

Figure S1

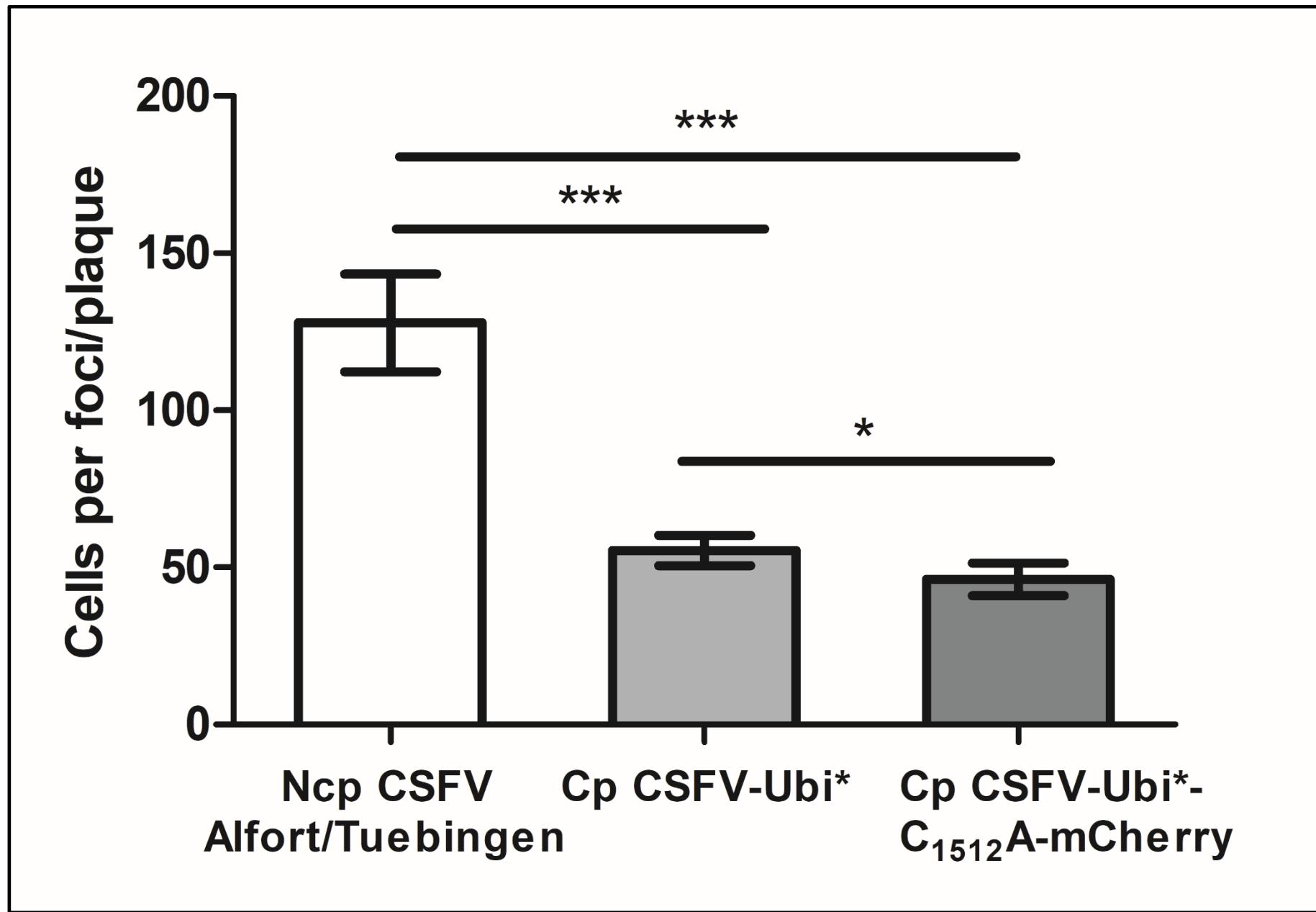
CSFV-7381\_for

5' -GCTCAGGGGGATGTGCAGAGATGTGTGGAAGCAGTGACCAATTATGCGAGAGAGGGCATCCAATTCATGAAGTCGCAGGCAGTGAA  
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TAGGGGCTGTAGCAGCCCACAATGCAATTGAAGCCAGTGAGCAAAAAAGAACACTACTTATGAAAGTCTTGTGAAAAACTTCTTAGAC  
CAGGCCGCCACCGACGAACTAGTCAAAGAGAGCCCTGAGAAAATAATAATGGCTTGTGAAGCGGTGCAAACGGTGGCAACCCCTCT  
TAGATTAGTGTACCACCTCTATGGAGTTTCTATAAAGGGTGGGAAGCAAAAGAGTTGGCCCAAAGAACAGCCGGCAGGAACCTTTCA  
CCTTGATAATGTTCGAGGCTGTGGAACACTGGGAGTAGACAGTGAGGGAAAAATTGCCAGCTA-3'

CSFV-8421\_rev

**Figure S1: Sequence of the NS4B gene from cp CSFV-Ubi\* after passage 5.** After virus infection, the surviving cells were trypsinized and total RNA extracted. An RT-PCR amplicon was generated using oligonucleotides CSFV-7381\_for and CSFV-8421\_rev. The PCR product was inserted into a T-vector and sequenced using M13 and M13 rev. A pinpoint reversion by homologous recombination to the parental ncp CSFV strain was documented.

Figure S2



**Figure S2: Focus and plaque size of the different CSFV strains.**

After infection of SK-6 cells, the supernatant was removed and replaced by fresh DMEM containing 1% carboxymethyl cellulose. The infected cells of 50 individual foci were counted for each virus strain, the mean focus size was calculated, and the 95% confidence interval was determined. The values were compared in the unpaired T-test.

**A** SVNA for positive serum 2016/01/0441/027

|          | Cp CSFV-Ubi*-C <sub>1215</sub> A-mCherryE2 |         |      | Ncp CSFV Alfort/Tuebingen |      |
|----------|--|---------|------|---------------------------|------|
|          | Bright field                               | mCherry | FITC | Bright field              | FITC |
| 1:4      |  |         |      |                           |      |
| 1:16     |  |         |      |                           |      |
| 1:64     |  |         |      |                           |      |
| 1:256    |  |         |      |                           |      |
| 1:1,024  |  |         |      |                           |      |
| 1:4,096  |  |         |      |                           |      |
| 1:16,384 |  |         |      |                           |      |
| 1:65,536 |  |         |      |                           |      |

**Figure S3: Documentation of the fluorescence-verified plaque reduction assay.** **(A)** Low positive serum 2016/01/0441/027. **(B)** Low positive serum 2008/03/0225/032. **(C)** High positive serum 2006/07/0056/086. **(D)** Negative control serum. The sera were tested against the viruses cp CSFV-Ubi\*-C1512A-mCherryE2 (left) and ncp CSFV Alfort/Tuebingen (right). Indicator cells were fixed 48 h post infection and stained with A18 anti E2 and FITC conjugate. A representative area of each 96-well was documented in brightfield, red and green fluorescence channels at 4x magnification. A scale bar (500 µm) is used for comparison between images.

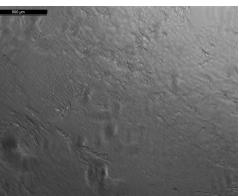
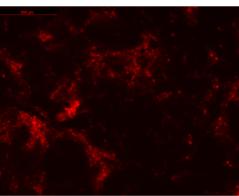
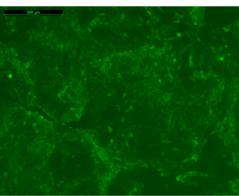
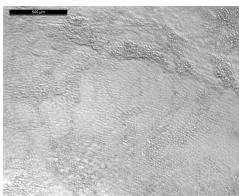
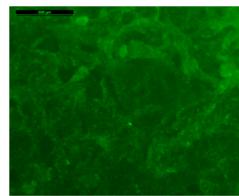
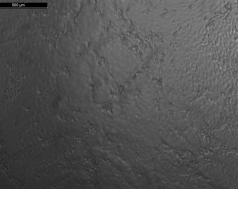
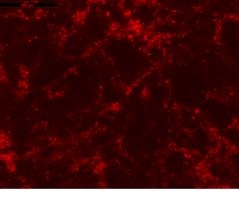
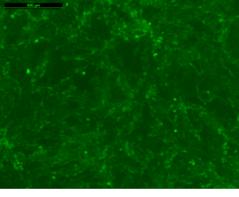
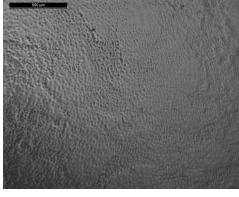
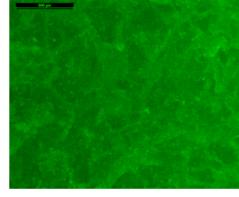
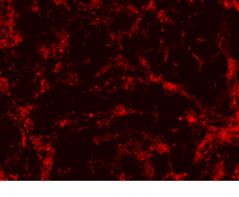
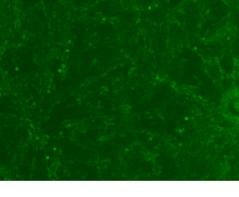
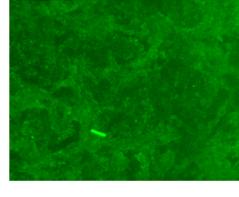
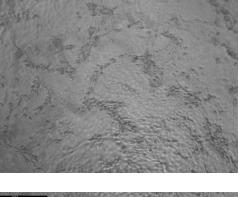
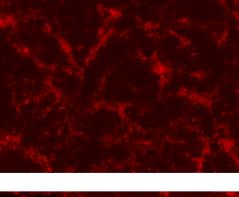
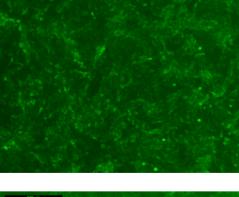
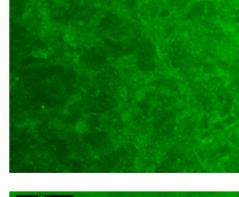
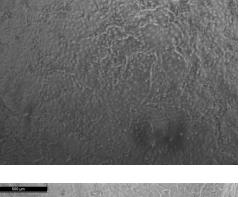
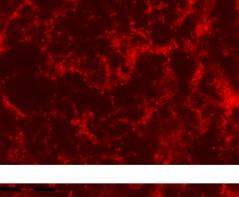
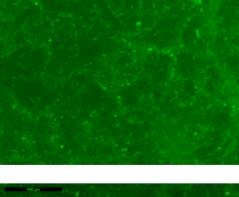
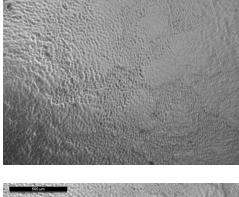
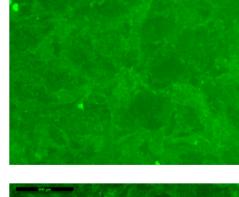
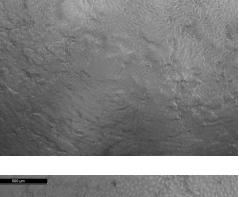
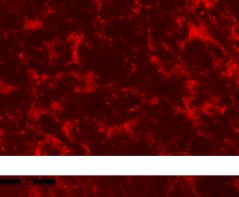
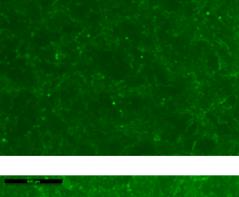
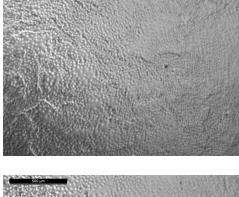
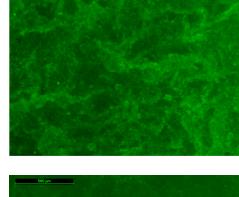
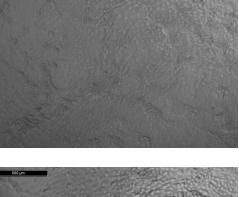
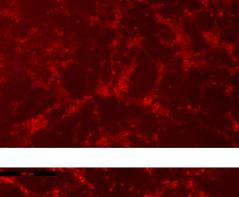
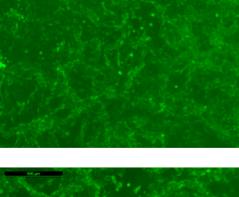
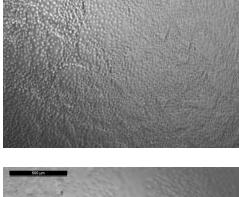
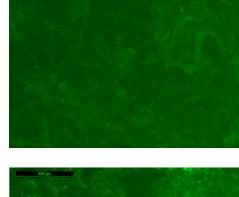
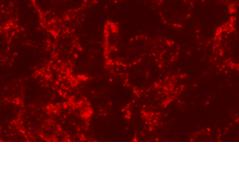
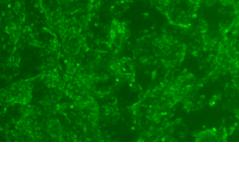
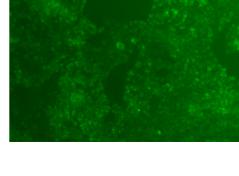
## B SVNA for positive serum 2008/03/0225/032

|          | Cp CSFV-Ubi*-C <sub>1215</sub> A-mCherryE2 |         | Ncp CSFV Alfort/Tuebingen |              |      |
|----------|--|---------|---------------------------|--------------|------|
|          | Bright field                               | mCherry | FITC                      | Bright field | FITC |
| 1:4      |  |         |                           |              |      |
| 1:16     |  |         |                           |              |      |
| 1:64     |  |         |                           |              |      |
| 1:256    |  |         |                           |              |      |
| 1:1,024  |  |         |                           |              |      |
| 1:4,096  |  |         |                           |              |      |
| 1:16,384 |  |         |                           |              |      |
| 1:65,536 |  |         |                           |              |      |

# C SVNA for positive serum 2006/07/0056/086

|          | Cp CSFV-Ubi*-C <sub>1215</sub> A-mCherryE2 |         |      | Ncp CSFV Alfort/Tuebingen |      |
|----------|--|---------|------|---------------------------|------|
|          | Bright field                               | mCherry | FITC | Bright field              | FITC |
| 1:4      |  |         |      |                           |      |
| 1:16     |  |         |      |                           |      |
| 1:64     |  |         |      |                           |      |
| 1:256    |  |         |      |                           |      |
| 1:1,024  |  |         |      |                           |      |
| 1:4,096  |  |         |      |                           |      |
| 1:16,384 |  |         |      |                           |      |
| 1:65,536 |  |         |      |                           |      |

## D SVNA for the negative control serum

|          | Cp CSFV-Ubi*-C <sub>1215</sub> A-mCherryE2  |   |  | Ncp CSFV Alfort/Tuebingen   |   |
|----------|---|---|--|---|---|
|          | Bright field  | mCherry   | FITC   | Bright field  | FITC  |
| 1:4      |    |    |    |    |    |
| 1:16     |    |    |    |    |    |
| 1:64     |    |    |    |    |    |
| 1:256    |   |   |   |   |   |
| 1:1,024  |  |  |  |  |  |
| 1:4,096  |  |  |  |  |  |
| 1:16,384 |  |  |  |  |  |
| 1:65,536 |  |  |  |  |  |