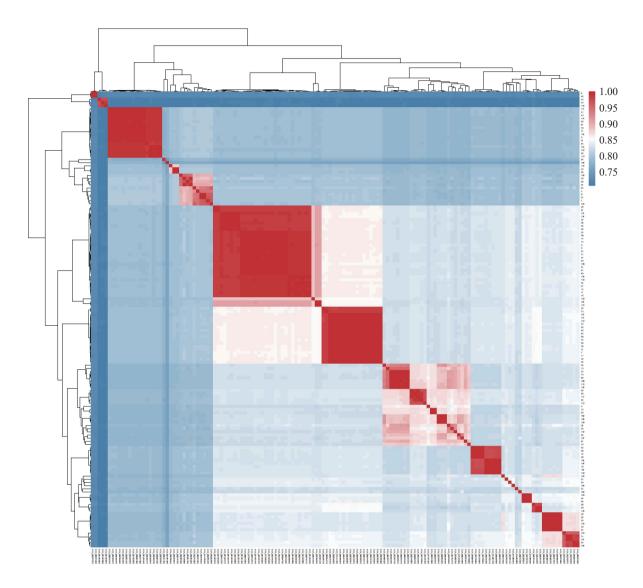
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SUPPLEMENTARY FIGURE S1. ANI analysis of the 144 whole-genome *Bartonella* spp. sequences from GenBank. Notes: The values generated by the Gegenees software Version 2 shown in the heat map indicate the percentage similarity between the analyzed genomes. The colors vary from blue (low similarity) to red (high similarity).