



Supplemental Digital Content 1. Phylogenetic tree of Coxsackie A16 (CA16) VP1 partial sequences (303 bp long) downloaded from GenBank (n=22, 28 June 2022), showing the relative position of the ‘Kerala tomato flu’ sequence (India 2022, red). Separate clades can be seen from Thailand (2000-2005, blue) and China (2011-2014, green). A single sequence from France (2006, purple) is shown as part of the China clade. The sequences were aligned and manually edited with BioEdit (v7.2.5), constructed under a general-reversible-time (GTR) model using FastTree (v2.1.11. double precision version) (<http://www.microbesonline.org/fasttree/>), and displayed using FigTree v1.4.4 (<http://tree.bio.ed.ac.uk/software/figtree/>).