

CSv2MF7-1

[illegible]

[illegible]

FIGURE S2 | Wrapped multiple sequence alignment of CSv24F/CSv24R targeting regions in 112 ciliates (16 classes) and 28 non-ciliate eukaryotes species viewed by Jalview v2.11.5.1. The sequences within the red dashed boxes in the first and last rows represent CSv24F and CSv24R, respectively. The aligned sequences from 112 ciliates (16 classes) and 28 non-ciliate eukaryotes are distinguished by different group colors. For all rows except the first and last (primer sequences), the leftmost columns show the 18S rRNA gene GenBank accession number and product length, and the rightmost column shows the sequence length of each row.