

Genetic Diversity of *Bartonella* spp. in Cave-Dwelling Bats and Bat Flies, Costa Rica, 2018

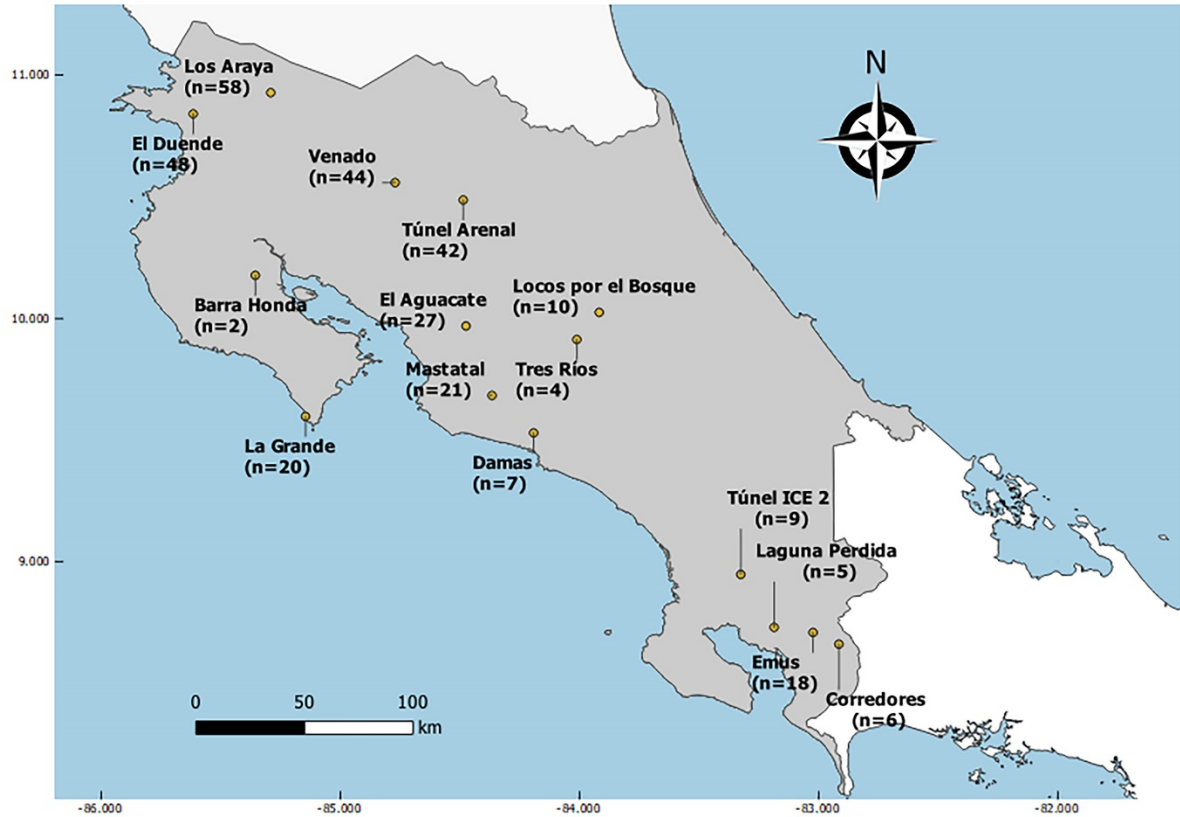
Appendix

Appendix Table. GenBank accession numbers and distribution of 11 genetic variants of *Bartonella* spp. found in bats and bat flies sampled from roost sites in Costa Rica

Accession no.	Roost	Host species	Clade
MW115627	Emus	<i>Carollia perspicillata</i>	IX
MW115628	Túnel Arenal	<i>Carollia perspicillata</i>	IX
MW115629	Emus	<i>Carollia perspicillata</i>	VII
MW115630	El Duende	<i>Carollia perspicillata</i>	VII
MW115631	El Duende	<i>Carollia perspicillata</i>	VII
MW115632	Mastatal	<i>Carollia perspicillata</i>	VIII
MW115624	Emus	<i>Carollia perspicillata</i>	IV
MW115625	Túnel ICE 2	<i>Carollia perspicillata</i>	IV
MW115626	Mastatal	<i>Trichobius pallidus</i> on <i>C. perspicillata</i>	IV
MW115635	Tres Ríos	<i>Megistopoda aranea</i> on <i>Artibeus jamaicensis</i> *	I
MW115637	Los Araya	<i>Trichobius lionycterdis</i> on <i>Lonchorhina aurita</i>	III
MW115636†	Emus	<i>Trichobius pallidus</i> on <i>Pteronotus mesoamericanus</i>	II
MW115633	Los Araya	<i>Trichobius pallidus</i> on <i>Pteronotus mesoamericanus</i>	V
MW115634†	Laguna Perdida	<i>Exastinion clovisi</i> on <i>Anoura cultrata</i> *	VI

*Blood sample not collected.

†Newly described genotype with a similarity value of 93.2%.



Appendix Figure. Roost sites and number of bats sampled per roost for *Bartonella* spp. in Costa Rica. Map was created using qGIS 3.13.8 software (<https://www.qgis.org/en/site>).