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Final Version

**Infection and transovarial transmission of rickettsiae in *Dermacentor variabilis*
acquired by artificial feeding**

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RH: INFECTION OF *DERMACENTOR VARILABILIS* WITH RICKETTSIAE

Abstract

In this study we examined the efficiency of an *in vitro* feeding technique using glass microcapillaries as a method of establishing rickettsiae-infected lines of ticks. To quantitate the volume ingested by ticks during microcapillary feeding, the incorporation of radiolabeled amino acids in tick gut and hemolymph was calculated. Fifteen of eighteen ticks consumed between 0.06 μ l and 6.77 μ l. However, ingestion of fluid was not correlated to weight gain during capillary feeding. Partially fed laboratory-reared females, *Dermacentor variabilis*, known uninfected, were exposed to either *Rickettsia montana*- or *R. rhipicephali*-infected Vero cells via microcapillary tubes, returned to rabbit hosts, and allowed to feed to repletion. All tissues collected from ticks allowed to feed overnight on rickettsia-infected fluids were found to be infected when examined by the immunofluorescence antibody assay (IFA). When rickettsia-infected and uninfected capillary-fed ticks were allowed to feed to repletion and lay eggs, no significant differences in mean engorgement weight or fecundity was observed. Assessing the efficiency of transovarial transmission of rickettsiae by ticks that imbibed rickettsia-infected solutions by the polymerase chain reaction (PCR) and IFA, infection was found in 85% of the eggs from ticks exposed to *R. montana* and 69% of the ticks exposed to *R. rhipicephali*. No rickettsial DNA was amplified in samples of the uninfected controls. 100% of egg samples from females exposed to rickettsia-infected fluids and 0% of the uninfected controls examined by IFA were found to be infected

Introduction

Rickettsiae are obligate intracellular bacteria that are transmitted to mammals by a number of arthropod vectors including mites, lice, fleas, and ticks. Most members of the spotted fever group (SFG) rickettsiae are transmitted by ixodid ticks. The SFG include both pathogenic and nonpathogenic rickettsiae. In most cases, the passage or development of tick-borne rickettsiae in a vertebrate host is not essential for survival of the bacteria because rickettsiae are maintained through transstadial and transovarial transmission (TOT) within the arthropod host. However, maintenance of rickettsiae in ticks via TOT may influence their virulence to vertebrate hosts.


To examine the tick-pathogen relationship, ticks must be deliberately infected with rickettsiae. Then, the spread of the bacteria throughout the tick's tissues can be investigated. Historically, the most widely used method to establish known rickettsia-infected ticks has been by allowing them to feed on rickettsemic animals (Burgdorfer 1988; Niebylski et al. 1999). However, important questions about the fate of ingested rickettsiae may be difficult to address due to uncertainty regarding the precise time when or number acquired during natural feeding. ←

Alternatively, ticks can be infected *in vitro* with cultured pathogens by artificial techniques. Artificial feeding offers important advantages, not the least of which is the ability to control the dose of microorganisms or the contents of the feeding medium. Several methods to artificially introduce microbial pathogens into ticks *in vitro* have been successful, including membrane feeding (Kemp et al. 1975; Howarth and Hokama 1983; Voigt et al. 1993; Abbasy et al. 1994; Young et al. 1996), percutaneous injection (Jongejan et al. 1980; Kocan et al. 1986), and capillary feeding (Chabaud 1950;

1 **Bacteria.** *R. montana* and *R. rhipicephali* were maintained in Vero cell monolayers.
2 Vero cells were grown in high glucose (4.5 grams per liter) Dulbecco's Modified eagle
3 Medium (DMEM; Gibco BRL, Grand Island, NY) containing 4% heat-inactivated fetal bovine
4 serum at 37° C in a humidified 5% CO₂ atmosphere. Vero cells were infected with rickettsiae
5 as previously described (Troyer et al. 1999). Briefly, frozen stock (~200 µl) of either *R.*
6 *montana* or *R. rhipicephali* was quickly thawed at 37°C and used to infect one 75-cm² flask
7 (Sarstedt Inc., Newton, NC) of a Vero cell monolayer. Infected monolayers were cultured for
8 7 d, and growth medium was changed daily. Prior to use in capillary feeding assay, cells were
9 harvested and resuspended in growth medium at a concentration of 2.8×10^6 cells per
10 milliliter. Staining with Diff-Quik (Dade Behring, New Castle, DE) showed that 90-95% of
11 the cells were infected with either *R. montana* or *R. rhipicephali*.

12 **Quantifying media ingested by capillary feeding.** In order to quantify the
13 volume of solution ticks were ingesting by capillary feeding, a ¹⁴C (uniformly labeled)
14 labeled amino acid mixture, 54.2 mCi/mmol (New England Nuclear Corp., Boston, MA) ←
15 was diluted in BSK with 6% rabbit serum (Sigma, St. Louis, MO) and fed to ticks in an
16 experiment separate from the Vero cell feeding experiment. Female *D. variabilis*, 5 d
17 post-attachment, were forcibly removed from the rabbit host and restrained ventral side
18 up on double-sided sticky tape in the bottom of a large petri dish. Microcapillary tubes
19 (10 µl) containing the feeding medium with the radiolabeled amino acids (0.0075 µCi/µl)
20 were placed over the entire mouthparts of each tick. The microcapillary tubes were
21 immobilized by embedding them in modeling clay that lined the perimeter of the petri
22 dish. Ticks were allowed to feed from microcapillary tubes for up to 24 h in an incubator
23 at 27° C and 92% RH. Following microcapillary tube feeding, the ticks were washed by

1 repeated immersion in water and subsequent washing of the mouthparts with brushes, to
2 remove coagulated BSK from these body structures. Washing was repeated until no
3 radioactive solution could be found contaminating the body surfaces of the treated ticks.
4 Ticks were then bled by severing the coxa-trochanteral joint of leg 2, far removed from
5 the mouthparts, and pressing gently. Hemolymph was collected with a 10 μ l Drummond
6 micropipette (Drummond Scientific Company, Broomall, PA), changing capillaries
7 between specimens. The amount of hemolymph collected from each tick specimen was
8 recorded. Each hemolymph sample was diluted in buffer and transferred to liquid
9 scintillation vials containing liquid scintillation cocktail and counted for radioactivity.
10 Following hemolymph collection, the ticks were dissected in 0.2 M PBS, and a sample of
11 midgut removed, minced and diluted to 100 μ l, then transferred to liquid scintillation
12 vials. The radioactivity in the hemolymph and midgut samples was recorded using a
13 liquid scintillation counter (LS-1700; Beckman Instruments Inc., Fullerton, CA) with
14 quench correction. To determine the loss of radioactivity due to quench (as a result of the
15 color from the midgut), a quench curve was prepared using midgut from ticks that were
16 not exposed to radiolabeled amino acids. In addition to quench correction, the sample
17 values were corrected for the differences between the ratio of the hemolymph sample
18 volume versus the estimated total body hemolymph volume (Kaufman and Phillips 1973)
19 and the ratio of the gut sample to the total weight of all midgut contents.

20 **Tick infection.** Part-fed (4 d) virgin female ticks were forcibly detached from the
21 rabbit host, individually weighed using a Cahn Microbalance (Cahn Instruments, 
22 Inc., Cerritos, CA) and restrained ventral side up in the same fashion as described
23 earlier. Three groups of ticks were fed a solution consisting of growth medium alone, or

1 growth medium containing Vero cells infected with either *R. montana* or *R. rhipicephali*.
2 Weights of partially fed ticks were taken prior to, and directly after, microcapillary
3 feeding, to determine if weight gain occurred during microcapillary feeding. Five
4 individuals from each treatment group were washed twice in 70% ethanol, dried, and
5 dissected under a stereoscopic microscope and their tissues examined for evidence of
6 rickettsial infection by IFA as described below. The remaining ticks from each treatment
7 were placed back on the rabbit hosts, using individual rabbits for each treatment, and
8 allowed to feed until repletion. After repletion, the ticks were weighed and held
9 individually in numbered vials in an incubator at 27°C and 92% R.H. Seven days after
10 the beginning of oviposition, the egg masses from five randomly-selected female ticks
11 from each group were selected and weighed. Engorgement and egg mass weights
12 recorded for ticks that were exposed to control treatment versus rickettsia-infected Vero
13 cells were compared.

14 **Detection and identification of rickettsiae.** Detection and identification of
15 rickettsial species in Vero cells and tick eggs was determined using PCR, and by analysis
16 of restriction fragment length polymorphism (RFLP) of PCR products. Also, presence of
17 rickettsiae in tick tissue was assessed by IFA. Detection of *R. montana*- and *R.*
18 *rhipicephali*-specific DNA sequences in Vero cells and tick eggs was done by PCR using
19 primers designed from the published sequences for *R. montana* (GenBank accession
20 number AJ238760) and *R. rhipicephali* (accession number AJ238761) *metK*. In addition,
21 amplification of *rompA* gene encoding the SFG-specific 190-kDa protein was also used
22 for species identification. Genomic DNA from rickettsia-infected Vero cells or tick eggs
23 (~100 eggs per tick) was obtained using a genomic DNA purification system (Promega,

1 Madison, WI). For the PCR of *metK*, 2 µl of purified genomic DNA extracted from Vero
2 cells or tick eggs was added to a solution containing 18 µl of PCR Master mix (Roche,
3 Mannheim, Germany), 1 µl each of forward [5' CGT GGA CAG TTA TGG AAG AAT
4 GG 3'] and reverse [5' CCG AAA TGA CCG TAA GAA GCC 3'] primers, and 3 µl of
5 sterile redistilled water. In a PRC Sprint thermal cycler (Hybaid), each 25 µl of sample
6 was heated to 94°C for 3 min, followed by 30 cycles of 94°C for 45 s, 60°C for 30 s, 72°C
7 for 45 s, with an additional incubation period of 72°C for 5 min on the final cycle. The
8 target PCR amplified DNA sequence was visualized by electrophoresis on a 1% agarose
9 gel and stained with ethidium bromide.



10 PCR amplification of *rompA* DNA utilized primers Rr190.70p (Regnery et al.
11 1991) and Rr190.701 (Roux et al. 1999), with the following thermal cycler conditions:
12 initial denaturation for 5 min at 94°C, followed by 35 cycles of 94°C for 30 s, 58°C for 1
13 min, 72°C for 2 min, with an additional incubation period of 72°C for 5 min on the final
14 cycle. The PCR product was visualized on a 1% agarose gel stained with ethidium
15 bromide, excised, and DNA recovered from the gel using a StrataPrep DNA extraction
16 kit (Stratagene, La Jolla, CA) according to manufacturer's protocol. Enzymatic digestion
17 of cleaned PCR product was accomplished by incubating 8 µl of DNA, 1 µl of enzyme
18 buffer, and 1 µl (15 U) of *Rsa*I (Stratagene) for 1 h at 37°C. Digested products were
19 visualized on 8% TBE gels (Novex, San Diego, CA) stained with ethidium bromide.

20 Detection of rickettsia in eggs or the tissues of microcapillary fed ticks by IFA
21 was done according to procedures described by Johns et al. (2000), with minor
22 modifications. Egg samples from each egg mass were smeared onto silylated glass slides
23 (CEL Associates, Houston, TX), air dried, acetone fixed, blocked overnight with 5%

1 dried milk/bovine serum albumin, and incubated with 10 μ l of diluted (1:500)
2 monoclonal antibody 7D11-A2 (kind gift of D. H. Walker, Galveston, TX) specific for
3 rOmpA for 1 h. FITC-conjugated goat anti-mouse (Kirkegaard & Perry, Gaithersburg,
4 MD) diluted 1:40 was added for 1 h, slides were washed with 0.2 M PBS between each
5 step, and viewed by fluorescence microscopy.

6 **Data analysis.** Tick biology parameters were evaluated for statistical
7 significance by 1-way ANOVA with Fisher's least significant difference mean separation
8 test (Steel et al. 1997). A P -value ≤ 0.05 was considered significant.

9 **Results**

10 **Quantification of ingested media.** Labeled amino acids diluted in BSK were
11 ingested by 88% (15/18) of the ticks tested. The volume taken averaged $1.35 \pm 2.15 \mu$ l,
12 including the 3 individuals that did not imbibe any fluid. For the 15 ticks that did 
13 imbibe fluid, the average \pm S.D. was $1.63 \pm 2.27 \mu$ l. There was a great deal of variation
14 in the amount taken up by different ticks, from 0.06 μ l to 6.77 μ l (Table 1). We followed
15 passage of labeled amino acids from the midgut into hemocoel during microcapillary
16 feeding. Of the ticks that did ingest the solution, $0.13 \pm 0.24 \mu$ l of solution, representing
17 ~9.6% of the total volume ingested during microcapillary feeding was in the hemolymph,
18 while $1.23 \pm 1.93 \mu$ l was present in the midgut. Based on the concentration of Vero cells 
19 used for tick infection, part fed ticks consumed between ~170 (0.06 μ l) and ~18,500
20 (6.77 μ l) Vero cells.

21 **Establishment of infected tick lines.** Changes in mean weights of ticks prior to
22 versus post-microcapillary feeding varied between the groups. Ticks exposed to growth
23 media alone had slightly decreased weights (-2.0%), while ticks exposed to *R. montana*-

1 infected Vero cells had slightly increased mean weights (+2.4%), after a 12 h period.
2 **These differences were not statistically significant.** However, the mean weight of ticks ←
3 exposed to *R. rhipicephali*-infected Vero cells (-4.6%) was significantly lower after
4 microcapillary feeding (Table 2).

5 IFA showed that 40.0% of ticks fed either *R. montana*- or *R. rhipicephali*-infected
6 Vero cells were positive for rickettsiae in gut, hemolymph, or salivary gland tissue.
7 Tissues from ticks fed on uninfected growth medium had no evidence of rickettsial
8 infection. Ticks exposed to rickettsiae-infected Vero cells had lower mean engorgement
9 weights (*R. montana* = -16.2%; *R. rhipicephali* = -9.0%), however, the differences
10 between the groups were not significantly different (Table 2). On day seven of
11 oviposition, average weight of the egg masses produced by five ticks of each group was
12 determined (Table 2). Although the ticks exposed to rickettsia-infected Vero cells had
13 decreased average egg mass weights (*R. montana* = -16.1%; *R. rhipicephali* = -21.2%)
14 compared to control ticks, the percent of egg mass weight relative to engorgement weight
15 did not differ significantly between control ticks (36.7 ± 5.2 mg), ticks fed *R. montana*- ←
16 (38.4 ± 12.0 mg), or *R. rhipicephali*-(32.4 ± 8.7 mg) infected Vero cells.

17 **Transovarial transmission of rickettsiae.** In order to determine if ticks exposed
18 to rickettsia-infected Vero cells during microcapillary feeding were capable of
19 transmitting rickettsiae to their progeny, a portion of eggs from individual ticks from each
20 group were assessed by PCR and IFA. The presence of rickettsial DNA was confirmed
21 by PCR using *metK* primers and/or *rompA* primers. Egg samples collected from
22 rickettsiae-infected adult ticks contained amplified rickettsial *metK* fragments with
23 different band sizes for *R. montana* (~350 bp) and *R. rhipicephali*, (~250 bp), indicating

1 that ticks exposed to rickettsiae via microcapillary feeding were capable of passing these
2 organisms to their progeny (Figure 1). Amplification of rickettsial *ompA* resulted in a
3 product for both rickettsial species (~630 bp), and *RsaI* digestion of the PCR products
4 yielded fragments unique to each species (Figure 2). Three bands were observed for *R.*
5 *montana* (~ 300, 200, and 100 bp), as compared to two pairs of bands for *R.*
6 *hipicephali* (200 and 100 bp). Transovarial transmission of *R. montana* was observed
7 in ~85% (11/13) of the ticks exposed to the *R. montana*-infected Vero cells versus
8 69% (11/16) of the ticks exposed to the *R. hipicephali*-infected Vero cells. Samples
9 of eggs from individual ticks were also examined for the presence of rickettsiae by IFA.
10 Egg pools from all the infected lines tested were positive for *R. montana* (11/11) and *R.*
11 *hipicephali* (3/3) by IFA. Control ticks remained negative throughout the
12 experimentation.

13 Discussion

14 Ticks, *D. variabilis*, readily imbibed liquids from glass microcapillaries applied to
15 their mouthparts, thereby resulting in rickettsial infection in these specimens. Moreover,
16 infection was established in the tick's internal tissues within as little as 15 hours, as
17 demonstrated by IFA of their salivary glands and hemolymph. This is the earliest
18 reported translocation of ingested rickettsiae from the midgut to the tick's tissues.

19 The use of radiolabeled amino acids provided a unambiguous measure of fluid
20 uptake by the feeding ticks. In contrast, measuring weight changes before and after
21 feeding does not allow for weight loss due to concomitant salivary secretion. As ticks
22 feed, they alternately suck fluid and salivate, often in an erratic pattern (Gregson 1960).
23 For example, *Theileria parva*-infected *Rhipicephalus appendiculatus* that were fed for 4

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1 days on rabbits secreted infective stage *T. parva* into capillary tubes after a period of 2
2 hours (Purnell and Joyner 1967). In addition, virgin female *Hyalomma asiaticum*,
3 detached after 7-8 days and forced to salivate into glass capillaries, secreted as much as 9
4 μl in 30 minutes (Balashov 1972). Such large volumes of salivary secretion would be
5 likely to compromise any attempt to measure fluid consumption by ticks feeding from
6 fluid-filled capillaries. Therefore, we believe that the volumes shown in Table 1 based on
7 radioisotopic assay are more reliable than similar measurements of artificial feeding
8 based solely on weight changes. Further, attempts to measure tick fluid uptake by
9 measuring fluid loss from the capillaries from which the ticks were imbibing must also
10 consider evaporation from the end of the capillaries. This may occur even in a
11 humidified environment. The use of radioassay measurements eliminates this concern.
12 However, other losses such as spillage of contents around the mouthparts would not have
13 been counted with either method.

14 The volume of fluid imbibed by *D. variabilis* females, as measured by radioassay,
15 is much less than that reported for *Amblyomma americanum* and *D. variabilis* by
16 Burgdorfer (1957) or Rechav et al. (1999), approximately 10 μl per tick, based on weight
17 determinations or fluid displacement. Direct comparisons cannot be made since the latter
18 workers used unfed females. However, 83% of the ticks used in this experiment imbibed
19 some fluid, slightly more than the 71.8% for *Amblyomma americanum* and considerably
20 more than the 61% for *D. variabilis* feeding success reported by Rechav et al. (1999).
21 Another noteworthy finding was the very great degree of variability in the amounts
22 imbibed by different ticks, from as little as 0.06 μl to as much as 6.77 μl ; only 6
23 individuals consumed more than 1 μl . Nevertheless, regardless of how much they

1 consumed, even very small amounts were sufficient to establish infection, as shown by
2 the very high percent success of TOT. Other disparities between the results reported here
3 and that of other workers could be due to the manner of capillary placement. We inserted
4 the capillaries over the mouthparts, including the palps, whereas others inserted them
5 over the hypostome, spreading the palps. However, differences in capillary placement
6 did not modify the feeding ability of *R. appendiculatus* (Purnell and Joyner 1967).

7 In addition to measuring weight gain, capillary feeding allowed us to examine the
8 influence of rickettsial infection on biological parameters of ticks, specifically
9 engorgement weight and fecundity. We observed no significant difference in either
10 engorgement weight or egg mass weight between control and infected female ticks. ←

11 Positive confirmation of rickettsial infection in the eggs post-oviposition supported the
12 hypothesis that the females were infected. These results differ from those reported by
13 Niebylski et al. (1999), in which reduced fecundity was observed in ticks naturally
14 infected with *R. montana* or *R. rhipicephali*, as compared to uninfected ticks. These
15 differences may be due to the fact that the ticks used in the experiments in this study were
16 infected during feeding, while the ticks studied by Niebylski et al. (1999) were infected
17 prior to feeding. Similar findings may arise upon engorgement and egg laying of the
18 progeny of ticks used in this study.

19 The volume of rickettsia-infected medium imbibed by the ticks did not appear to
20 influence infection success. The percent of egg mass infection for *R. montana* (85%) as
21 determined by PCR was almost identical to the percent of ticks that fed (88%), even
22 though 60% imbibed less than 1 μ l. Monitoring of rickettsial infection directly after
23 microcapillary feeding identified a minimum infection rate of 40%. Compared to the

1 higher percentage of TOT observed in this study, the data suggest that infection due to
2 co-feeding on the rabbit host likely occurred. Clearly, ticks will become infected even if
3 they only imbibe very small amounts of rickettsia-infected media. The reason for the
4 lower percentage success with *R. rhipicephali*, 69%, is unknown, but it may be because
5 *D. variabilis* has not been proven to be a natural vector for this rickettsial species
6 (Schreifer and Azad 1994) originally isolated from the brown dog tick, *Rhipicephalus*
7 *sanguineus* (Hayes and Burgdorferi 1979).

8 The association between arthropods and rickettsiae is the result of an evolutionary
9 relationship in which highly adapted rickettsiae coexist with their arthropod host. Tick-
10 borne rickettsiae are somewhat unique in that the bacteria are maintained through
11 transstadial and transovarial transmission, and typically have no significant deleterious
12 effects on the arthropod host itself. Exceptions exist in which the bacteria are capable of
13 manipulating cellular functions of the host (Werren 1997), or even result in mortality of
14 the arthropod host, as in the case of *R. prowazekii* in its vector, the human body louse
15 (Azad 1988).

16 The presence of SFG rickettsia in many tick surveys is to be expected due to the
17 mechanisms of maintenance that occurs within tick populations. One aspect of this
18 relationship that is of interest is the higher occurrence of nonpathogenic rickettsia
19 reported in ticks, compared to *R. rickettsii* in the U.S. The relationship, or competition,
20 of rickettsiae within the arthropod vector is of interest not only from an evolutionary
21 standpoint, but from a pathogen control aspect as well. To date, the mechanisms of
22 intraspecific competition between rickettsiae within the tick vector is unclear.

1 Considering the need for increased knowledge concerning the interaction of rickettsiae
2 within the arthropod vector, known rickettsia-infected lines of ticks must be established.

3 We report here a capillary technique to infect partially-fed *D. variabilis* ticks with
4 either *R. montana* or *R. rhipicephali*, and furthermore report TOT of rickettsiae in the
5 eggs as detected by PCR and IFA. Rickettsiae-infected tick lines have now been
6 established and propagation of these lines is in progress. Having known infected lines of
7 ticks, a technique to infect ticks with a relatively high amount of efficiency, and the
8 ability to differentiate between species using PCR and RFLP analysis, will allow for
9 studies to examine intraspecific competition within the tick vector.

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12
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Table 1. Amount (μ l) of radiolabeled (^{14}C) amino acids in BSK medium ingested by *D. variabilis* via microcapillary feeding*

Tick #	Hemolymph	Midgut	Total
1	0.37	2.32	2.69
2	0.07	0.78	0.84
4	0.02	0.10	0.13
5	0.04	0.57	0.61
6	0.32	2.90	3.22
7	0.00	0.06	0.06
8	0.00	0.08	0.08
9	0.41	6.10	6.51
10	0.01	0.08	0.08
11	0.02	0.16	0.18
12	0.01	0.12	0.14
13	0.93	5.84	6.77
15	0.00	0.06	0.06
17	0.03	1.86	1.89
18	0.07	1.05	1.11
Mean \pm S.D.	0.15 \pm 0.26	1.47 \pm 2.04	1.63 \pm 2.27

* Partially fed ticks were exposed to ^{14}C -labeled amino acids in BSK for ~24 h.

radioactivity present in the hemolymph and midgut was measured, and volume of solution ingested was calculated for individual ticks.

TABLE 2. Mean weight (mg) \pm standard deviation for the biological parameters of *D. variabilis* used in capillary feeding experiment*

Group	Capillary feeding ^a		Engorgement ^b	Egg mass ^b
	Pre-fed weight	Post-fed weight	weight	weight
<i>R. montana</i>	23.9 \pm 8.6	24.5 \pm 9.3	521.0 \pm 141.6 a	189.1 \pm 51.7 a
<i>R. rhipicephali</i>	25.1 \pm 7.8	23.9 \pm 7.4	565.4 \pm 182.0 a	177.6 \pm 59.0 a
Control	19.5 \pm 8.4	19.1 \pm 8.1	621.5 \pm 107.8 a	225.3 \pm 31.3 a

* Partially fed ticks were exposed to growth medium alone (control), or growth medium containing either *R. montana*- or *R. rhipicephali*-infected Vero cells via microcapillary feeding for ~12 h. Weights of ticks were taken prior to, and directly after, microcapillary feeding, and ticks were returned to rabbit host. Post-microcapillary fed weight values in bold are significantly different than pre-fed weights ($P \leq 0.05$). After feeding to repletion, weight of engorged ticks and egg mass (seven day post-oviposition) were recorded. In each column, means followed by the same letter are not statistically significant ($P \geq 0.05$).



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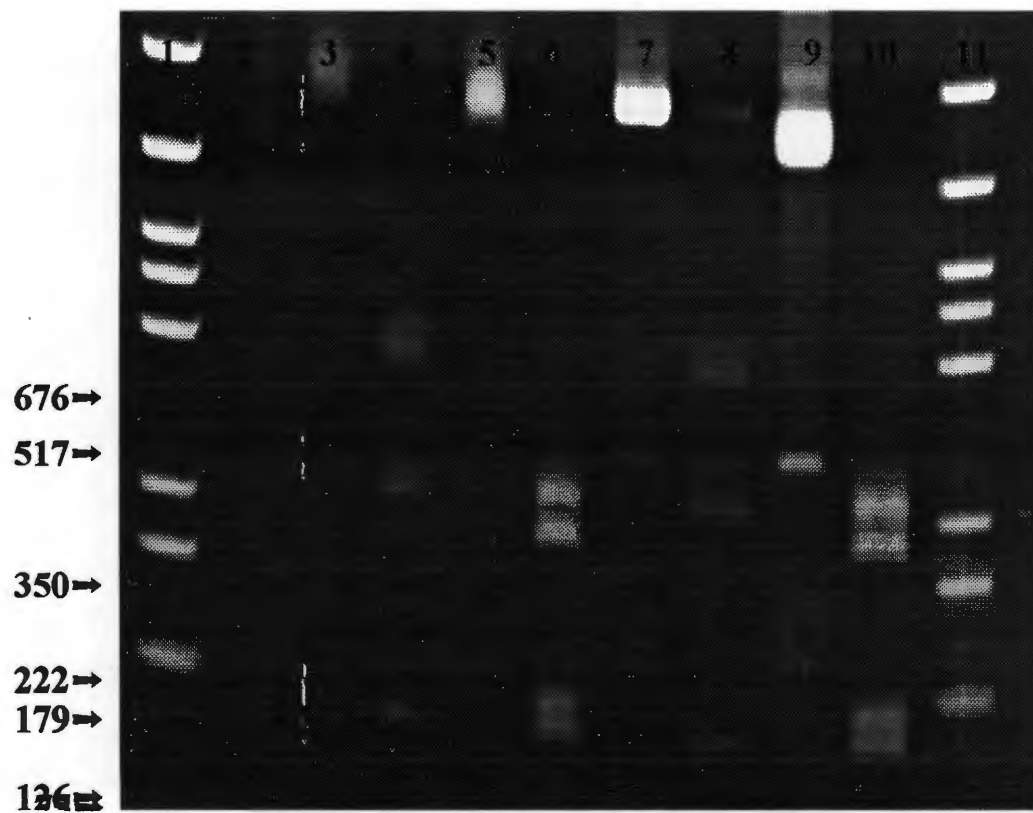


Figure Legends

Figure 1. Representative polymerase chain reaction (PCR) of rickettsial *metK* in eggs from capillary fed *Dermacentor variabilis* ticks. Lane 1, 100-bp marker ladder; Lane 2, uninfected *D. variabilis* eggs; Lane 3, *R. montana*-infected *D. variabilis* eggs; Lane 4, *R. rhipicephali*-infected *D. variabilis* eggs; Lane 5, *R. rhipicephali*-infected Vero cells; Lane 6, *R. montana*-infected Vero cells; Lane 7, H₂O PCR negative control. Numbers on the left are in base pairs.

Figure 2. Representative polymerase chain reaction (PCR) of a 630-bp amplification product of rickettsial *rompA*, and digest with *Rsa*I, in eggs from capillary fed *Dermacentor variabilis* ticks. Lane 1 and 11, D-15 DNA marker (Novex); Lane 2, uninfected *D. variabilis* eggs; Lanes 3 and 4, purified *rompA* amplification product from *R. montana*-infected *D. variabilis* eggs and *Rsa* I digest, respectively; Lanes 5 and 6, purified *rompA* amplification product from *R. rhipicephali*-infected *D. variabilis* eggs and *Rsa* I digest, respectively; Lanes 7 and 8, *rompA* amplification product from *R. montana*-infected Vero cells and *Rsa* I digest, respectively; Lanes 9 and 10, *rompA* amplification product from *R. rhipicephali*-infected Vero cells and *Rsa* I digest, respectively. Numbers on the left are in base pairs.