

**Supplementary Table S1.** The number of carbapenemase-encoding genes in carbapenem-resistant *Citrobacter* spp. isolates

<b>Isolates</b>	<b><i>bla</i><sub>KPC</sub></b>	<b><i>bla</i><sub>NDM</sub></b>	<b><i>bla</i><sub>IMP</sub></b>	<b><i>bla</i><sub>VIM</sub></b>	<b><i>bla</i><sub>OXA-48</sub></b>	<b><i>bla</i><sub>KPC</sub>+<i>bla</i><sub>NDM</sub></b>	<b><i>bla</i><sub>KPC</sub>+<i>bla</i><sub>NDM</sub>+<i>bla</i><sub>OXA-48</sub></b>	<b><i>bla</i><sub>NDM</sub>+<i>bla</i><sub>OXA-48</sub></b>	<b><i>bla</i><sub>NDM</sub>+<i>bla</i><sub>VIM</sub></b>
<i>C. freundii</i> (N=32)	4	19	1	1	1	3	1	1	1
<i>C. koseri</i> (N=3)	3	0	0	0	0	0	0	0	0
<i>C. braakii</i> (N=3)	0	2	0	0	0	1	0	0	0
<i>C. amalonaticus</i> (N=2)	0	2	0	0	0	0	0	0	0
<i>C. youngae</i> (N=2)	1	1	0	0	0	0	0	0	0

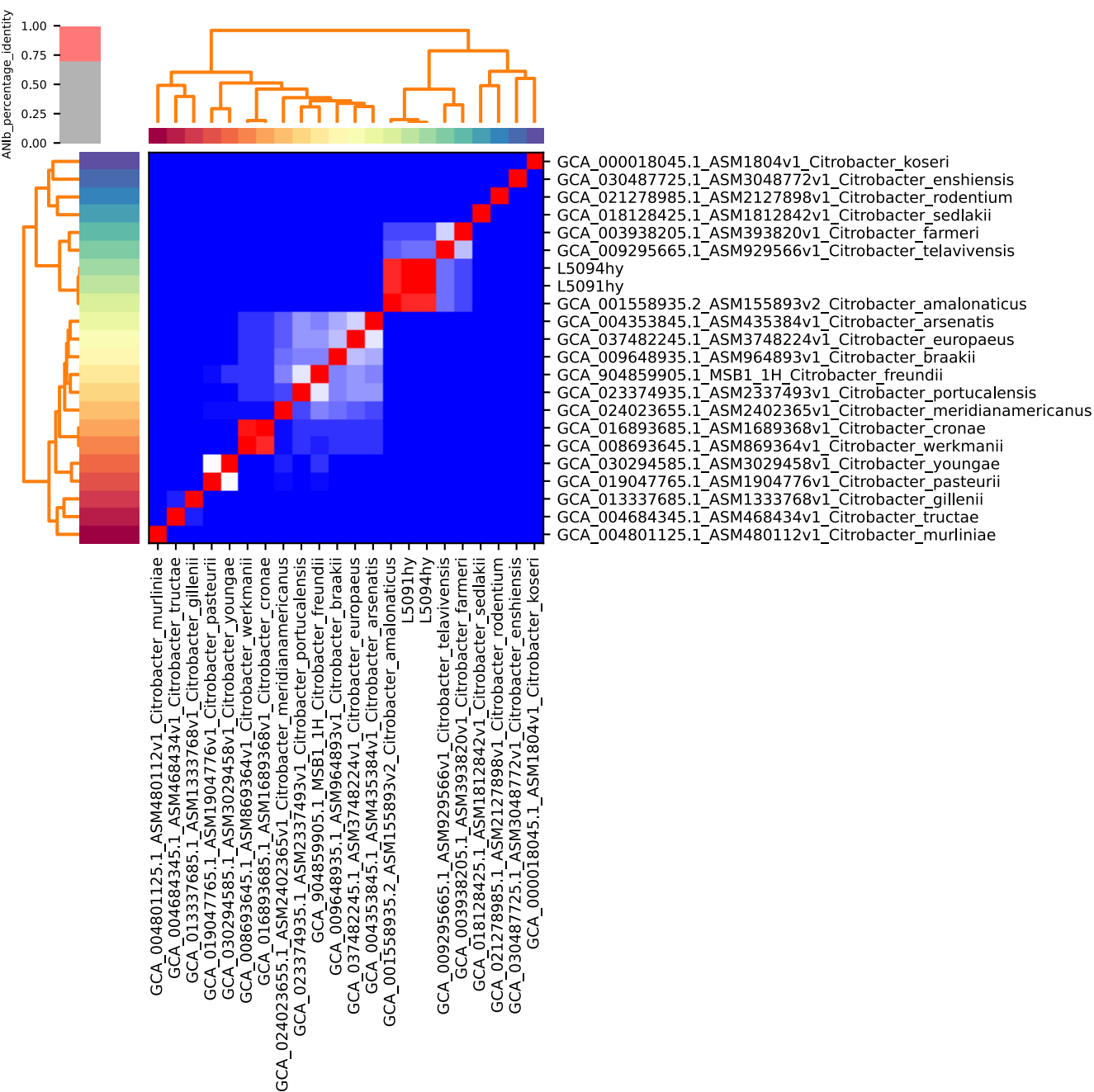


Figure S1. Average nucleotide identity analysis of two *C. amalonaticus* isolates (L5091hy and L5094hy) and type strains of the genus *Citrobacter*.