

# Supplementary Information

1	AGAGTTTGATCATGGCTCAGATTGAAACGCTGGCGCAGGCCTAACACATGCAAGTCGAACGGAGTTATAT AGAGTTTGATCCTGGCTCAGAATGAAACGCTGGCGCAGGCCTAACACATGCAAGTCGAACGGAGTTATAT	70
71	TGTAGCTTGTATGGTATAACTTAGTGGCAGACGGGGGGTAATGTATAGGAATCTACCTAATAGTGGG TGTAGCTTGTATGGTATAACTTAGTGGCAGACGGGGAGTAATGTATAGGAATCTACCTAGTAGTACGG	140
141	AATAATTGTTGAAACGGCAACTAACCGTATACGCCCTACGGGGAAAAATTATTGCTATTAGATGA AATAATTGTTGAAACGGCAACTAACCGTATACGCCCTACGGGGAAAAATTATTGCTATTAGATGA	210
211	GCCTATATTAGATTAGCTAGTTGGGGATAATAGCCCTACCAAGGCAATGATCTAGCTGATCTGAGAG GCCTATATTAGATTAGCTAGTTGGGGATAATAGCCCTACCAAGGCAATGATCTAGCTGATCTGAGAG	280
281	GATGATCAGCCACACTGGAACCTGAGATAACGGTCCAGACTCTACGGGAGGCAGCAGTGGGAATTGGA GATGATCAGCCACACTGGAACCTGAGATAACGGTCCAGACTCTACGGGAGGCAGCAGTGGGAATTGGA	350
351	CAATGGCGAAGCGCTGATCCAGCCATGCCGATGAGTAAGAAGGGCTTGGGGTAAAGCTTTTA CAATGGCGAAGCGCTGATCCAGCCATGCCGATGAGTAAGAAGGGCTTGGGGTAAAGCTTTTA	420
421	GTGAGGAAGATAATGACGGTACTCACAGAAGAAGTCTGGCTAACCTCGTGCAGCAGCCGCGTAATAC GTGAGGAAGATAATGACGGTACTCACAGAAGAAGTCTGGCTAACCTCGTGCAGCAGCCGCGTAATAC	490
491	GGAGAGGGCTAGCGTTTCCGAAATTGGGGCTAAAGGGCGCTAGGGGGATTAGTAAGTAAAAGTG GGAGAGGGCTAGCGTTTCCGAAATTGGGGCTAAAGGGCGCTAGGGGGATTAGTAAGTAAAAGTG	560
561	AAATCCAAGGCTAACCTTGAATTGCTTTAAACTGCTAATCTAGAGATTGAAAGAGGATAGAGGAA AAATCCAAGGCTAACCTTGAATTGCTTTAAACTGCTAATCTAGAGATTGAAAGAGGATAGAGGAA	630
631	TTCTAGTGTAGAGGTGAAATTGCTAAATTAGGAGGAACACCAGTGGCGAAGGGCTCTATCTGGTTCA TTCTAGTGTAGAGGTGAAATTGCTAAATTAGGAGGAACACCAGTGGCGAAGGGCTCTATCTGGTTCA	700
701	AATCTGACGCTGAGGCCGAAGGGTGGGAGCAAACAGGATTAGATAACCTGGTAGTCCACGCTGTAAA AATCTGACGCTGAGGCCGAAGGGTGGGAGCAAACAGGATTAGATAACCTGGTAGTCCACGCTGTAAA	770
771	CGATGAATGTTAAATATGGAAAGTAACTTCTGTATTACAGCTAACCGTAAACATTCCGCTGGGG CGATGAATGTTAAATATGGAAAGTAACTTCTGTATTACAGCTAACCGTAAACATTCCGCTGGGG	840
841	ACTACGGTCGAAGATAAAACTCAAAGGAATTGACGGGGACCCGCACAAGCGGTGGACATGTGGTTA ACTACGGTCGAAGATAAAACTCAAAGGAATTGACGGGGACCCGCACAAGCGGTGGACATGTGGTTA	910
942	CCCCGCACAAGCGGTGGACATGTGGTTA CCCCGCACAAGCGGTGGACATGTGGTTA	970
911	ATTCGACGCCAACCGGAAGAACCTTACCTGGTCTGACATCCATAGAATTATAGAAATAGAAGTGCC ATTCGATGCAACCGGAACACCTTACCA	980
971	ATTCGATGCAACCGGAACACCTTACCTGGTCTGACATCCATAGAATTATAGAAATAGAAGTGCC ATTCGATGCAACCGGAACACCTTACCA	1040
981	TTCGGGAACATGAGACAGGTGCTGCATGGCTGCTCAGCTCGTGTGAAATGTGGGTTAAGTCCC	1050
1041	TTCGGGAACATGAGACAGGTGCTGCATGGCTGCTCAGCTCGTGTGAAATGTGGGTTAAGTCCC	1110
1051	GCAACGAGCGAACCCCTATCCTTGTGACATCGTGTGAAATGTGGGTTAAGTCCC	1120
1111	GCAACGAGCGAACCCCTATCCTTGTGACATCGTGTGAAATGTGGGTTAAGTCCC	1180
1121	AAACCGAGGAAGGTGGGACGACGTCAGTCATCATGCCCTACGACCAGGGCTACACACGTGTACA	1190
1181	AAACCGAGGAAGGTGGGACGACGTCAGTCATCATGCCCTACGACCAGGGCTACACACGTGTACA	1250
1191	ATGGTATACAAAGAGAAGCGACTCCGTAAGATAAGCAACCTCATAAAGTATATGTAGTCCGGACT	1260
1251	ATGGTATACAAAGAGAAGCAACTCTGTAAAGATAAGCAACCTCATAAAGTATATGTAGTCCGGACT	1320
1261	GGAGTCTGCACTCGACTCCACGAAGTCGAATCGTAGTAATCGGGATCAGAATGCCACGGTAATAC	1330
1321	GGAGTCTGCACTCGACTCCACGAAGTCGAATCGTAGTAATCGGGATCAGAATGCCACGGTAATAC	1390
1331	GTTCCCGGGCTTGTACACGCCGCCGTACACCATGGGAGTGGGTTGCAAAAGAAGCAGATATCTAAC	1400
1391	GTTCCCGGGCTTGTACACGCCGCCGTACACCATGGGAGTGGGTTGCAAAAGAAGCAGATATCTAAC	1460
1401	TGTTTAAACAA -- GAGGGAAATCTACTTGTGATTGACTGGGTGAAGTCGAAACAGT 1463	
1461	CGTTTAAAAACGGGGAAATCTACTTGTGATTGACTGGGTGAAGTCGAAACAGT	1525

**Figure S1.** The composition of the chimeric sequence from the aphid *Stomaphis sinisalicis*.

Black represents the chimeric sequence, green represents the *Wolbachia* sequence, and blue represents the *Buchnera* sequences. The red bar indicates different bases in alignment. In the region of the pink square frame, the sequences of *Wolbachia* and *Buchnera* are the same.

The screenshot displays the NCBI BLAST search interface. At the top, it shows the search parameters: Query ID (Id|11635), Description (None), Molecule type (nucleic acid), and Query Length (1504). The Database Name is nr, and the Program used is BLASTN 2.2.28+. The results table lists 21 entries of significant alignments, each with a link to the GenBank record. The columns include Description, Max score, Total score, Query cover, E value, Ident, and Accession number.

Description	Max score	Total score	Query cover	E value	Ident	Accession
Buchnera aphidicola ( <i>Lachnus roboris</i> ) partial 16S rRNA gene	2287	2287	100%	0.0	94%	AJ296756.1
Buchnera aphidicola ( <i>Tuberolachnus salignus</i> ) partial 16S rRNA gene	2270	2270	100%	0.0	94%	AJ296754.1
Buchnera aphidicola ( <i>Lachnus roboris</i> ) 16S ribosomal RNA gene, partial sequence	2261	2261	99%	0.0	94%	FJ655507.1
Buchnera aphidicola ( <i>Nippolachnus piri</i> ) 16S ribosomal RNA gene, partial sequence	2222	2222	99%	0.0	93%	FJ655512.1
Buchnera aphidicola ( <i>Pterochloroides sp.</i> ) 16S ribosomal RNA gene, partial sequence	2218	2218	99%	0.0	93%	FJ655511.1
Buchnera aphidicola ( <i>Nippolachnus piri</i> ) 16S ribosomal RNA gene, partial sequence	2209	2209	99%	0.0	93%	FJ655513.1
Buchnera aphidicola ( <i>Stomaphis fagi</i> ) 16S ribosomal RNA gene, partial sequence	2207	2207	99%	0.0	93%	FJ655501.1
Buchnera aphidicola ( <i>Stomaphis takahashii</i> ) 16S ribosomal RNA gene, partial sequence	2200	2200	99%	0.0	93%	FJ655500.1
Buchnera aphidicola ( <i>Stomaphis sp.</i> ) 16S ribosomal RNA gene, partial sequence	2193	2193	99%	0.0	93%	FJ655498.1
Buchnera aphidicola ( <i>Stomaphis yanonis</i> ) 16S ribosomal RNA gene, partial sequence	2189	2189	99%	0.0	93%	FJ655497.1
Buchnera aphidicola ( <i>Stomaphis sp.</i> ) 16S ribosomal RNA gene, partial sequence	2178	2178	99%	0.0	93%	FJ655499.1
Buchnera aphidicola ( <i>Macrolachnus submacula</i> ) partial 16S rRNA gene	2178	2178	100%	0.0	93%	AJ296755.1
Buchnera aphidicola ( <i>Stomaphis querqus</i> ) 16S ribosomal RNA gene, partial sequence	2169	2169	99%	0.0	93%	FJ655503.1
Buchnera aphidicola ( <i>Stomaphis aphananthe</i> ) 16S ribosomal RNA gene, partial sequence	2154	2154	99%	0.0	92%	FJ655495.1
Buchnera aphidicola ( <i>Stomaphis querqus</i> ) partial 16S rRNA gene	2154	2154	100%	0.0	92%	AJ296753.1
Buchnera aphidicola ( <i>Stomaphis pini</i> ) 16S ribosomal RNA gene, partial sequence	2145	2145	99%	0.0	92%	FJ655503.1
Buchnera aphidicola ( <i>Cinara tujafilina</i> ), complete genome	2141	2141	100%	0.0	92%	CP001817.1

**Figure S2.** The webpage interface that indicates the results of the BLAST search of the chimeric sequence from *Lachnus quercihabitans*.

**Table S1.** The detailed host information and GenBank accession numbers of *Buchnera* download from GenBank.

Subfamily	Species	GB. Number
	<i>Aphis nerii</i>	JQ269567
	<i>Aphis nasturtii</i>	JQ269568
	<i>Aphis craccivora</i>	JQ269569
	<i>Rhopalosiphum maidis</i>	M63247
	<i>Schizaphis_graminum</i>	NR074512
	<i>Acyrtosiphon pisum</i>	NR074159
Aphidinae	<i>Aspidophorodon longicaudus</i>	JQ269566
	<i>Capitophorus hudsonicus</i>	JQ269564
	<i>Diuraphis noxia</i>	M63251
	<i>Macrosiphoniella sanborn</i>	JQ269571
	<i>Myzus persicae</i>	AY849937
	<i>Sitobion miscanthi</i>	HM156635
	<i>Uroleucon sonchi</i>	M63250
Chaitophorinae	<i>Chaitophorus viminalis</i>	M63252
	<i>Sipha elegans</i>	JQ269587
Drepanosiphinae	<i>Yamatocallis tokyoensis</i>	AB064514

**Table S1.** Cont.

<b>Subfamily</b>	<b>Species</b>	<b>GB. Number</b>
	<i>Eutrichosiphum sinense</i>	JQ269580
Greenideinae	<i>Mollitrichosiphum nigrofasciatum</i>	JQ269579
	<i>Greenideinae</i> sp.	JQ269578
Hormaphidinae	<i>Hamamelistes spinosus</i>	JQ269574
	<i>Cinara edulis</i>	FJ655493
	<i>Cinara gudaris</i>	EU334771
	<i>Cinara maghrebica</i>	EU334772
	<i>Cinara pinea</i>	FJ655490
	<i>Cinara tujafilina</i>	EU334773
	<i>Cinara wahtolca</i>	FJ655492
Lachninae	<i>Lachnus roboris</i>	FJ655507
	<i>Nippolachnus piri</i>	FJ655513
	<i>Pterochloroides</i> sp.	FJ655511
	<i>Stomaphis aphanantheae</i>	FJ655495
	<i>Stomaphis fagi</i>	FJ655501
	<i>Stomaphis quercus</i>	FJ655505
	<i>Tuberolachnus salignus</i>	AJ296754
	<i>Myzocallis agrifollicola</i>	JQ269546
Myzocallidinae	<i>Hoplocallis pictus</i>	AJ296759
	<i>Panaphis juglandis</i>	AJ296758
Mindarinae	<i>Mindarus victoria</i>	M63253
	<i>Baizongia pistaciae</i>	NR074609
	<i>Eriosoma lanigerum</i>	DQ418492
Pemphiginae	<i>Melaphis rhois</i>	JQ269582
	<i>Pemphigus populi</i>	AJ296750
	<i>Prociphilus caryae</i>	JQ269572
	<i>Tetraneura caerulescens</i>	AJ296749
Pterocommatinae	<i>Pterocomma populeum</i>	AJ296747
Thelavinae	<i>Thelaxes suberi</i>	AJ296757

**Table S2.** The detailed information and GenBank accession numbers of *Serratia* download from GenBank as outgroups.

<b>Subfamily</b>	<b>Species</b>	<b>GB. Number</b>
Aphidinae	<i>Aphis fabae</i>	GU394001
	<i>Cinara cedri</i>	CP002295
Lachninae	<i>Cinara cedri</i>	EU348324
	<i>Cinara schimitscheki</i>	EU348318