Gut cancer increases the risk for *Drosophila* to be preyed upon by hunting spiders

David Duneau[1,2*] and Nicolas Buchon[3]

[1]Université Toulouse 3 Paul Sabatier, CNRS, IRD; UMR5174 EDB (Laboratoire Évolution &

Diversité Biologique); Toulouse, France.

[2]Instituto Gulbenkian de Ciência, Rua da Quinta Grande 6, P-2780, Oeiras, Portugal

[3]Cornell Institute for Host Microbe Interactions and Disease, Department of Entomology, Cornell

University, Ithaca, NY 14853, USA.

* Corresponding author: David Duneau. david.duneau@gmail.com

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Abstract

Predators are thought to prey on individuals that are in poor physical condition, although the evidence supporting this is ambiguous. We tested if sick individuals where more predated using *Drosophila melanogaster* flies as manipulable preys. We asked whether hunting spiders, trapped from the wild, would selectively prey upon flies with compromised health (i.e. chronically infected or cancerous) versus healthy flies under laboratory conditions. Flies chronically infected with the bacterium *Providencia rettgeri*, a natural *Drosophila* pathogen, were not selectively preyed upon by jumping spiders. We strengthened and confirmed our finding with another hunting spider species, small wolf spiders. We discuss that this result supports the hypothesis that chronic infection is associated with reduced symptoms notably to avoid the potentially deadly consequences of host predation on pathogens. We then induced colon cancer in some of the flies and asked whether the presence of cancer led to selective predation; there is little evidence for this, even in vertebrates. As the cancer developed, the incidence of predation by jumping spiders on the afflicted flies increased. We conclude that disease can have different lethal consequences through predation, even in invertebrate species, and that cancer is a factor in selective predation. Our results may explain why early tumours, but not metastasized cancers, are commonly detected in organisms in the wild, as cancer bearing individuals are rapidly eliminated due to the strong selective pressure against them.

Introduction

In vertebrates, individuals that are injured, or diseased are often preyed upon, as are juveniles who have not reached adult speed or reflex (Furey et al., 2021; Genovart et al., 2010; Mesa, Poe, Gadomski, & Petersen, 1994; Møller, 2008; Murray, 2002). This is especially true when a predator's preferred type of prey is difficult to catch at the prime of its performance (Temple, 1987), however, there are exceptions (Penteriani et al., 2008). Although this conventional wisdom of selective predation on substandard individuals is largely accepted, only a few studies have tested it empirically. Selective predation can only be said to exist when the relative frequencies of the types of prey in a predator's diet differ from the frequencies of the prey in the environment (Chesson, 1978). Hence, to test the selective prey hypothesis in the wild, data on the types of prey sought by many of the predators and the health status of the whole prey population would be determined. This is obviously difficult to obtain. Furthermore, a prey individual may suffer from different conditions which influence each other and lead to spurious effects. For example, exhaustion from care for large brood increases susceptibility to parasites (Oppliger, Christe, & Richner, 1996). Thus, the fittest hosts are the ones classified as being sick. Although those information may be vitally important in the context of eco-evolutionary dynamics (Brunner, Anaya-Rojas, Matthews, & Eizaguirre, 2017; Brunner, Deere, Egas, Eizaguirre, & Raeymaekers, 2019), the topic has been subjected more to speculation than to empirical study.

Probably the most studied question regarding selective predation is whether predators prey more on individuals with suboptimal conditions over healthy individuals. Even if hosts predators may avoid infected prey to avoid the risk of getting sick themselves (Goren & Ben-Ami, 2017), it has been suggested, based on field observations, that infections could increase the vulnerability to predation (Adelman, Mayer, & Hawley, 2017; Duffy, Hall, Tessier, & Huebner, 2005; Gooding et al., 2020; Miller et al., 2008; Moller & Erritzoe, 2000; Moller, Erritzoe, & Tottrup, 2010; Møller & Nielsen, 2007; Murray, Cary, & Keith, 2006) and there is some experimental support for this (DeBlieux & Hoverman, 2019; Gallagher et al., 2019; Johnson, Stanton, Preu, Forshay, & Carpenter, 2006; Murray, 2002). Consequently, by mediating selective predation, parasites can mediate the relationship between hosts and their prey (Hall, Cáceres, Duffy, & Cáceres, 2005; Møller, 2008). For instance, removing

predators can reduce vertebrate prey populations (Sih, Crowley, Mcpeek, Petranka, & Strohmeier,

1985), which seems counterintuitive, as predators are expected to remove preys. However,

mathematical models predict that virulent parasites may be selected against if predation

disproportionately removes them before they can be transmitted to the host (Packer, Holt, Hudson,

Lafferty, & Dobson, 2003). Without this check on virulent parasites by the predator, more prey may

die from lethal infections than from predation. Hence, if selective predation for infected individuals is

a common pattern, it may also be a major driver of host and parasite evolution and cannot be ignored

any longer (Møller, 2008).

It is also relevant to consider that not all sickness come from infections and that non-infectious

diseases could also increase the risk of being predated. In fact, understanding the role of cancer on

ecosystem function has been identified as a key endeavour (Dujon, Aktipis, et al., 2021). So far, most

of our knowledge is based on the consequence of transmissible cancer on an apex predator (i.e.

Tasmanian devil (Cunningham, Johnson, & Jones, 2020; Hollings, Jones, Mooney, & Mccallum,

2014; Hollings, Jones, Mooney, & McCallum, 2016; Woods et al., 2018)), and few to nothing is

known about the potential selective predation of prey that are sick due to a non-infectious disease (see

(Perret, Gidoin, Ujvari, Thomas, & Roche, 2020) for a theoretical study). Mutations can occur during

the cell replication and division required to grow and maintain multicellular animals, and certain types

of mutation, said oncogenic mutations, can lead to the formation of tumours (Aktipis et al., 2015;

Albuquerque, Drummond do Val, Doherty, & de Magalhães, 2018; Hanahan & Weinberg, 2011;

National Cancer Institute, 2020a). In most cases, tumours are benign (Bissell & Hines, 2011), but they

can be malignant, and even fatal (Aktipis et al., 2015; Albuquerque et al., 2018; Hanahan & Weinberg,

2011; National Cancer Institute, 2020b). Clinical categorization of tumours has been established in

unchallenging, even 'optimal' environments (i.e. in the laboratory or clinic), usually to the point of

death via organ failure. The impact of tumour progression is likely to influence tremendously

individual's life history traits (e.g. lifespan and reproductive output but also competitive and dispersal

ability, and pathogen susceptibility), yet, its impact under more natural conditions has largely been

neglected (Roche, Møller, DeGregori, & Thomas, 2017; Marion Vittecoq et al., 2013). Furthermore,

depending on the relationship between tumour development and the health of the prey, predation could

affect the selection of oncogenic mutations and thus the risk to develop cancer. To our knowledge, the

possible interactions between a predator and a prey that is afflicted with cancer are poorly understood

and have not been empirically studied to date (Marion Vittecoq et al., 2013).

Here, we induced chronic bacterial infection and colon cancer in prey to understand their role

in increasing the likelihood of being predated. We use the fruit fly *Drosophila melanogaster*, a model

which has started recently to be used to study the effect of cancer from an ecological perspective

(Arnal et al., 2017; Dawson et al., 2018) and is already a well-characterized model for infectious and

non-infectious diseases, as a novel model for sick prey. We assess the predation of flies by hunting

spiders, using flies chronically infected by a wild-caught bacterial pathogen (*Providencia rettgeri*),

and have genetically induced cancer, under laboratory-controlled conditions. These conditions allow

for the ideal experimental setting to study predation and the role of prey well-being in a population

context.

Materials and methods

Prey and predators.

We used *Drosophila melanogaster* flies as prey in this study as they are tractable to genetic induction

of tumours, are well-defined models for bacterial infection, and allow for control of environmental and

genetic differences. Only one sex (males) were used, to avoid sex effects. We induced chronic

infection by injecting 23 nL of a suspension of few thousand cells of the bacterium *Providencia*

rettgeri (strain *Dmel*, isolated from wild-caught *Drosophila melanogaster;* Juneja and Lazzaro 2009)

in phosphate-buffered saline (PBS) into the abdomen of flies (Canton S line, a wildtype genotype kept

in the laboratory since several decades). The same volume of sterile PBS without bacteria was injected

into another group of flies that served as the control. Individuals surviving for three days or more after

injection of the bacteria have an established chronic infection, as we have previously demonstrated

(Duneau et al., 2017).

To induce tumours in the flies, we took advantage of signalling pathways that regulate cell growth in

mammals, which have a conserved function in *Drosophila*, the EGFR and Wnt pathways (Millburn et

al., 2016; Mirzoyan et al., 2019; Rudrapatna, Cagan, & Das, 2012; Villegas, 2019). We used targeted

induction of these pathways (Rasv12 for EGFR, APC RNAi for the Wnt pathway) to induce two types

of enlargement of the flies digestive tract (i.e. hyperplasia, often called benign tumour, and cancer).

Those types of colon cancer is appropriate for this study, as it does not overtly affect physical

performance including locomotion, and will thus allow for the study of the interaction of oncogenic

phenomena and predation (see Dawson et al. 2018 for locomotion assays in gut cancerous flies in

another genetic background). To generate tumours in the gut of *Drosophila*, flies *esg^{TS}* (*esgGal4*,

Gal80^{ts}, *UAS-GFP*) were crossed to either background control flies (Trip stock 35786), flies carrying a

UAS-Ras^{v12} (Bloomington 64196) or flies carrying a combination of *UAS-Ras^{v12}* (Bloomington

641926) and *UAS-APC-RNAi* (Bloomington Trip stock 28582) combined. The expression of Ras^{v12} in

progenitor cells stimulates an accelerated turnover and a mild accumulation of epithelial cells in the

gut, akin to a dysplastic tissue (Buchon, Broderick, Kuraishi, & Lemaitre, 2010; Houtz, Bonfini, Bing,

& Buchon, 2019), later also referred as hyperplasia. The overexpression of *Ras^{v12}* combined with the

knock-down of *APC* leads to increased proliferation, and accumulation of progenitor cells and

epithelial cells in the gut, reminiscent of an intestinal disseminated tumour (Wang et al., 2013), later

also referred as cancerous. F1 flies from these crosses were raised at 18°C (Gal4 off, transgene

expression off, normal development and emergence), and 3 days after emergence were switched to

29°C (Gal4 on, transgene expression on) for 10 or 20 days before performing the experiment. As in

(Wang et al., 2013), crosses with WT flies showed mild tissue renewal (restricted GFP signal), flies

with expressing *Ras^{v12}* showed disseminated GFP in their gut and flies expressing *Ras^{v12}* and *APC*

RNAi showed gut enlargement and GFP accumulation. The longest the induction is done and the

strong is the phenotypes leading to a gradient from induced hyperplasia for 10 days (i.e. mild colon

enlargement) to induced cancer for 20 days (i.e. limit to the complete obstruction of the colon).

We used two families of hunting spiders as predators: jumping spiders of several species from

the family of Salticidae (see illustration in Fig. 1) and small wolf spiders from the family of Lycosidae

(likely *Pardosa lugubris*). Both jumping and wolf spiders hunt by wandering rather than by using

webs as traps.

Jumping spiders have the sharpest vision of any arthropod, even surpassing many vertebrates

(Land & Nilsson, 2012). They hunt during the day and rely on this astonishing vision when catching a

prey. Similar to the way a cat stalks its prey, jumping spiders turn towards their prey, directed by a

pair of lateral eyes that provide a nearly panoramic field of view with the ability to discern motion

(Zurek & Nelson, 2012). Then, the spiders track, approach, and jump on the prey. The vision accuracy

is conferred by two pairs of forward-facing eyes that also provide a precise perception of depth

(Nagata et al., 2012; Zurek, Taylor, Evans, & Nelson, 2010).

After having tested jumping spiders as predator, we decided to test whether infected

individuals would still not be more likely to be eaten by another type of hunting spider species with

different hunting behaviour, wolf spiders. This was additionally performed to strengthen our

"negative" result. Wolf and jumping spiders have already successfully been used to test for selective

predation (Holmberg & Turnbull, 1982; Vickers & Taylor, 2018). Wolf spiders are generally nocturnal

hunters and their eyes function mainly as low-light movement detectors, however *Pardosa* species are

mostly hunting during the day (Edgar, 1969). Their vision is not as acute as that of jumping spiders

(Land & Nilsson, 2012). The use of visual cues in prey detection and orientation have only been well-

studied in jumping spiders, and empirical evidence suggests that wolf spiders rely more on vibrations

to capture their prey than on vision (Lizotte & Rovner, 1988). Hence, the differences between these

two types of hunting spiders primarily lie in their ability to prey on flies that are at rest. In an

environment where there is no place to hide, jumping spiders would likely prey equally well on

moving and non-moving prey, and selective predation in this case should depend largely on the prey's

capacity to escape.

Predation trials

Predation trials were performed during the day by incubating five healthy and five sick flies, age-

matched, at 20°C with a spider in a round plastic box (11.5cm x 8.5cm, see illustration in fig. 1B)

172 containing a wet piece of cotton or foam. The flies were first added to the box to settle before the

173 introduction of the predators. After about 30 min, spiders were put in the box. The trials generally

174 lasted for about 4 hours or were stopped when 50% of the flies had been preyed upon, as, when

175 scoring only at the end of the experiment, it allows to score the most extreme result (i.e. all individuals

176 from one treatment have been eaten before the other treatment). Even if no spiders were starved before

177 the experiment, this goal was sometimes exceeded as some spiders ate faster than expected. Individual

178 spiders have never been used twice as predators within an experiment.

179 To determine the number of surviving flies that were infected, we ground the remaining flies

180 at the end of the predation period and suspended the mixture in 250 µL of Luria-Bertani (LB) medium.

181 A droplet of five microliters of the resulting suspension was spread on a petri dish filled with agar

182 containing LB medium, and the plates were incubated overnight at 37°C to assess for the presence of

183 *P. rettgeri*. We calculated the proportion of infected flies at the end of predation period by determining

184 the number of droplets in which *P. rettgeri* were present. For a control comparison, we performed the

185 same process with infected and healthy flies, but did not expose them to predatory spiders. Our

186 protocol consistently led to the induction of tumour, even if interindividual variation in tumour size

187 exist which can be a source of variation in our trials. In the unlikely event that some individuals were

188 considered has carrying a tumour but did not, those individuals would reduce the strength of our signal

189 as they would be like control individuals. The number of surviving flies that had cancer were counted

190 by dissecting all surviving flies and observing their guts under a fluorescent microscope, which reveals

191 the presence/absence of cancer.

192 **Statistical analysis**

193 Statistical analysis were performed using R, version 4.1 (R Core Team, 2020). The use of the Fisher

194 exact test or a chi-squared test would have permitted us to test for every trial regardless of whether the

195 number of flies from a particular category was more or less preyed upon than those from another

196 category. However, we used a hypergeometric distribution instead of a Chi-squared distribution

197 because, as flies were preyed upon, the population size decreased and we only had information on

198 predation rate per treatment at the end of the experiment. The *p*-value for each experimental trial was

added up and compared to the summed *p*-value obtained by simulating random trials. We assessed the

overall *p*-value by determining the number of simulated trials required to have an observed result

(summed *p*-value observed) as probable as the simulation (summed *p*-value simulated). Hence, a *p*-

value of 0.05 indicated that 20 trials would be sufficient to have the same result as the empirical data

by chance, whereas a *p*-value of 0.0001 meant that 10,000 trials would be required to achieve the same

result as the empirical data by chance. We used the Manly Alpha Index as the preference index to

illustrate the selective predation on sick individuals (index 0: preference for healthy individuals, index

0.5: no preference, index 1: preference for sick individuals). It was calculated as log (total flies sick /

flies sick but not eaten) / [(log(total flies sick / flies sick but not eaten) + log(total flies healthy / flies

healthy but not eaten)] as established by Manly (1972). To test whether the preference index for sick

individuals increased with the severity of the cancer, we considered hyperplasia after 10 and 20 days

of induction and cancer after 10 days and 20 days as ordinal variables. This was used because although

we knew the rank of severity (i.e. from inducing hyperplasia for 10 days to inducing cancer for 20

days), we could not know the size of the differences among them. This allowed us to test in a single

model (*i.e.* using an ordinal logistic regression) for the increase of the preference index with the

increase in cancer severity.

Ethical note

All spiders in the study were obtained from the wild during their daily activity and kept for less than a

week, eventually fed with healthy *Drosophila*. Spiders used for testing the effect of cancer were

sampled and tested on the Cornell University campus (Ithaca, NY, USA) and those used for testing the

effect of infectious diseases were sampled and tested on the campus of the University of Toulouse 3

(Toulouse, France). All individuals have been released in the wild at the area they have been caught

shortly after the experiments were performed.

Results

Predation of chronically infected flies by hunting spiders. We tested whether chronic infection of the male flies by *P. rettgeri* affected the risk of being preyed upon by the spiders. The flies had been infected for several days before exposure to the spiders, and their immune response was expected to be activated (Chambers, Jacobson, Khalil, & Lazzaro, 2019). Despite using two predator species, we did not detect selective predation by either of the hunting spider types (Fig. 2A and B for jumping and wolf spiders, respectively).

Predation of flies with cancer by hunting spiders. We asked if cancer of the gut, which we induced, affected the risk for predation of male flies by the hunting spiders. Hyperplasia is the enlargement of an organ or tissue caused by increased cell division and is often an initial stage in the development of cancer. We observed that flies in this early hyperplastic stage ($esg^{ts} > UAS\text{-}Ras^{v12}$) did not show an increased likelihood of getting preyed upon by the jumping spiders, which are active diurnal hunters (Fig. 3A). However, jumping spiders did catch more successfully flies with an advanced stage of cancer ($esg^{ts} > UAS\text{-}Ras^{v12}, UAS\text{-}APC\text{-}IR$) compared with healthy flies. Furthermore, the preference index for sick individuals increased with the length of time after the tumour had been induced (Fig. 3B).

Discussion

Many diseases do not kill their hosts and are considered benign in optimal conditions, such as in a clinic or laboratory. However, it is very likely that non-lethal illnesses still increase the risk of mortality through indirect interactions with other environmental factors. Indeed, being sick may reduce foraging and mating abilities, increase the risk for lethal superinfections, or of being preyed upon. Even though many biological questions are presently being addressed using an integrative approach, the question of whether a non-lethal disease is truly benign in natural conditions seems to have been overlooked in many cases. We hope to generate interest in evaluating the lethality of diseases when their interaction with other risks, such as predation, is considered. Chronic infection by

a wild-caught *Drosophila* pathogen did not significantly affect the risk of being preyed upon in the

laboratory. In contrast, we observed that while the presence of early stage tumours did not change

predation risk, the presence of advanced cancer did compared to age-matched controls.

Little is known about selective predation in individuals with non-infectious diseases. The

prevailing opinion that general health, which can be affected by non-infectious sickness, correlates

with the risk of being preyed upon, is largely observational (Genovart et al., 2010; Hoey &

McCormick, 2004). Nonetheless, although we should not be jumping to hasty conclusions, our results

on the effects of non-infectious disease on selective predation suggests a stronger selective pressure

than natural chronic infections.

Cancers are common in multicellular organisms and can occur whenever cells evade the

normal cell checkpoints that control cell division, proliferation, and apoptosis, thus leading to

uncontrolled cell division (Aktipis et al., 2015). Individuals always have some tumours, most of which

never become lethal (Abu-Helil & van der Weyden, 2019; Bissell & Hines, 2011). Our understanding

of oncogenic phenomena in wild populations of animals is limited, in particular whether tumours that

are clinically non-lethal affect the well-being of organisms in the wild (McAloose & Newton, 2009;

Pesavento, Agnew, Keel, & Woolard, 2018; M. Vittecoq et al., 2015; Marion Vittecoq et al., 2013).

Theoretical modelling further suggests that biotic interactions complicate the predictions regarding the

impact of cancer on populations and that prey and predator populations are likely to suffer differently

(Perret et al., 2020). If tumour formation is inevitable, it is likely that multicellular organisms have

evolved ways to control and tolerate them, perhaps resulting in a trade-off between surviving the

tumour and surviving other perils, such as infections or predation (Pavard & Metcalf, 2019). In some

cases, selection for tumour tolerance may have arisen from selection for tolerance to other factors,

perhaps explaining why early stages of cancer, such as the colon hyperplasia that we induced in

Drosophila, do not change the likelihood of being preyed upon. In our study, when the tumour

developed into a cancer, predation of the flies increased, suggesting that uncontrolled tumours strongly

increase the likelihood of being preyed upon. This conclusion supports the idea that individuals with

uncontrolled tumours will suffer the risk of increased predation compared with healthy individuals.

Predation is a critical force shaping natural selection (Wade & Kalisz, 1990). By removing prey

genotypes susceptible to tumours, predators are likely to have selected for the tumour tolerance largely

observed in nature (Bissell & Hines, 2011). Furthermore, by removing individuals with advanced

tumours, predation may explain why cancerous animals are rarely recorded in autopsies of wild animal

populations (Marion Vittecoq et al., 2013).

279   We do not know why cancerous flies are more predated than their age-matched controls or

even if this result could be to some extend dependent on the sex of the fly. Jumping spiders have been

shown to be able to choose prey based on odour and coloration (Vickers & Taylor, 2018). It would be

difficult to interpret why such behaviour would have evolved in the context of cancer but we cannot

exclude that, spiders recognized and preferred eating cancerous flies. Assays in gut cancerous flies

from another study in another genetic background suggests that induction of colon cancer does not

affect strongly fly locomotion (Dawson et al., 2018). However, there is still the possibility that

cancerous flies have worst reflex than healthy flies or that subtle locomotion differences may be

enough to explain a difference in the chance to escape predation. Our results lay the premise of future

studies on understanding the mechanisms explaining the reasons for this difference.

289   The progression of cancer can, in many ways, be compared with infectious diseases. In fact,

cancer cells have already been considered as a parasitic species consuming the host's resources after

having emerged from healthy cells (Capp & Thomas, 2020; Duesberg, Mandrioli, McCormack, &

Nicholson, 2011). However, if both reduce viability and survival, infectious diseases are, unlike

oncogenic processes, under selective pressure to ensure transmission of the pathogen to another host.

For both non-infectious diseases and cancer, selective predation could increase the selection against

the sickness, as diseases that are moderate and non-lethal could lead to a higher risk of the host being

preyed upon, resulting in mortality rates that are similar to those of infectious diseases. With infectious

diseases, there may be additional consequences. First, selective predation could affect the transmission

of pathogens, spreading them through predator feces over long distances. Second, it might affect

epidemics, either preventing them by removing parasite spreaders from the host population (Duffy et

al., 2005) or by increasing their likelihood of occurrence by dispersing parasites instead of containing

them in dead hosts (Strauss et al., 2016). An additional consideration is that predation may affect the

evolution of parasite virulence. It is generally assumed that predation can reduce virulence because

hosts with rapidly proliferating parasites would suffer more from the infection and may be eaten

before transmitting the parasite to another host, whereas the less rapidly proliferating parasite would

impact less host health and would be transmitted before the host is eaten (Moller et al., 2010).

However, lower virulence could evolve also through lower pathogenicity, that is by inducing less

damage to the host, and not only through reduced proliferation. The most important factor being that

bacteria can proliferate within their host as long as the symptoms do not increase the risk of predation.

The bacteria studied here, *P. rettgeri*, was obtained from the wild and is prevalent in populations of *D.*

melanogaster (Juneja & Lazzaro, 2009). Despite being chronic and the fact that the pathogen can

trigger an immune response through the course of the infection (Chambers et al., 2019; Duneau et al.,

2017), the chronic infection did not increase the predation by the two species of predator we used in

our study. This result supports the hypothesis that chronic infections are selected for such that the

symptoms that enhance predation are reduced.

It is reasonable to assume that the success of predators is not simply due to favourable luck. In

fact, selective predation may be more the rule than the exception and is likely to have a role on the

evolution of diseases, infectious or not (Møller, 2008). On the one hand, selective predation on sick

individuals is likely to counter-select against parasites or genetic diseases more strongly than if

selection occurred upon disease-mediated host death later in life in absence of predation. On the other

hand, in the same way that some parasites evolved to manipulate their host to increase transmission to

intermediate hosts (Hughes & Libersat, 2019), parasites may have evolved to make their hosts less

conspicuous to predation, for example, by interfering with the search for mates or by lessening the

effects of the disease on the host; for example, by reducing symptoms. Because cancer progression

generally results in the death of the organism, we cannot expect the incidence of lethal cancer to have

evolved in the same way as parasites, except for transmissible cancers such as in Tasmanian devils

(Dujon, Bramwell, Roche, Thomas, & Ujvari, 2021). We argue that selective predation on prey with

cancer may be one of the reasons why tumours are commonly tolerated in animals, where advanced

cancer is only rarely detected.

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Figure legends

Figure 1. Photograph of a jumping spider consuming *Drosophila melanogaster* flies in the laboratory.

Figure 2. Predation by hunting spiders on a population of *Drosophila melanogaster* Canton S, of either healthy flies or flies chronically infected with *P. rettgeri*. A. Predation by jumping spiders (Hypergeometric test: n= 27 trials, *p*-value > 0.05, Median pref. index = 0.64) B. Predation by wolf spiders (Hypergeometric test: n= 30 trials, *p*-value > 0.05, Median pref. index = 0.43). Each dot represents the proportion of individuals eaten over five flies and the lines connect the groups of individuals from a same trial. Each trial represents ten flies (five from each type) presented to one spider at once. Preference index is calculated as in Manly (1972) and ranges from 0 to 1, from a preference to healthy flies to a preference for sick flies, respectively.

Figure 3. Predation by jumping spiders in a population of *Drosophila melanogaster* that were either healthy or in which colon tumours (hyperplasia) had been induced. Tumour size increased with induction time, during which uncontrolled cell division occurred. A. Uncontrolled cell division was triggered for 10 or 20 days in *esg^{TS} > Ras^{v12}* flies to lead to hyperplasia (respectively with n= 17 and 15 trials). In both cases, individuals with hyperplasia were not differently predated upon compared with the healthy individuals. B. Fast uncontrolled cell division was triggered for 10 or 20 days in *esg^{TS} > Ras^{v12}, APC-RNAi* flies to lead to cancer (respectively with n= 16 and 20 trials). In both cases, individuals with cancer were more likely to be predated upon than individuals which did not have cancer. This pattern tended to be stronger as the tumour grew larger (Ordinal logistic regression: LRT=8.07, df=1, p=0.004). Each dot in A and B represents the proportion of individuals eaten over five flies and the lines connect the groups of individuals from a same trial. Each trial represents ten flies (five from each type) presented to one spider at once. Preference index is calculated as in Manly (1972) and ranges from 0 to 1, from a preference to healthy flies to a preference for sick flies, respectively.