

Table S2. Genotyping result per species. Samples denoted as NA could not be genotyped by conventional PCR.

<b>Species</b>	<b>NA</b>	<b>Tcl</b>	<b>Tcll</b>
<i>Eratyrus mucronatus</i>	1	3	-
<i>Panstrongylus geniculatus</i>	3	14	1
<i>Psammolestes arthuri</i>	11	2	-
<i>Rhodnius colombiensis</i>	10	1	-
<i>Rhodnius pallescens</i>	8	5	-
<i>Rhodnius prolixus</i>	13	18	-
<i>Triatoma dimidiata</i>	-	30	-
<i>Triatoma dispar</i>	1	1	-
<i>Triatoma maculata</i>	1	1	-
<i>Triatoma venosa</i>	20	4	-
<b>Total</b>	<b>66</b>	<b>79</b>	<b>1</b>