------|------|----------|--------------------------|
| Khushab  | Khushab                     | 32.2955° N, 72.3489° E    | 2017 | Caroda   | 3                        |
| Khushab  |                             | 32.2955° N, 72.3489° E    | 2018 | Caroda   | 3                        |
| Naushehra|                             | 32.5641° N, 72.1594° E    | 2017 | Barsenna | 4                        |
| Naushehra|                             | 32.5641° N, 72.1594° E    | 2018 | Barsenna | 6                        |
| Jauharabad|                            | 32.2899° N, 72.2719° E    | 2017 | Caroda   | 3                        |
| Jauharabad|                            | 32.2899° N, 72.2719° E    | 2018 | Caroda   | 3                        |
| Sargodha | Sargodha                    | 32.0740° N, 72.6861° E    | 2017 | Caroda   | 3                        |
| Sargodha |                             | 32.0740° N, 72.6861° E    | 2018 | Caroda   | 3                        |
| Shahpur  |                             | 32.2878° N, 72.4288° E    | 2017 | Caroda   | 4                        |
| Shahpur  |                             | 32.2878° N, 72.4288° E    | 2018 | Caroda   | 4                        |
| Sahiwal  |                             | 31.9765° N, 72.3282° E    | 2017 | Caroda   | 5                        |
| Sahiwal  |                             | 31.9765° N, 72.3282° E    | 2018 | Caroda   | 5                        |
| Sahiwal  | Chak Jeevan Shah            | 30.6682° N, 73.1114° E    | 2017 | Mazika   | 3                        |
| Sahiwal  | Chak Jeevan Shah            | 30.6682° N, 73.1114° E    | 2018 | Mazika   | 3                        |
| Saleem Kot|                             | 30.3708° N, 73.2397° E    | 2017 | Mazika   | 4                        |
| Saleem Kot|                             | 30.3708° N, 73.2397° E    | 2018 | Mazika   | 4                        |
| Chak Sundhey Khan|                    | 30.6682° N, 73.1114° E    | 2017 | Mazika   | 6                        |
| Chak Sundhey Khan|                       | 30.6682° N, 73.1114° E    | 2018 | Mazika   | 7                        |
| Okara    | Okara                       | 30.8138° N, 73.4534° E    | 2017 | Sante    | 4                        |
| Okara    |                             | 30.8138° N, 73.4534° E    | 2018 | Sante    | 4                        |
| Renala Khurd|                            | 30.8782° N, 73.5954° E    | 2017 | Sante    | 4                        |
| Renala Khurd|                            | 30.8782° N, 73.5954° E    | 2018 | Sante    | 4                        |
| Deepalpur|                             | 30.6770° N, 73.6477° E    | 2017 | Mazika   | 6                        |
| Deepalpur|                             | 30.6770° N, 73.6477° E    | 2018 | Mazika   | 7                        |
| Chiniot  | Chiniot                     | 31.7292° N, 72.9822° E    | 2017 | Asterix   | 4                        |
| Chiniot  |                             | 31.7292° N, 72.9822° E    | 2018 | Asterix   | 4                        |
| Bhawana  |                             | 31.5659° N, 72.6447° E    | 2017 | Asterix   | 4                        |
| Bhawana  |                             | 31.5659° N, 72.6447° E    | 2018 | Asterix   | 4                        |
| Laliyan  |                             | 31.49.21° N 72.47.50° E   | 2017 | Asterix   | 3                        |
| Laliyan  |                             | 31.49.21° N 72.47.50° E   | 2018 | Asterix   | 4                        |
| Jhang    | Jhang                       | 31.2781° N, 72.3317° E    | 2017 | Burana    | 4                        |
| Jhang    |                             | 31.2781° N, 72.3317° E    | 2018 | Burana    | 4                        |
| Shorkot  |                             | 30.8335° N, 72.0883° E    | 2017 | Burana    | 4                        |
| Shorkot  |                             | 30.8335° N, 72.0883° E    | 2018 | Burana    | 4                        |
| Ahmad Pur Sial|                     | 30.4042° N, 71.4643° E    | 2017 | Burana    | 4                        |
| Ahmad Pur Sial|                     | 30.4042° N, 71.4643° E    | 2018 | Burana    | 4                        |
mycelium from the actively growing edge of a colony and transferring to a pea agar plate (Hollomon 1965).

**Determination of Mating Types**

The mating type of each unknown isolate was determined by dual culture on rye agar medium; a small section cut from an agar culture of the test isolate was placed approx. 2 cm from a similar section of a reference A1 or A2 isolate (reference isolates were provided by Professor Shan, Northwest Agriculture and Forestry University, China). After 2 weeks’ incubation (16–18 °C in darkness), plates were examined microscopically for oospore formation at the hyphal interfaces in the contact zone. Isolates which produced sexual oospores with the reference A1 and A2 mating type isolates were categorised as A2 and A1, respectively. When oospores with antheridia formed in the A1 × A1 or A2 × A2 paired cultures, the isolate was considered likely to be self-fertile. This was confirmed by single spore isolation and culture of the resulting isolates alone on rye agar; if oospores formed in these single cultures, the isolate was categorised as self-fertile. This was done to eliminate the possibility of mixed mating type cultures (Tian et al. 2015).

**Morphology of Oospores**

The micromorphology of oospores was examined microscopically; the presence of antheridia, the shape of the oogonia, the oospore diameter and the wall thickness were recorded (Mosa et al. 1989; Cohen et al. 1997).

**Foliar Aggressiveness Testing of Selected Phytophthora infestans Isolates from 2017 to 2018**

Isolates (104, 50 from 2017 and 54 from 2018) were selected on the basis of their pathogenicity, as reported by (Raza et al. 2019), from among the 149 collected in 2017–2018 and tested for mating type (as described above); isolates that caused lesions ≥ 1 cm in length were chosen for aggressiveness testing. Newly formed sporangia from cultures of each isolate on rye agar were collected by adding 10 mL of sterile, distilled water to each Petri dish, dislodging the sporangia using a needle. Spore suspensions were filtered through cheese cloth, and the concentration was adjusted to $4 \times 10^4$ sporangia/mL using a haemocytometer. The sporangia were chilled at 4 °C for 2 h to encourage release of zoospores for use in inoculation.
A protocol adapted from Majeed et al. (2014) was used to assess aggressiveness. Uniform-sized leaves were harvested from 6-week-old glasshouse-grown potato plants cv. Caroda, washed with tap water and dried on tissue paper. For each isolate, three detached leaflets were placed adaxial side up in moist chambers and each leaflet was inoculated with a single drop (15 μL) of the test isolate applied to the midrib. The boxes of leaflets were incubated at 18–20 °C in an illuminated incubator (12 h photoperiod) for 10 days. A completely randomised design was used.

Leaflets were assessed daily for 10 days after inoculation, and lesion area (LA), latent period (LP) and infection frequency (IF) were determined. The length and width of each lesion was measured daily using a ruler and used to calculate the LA. Leaflets were viewed under a low-power microscope to check for sporulation; the LP was defined as the time in hours after inoculation until development of sporulating lesions. The infection frequency (IF) was calculated from the percentage of infected leaflets (Carlisle et al. 2002). A composite aggressiveness index (CAI) was calculated using the formula of Montarry et al. (2007):

$$\text{CAI} = \frac{\text{IF} \times \text{LA}}{\text{LP}}$$

The experimental data (LA, LP, IF and CAI) were subjected to ANOVA using SPSS software v. 19. Parameter means for the two mating types were compared using Student’s t test.

Genotyping and Genetic Analysis of Samples of P. infestans

Lesions from 2019 and 2020 were genotyped using a 12-plex simple sequence repeat (SSR) assay according to published methods (Li et al. 2013). Briefly, the abaxial sides of foliar lesions were pressed against FTA cards and a 3-mm disc of dried card was cut from the interface of green leaf tissue and brown lesion. The disc was washed according to the manufacturer’s protocols and subjected to PCR to amplify 12 SSR loci (Li et al. 2013). After running on an ABI3730 capillary sequencer, the peak profiles were edited in GeneMapper (Applied Biosystems) version 5.0 and the profiles were exported. A data file of SSR data comprising 13_A2 samples from Pakistan (25), India (88) and Europe (12) and 2 samples each of another four clonal lineages (EU_5_A1, EU_6_A1, EU_8_A1 and EU_2_A1) was compiled and Bruvo’s distances were calculated and used to generate a minimum spanning network (MSN) in poppr 2.8.5 (Kamvar et al. 2015) in R 3.6.3 (R core team 2020).

Results

Mating Type

A total of 149 isolates of P. infestans were obtained from samples of blighted potato leaves collected from fields in the potato-growing districts of the Punjab Province of Pakistan during 2017–2018 (Table 1). Of these isolates, 73% were A1 mating type, 23% were A2 mating type, while 4% were self-fertile. In 2017 and 2018, the ratio of
A1 to A2 mating types was 76:21 and 70:25, respectively (Table 3). Isolates were obtained from six potato cultivars in total; there did not appear to be any association between mating type and cultivar as A2 isolates were obtained from all six cultivars sampled (Table 4). Just under half of the fields sampled (47%) yielded isolates of both mating types, and self-fertile isolates were also present in three of these fields (Table 5).

**Micromorphology of Oospores**

Antheridia that developed in A1 × A2 paired cultures were amphigynous and varied from hemispherical to almost spherical in shape, hyaline and from 17–22 μm × 11–14 μm in size. The oogonia were nearly spherical, and their colour ranged from red-orange to brown in colour. The oospores varied from brown to dark-brown in colour, and they were spherical (21–33 μm diameter) with two walls 2.3–5.3 μm thick.

**Foliar Aggressiveness**

The three parameters of aggressiveness (LA, LP and IF) were very highly significantly \((P < 0.001)\) correlated with each other and with the CAI (Table 6); the CAI was therefore used for subsequent comparisons. There were very highly significant differences in terms of CAI between individual isolates in both 2017 and 2018 (Figs. 1 and

| Year and district | Total number of isolates | Number of isolates of mating type |
|------------------|------------------------|----------------------------------|
|                  |                        | A1 | A2 | Self-fertile |
| 2017             |                        |    |    |              |
| Khushab          | 10                     | 9  | 1  | 0            |
| Sargodha         | 12                     | 10 | 2  | 0            |
| Sahiwal          | 13                     | 8  | 4  | 1            |
| Okara            | 14                     | 7  | 6  | 1            |
| Jhang            | 12                     | 12 | 0  | 0            |
| Chiniot          | 11                     | 9  | 2  | 0            |
| Total            | 72                     | 55 (76)a | 15 (21)a | 2 (3)a |
| 2018             |                        |    |    |              |
| Khushab          | 12                     | 12 | 0  | 0            |
| Sargodha         | 13                     | 10 | 2  | 1            |
| Sahiwal          | 13                     | 7  | 5  | 1            |
| Okara            | 15                     | 6  | 8  | 1            |
| Jhang            | 12                     | 10 | 2  | 0            |
| Chiniot          | 12                     | 9  | 2  | 1            |
| Total            | 77                     | 54 (70) | 19 (25) | 4 (5) |
| Total 2017–2018  | 149                    | 109 (73)a | 34 (23)a | 6 (4)a |

a Percentage of isolates of each mating type
but overall, there was no significant difference in the mean CAI for the 50 isolates from 2017 (974) and for the 54 isolates from 2018 (1185). However, a tendency was noted for the more aggressive isolates to be those of the A2 mating type. Comparison of the CAIs for the A1 and A2 isolates (self-fertile isolates were excluded as only two from 2017 and four from 2018 were tested for aggressiveness) by ANOVA showed that the A2 isolates had a significantly greater mean CAI than the A1 isolates for both 2017 and 2018 isolates (Table 7).

There were no significant differences in CAI between isolates of *P. infestans* from different districts of the Punjab (Table 8). There were some significant differences between isolates obtained from different cultivars, with those from cv. Barsenna being more aggressive than those from cvs. Mazika, Caroda and Sante; this was not associated with a larger proportion of A2 isolates from this cultivar (Table 9). However, as there were only four isolates from cv. Barsenna tested for aggressiveness, this could be a sampling artefact.

### Table 4

| District | Number of isolates | Cultivar(s) yielding isolates of each mating type |
|----------|--------------------|--------------------------------------------------|
| Khushab  | 22                 | Caroda, Barsenna                                 |
| Sargodha | 24                 | Caroda                                           |
| Sahiwal  | 27                 | Mazika                                           |
| Okara    | 29                 | Mazika, Sante                                    |
| Jhang    | 24                 | Burana                                           |
| Chiniot  | 23                 | Asterix                                          |

### Table 5

| District | Number of fields sampled | A1 only | A2 only | A1 + A2 |
|----------|--------------------------|---------|---------|---------|
| Khushab  | 6                        | 5       | 0       | 1       |
| Sargodha | 6                        | 2*      | 0       | 4       |
| Sahiwal  | 6                        | 1       | 0       | 5**     |
| Okara    | 6                        | 1       | 2*      | 3*      |
| Jhang    | 6                        | 5       | 0       | 1       |
| Chiniot  | 6                        | 3*      | 0       | 3       |
| Total    | 36                       | 17      | 2       | 17      |

*One field also yielded one self-fertile isolate*

**One field also yielded two self-fertile isolates*
Genotyping

Technical problems led to the loss of the living isolates collected in 2017–2018. Genotyping of pathogen DNA from 25 lesions collected during December 2019 and January 2020 (Table 2) revealed a single multi-locus genotype (MLG). Comparisons against a large database of published (Chowdappa et al. 2013; Chowdappa et al. 2015; Dey et al. 2018; Martínez et al. 2019) and unpublished SSR profiles indicated that this MLG was a sub-clonal variant of the 13_A2 clonal lineage. This single MLG was collected from five districts in Pakistan. A sample of MLGs from India and Europe was examined using poppr and the MSN demonstrated a cluster of closely related 13_A2 MLGs genetically distinct from each of the other four clonal lineages (Fig. 3). The MLG sampled in Pakistan was identical to one sampled in both India and Europe.

Table 6  Correlation coefficients of the parameters of aggressiveness derived from inoculation onto detached potato leaflets of isolates of Phytophthora infestans obtained from the Punjab, Pakistan, in 2017 and 2018

|                         | Lesion area (mm²) | Latent period (h) | Infection efficiency (%) | Composite aggressiveness index |
|-------------------------|-------------------|-------------------|--------------------------|-------------------------------|
| Lesion area (mm²)      | 1                 |                   |                          |                               |
| Latent period (h)      | -0.9233***        | 1                 |                          |                               |
| Infection efficiency (%)| 0.8821***         | -0.9288***        | 1                        |                               |
| Composite aggressiveness index | 0.8797***         | -0.9027***        | 0.8828***    | 1                              |

***P < 0.001

Fig. 1 Composite aggressive index (CAI) for Phytophthora infestans isolates collected in 2017 from the Punjab, Pakistan. A1 isolates are indicated by cross-hatched bars, A2 isolates by black bars and self-fertile isolates by grey bars.
Discussion

The oospores observed in this study, when isolates were crossed with the opposite mating type, were spherical and had 21–33 μm diameter, which is similar in size to those reported by others; Erwin and Ribeiro (1996) state that in culture oospores’ diameters range from 24 to 56 μm, while Gallegly and Hong (2008) give a figure of about 30 μm as the mean oospore diameter. Singh et al. (1994) reported that they obtained slightly larger oospores (32–44 μm diameter) when A2 *P. infestans* isolates from India were paired in culture with an A1 isolate of *Phytophthora capsici*.

In the present study, both mating types of *P. infestans* were found to occur in all six districts of the Punjab, Pakistan, from which isolates were obtained either in 2017 or in 2018. However, in every case, the A1 mating type predominated, and overall, 73% of isolates were A1. In the districts of Jhang and Khushab, only A1 isolates were detected in 2017 and 2018, respectively. In contrast, recent studies in India showed exclusively

### Table 7  Comparison of composite aggressiveness indices for A1 and A2 isolates of *Phytophthora infestans* obtained from the Punjab, Pakistan, in 2017 and 2018

| Year of isolate collection | Mean CAI | Significance |
|---------------------------|----------|--------------|
|                           | A1 only  | A2 only      | ***          |
| 2017                      | 694      | 1647         | ***          |
| 2018                      | 859      | 1819         | ***          |
| 2017 and 2018             | 774      | 1743         | ***          |

***P<0.001
A2 populations of the aggressive European 13\_A2 genotype in southern India on potato and tomato during 2010–2012 (Chowdappa et al. 2015) and in eastern and north-eastern India on potato and tomato during 2012–2014 (Dey et al. 2018). Published studies of the *P. infestans* population on potatoes in north-west India appear to be lacking. Genotype 13\_A2 is reported to be widespread in Bangladesh (Kessel et al. 2017) and has also been found in Nepal (Sharma Adhikari 2017). When genotype 13\_A2 is initially introduced into a region, it tends to displace the existing *P. infestans* population, often rapidly (Cooke et al. 2012). The predominance of the A1 mating type of *P. infestans* found in the present study in the Punjab, Pakistan, in 2017–2018 suggests that genotype 13\_A2, detected in 2019–2020, has only recently been introduced.

Aggressiveness, defined as the quantity of disease induced by a pathogenic strain on a susceptible host (Andrivon 1993) is often invoked to explain changes in *P. infestans* populations. In reality, pathogenic fitness (the expected contribution of a phenotype to the subsequent generation; Antonovics and Alexander 1989), of which aggressiveness is only one aspect, is of greater epidemiological significance. However, while fitness is environmentally dependent and difficult to measure, one of the attributes of

| District    | Number of isolates | A2 isolates (%) | Mean CAI* |
|-------------|--------------------|-----------------|-----------|
| Khushab     | 14                 | 7               | 1145      |
| Sargodha    | 14                 | 29              | 938       |
| Sahiwal     | 19                 | 47              | 949       |
| Okara       | 24                 | 58              | 1076      |
| Jhang       | 17                 | 12              | 1226      |
| Chiniot     | 16                 | 25              | 1174      |
| Total/mean  | 104                | 33              | 1085      |

*There were no significant differences (*P* > 0.05) between CAIs for isolates from the six sampled districts of the Punjab

| Cultivar    | Number of isolates | A2 isolates (%) | Mean CAI* |
|-------------|--------------------|-----------------|-----------|
| Asterix     | 16                 | 25              | 1174ab    |
| Barsenna    | 4                  | 25              | 1710a     |
| Burana      | 17                 | 12              | 1226ab    |
| Caroda      | 24                 | 17              | 930b      |
| Mazika      | 31                 | 52              | 1059b     |
| Sante       | 12                 | 58              | 919b      |
| Total/mean  | 104                | 33              | 1170      |

*CAIs for isolates from the sampled cultivars followed by the same letter are not significantly different (*P* < 0.05)
aggressiveness is that it can be measured under standard conditions allowing comparison between isolates (Andrivon 1993). Studies of *P. infestans* have therefore frequently focussed on aggressiveness in order to try to understand the behaviour of components of populations of the pathogen (e.g. Carlisle et al. 2002).

In the present study, the foliar aggressiveness of all isolates collected in 2017 and 2018, except for the 45 which were only weakly pathogenic (which were all of the A1 mating type), was evaluated and this was done using detached potato leaflets. Although this technique does not completely mimic infection in the field, it allows for the comparison of substantial numbers of isolates under controlled conditions and is therefore widely used (e.g. Majeed et al. 2014). Because the components of aggressiveness (latent period, lesion area and infection frequency) measured in the present study were closely correlated, a composite aggressiveness index (CAI) was calculated and used to compare the isolates. Majeed et al. (2014), in their study using 10 isolates of *P. infestans* collected in the Khyber Pakhtunkhwa Province of Pakistan in 2011, defined isolates possessing a CAI of 1000 or greater as being highly aggressive and reported that 40% of their isolates belonged to this class. In the present study of 104 isolates from the Punjab, 46% possessed CAIs of 1000 or more. However, as different cultivars were used (cv. Desiree by Majeed et al. 2014 and cv. Caroda in the present study) and only a small number of isolates were evaluated in the earlier study, the results may not be strictly comparable. The most striking finding in terms of aggressiveness in the present study was that, overall, isolates of the A2 mating type had a very highly significantly greater CAI than A1 isolates. While some individual A1 isolates were also highly aggressive, others were only weakly aggressive (CAI < 500), whereas no A2 isolates fell into the weakly aggressive class. Nonetheless, within some fields, aggressive

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**Fig. 3** Minimum spanning network (MSN) of samples of *Phytophthora infestans* DNA collected from Pakistan, India and Europe. Each node represents an MLG with edge width and shading reflecting genetic relatedness based on Bruvo’s genetic distance. This MSN indicates a cluster of clonal MLG variants of EU_13_A2 that are genetically closely related in comparison with the more genetically distinct pairs of isolates of other clonal lineages. The latter clones are delineated within the dashed line.
isolates of both mating types occurred together. It should be emphasised that there is no genetic link between aggressiveness and mating type of \textit{P. infestans}; because of its clonal reproduction, strong associations between mating type and phenotypic characters such as aggressiveness and fungicide resistance can arise in particular places and times, but this is a result of selection of specific genotypes. The generally greater aggressiveness of A2 compared with A1 isolates is most probably an effect of the particular A2 genotype(s) of the pathogen present in the Punjab. Such an association, as noted above, occurred in the case of the aggressive genotype 13\_A2 (Cooke et al. 2012). The finding of only the 13\_A2 type in each of 25 samples from five districts of the Punjab Region in Pakistan collected in 2019–2020 (Fig. 3) suggests that this lineage, which has not previously been reported from Pakistan, is widely distributed. This supports the premise that the greater aggressiveness of the A2 isolates obtained in 2017–2018 might be because they belong to the 13\_A2 genotype, recently introduced to Pakistan. Unfortunately, it was not possible to genotype the 2017–2018 isolates themselves.

The occurrence of both mating types within a single field in 17 cases indicated a potential for sexual reproduction by \textit{P. infestans}. There could be a risk of oospore-derived infections in such fields (Widmark et al. 2007), which could lead to earlier attacks and more diverse pathogen populations as has been reported from the Nordic countries (Hannukkala et al. 2007; Brurberg et al. 2011). It is therefore important that, as well as using fungicides to prevent pathogen infection and, where possible, growing potato cultivars with some resistance to late blight, growers should avoid continuous cropping, which increases the risk of infection by soil-borne oospores and may cause earlier attacks (Runno-Paurson et al. 2009). Turkensteen et al. (2000) showed that oospores can survive in field soil for up to 4 years, so using rotations with 5 or more years between potato crops should reduce this risk.

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