Non-host and poor-host resistance against aphids may reside in

different plant cell layers depending on the plant species-aphid species

interaction

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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. Here we use the Electrical Penetration Graph technique to study aphid probing and feeding behaviour on poor- and non-host plants for the broad host range pest *Myzus persicae* and the cereal pest *Rhopalosiphum padi*. 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 to be 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.

Introduction

Aphids are important insect pests which cause significant yield losses to crops globally [1]. There are approximately 5000 aphid species described and around 250 of these are important agricultural and horticultural pests which vary in their host range – the ability to successfully infest different plant species. This host range variation generally applies to secondary hosts during summer

months, where aphid populations increase rapidly due to asexual reproduction

[2]. Whilst the majority of aphid species exhibit a limited host range, dedicated

to few closely related plant species, some aphid species, like *Myzus persicae*

(green peach aphid), have an exceptionally broad host range which includes

representatives from more than 40 plant families [1,3]. The evolutionary drivers

and molecular determinants of such exceptionally broad host ranges in aphids

remain to be elucidated.

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

the plant surface or within plant tissues and cells [3]. Prior to probing the leaf

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

including leaf colour, emitted volatile organic compounds and leaf surface

components, such as epicuticular waxes or trichomes [4-6]. Regardless of

whether the aphid encounters a host or non-host plant species their

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

tissue [3,7,8]. This probing behaviour is associated with the transmission of

important plant viruses during both host and non-host interactions [3,9-11] which

can substantially reduce crop yields [12]. 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 into probed cells along the stylet-pathway [13,14]. During compatible

plant-aphid interactions the aphid stylets are able to successfully puncture the

sieve-tube elements to facilitate ingestion of phloem sap [13,15].

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 [14,16-18]. 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 [17-20]. 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 [14]. The biological

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

have been extensively analysed [16-18,20]. 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 [21]. 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. By

characterising aphid probing and feeding behaviour on non-/poor-host species

we will 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 [7]. On the contrary, *Rhopalosiphum padi* (bird cherry-oat aphid) is

a pest of barley but is unable to feed from, and therefore survive, on

Arabidopsis [8]. 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 [7,8]. Here, we made use of the different host and non-/poor-host

combinations of *M. persicae* and *R. padi* with Arabidopsis and barley to

explore aphid probing and feeding behaviour during these different

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.

Results

The Arabidopsis-*R. padi* non-host interaction is characterised by long

no-probing periods and difficulties to locate the vascular tissues

We used the EPG technique to monitor *R. padi* probing and feeding behaviour

for the first 6 hours on either Arabidopsis (non-host) or barley cv. Golden

Promise (host) and found significant differences with regards to parameters

relating to probing and interactions with the plant vasculature (Fig. 2). The

statistical results for all EPG parameters, which were significantly different in

host vs non/poor-host interactions (71/97 tested parameters) are displayed in

Table S1, with the statistical results for all non-significant parameters (26/97

tested) shown in Table S2.

In general, probing parameters which differed for *R. padi* when interacting

with non-host versus host plants were non-probing periods, number of probing

events, and time spent in the epidermal/mesophyll cells (C phase) (Fig. 2a;

Table S1). In the non-host interaction, the total time the aphids were not

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

host interaction (1767s) (Fig. 2a; Table S1; W = 33.00; p = <0.001). Also,

aphids probed non-host plants more frequently (18 attempts) than host plants

(8 attempts) (Fig. 2a; Table S1; W = 52.50, p = 0.001). Although the total

number of C phases (stylet activity at the epidermis/mesophyll) was not

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

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

(14128s) compared with the host interaction (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

the non-host compared with the host interaction (Fig. 2b; Table S1).

Additionally, the number of xylem events was halved during the non-host

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

2b; Table S1; W = 2.28.50; p = 0.001). The total length of xylem ingestion (G

phase) was significantly shorter on the non-host (1021s) compared with the

host plants (1483s) (Fig. 2b; Table S1; W = 221.50; p = 0.003). We 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), and salivation events were five-fold shorter during the non-host

(18s) versus host (93s) interactions (Fig. 2b; 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 (19s) versus host plants (10030s, or 2.78 hours)

(Fig. 2b; Table S1; W = 288.00; p = <0.001).

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

of sustained phloem ingestion

Similar to the EPG assays performed with *R. padi*, we also assessed probing

and feeding of *M. persicae* on host plants (Arabidopsis) and poor-host plants

(barley cv. Golden Promise) over a 6-hour period (Fig. 3). Within the probing

parameters, we observed significant differences in the number of probing

attempts on poor-host compared with host plants, while the non-probing time,

and number and time of events associated with stylet activity in the epidermal

and mesophyll tissue (C phase) were similar (Table S2; Fig. 3b). In contrast,

the differences between *M. persicae* interactions with poor-host versus host

plants were primarily at the vascular level with aphids spending extensive

periods of time in the xylem and showing reduced phloem ingestion on poor-

host plants (Table S1; Fig. 3b). During the poor-host interaction, there was a

significant increase in the number of aphid probing attempts (19 attempts)

compared with the host interaction (16 attempts) (Fig. 3a; Table S1; W =

186.00; p = 0.024). The time spent in the non-probing phase was longer in the

poor-host interaction (3130s) than the host interaction (2275s), but this

difference was not statistically significant (Fig. 3a; Table S2). The aphids

spent a similar amount of time in the C phase, which lasted 13328s during the

poor-host and 11879s during the host interaction. 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. 3b; 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 time of interaction 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 in the xylem

(Fig. 3b; Table S1).

For the E1 salivation phase the number and duration of events was reduced

during the poor-host interaction, with 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. 3b; 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, with 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. 3b; 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. 3b; Table 1). Therefore, the *M. persicae* poor-host

interaction features substantially reduced phloem ingestion.

Discussion

The overall aim of this study was to characterise aphid probing and feeding

behaviour during host versus non/poor-host interactions in order to gain

insight into where resistances against aphids may reside within the plant

tissue. Our EPG analyses revealed that common features of the non-host and

poor-host interactions were an increased number of probing events 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 reached the phloem but were

unable to ingest sap for prolonged periods of time. 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. 4).

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

time probing and in the pathway phase compared to the non-host interaction

with Arabidopsis, and readily reach the phloem where salivation and phloem

sap ingestion takes place for several hours (Fig. 4a). Occasionally, aphids will

drink from the xylem, which is thought to be important in coping with osmotic

effects associated with ingestion of large amounts of phloem sap [22,23]. In

contrast, *R. padi* shows increased probing behaviour on the non-host plant

Arabidopsis, as well as an extended stylet pathway phase, and only rarely

does the aphid reach the Arabidopsis phloem or xylem (Fig. 4b). On the

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

interactions the ingestion of sap on these occasions is not effective, in line

with this aphid being unable to survive on Arabidopsis [8]. Interestingly, *R. padi*

encountered more frequent stylet penetration difficulties when interacting with

Arabidopsis, as reflected by the F phase. This F phase is known to occur

exclusively at the mesophyll cell layers [20], suggesting that the non-host

resistance could reside there (Fig. 4b). 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* [8]. 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. 4c). 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. 4c and d). It is likely that this reduced phloem sap

ingestion is responsible for the reduced *M. persicae* performance on barley

[7,24]. 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. 4d) [24].

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

parameters, and in particular ingestion phases shorter than 10 minutes [21,25].

Phloem-mediated defences against aphids include the occlusion of sieve

elements, which prevents aphids from ingesting phloem sap [26-28]. 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 [13,28]. Interestingly, PAD4 was found to be a component of

phloem-based immunity against *M. persicae* in Arabidopsis [29]. However, no

barely PAD4 (MLOC_1340) or PAD4-related genes were up-regulated during

the barley-*M. persicae* interaction [7]. 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 [7]. Lectins have

carbohydrate-binding properties and function in cell communication,

development, and plant defence [30].   PP2 is a lectin highly abundant in the

phloem and accumulates in damaged phloem sieve pores to form protective

plugs [31]. Overexpression of *AtPP2* in Arabidopsis leads to reduced *M.*

persicae feeding suggesting PP2 may contribute to defences against aphids

[32], possibly by interfering with aphid digestion in the midgut [33]. 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 [34]. Indeed, lectins, including PP2-like proteins, have been shown

to have deterrent activities and insecticidal activities against *M. persicae*

[32,35,36]. 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 [3,7]. For example, we previously showed that the

black cherry aphid (*Myzus cerasi*), 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 [7], pointing to a potential role of

barley defences that act at the pre-probing level against this aphid species [37].

In addition, some plant induced volatile compounds have been reported to be

repellent to aphid pests and attractants of their natural enemies [38-40].

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.

Methods

Aphid rearing

R. padi (JHI-JB, genotype G) [41,42] was maintained on *Hordeum vulgare* L. 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, 16 h of light).

Plant growth

Barley plants (cv. OpticGolden Promise) were pre-germinated in Petri dishes

with wet filter paper for three days in the dark. Then, they were moved to a

cabinet under controlled conditions and grown for 7 days (growth stage 1.10,

determined using the staging key [43]) until the EPG experiments. Arabidopsis

plants were sown directly in soil; the seeds were stratified for 3 days at 4ºC

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 [44]). 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).

312   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

[15] on a Giga-4 DC-EPG device with 1 Giga Ω resistance (EPG Systems, The

Nethelands). 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)

[15,21]. 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. (2014)

[45], 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) [46] using the Wilcoxon rank test.

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353 **Author contributions**
354
355 JIBB, CEM and DJL conceived and designed the experiments, CEM and DJL

356 performed the experiments, JIBB, CEM and DJL analysed the data, JIBB and

357 CEM wrote the manuscript with input from DJL. All authors read and approved

358 the final manuscript.

Competing interests

The author(s) declare no competing interests.

Figure Legends

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

a)   Example of obtained EPG waveform, top panel shows waveform over a 1 hour period and lower panel is the magnified waveform for the highlighted section over a 60s period. np = non-probing, C = C phase (stylet activity in epidermis/mesophyll), pd = potential drop (cellular puncture) E1/E2 indicates phloem phases, saliva secretion (E1) and phloem ingestion (E2).

b)   Example of aphid activity during np waveform, stylet is not in contact with leaf tissue therefore voltage is approximately 0. The magnified panels cover only a 30s period.

c)   Initiation of pathway (C) phase - aphid stylet pierces leaf epidermis, voltage becomes positive as aphid stylet moves through epidermis. Waveform characterised by large initial waves, voltage is generally between 0 and +4 volts.

d)   Potential drop (pd) – aphid stylet penetrates adjacent plant cell leading to rapid decrease in voltage. Waveform characterised by a reduction in voltage to around – 3 volts for 2 – 3s before stylet retraction from pierced cell and return to C phase.

e)   Stylet penetration difficulties (F phase) – waveform characterised by

rapid oscillations in positive voltage between +1 and +5 volts, waveforms

generally appear smooth.

f)   Extracellular saliva secretion (E1e) phase – salivation into extracellular

space, waveform is characterised by short fin-shapes waves in a positive

voltage, around +2 to +4 volts.

g)   Xylem ingestion (G phase) – stylet penetrates vascular xylem cells to

initiate xylem drinking. Waveform is characterised by oscillating positive

waveforms between +2 and +4 volts.

h)   Salivation into phloem (E1 phase) – stylet penetrates sieve tube

element and aphid initiates salivation into phloem sap. Waveform is preceded

by a series of rapid potential drops (cellular probes) with a final probe into a

sieve element, salivation waveform characterised by fin-shaped waveforms in

the negative voltage.

i)   Phloem ingestion (E2 phase) – aphid begins passive ingestion of

phloem sap. Waveform always follows on from E1 waveform and is

characterised by brief and sharp negative peaks.

Figure 2. Box plots showing different EPG parameters associated with

***Rhopalosiphum padi*-barley (host) and *Rhopalosiphum padi*-Arabidopsis**

(non-host) interactions.

a) Probing-related parameters: non-probing period length, number of probing

events, number of pathway (C phase) events and pathway phase length.

b) Vascular-related parameters: number of xylem (G phase) events, xylem

phase length, number salivation (E1 phase) events, salivation phase length,

number of ingestion (E2 phase) events and ingestion phase length. Green

boxes indicate the host interaction and red boxes represent the non-host

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 ≤0.05 and

*** = p ≤0.01).

Figure 3: Box plots showing different EPG parameters in *Myzus persicae*

interaction with a host (Arabidopsis) and a poor-host plant (barley).

a) *M. persicae* probing-related parameters: number of probing events, no--

probing period length, number of pathway (C phase) events and pathway

phase length.

b) *M. persicae* vascular-related parameters: number of xylem (G phase)

events, xylem phase length, number of salivation (E1 phase) events,

salivation phase length, number of ingestion (E2 phase) events and ingestion

phase length. Green boxes indicate the host interaction and red boxes

represent the poor-host 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 4. 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 variables 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 variables was

taken from EPG systems:

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Figure 1

Figure 2

Figure 3

Figure 4