

1 ***Ixodes scapularis* does not harbor a stable midgut microbiome**

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1 **Supplementary Figures**

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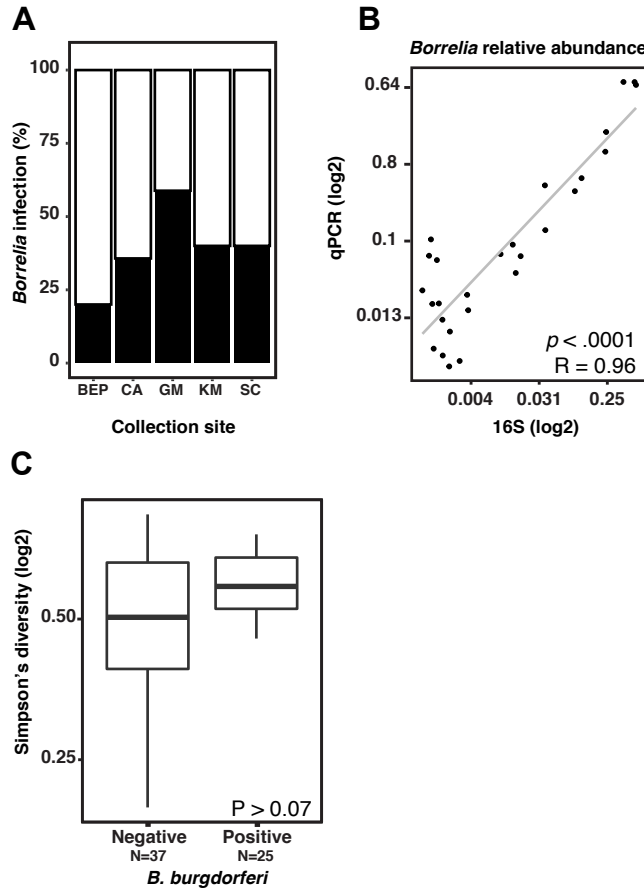
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16 **Figure S1. Characterization of *Borrelia burgdorferi* infection in wild *I. scapularis*.**

17 **(A)** *Borrelia* infection frequency as determined by qPCR for *B. burgdorferi* *flaB* gene, plotted by

18 collection site. **(B)** The relative abundance of *B. burgdorferi* as measured by qPCR with *flaB*

19 primers calculated as a percent of total 16S rRNA gene counts exhibits a strong positive

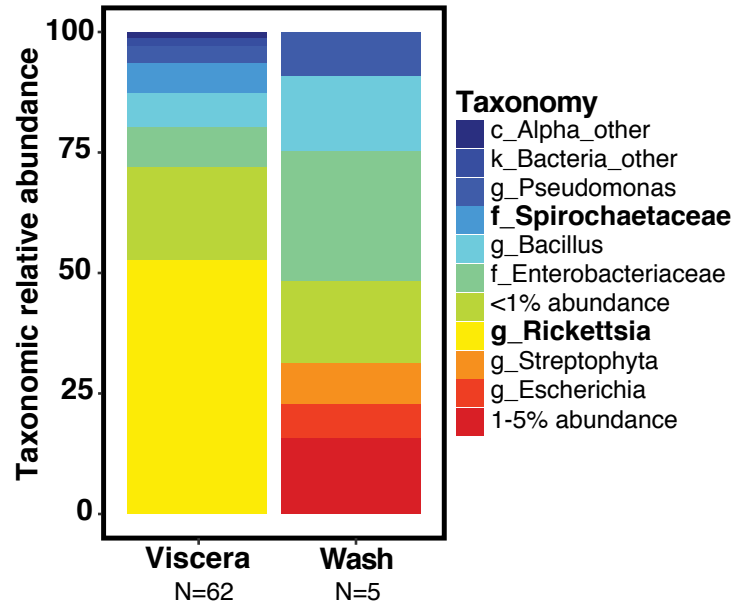
20 correlation between the relative abundance of *Borrelia* by 16S rRNA gene sequencing (Pearson

21 $p < 0.0001$, $r = 0.96$). **(C)** Alpha diversity as quantified by the Simpson's diversity metric,

22 plotted for *B. burgdorferi* infected and uninfected ticks. No significant difference is observed (t-

23 test, $p > 0.07$).

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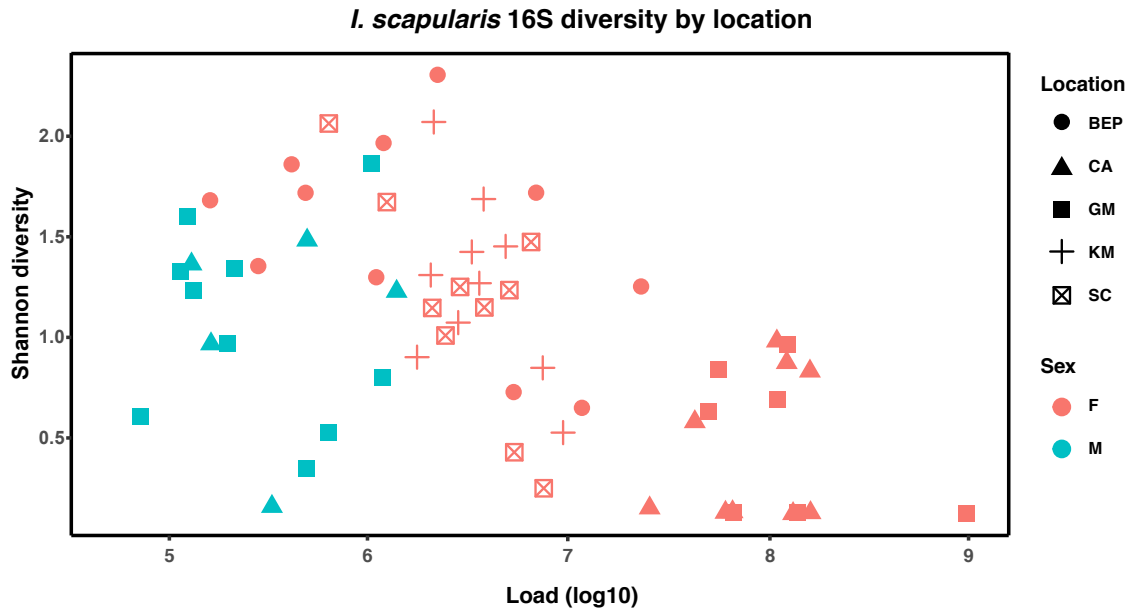
Figure S2. Taxonomic characterization of the *I. scapularis* microbiota.

Relative taxonomic abundance for bacteria detected by 16S rRNA gene sequencing averaged across all adult *I. scapularis* internal viscera samples and external washes. Taxa previously known to be associated with *I. scapularis* are bolded in legend.

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5 **Figure S3. Bacterial load and diversity in adult female *I. scapularis* ticks across sample**
 6 **collection sites.**

7 Shannon diversity (y-axis) and bacterial load as measured by qPCR (x-axis) of adult female *I.*
 8 *scapularis* ticks. Samples cluster by sex, but not by geographic collection site. Big Eau Pleine
 9 (BEP), Carlos Avery (CA), Gordie Mikkelson (GM), Kettle Moraine (KM), Sandberg
 10 Conservancy (SC).

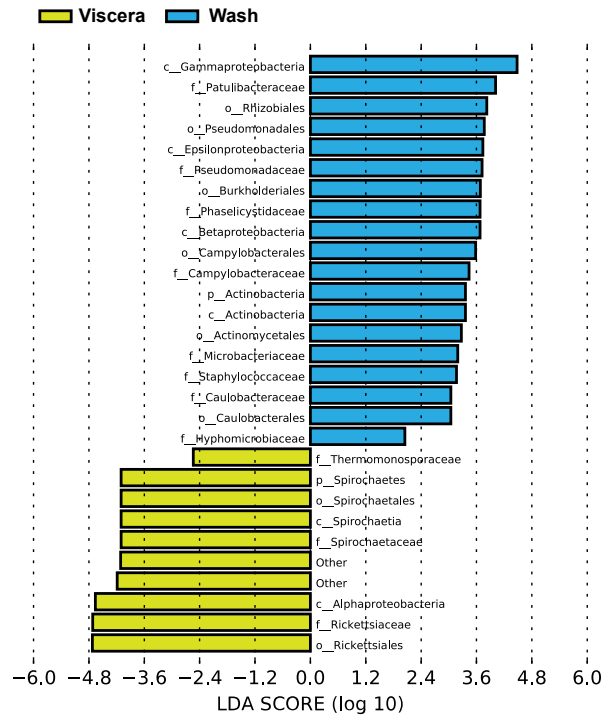
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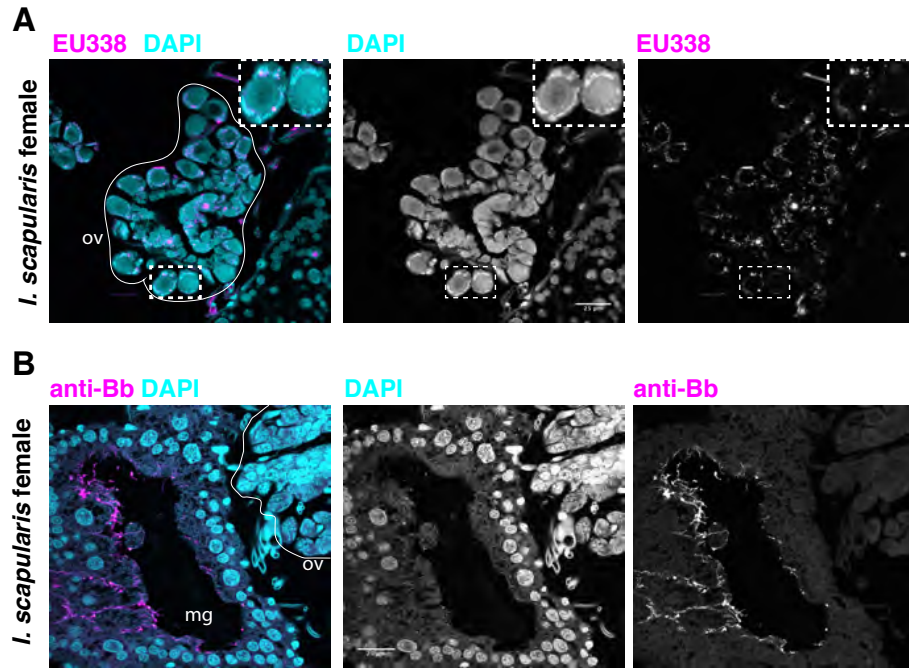
5 **Figure S4. LefSe plot of *I. scapularis* samples.**

6 Taxa enriched in either the viscera (yellow) or the wash samples (blue) as measured by the LDA
7 score are indicated.

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5 **Figure S5. Detection of internal bacteria within *I. scapularis* by confocal microscopy.**

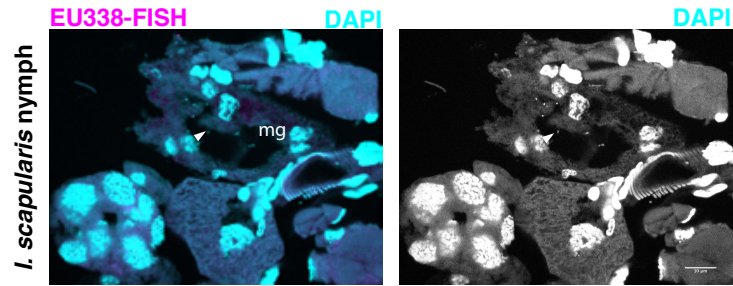
6 **A)** DAPI (blue) and EU338-TSA-ISH (magenta) staining of the central section of the ovary of a
7 female adult *I. scapularis* tick. The ovary is outlined in white and region of inset zoom is
8 indicated by a dashed white rectangle. **(B)** Anti-*B. burgdorferi* (magenta) and DAPI (blue)
9 staining of a midgut cross-section and ovary of an infected *I. scapularis* adult female tick. The
10 midgut lumen is indicated by “mg”, and ovary is outlined in white. Scale bars are 25 microns.

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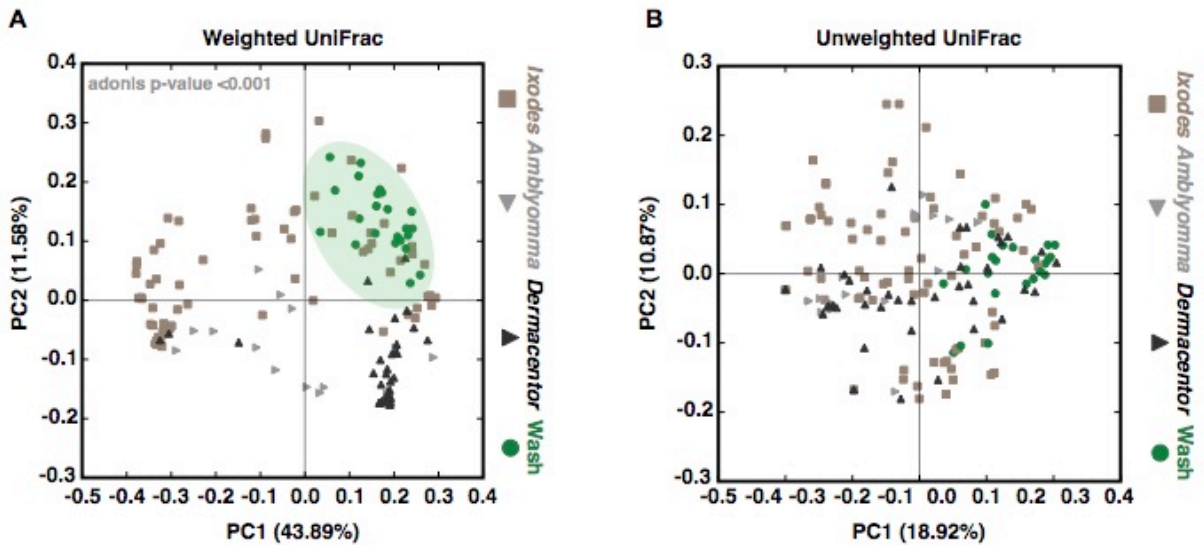
5 **Figure S6. Confocal microscopy of *I. scapularis* nymph.**

6 DAPI (blue) and EU338-TSA-ISH (magenta) staining of an *I. scapularis* nymph. Midgut is
7 indicated by an arrowhead. Scale bar is 10 microns.

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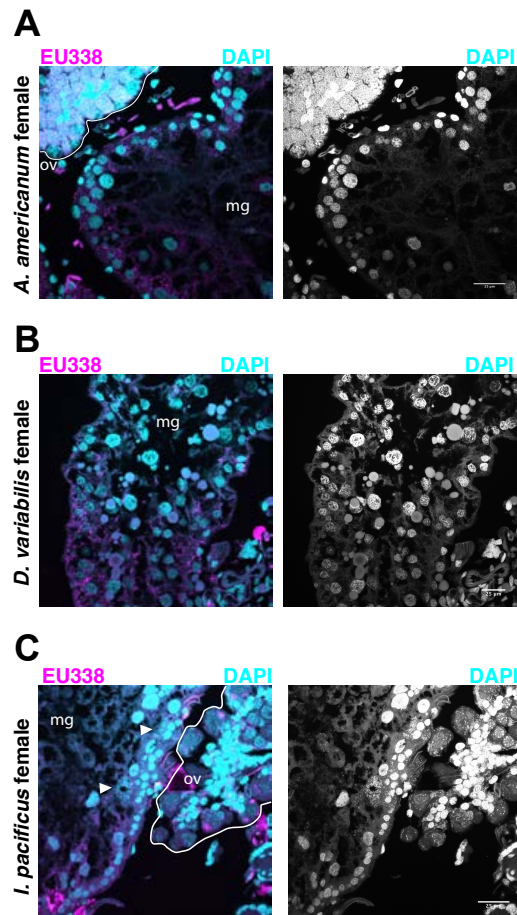
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3 **Figure S7. Beta diversity analysis of adult hard tick samples.**

4 **A)** Beta diversity analysis using weighted UniFrac (which accounts for taxon relative abundance
5 within samples) reveals clustering by external wash samples and by tick species. **B)** Beta
6 diversity analysis using the unweighted UniFrac metric (in which the relative abundance of taxa
7 within samples is not taken into account) shows a lack of species-specific clustering.

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3 **Figure S8. Confocal microscopy of non-*I. scapularis* hard ticks.**

4 **(A)** *Amblyomma maculatum* adult female, **(B)** *Dermacentor variabilis* adult female, **(C)** and

5 *Ixodes pacificus* adult female. All ticks stained with DAPI (blue) and EU338-TSA-ISH

6 (magenta), with the midgut lumen indicated with “mg” and ovary with “ov”. Arrowheads

7 indicate bacterial micro-colonies within midgut. Ovaries are outlined in white in A and C. Scale

8 bars are 25 microns.

Table S1. 84 Spirochaetes, Deferribacteres, tick endosymbiont, and pathogen genomes analyzed in this study.

Genome	Strain	Assembly
<i>Alkalispirochaeta alkalica</i>	DSM 8900	GCF_000373545.1
<i>Anaplasma phagocytophilum</i>	HZ2	GCF_000439755.1
<i>Anaplasma phagocytophilum</i>	HZ	GCF_000013125.1
<i>Anaplasma phagocytophilum</i>	JM	GCF_000439775.1
<i>Anaplasma phagocytophilum</i>	Norway_variant2	GCF_000689635.2
<i>Borrelia afzelii</i>	HLJ01	GCF_000304735.1
<i>Borrelia afzelii</i>	K78	GCF_000962775.1
<i>Borrelia afzelii</i>	PKo	GCF_000222835.1
<i>Borrelia afzelii</i>	Tom3107	GCF_000741005.1
<i>Borrelia burgdorferi</i>	B31	GCF_000008685.2
<i>Borrelia burgdorferi</i>	CA-11.2A	GCF_000172315.2
<i>Borrelia duttonii</i>	Ly	GCF_000019685.1
<i>Borrelia garinii</i>	BgVir	GCF_000239475.1
<i>Borrelia garinii</i>	NMJW1	GCF_000300045.1
<i>Borrelia garinii</i>	SZ	GCF_000691545.1
<i>Borrelia hermsii</i>	CC1	GCF_000956315.1
<i>Borrelia hermsii</i>	HS1	GCF_001660005.1
<i>Borrelia miyamotoi</i>	CT14D4	GCF_000807295.1
<i>Borrelia miyamotoi</i>	LB-2001	GCF_000445425.4
<i>Borrelia persica</i>	No12	GCF_000500045.1
<i>Borrelia parkeri</i>	HR1	GCF_000512145.1
<i>Borrelia turicatae</i>	91E135	GCF_000012085.1
<i>Borrelia recurrentis</i>	A1	GCF_000019705.1
<i>Brachyspira hyodysenteriae</i>	ATCC_27164	GCF_001676785.2
<i>Brachyspira hyodysenteriae</i>	WA1	GCF_000022105.1
<i>Brachyspira intermedia</i>	PWS_A	GCF_000223215.1
<i>Brachyspira murdochii</i>	DSM_12563	GCF_000092845.1
<i>Brachyspira pilosicoli</i>	95_1000	GCF_000143725.1
<i>Brachyspira pilosicoli</i>	B2904	GCF_000296575.1
<i>Brachyspira pilosicoli</i>	P43_6_78	GCF_000325665.1
<i>Caldithrix abyssi</i>	DSM_13497	GCF_000241815.1
<i>Coxiella-like endosymb. of Amblyomma</i>	C904	GCF_000815025.1
<i>Coxiella burnettii</i>	RSA 493	GCF_000007765.1
<i>Ehrlichia chaffeensis</i>	Arkansas	GCF_000013145.1
<i>Ehrlichia chaffeensis</i>	Heartland	GCF_000632815.1
<i>Ehrlichia chaffeensis</i>	Jax	GCF_000632865.1
<i>Ehrlichia chaffeensis</i>	Liberty	GCF_000632885.1
<i>Ehrlichia chaffeensis</i>	Osceola	GCF_000632905.1

<i>Ehrlichia chaffeensis</i>	Saint Vincent	GCF_000632925.1
<i>Ehrlichia chaffeensis</i>	Wakulla	GCF_000632945.1
<i>Ehrlichia chaffeensis</i>	West Paces	GCF_000632965.1
<i>Geovibrio sp. L21</i>	Ace-BES	GCF_000421105.1
<i>Leptospira alstonii</i>	GWTS #1	GCF_001729245.1
<i>Leptospira biflexa</i>	serovar Patoc 1 (Ames)	GCF_000017605.1
<i>Leptospira biflexa</i>	serovar Patoc 1 (Paris)	GCF_000017685.1
<i>Leptospira borgpetersenii</i>	serovar Ballum	GCF_001444465.1
<i>Leptospira borgpetersenii</i>	serovar Hardjo-bovis str. JB197	GCF_000013965.1
<i>Leptospira borgpetersenii</i>	serovar Hardjo-bovis str. L550	GCF_000013945.1
<i>Leptospira interrogans</i>	serovar Bratislava	GCF_001010765.1
<i>Leptospira interrogans</i>	serovar Copenhageni str. Fiocruz_L1-130	GCF_000007685.1
<i>Leptospira interrogans</i>	serovar Hardjo str. Norma	GCF_001293065.1
<i>Leptospira interrogans</i>	serovar Lai str. 56601	GCF_000092565.1
<i>Leptospira interrogans</i>	serovar Lai str. IPAV	GCF_000231175.1
<i>Leptospira interrogans</i>	serovar Linhai str. 56609	GCF_000941035.1
<i>Leptospira interrogans</i>	serovar Manilae	GCF_001047655.1
<i>Leptospira interrogans</i>	Grippotyphosa UI 12764	GCF_000244315.2
<i>Leptospira kirschneri</i>	H1	GCF_000243915.1
<i>Leptospira santarosai</i>	serovar Arenal str. MAVJ 401	GCF_000243835.1
<i>Leptospira santarosai</i>	serovar Shermani str. LT 821	GCF_000313175.2
<i>Rickettsia endosymb. of I. scapularis</i>		GCF_000160735.1
<i>Rickettsia peacockii</i>	Rustic	GCF_000021525.1
<i>Rickettsia belli</i>	RML369-C	GCF_000012385.1
<i>Salinispira pacificus</i>	L21-RPul-D2	GCF_000507245.1
<i>Sediminispirochaeta bajacaliforniensis</i>	DSM 16054	GCF_000378205.1
<i>Sediminispirochaeta smaragdinae</i>	DSM 11293	GCF_000143985.1
<i>Sphaerochaeta globosa</i>	Buddy	GCF_000190435.1
<i>Sphaerochaeta coccoides</i>	DSM 17374	GCF_000208385.1
<i>Sphaerochaeta pleomorpha</i>	Grapes	GCF_000236685.1
<i>Spirochaeta lutea</i>	JC230	GCF_000758165.1
<i>Spirochaeta africana</i>	DSM_8902	GCF_000242595.2
<i>Spirochaeta aurantia</i>	M1	IMG: 2579779150
<i>Spirochaeta thermophila</i>	DSM_6192	GCF_000147075.1
<i>Spirochaeta thermophila</i>	DSM_6578	GCF_000184345.1
<i>Treponema brennaboreense</i>	DSM_12168	GCF_000212415.1
<i>Treponema caldarium</i>	DSM_7334	GCF_000219725.1
<i>Treponema denticola</i>	ATCC 35405	GCF_000008185.1
<i>Treponema pallidum subsp. pallidum</i>	Nichols	GCF_000008605.1
<i>Treponema pallidum subsp. pallidum</i>	Sea 81-4	GCF_000604125.1
<i>Treponema pedis</i>	T A4	GCF_000447675.1
<i>Treponema phagedenis</i>	4A	GCF_000513775.1

<i>Treponema primitia</i>	ZAS-2	GCF_000214375.1
<i>Treponema saccharophilum</i>	DSM 2985	GCF_000255555.1
<i>Treponema succinifaciens</i>	DSM 2489	GCF_000195275.1
<i>Turneriella parva</i>	DSM 21527	GCF_000266885.1

Table S2. Wild-collected hard ticks analyzed in this study.

Genus	Species	Life Stage	Sex	Collected	Sequenced	Analyzed by Microscopy
<i>Amblyomma</i>	<i>maculatum</i>	Adult	Female	20	10	10
<i>Amblyomma</i>	<i>maculatum</i>	Adult	Male	20	10	10
<i>Amblyomma</i>	<i>maculatum</i>	Nymph	N/A	17	N/A	10
<i>Dermacentor</i>	<i>andersoni</i>	Adult	Female	10	10	N/A
<i>Dermacentor</i>	<i>andersoni</i>	Adult	Male	10	10	N/A
<i>Dermacentor</i>	<i>variabilis</i>	Adult	Female	22	10	10
<i>Dermacentor</i>	<i>variabilis</i>	Adult	Male	19	10	9
<i>Dermacentor</i>	<i>variabilis</i>	Nymph	N/A	14	N/A	N/A
<i>Ixodes</i>	<i>pacificus</i>	Adult	Female	37	10	10
<i>Ixodes</i>	<i>pacificus</i>	Adult	Male	25	10	10
<i>Ixodes</i>	<i>scapularis</i>	Adult	Female	14	9	2
<i>Ixodes</i>	<i>scapularis</i>	Adult	Male	16	5	2
<i>Ixodes</i>	<i>scapularis</i>	Nymph	N/A	1	N/A	N/A
<i>Ixodes</i>	<i>scapularis</i>	Adult	Female	13	7	2
<i>Ixodes</i>	<i>scapularis</i>	Adult	Male	22	10	2
<i>Ixodes</i>	<i>scapularis</i>	Nymph	N/A	15	N/A	10
<i>Ixodes</i>	<i>scapularis</i>	Adult	Female	20	10	10
<i>Ixodes</i>	<i>scapularis</i>	Adult	Female	20	10	10
<i>Ixodes</i>	<i>scapularis</i>	Adult	Female	20	10	10
<i>Ixodes</i>	<i>scapularis</i>	Adult	Female	5	N/A	N/A
<i>Ixodes</i>	<i>scapularis</i>	Adult	Male	9	N/A	3

Table S2. Wild-collected hard ticks analyzed in this study.

Genus	Species	Site of Collection	Latitude/Longitude
<i>Amblyomma</i>	<i>maculatum</i>	USA: McPherson Preserve, Stillwater, OK	36.13 N 97.20 W
<i>Amblyomma</i>	<i>maculatum</i>	USA: McPherson Preserve, Stillwater, OK	36.13 N 97.20 W
<i>Amblyomma</i>	<i>maculatum</i>	USA: McPherson Preserve, Stillwater, OK	36.13 N 97.20 W
<i>Dermacentor</i>	<i>andersoni</i>	USA: Klickitat River Canyon, WA	45.78 N 121.20 W
<i>Dermacentor</i>	<i>andersoni</i>	USA: Klickitat River Canyon, WA	45.78 N 121.20 W
<i>Dermacentor</i>	<i>variabilis</i>	USA: Carlos Avery WMA, Columbus, MN	45.29 N 93.13 W
<i>Dermacentor</i>	<i>variabilis</i>	USA: Carlos Avery WMA, Columbus, MN	45.29 N 93.13 W
<i>Dermacentor</i>	<i>variabilis</i>	USA: Carlos Avery WMA, Columbus, MN	45.29 N 93.13 W
<i>Ixodes</i>	<i>pacificus</i>	USA: Klickitat River Canyon, WA	45.78 N 121.20 W
<i>Ixodes</i>	<i>pacificus</i>	USA: Klickitat River Canyon, WA	45.78 N 121.20 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Carlos Avery WMA, Columbus, MN	45.29 N 93.13 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Carlos Avery WMA, Columbus, MN	45.29 N 93.13 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Carlos Avery WMA, Columbus, MN	45.29 N 93.13 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Gordie Mikkelson WMA, East Bethel, MN	45.37 N 93.13 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Gordie Mikkelson WMA, East Bethel, MN	45.37 N 93.13 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Gordie Mikkelson WMA, East Bethel, MN	45.37 N 93.13 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Sandburg Woods Conservancy, Madison, WI	43.13 N 89.31 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Kettle Moraine State Forest Southern Unit, Kewaskum, WI	43.49 N 88.18 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Big Eau Pleine County Park, Mosinee, WI	44.76 N 89.87 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Wolf Creek State Park, Windsor, IL	39.48 N 88.69 W
<i>Ixodes</i>	<i>scapularis</i>	USA: Wolf Creek State Park, Windsor, IL	39.48 N 88.69 W