Table S4. Clinical *Vibrio cholerae* strains used as amplification controls in PCR typing of virulence genes.

Strain	Species	Year	Geographical origin	Type of infection	Reference	Genes/Features[1]
VN-00211	*Vibrio cholerae* non-O1, non-O139	2011	Italy	Diarrhea	RKI	*rtxA*[2]
VN-00300	*Vibrio cholerae* non-O1, non-O139	1999	Germany	Diarrhea, peritonitis	BfR	*chxA*
VN-00303	*Vibrio cholerae* non-O1, non-O139	2012	Hungary	Otitis	RKI	negative for $hlyA^{CL}$, $hlyA^{ET}$
VN-00308	*Vibrio cholerae* non-O1, non-O139	2012	Romania	Diarrhea	RKI	TTSS *vcsC2*/*vcsN2*/*vspD*/*vcsV2*; *rtxC*
VN-00315	*Vibrio cholerae* non-O1, non-O139	2011	Cuba	Diarrhea	AGES	VSP-2
1576	*Vibrio cholerae* O1	1959	Thailand	Diarrhea	BfR	classical biotype; $rstR^{CL}$
1360	*Vibrio cholerae* O1	1959	Thailand	Diarrhea	BfR	classical biotype; *toxR*; *ctxA*; O1 *rfb*; $hlyA^{CL}$; *ompU*; $tcpA^{CL}$, $tcpA^{ET}$; negative for VSP-1, VSP-2
MO45	*Vibrio cholerae* O139	1992	India	Diarrhea	BfR	El Tor biotype; *toxR*; *ctxA*; O139 *rfb*; $hlyA^{ET}$; *mshA*; $tcpA^{ET}$; VSP-1; $rstR^{ET}$

AGES: Austrian Agency for Health and Food Safety, Vienna, Austria
BfR: Federal Institute for Risk Assessment, Berlin, Germany
RKI: Robert Koch Institute, Berlin, Germany

CL, Classical; ET, El Tor.

[1] Only genes/features relevant to PCR typing of virulence genes are given.
[2] VC1451 of *Vibrio cholerae* O1 biovar El Tor str. N16961.