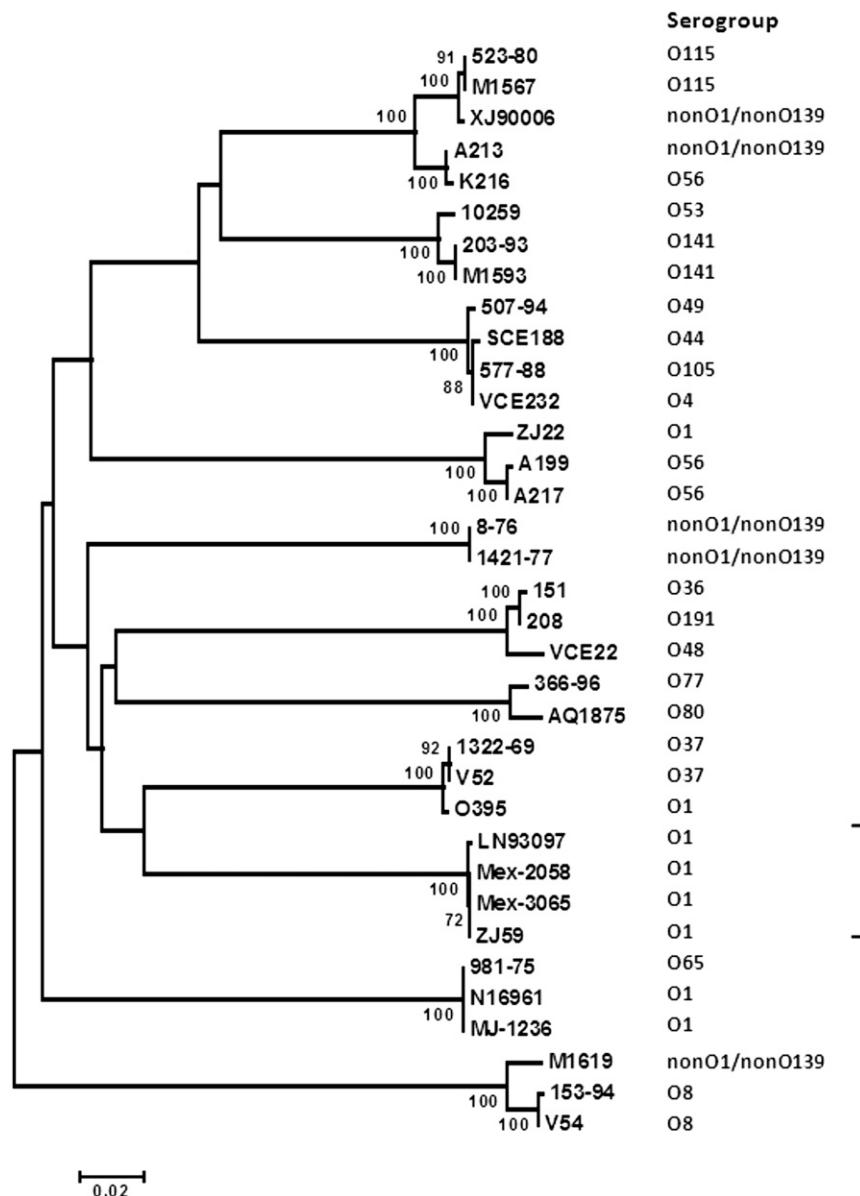


# Supporting Information

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**Fig. S1.** The *tcpA* phylogenetic tree was constructed using neighbor joining. Mex-2058 and Mex-3065 strains represent the TcpA variant of *Vibrio cholerae* O1 biotype El Tor isolated in Mexico during 2000–2004 and share the cluster with TcpA of *V. cholerae* O1 strains isolated in China obtained from GenBank. All other TcpA sequences were from GenBank. Bootstrap values are percentages of 1,000 replications and, if 50% or greater, are indicated at the nodes.

**Table S1. Genetic arrangement of CTXΦ and flanking region of representative *ctx*<sup>+</sup> strains (2004–2008)**

Year of isolation	No. of strains (representative)	RS1		Primer sets targeting TLC-CTXΦ-RTX region							
		<i>rstC</i> (238 bp)	<i>rstAF-rstCR</i> (~1.6 kb)	CIIF-CIIR (0.75 kb)	TLC3F-rstAR (~1.8 kb)	<i>ctxAF-RTX5R</i> (~1.4 kb)	<i>ctxBF-cepR</i> (~3.0 kb)	<i>rstAF-rstR</i> <sup>ET</sup> R (~2.8 kb)	<i>cepF-rstR</i> <sup>ET</sup> R (~1.2 kb)	<i>cepF-rstAR</i> (~2.2 kb)	<i>cepF-rstBR</i> (~2.7kb)
2004	2	–	–	+	+	+	–	+	+	+	+
2005	2	–	–	+	+	+	–	+	+	+	+
2006	2	–	–	+	+	+	–	+	+	+	+
2007	2	–	–	+	+	+	–	+	+	+	+
2008	2	–	–	+	+	+	–	+	+	+	+
N16961 (ET Reference)	+	+	+	+	+	–	–	–	–	–	–
O395 (CL Reference)	–	–	–	+	+	–	–	–	–	+	+
B-33 (Mozambique Variant)	–	–	–	–	–	+	–	–	–	–	–

+, positive; –, negative.