Plant resistance in different cell layers affects aphid probing and feeding behaviour during poor- and non-host interactions

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Abstract

Aphids are phloem-feeding insects that cause economic losses to crops globally. Whilst aphid interactions with susceptible plants and partially resistant genotypes have been well characterised with regards to aphid probing and feeding behaviour, the interactions with non-natural host species are not well understood. Using aphid choice assays with the broad host range pest *Myzus persicae* and the cereal pest *Rhopalosiphum padi* we show that about 10% of aphids settle on non-/poor-host species over a 24h time period. We used the Electrical Penetration Graph technique to assess aphid probing and feeding behaviour during the non-/poor-host interactions. In the Arabidopsis non-host interaction with the cereal pest *R. padi* aphids were unable to reach and feed from the phloem, with resistance likely residing in the mesophyll cell layer. In the barley poor-host interaction with *M. persicae*, resistance is likely phloem-based as aphids were able to reach the phloem but ingestion was reduced compared with the host interaction. Overall our data suggests that plant resistance to aphids in non-host and poor-host interactions with these aphid species likely resides in different plant cell layers. Future work will take into account specific cell layers where resistances are based to dissect the underlying mechanisms and gain a better understanding of how we may improve crop resistance to aphids.

Keywords: aphid, EPG analyses, nonhost, plant resistance, probing, stylet pathway.

1 **Introduction**

2 Aphids are important insect pests which cause significant yield losses to crops

3 globally (Blackman R, 2000). There are approximately 5000 aphid species described

4 and around 250 of these are important agricultural and horticultural pests which vary

5 in their host range – the ability to successfully infest different plant species. This host

6 range variation generally applies to secondary hosts during summer months, where

7 aphid populations increase rapidly due to asexual reproduction (Moran, 1992). Whilst

8 the majority of aphid species exhibit a limited host range, dedicated to few closely

9 related plant species, some aphid species, like *Myzus persicae* Sulzer (green peach

10 aphid), have an exceptionally broad host range which includes representatives from

11 more than 40 plant families (Blackman R, 2000, Powell et al., 2006). The

12 evolutionary drivers and molecular determinants of such exceptionally broad host

13 ranges in aphids remain to be elucidated.

14 Host suitability relies on a number of factors, which could be based either at the plant

15 surface or within plant tissues and cells (Powell et al., 2006). Prior to probing the leaf

16 surface aphid behaviour can be influenced by a range of these factors including leaf

17 colour, emitted volatile organic compounds and leaf surface components, such as

18 epicuticular waxes or trichomes (Doring, 2014, Doring & Chittka, 2007, Neal et al.,

19 1990). Regardless of whether the aphid encounters a host or non-host plant species

20 their specialised mouthparts, known as stylets, are utilised to probe into the plant

21 tissue (Escudero-Martinez et al., 2017, Jaouannet et al., 2015, Powell et al., 2006).

22 This probing behaviour is associated with the transmission of important plant viruses

23 during both host and non-host interactions (Debokx & Piron, 1990, Katis & Gibson,

24 1985, Powell et al., 2006, Verbeek et al., 2010) which can substantially reduce crop

25 yields (Perry et al., 2000). During interactions with susceptible plant species the

aphid stylets penetrate the plant epidermis and move through the plant tissue

towards the vascular bundle. During this process the stylets probe into adjacent plant

cells, and saliva is secreted both in the apoplast and into probed cells along the

stylet-pathway (Tjallingii, 2006, Tjallingii & Esch, 1993). During compatible plant-

aphid interactions the aphid stylets are able to successfully puncture the sieve-tube

elements to facilitate ingestion of phloem sap (Tjallingii, 1995, Tjallingii, 2006).

The aphid stylet-pathway through the plant tissue has been well-characterised during

interactions with susceptible plants using the Electrical Penetration Graph (EPG)

technique. This technique uses an electrical circuit to connect the aphid to the plant

via a series of electrical probes, allowing distinction between different phases of the

stylet pathway from obtained electrical waveforms which correlate with the position of

the aphid stylet within plant tissue in real-time (Prado & Tjallingii, 1994, Tjallingii,

1985a, Tjallingii, 1985b, Tjallingii & Esch, 1993). Briefly, the aphid is attached to an

electrical probe with gold wire, and a copper electrode is placed into the soil to

incorporate the plant into the electrical system. Both the plant and the aphid

electrodes are attached to a data-logger, which is read by computational software

and the whole set-up is contained in a grounded Faraday cage (Mclean & Kinsey,

1968, Tjallingii, 1978, Tjallingii, 1985a, Tjallingii, 1985b). Once the aphid probes the

plant tissue the circuit closes and changes in electrical voltage are displayed as

alternating waveforms which can be manually annotated using computational

software and translated into time-series data (Tjallingii & Esch, 1993). The biological

relevance of the different waveforms that are detected by the EPG technique have

been extensively analysed (Prado & Tjallingii, 1994, Tjallingii, 1978, Tjallingii, 1985a,

Tjallingii, 1985b). Waveforms associated with aphid probing are: waveform np,

representing non-probing behaviour where the stylets are not in contact with the leaf

surface; waveform C, which begins upon stylet penetration of leaf tissue and is correlated with the intercellular apoplastic stylet pathway located at the epidermis or the mesophyll cell layers; waveform pd, associated with piercing of a plant cell which leads to a signal potential drop; waveform F, which reflects stylet mechanical/penetration difficulties; and waveform E1e, which represents extracellular saliva secretion into plant tissues other than phloem. Waveforms associated with vascular interactions and which provide intricate information at the aphid feeding site are: waveform G, which represents aphids drinking from the xylem sap; waveform E1, which is linked to aphid salivation into phloem before ingestion; and waveform E2, which corresponds to phloem sap ingestion (Alvarez et al., 2006). A graphical representation of examples of these waveforms, alongside the stylet activity during each, is shown in Fig. 1.

Although the EPG technique has mainly been used to study aphid interactions with susceptible and (partially-)resistant genotypes of host plant species, it also represents a suitable tool to explore how aphids interact with plants which are not natural hosts, including non-host and poor-host species. Indeed, EPG analyses of *Brevicoryne brassicae* Linnaeus (cabbage aphid) on host Brassicaceae and non-host *Vicia faba* showed that this aphid species was unable to reach the phloem when feeding on the non-host *V. faba*, despite probing the leaf surface (Garbys & Pawluk, 1999). Also, epidermis and phloem factors contributed to resistance in different legume species to different pea aphid biotypes (Schwarzkopf et al., 2013). By characterising aphid probing and feeding behaviour across different aphid interactions with non-/poor-host species we aim to generate a better understanding of where associated resistance mechanisms reside. This in turn will facilitate

important mechanistic studies to reveal the molecular determinants of plant immunity to aphids.

We previously showed that *M. persicae*, which is not a pest of barley, is able to feed and reproduce on this crop under controlled environment conditions, but to a lower extent than on a host species such as oil seed rape or Arabidopsis (Escudero-Martinez et al., 2017). On the contrary, *Rhopalosiphum padi* Linnaeus (bird cherry-oat aphid) is a pest of barley but is unable to feed from, and therefore survive, on Arabidopsis (Jaouannet et al., 2015). However, in both the *M. persicae*-barley poor-host interaction and the *R. padi*-Arabidopsis non-host interaction probing of the leaf surface takes place (Escudero-Martinez et al., 2017, Jaouannet et al., 2015). In line with our previous findings, choice assays showed that both aphid species will settle on and interact with non-/poor-host plant species if given a choice, with 10% of aphids found on non-/poor-hosts after 24h. Using EPG analyses of *M. persicae* and *R. padi* on Arabidopsis and barley we explored differences in aphid probing and feeding behaviour during non-/poor-host versus host interactions. We show that resistance in the non-/poor-host interactions can reside in different plant cell layers, suggesting complex mechanisms may underlie plant immunity to aphids.

Materials and Methods

Aphid rearing

R. padi (JHI-JB, genotype G) (J. et al., Thorpe et al., 2018) was maintained on *Hordeum vulgare* cv Optic and *M. persicae* (JHI_genotype O) was maintained on *Brassica napus* (oilseed rape). All aphid species used in the experiments were maintained in growth chambers under controlled conditions ($18°C \pm 2°C$, 16 h of light).

Plant growth

Barley plants (cv. Golden Promise) were pre-germinated in Petri dishes with wet filter paper for three days in the dark. Then, they were moved to a plant growth cabinet under controlled conditions and grown for 7 days (growth stage 1.10, determined using the staging key (Zadoks et al., 1974)) until the EPG experiments. *Arabidopsis thaliana* Col-0 plants were sown directly in soil; the seeds were stratified for 3 days at 4°C and placed in the growth cabinet for 4-5 weeks before use in experiments (growth stage 1.10 to 3.90, determined using the Boyes growth key (Boyes et al., 2001)). The cabinet conditions for Arabidopsis were 8 hours of light (125 μmol photons/m^2.s), at 22 °C and 70% humidity. The cabinet conditions for barley were 8 hours of light (150 μmol photons/m2.s), at 20 °C (+-2°C).

Aphid choice experiment

Aphid choice tests were devised to investigate the host plant preference of *R. padi* and *M. persicae.* Three choice test assays were developed: one using 50 *R. padi* aphids, a second using 50 *M. persicae* aphids, and a third using a mixed species population (25 *R. padi,* 25 *M. persicae*). For each assay, fifty aphids (mixed aged: 2nd instar – apterous adult) were placed on a sheet of tissue paper and were placed in the centre of a Perspex cage halfway between two plants (one Arabidopsis, one barley). Aphids were 90 mm away from both plants and the two plants were 180 mm apart. Bamboo sticks served as bridges from the cage bottom (where the aphids were placed) to each plant, with additional bamboo sticks acting as bridges between the two plants, similar to the set-up used by Nowak and Komor (Nowak & Komor, 2010). Once the aphids were placed between the plants and the ladders were positioned, the cages were closed and the proportion of aphids present on the host, non-/poor-host, or which had not settled were scored three and 24 hours later.

Choice assays were carried out in growth chambers under controlled conditions

(18°C ± 2°C, 16 h of light).

Choice tests were carried out simultaneously in separate Perspex cages (440 mm x

340 mm x 390 mm). For each replicate the assignment of aphid mixture (*R. padi, M.*

persicae, or mixed) to cage (1, 2, or 3) and the position (1 or 2) of Arabidposis and

barley within each cage was randomly assigned. Seven replicates were collected for

each aphid mixture. The proportion of aphids detected on each plant were modelled

in response to plant type (Host, non-/poor-host, or not settled), aphid mixture (*R.*

padi, *M. persicae*, mixed species), time-point (three hours and 24 hours) and all

interactions using a linear mixed effects model. Cage and block were included as

random factors, the model was simplified using manual backward stepwise model

selection, and fitted-residual plots were observed at each stage to assess model

suitability. Models were analysed using a χ^2 Analysis of Deviance Test. Differences in

the Least Squares Mean with Tukey correction for multiple comparison was used as

a post-hoc test. Data were analysed in R Studio v. 1.0.143 running R v. 3.4.3 (R Core

Team, 2017) with additional packages car v.2.1-4 (Weisberg & Fox, 2011), lme4

v.1.1-13, and lsmeans v.2.27-62 (Lenth, 2016).

Electrical penetration graph (EPG) analyses

The probing and feeding behaviour of *R. padi* and *M. persicae* on different plant

species was assessed using the Electrical Penetration Graph technique (Tjallingii,

1995) on a Giga-4 DC-EPG device with 1 Giga Ω resistance (EPG Systems, The

Netherlands). We used a randomized block design for all EPG experiments

performed here. Aphids were connected to a copper electrode with a golden wire (20

μm diameter), attached at the aphid dorsum and connected to the electrode with water-based silver glue. Aphids were lowered onto either an Arabidopsis or barley leaf approximately 1-1.5 hr after being removed from culture, depending on the treatment, and feeding behaviour was recorded over a 6h period. Three recordings were taken simultaneously. Each experiment was initiated between 10-12 am and the experiment was performed over a 6-month period, with 18 host and 17 non-host replicates for *R. padi* and 23 host and 28 poor-host replicates for *M. persicae*. Data were acquired using the Stylet+ D software package version v.01.28 and annotated manually using the Stylet+ A v.01.30 software (EPG-Systems, The Netherlands). Obtained waveforms were annotated with one of the following signals: no penetration (np), stylet penetration into the epidermal and mesophyll tissue (pathway/C phase), cellular punctures during the C phase (pd), watery salivation into sieve elements (E1), ingestion of phloem sap (E2), derailed stylet mechanics/stylet penetration difficulties (waveform F), xylem ingestion (waveform G), or extracellular saliva secretion into mesophyll (E1e) (Alvarez et al., 2006, Tjallingii, 1995). Annotated waveforms were converted into time-series data using the excel macro developed by Dr Schliephake (Julius Kühn-Institut); these converted parameters were used for statistical analysis. Parameters used for comparisons in these experiments are described by Giordanengo et al. (Giordanengo, 2014), and include total time of probing, number of probes, duration of phloem sap ingestion, and duration of xylem sap ingestion, a total of 97 parameters were measured. Statistical analyses were performed in R Studio running R v. 3.2.3. (R Core Team, 2017) using the Wilcoxon rank test, a significance threshold of 0.05 was used.

Results

Aphids preferentially settle on their host plant

We used aphid choice assays to examine the host plant preference of

Rhopalosiphum padi and *Myzus persicae*. We monitored the settling behaviour of *R.*

padi when provided with a choice between barley (host) and Arabidopsis (non-host),

of *M. persicae* when provided with a choice between Arabidopsis (host) and barley

(non-host), and of a mixed species population containing *R. padi* and *M. persicae*.

The majority of aphids preferentially settled on the host plant, *c.* 50% of aphids

settled on the host plant within three hours (Table 1; Fig. 2). The number of aphids

that settled on the host plant increased to around 80% after 24 hours for all aphid

populations assessed (t = -9.48; p = <0.001) with the number of unsettled aphids

decreasing (t = 8.30; p = <0.001). However, approximately 10% of aphids were found

on either the non-host or the poor-host plant at both time-points. No effect of aphid

mixture was observed (Table 1), indicating that the presence of additional aphid

species did not influence aphid behaviour.

The Arabidopsis-*R. padi* non-host interaction is characterised by long no-

probing periods and difficulties in locating the vascular tissues

We employed the Electrical Penetration Graph (EPG) technique to compare the

feeding behaviour of *R. padi* on barley (host) with Arabidopsis (non-host) and of *M.*

persicae on Arabidopsis (host) with barley (poor-host) over a six hour period in order

to identify the tissue layers involved in non-host and poor-host resistance against

aphids. We assessed 97 feeding parameters in total, 71 of these were altered during

feeding on non/poor-host plants compared with feeding patterns on host plants

(Table S1) with 26 parameters remaining unaffected (Table S2).

The majority of feeding parameters that differed between *R. padi* feeding on host

compared with non-host plants were related to stylet probing of the plant tissue and

interactions with the plant vasculature (Fig. 3). In general, probing parameters that

differed for *R. padi* when interacting with non-host versus host plants were non-

probing periods, number of stylet probes into plant tissue, and time spent in the

epidermal/mesophyll cells (C phase) (Fig. 3A; Table S1).

During non-host interactions with Arabidopsis, the total time the aphids were not

probing plant tissue during the 6 h recording was 2.5 times greater (4889s) than the

host interactions (1767s) (Fig. 3A; Table S1; W = 33.00; p = <0.001). However, the

overall number of stylet probes into plant tissue was higher on non-host plants (18)

than host plants (8) (Fig. 3A; Table S1; W = 52.50, p = 0.001). Although the total

number of C phases (stylet activity at the epidermis/mesophyll, including a return to

C phase following stylet interactions in the vasculature) was not significantly different

between non-host and host interactions, the overall time spent in the

epidermis/mesophyll (C phase) was over two times longer for the non-host (14128s)

compared with host interactions (6237s) (Fig. 2A; Table S1; W = 37.00; p = <0.001).

All the vascular-related parameters (G, E1 salivation and E2 ingestion phases)

measured for *R. padi* were significantly reduced during non-host interactions

compared with host interactions (Fig. 3B; Table S1). This included a two-fold

reduction in the number of xylem ingestion (G phase) events during the non-host

interaction (0.24 times) compared with the host interaction (0.50 times) (Fig. 3B;

Table S1; W = 2.28.50; p = 0.001) alongside a significant decrease in the total length

of xylem ingestion, 1021s for non-host compared with 1483s for host plants (Fig. 3B;

Table S1; W = 221.50; p = 0.003). We also observed significantly fewer salivation

events (E1 phase) during the non-host interaction (0.18 events) compared with the

host interaction (3.67 events; W = 282.00; p = <0.001), with salivation events five-fold

shorter during the non-host interaction (18s) compared with the host interaction (93s)

(Fig. 3B; Table S1; W = 278.00; p = <0.001). Ingestion of phloem sap (E2 phase)

was rarely observed during the non-host interaction (0.06 times) compared with the

host interaction (3 times; W = 285.00; p = <0.001), and the total duration of this

ingestion period was greatly reduced on non-host plants (19s) compared with host

plants (10030s, or 2.78 hours) (Fig. 3B; Table S1; W = 288.00; p = <0.001).

The barley-*M. persicae* poor-host interaction is characterised by a lack of

sustained phloem ingestion

The majority of feeding parameters that differed between *M. persicae* feeding on host

compared with poor-host plants were primarily related to interactions within the plant

vasculature, specifically a decrease in interactions with the phloem and an increase

in interactions with the xylem (Fig. 4; Table S1). In general, this involved a decrease

in the ability to locate the phloem and initiate ingestion of phloem sap. When feeding

on poor-host plants there was a significant increase in the number of probes made

into the plant tissue by aphids (19) compared with the number of probes made into

host plants (16) (Fig. 3A; Table S1; W = 186.00; p = 0.024). However, the total length

of time aphids probed into plant tissue, the number of pathway (C) phase events, and

the total time spent within the pathway (C) phase was similar for the host and poor-

host interactions (Fig. 4A)

Aphid stylet activities related to the vascular parameters (G – xylem, E1 – phloem

salivation, and E2 – phloem ingestion) were different between host and poor-host

interactions (Fig. 4B; Table S1). The number of times that *M. persicae* reached the

xylem (G phase) during the poor-host interaction was higher (1.33 times; W =

133.50; p = <0.001) and the total time of xylem ingestion was longer (2321s; W =

142.50; p = <0.001) than during the host interaction, where aphids reached the xylem

0.30 times and spent a total of 691s ingesting xylem sap (Fig. 4B; Table S1). For the

E1 salivation phase the number and duration of events was reduced during the poor-

host interaction, 1.73 events (W = 5.28; p = <0.001) with a total length of time spent

salivating into the phloem of 562s (W = 500.00; p = <0.001), compared with the host

interaction (7 events with a time length of 652s) (Fig. 4B; Table S1).

M. persicae showed limited ingestion periods during the poor-host compared with

host interactions. The number of E2 phases and their length was greatly reduced on

poor-host plants, 0.53 events (W = 552.50; p = <0.001) with a 40-fold decrease in the

total time spent ingesting phloem (126s; W = 573.50; p = <0.001), compared with

host plants (5.7 events with a total length of 5064s) (Fig. 4B; Table S1). Moreover, on

the poor-host sustained phloem ingestion was severely lacking, and aphids spent

only 49s in the E2 ingestion phase on poor-host plants (W = 520.00; p= <0.001) with

events being nearly absent, 0.07 events (W = 515.00; p = <0.001). In contrast,

aphids spent 4322s in the E2 sustained ingestion phase on host plants over 2.1

events during the 6h recording (Fig. 4B; Table 1). Therefore, the *M. persicae* poor-

host interaction features substantially reduced phloem ingestion.

Discussion

The overall aim of this study was to gain insight into where resistances against

aphids may reside within the plant tissue during host versus non/poor-host

interactions by analysing aphid probing and feeding behaviour. We showed that

when given a choice aphids do interact with non-/poor-host plants under controlled

conditions, and we further explored these interactions using EPG analyses. Common

features of the non-host and poor-host interactions were an increased number of

probes and longer no-probing periods. Importantly, our data showed differences

between *R. padi* and *M. persicae* probing and feeding behaviour on the non-/poor-

host plants. During the *R. padi*-Arabidopsis (non-host) interaction the aphids only

occasionally reached the vascular tissues. On the contrary, during the *M. persicae*-

barley interaction (poor-host) aphids successfully reached the vascular tissue and

could ingest xylem and phloem, however prolonged periods of phloem ingestion were

inhibited. Based on the data generated here for *M. persicae* and *R. padi* we propose

a model wherein poor- and non-host plant resistances against these aphid species

may reside within the phloem and mesophyll cell layers, respectively (Fig. 5).

During the *R. padi*-barley interaction (host interaction) the aphids spend less time

probing and in the pathway (C) phase and readily reach the phloem where salivation

and phloem sap ingestion occurs for several hours (Fig. 5A). Occasionally, aphids

ingest xylem, which is thought to be important in coping with osmotic effects

associated with ingestion of large amounts of phloem sap (Pompon et al., 2010,

Spiller et al., 1990). In contrast, during the *R. padi* – Arabidopsis interaction (non-host

interaction) aphids exhibit altered probing behaviour, including an increase in the

number of plant probes alongside a decrease in the total time probing into plant

tissue. Additionally, *R. padi* shows an extended stylet pathway phase, and only rarely

does the aphid reach the Arabidopsis phloem or xylem (Fig. 5B). On the occasions

where the *R. padi* stylets reach the vascular tissue during non-host interactions the

ingestion of phloem and xylem sap is ineffective, in line with this aphid being unable

to survive on Arabidopsis (Jaouannet et al., 2015).

Interestingly, *R. padi* spent less time probing into plant tissue during the non-host

interaction. However, during these probes aphids spent an increased time interacting

with the mesophyll tissue during the non-host interaction than the host interaction,

including an increase in the total time spent in the pathway (C) phase. This indicates

that non-host resistance could potentially reside in the mesophyll tissue as the aphids

struggled to probe beyond this layer and access to the vascular tissue was limited

(Fig. 5B), as further indicated by the increased time required for aphids to reach the

phloem during non/poor-host interactions compared with the host interactions.

Further research will be needed to further understand the mechanisms underlying

Arabidopsis non-host resistance to *R. padi*, and to investigate the potential

involvement of specific recognition receptors within the mesophyll cell layer.

Interestingly, the NADPH oxidase *AtRbohF*, involved in ROS (Reactive Oxygen

Species) production, a member of the *LEA* (*Late Embryogenesis Abundant*) family,

implicated in abiotic and biotic stress, as well as the *VSP1* (*Vegetative Storage*

Protein 1), which is activated by jasmonate signalling, contribute to Arabidopsis non-

host resistance against *R. padi* (Jaouannet et al., 2015). Whether these genes act

within the mesophyll cell layer to activate defences against aphids remains to be

determined.

The *M. persicae*-Arabidopsis (host) interaction, features short probing and pathway

times, and prolonged salivation and ingestion once the phloem is reached, as well as

occasional xylem drinking (Fig. 5C). In contrast, during the *M. persicae*-barley

interaction (poor-host interaction) aphids show increased probing but spend a similar

time in the stylet pathway phase as aphids on host Arabidopsis plants. The main

differences between the Arabidopsis (host) and barley (poor-host) interactions with

M. persicae are reduced salivation in the phloem and relatively short periods of

phloem ingestion (less than 10 minutes) on barley (Fig. 5C and 5D). It is likely that

this reduced phloem sap ingestion is responsible for the reduced *M. persicae*

performance on barley (Escudero-Martinez et al., 2017, Ramirez & Niemeyer, 2000).

It is possible that *M. persicae* attempts to compensate for this reduced ingestion of

phloem sap with increased xylem drinking, in line with the observation that aphid

starvation increases the xylem phase (Fig. 5D) (Ramirez & Niemeyer, 2000).

Phloem resistance factors are related to the E1 salivation and E2 ingestion

parameters, and in particular ingestion phases shorter than 10 minutes (Alvarez et

al., 2006, Prado & Tjallingii, 1997). Phloem-mediated defences against aphids

include the occlusion of sieve elements, which prevents aphids from ingesting

phloem sap (Dreyer & Campbell, 1987, Medina-Ortega & Walker, 2015, Will & van

Bel, 2006). This phloem occlusion occurs upon callose deposition and formation of P-

protein plugs. The latter is thought to seal off the phloem upon damage and/or to

block the aphid food canal (Tjallingii, 2006, Will & van Bel, 2006). Interestingly, PAD4

was found to be a component of phloem-based immunity against *M. persicae* in

Arabidopsis (Pegadaraju et al., 2007). However, no barley PAD4 (MLOC_1340) or

PAD4-related genes were up-regulated during the barley-*M. persicae* interaction

(Escudero-Martinez et al., 2017). However, our previous transcriptome analyses

showed induction of a barley gene encoding Phloem Protein 2-like (PP2), which is a

phloem specific lectin, with the induction being most pronounced during the barley-*M.*

persicae interaction (Escudero-Martinez et al., 2017). Lectins have carbohydrate-

binding properties and function in cell communication, development, and plant

defence (Bellande et al., 2017).   PP2 is a lectin highly abundant in the phloem and

accumulates in damaged phloem sieve pores to form protective plugs (Read &

Northcote, 1983). Overexpression of *AtPP2* in Arabidopsis leads to reduced *M.*

persicae feeding suggesting PP2 may contribute to defences against aphids (Zhang

et al., 2011), possibly by interfering with aphid digestion in the midgut (Kehr, 2006).

The very infrequent phloem sap ingestion we observed might reflect a rejection of the

sieve element, possibly due to the presence of a deterrent factor in the phloem sap

(Mayoral et al., 1996). Indeed, lectins, including PP2-like proteins, have been shown

to have deterrent activities and insecticidal activities against *M. persicae* (Jaber et al.,

2010, Sauvion et al., 1996, Zhang et al., 2011). Whether barley phloem-lectins like

PP2 indeed contribute to phloem-based defences of barley against *M. persicae*

needs to be further tested.

It is important to note that the EPG experimental set-up was of a no-choice nature

(i.e. aphids were placed on the plants) and that additional plant resistance

components that affect aphid choice may play a role in the interactions studied here

(Escudero-Martinez et al., 2017, Powell et al., 2006). For example, we previously

showed that the black cherry aphid (*Myzus cerasi* Fabricius), which infests cherry

trees as well as several herbaceous plants, displays only limited probing on non-host

barley plants, and does not settle on barley leaves (Escudero-Martinez et al., 2017),

pointing to a potential role of barley defences that act at the pre-probing level against

this aphid species (Nottingham et al., 1991). In addition, some plant induced volatile

compounds have been reported to be repellent to aphid pests and attractants of their

natural enemies (Dreyer & Jones, 1981, Mallinger et al., 2011, Turlings & Ton, 2006).

With limited genetic crop resistance available against aphids, identifying the

determinants of non/poor-host resistance is an important area of research that may

help the development novel crop protection strategies. Using a detailed assessment

of aphid probing and feeding behaviour on different natural host and non-host

species we show that resistances may reside in different cell layers depending on the plant species-aphid species interaction.

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Competing interests

The author(s) declare no competing interests.

Author contributions

JIBB, CEM and DJL conceived and designed the experiments, CEM and DJL performed the experiments, JIBB, CEM and DJL analysed the data, JIBB and CEM wrote the manuscript with input from DJL. All authors read and approved the final manuscript.

Tables

Table 1: Statistical results of the choice test assay

Response variable	Test Statistic (degrees of freedom)	p-value
Plant Type	$X^2_{(2)} = 532.65$	P = <0.001
Aphid Mixture	$X^2_{(2)} = 0.01$	P = 0.996
Time-point	$X^2_{(1)} = 0.01$	P = 0.949
Plant Type x Aphid mixture	$X^2_{(4)} = 5.43$	P = 0.245
Plant Type x Time-Point	$X^2_{(2)} = 162.06$	P = <0.001
Aphid Mixture x Time-Point	$X^2_{(2)} = 0.01$	P = 0.996
Plant Type x Aphid Mixture x Time-Point	$X^2_{(4)} = 0.34$	P = 0.986

Figures

Figure 1. Graphical representation of aphid/stylet activities associated with each EPG waveform.

(A)   Example of aphid activity during np (non-probing) period, stylet is not in contact with leaf tissue.

(B)   Initiation of pathway (C) phase - aphid stylet pierces leaf epidermis,

(C)   Potential drop (pd) – aphid stylet penetrates adjacent plant cell

(D)   Stylet penetration difficulties (F phase)

(E)   Extracellular saliva secretion (E1e) phase – salivation into extracellular space.

(F)   Xylem ingestion (G phase) – stylet penetrates vascular xylem cells to initiate xylem drinking.

(G)   Salivation into phloem (E1 phase) – stylet penetrates sieve tube element and aphid initiates salivation into phloem sap.

(H)   Phloem ingestion (E2 phase) – aphid begins passive ingestion of phloem sap. Also includes sustained phloem ingestion (sE2 phase) - a period of phloem sap ingestion lasting > 10 mins.

Figure 2. Stacked bar charts showing the settling behaviour of aphids in the choice experiment.

(A) Aphid settling after three hours

(B) Aphid settling after 24 hours

Graphs show the mean proportion of aphids from the *R. padi* (Rp)*, M. persicae* (Mp)*,* and the mixed species population (Mix) which had settled on the host plant (H; green), the non-host plant (NH; red), the poor-host plant (PH; yellow), the non/poor-host plant (NH.PH; orange) or which has not settled (NS; grey). Letter under each

bar indicate differences based on Least Squares Mean post-hoc analysis with Tukey correction.

Figure 3. Box plots showing different EPG parameters associated with *Rhopalosiphum padi*-**barley (host) and** *Rhopalosiphum padi*-**Arabidopsis (non-host) interactions.**

(A) Probing-related parameters: total number of probing events, total length of no probing time, total number of pathway (C) phase events, total length of pathway (C) phase time.

(B) Vascular-related parameters: number of xylem ingestion (G phase) events, total length of xylem ingestion, number of salivation (E1 phase) events where aphid saliva is secreted into phloem sap, total length of salivation (E1 phase), number of phloem sap ingestion (E2 phase) events and total length of phloem sap ingestion (E2 phase). Green boxes indicate the host (H) interaction and red boxes represent the non-host (NH) interaction. *R. padi* on host plants was replicated 18 times and *R. padi* on non-host plants was replicated 17 times. Significant differences between interactions were assessed by Wilcoxon non-parametric t-test (*= $p \leq 0.05$ and *** = $p \leq 0.01$).

Figure 4: Box plots showing different EPG parameters in *Myzus persicae* interaction with a host (Arabidopsis) and a poor-host plant (barley).

(A) Probing-related parameters: total number of probing events, total length of no probing time, total number of pathway (C) phase events, total length of pathway (C) phase time. (B) Vascular-related parameters: number of xylem ingestion (G phase) events, total length of xylem ingestion, number of salivation (E1 phase) events where

aphid saliva is secreted into phloem sap, total length of salivation (E1 phase), total length of phloem sap ingestion (E2 phase) and total length of sustained phloem sap ingestion (sE2 phase).

Green boxes indicate the host (H) interaction and yellow boxes represent the poor-host (PH)interaction. *M. persicae* on host plants was replicated 23 times and *M. persicae* on poor-host plants was replicated 28 times. Significant differences between interactions were assessed statistically by Wilcoxon non-parametric t-test (*= p ≤0.05 and *** = p ≤0.01).

Figure 5. Model showing *R. padi* and *M. persicae* probing and feeding during host, poor-host and non-host plant interactions.

(A) During the host interaction (*R. padi*-barley)*,* the aphids will probe the epidermal and mesophyll cells (pathway C phase), then will drink from the xylem or salivate and feed from the phloem, with feeding lasting for hours.

(B) During the non-host interaction (*R. padi*-Arabidopsis), the aphids will spend a long time not probing, and when probing eventually occurs the aphids remain in stylet pathway phase (in epidermis and mesophyll cell layers) most of the time and only occasionally will reach the vascular tissue, either xylem or phloem. No sustained ingestion of phloem sap takes place.

(C) During the host interaction (*M. persicae*-Arabidopsis)*,* the aphids will probe the epidermal and mesophyll cells (pathway C phase), then will drink from the xylem or salivate and feed from the phloem, with feeding taking place for hours.

(D) During the poor-host interaction (*M. persicae*-barley), the aphids show increased probing compared to the host interaction, while the stylet pathway phase (in

epidermis and mesophyll cell layers) is similar to the interaction with the host plant. At the vascular level, long periods of time will be spent in the xylem, and eventually aphid will reach the phloem, salivate and ingest phloem sap. However, contrary to the host interaction, no sustained (>10 minutes) ingestion of phloem sap takes place.

Supplementary Data

Table S1. Results for all obtained Electrical penetration graph (EPG) parameters which were significantly different between host and non/poor-host feeding. Table displays the EPG parameter assessed, a description of the parameter, and the plant tissue layer involved. Results displayed are the mean and standard deviation (SD) for each aphid-plant combination for each parameter alongside the Wilcoxon test statistic (W value) and p value for each pairwise host vs non/poor host comparison. p values in bold represent values significantly different in both host vs non-host and host vs poor-host interactions, italicised p values represent parameters which only differed in one combination. Average and standard deviation of the 97 electrical EPG parameters calculated for *R. padi* host (Rp_Hv) and non-host (Rp_At). Average and standard deviation of the 97 electrical EPG parameters calculated for *M. persicae* host (Mp_At) and poor-host (Mp_Hv). Calculations were made with summary statistics in Rstudio. The EPG list of parameters was taken from EPG systems: www.epgsystems.eu/files/List%20EPG%20variables.xls

Table S2: Results for all obtained Electrical penetration graph (EPG) parameters which were not significantly different between host and non/poor-host feeding. Table displays the EPG parameter assessed, a description of the parameter, and the plant tissue layer involved. Results displayed are the mean and standard deviation (SD) for each aphid-plant combination for each parameter alongside the Wilcoxon test

statistic (W value) and p value for each pairwise host vs non/poor host comparison. p values in bold represent values significantly different in both host vs non-host and host vs poor-host interactions, italicised p values represent parameters which only differed in one combination. Average and standard deviation of the 26 electrical EPG parameters calculated for *R. padi* host (Rp_Hv) and non-host (Rp_At). Average and standard deviation of the 97 electrical EPG parameters calculated for *M. persicae* host (Mp_At) and poor-host (Mp_Hv). Calculations were made with summary statistics in Rstudio. The EPG list of parameters was taken from EPG systems: www.epgsystems.eu/files/List%20EPG%20variables.xls

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