Title: Slow viral propagation during initial phase of infection leads to viral persistence in mice

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IFN-γ



Supplementary Figure 1: TNF-α and IL-β production *in vivo* is limited during LCMV infection.

C57BL/6 mice were infected with LCMV-WE, LCMV-Docile, or co-infected. (a) T cell tetramer gating strategy of Figure 1a-d. Representative FACS blot of Figure 1F are shown for IFN- γ (b) and (c)TNF- α staining. At the indicated time points post-infection, (d) blood cells, (e) singly cell suspended splenocytes were re-stimulated with LCMV-specific CD4⁺ T cell epitopes as indicated followed by staining for IFN- γ and TNF- α (n=9-11). (f) Serum neutralizing antibodies against LCMV were determined at day 20 post-infection (n=4). (g) Serum LCMV GP specific binding antibodies were quantified at day 20 post-infection (n=4). (Error bars show SEM, **p <0.01, ***p < 0.001, ####p<0.001, and ns indicates statistically not significant between the indicated groups).



Supplementary Figure 2: Viral presence positively correlates with T cell exhaustion.

(a) PD-1 or TIM3 expression of splenic tet-gp33 from Figure 1E were plotted against spleen LCMV titers from Figure 1H. (b) IFN- γ or TNF- α producing CD8⁺ T cells in response to LCMV gp33 peptides (Figure 1g) were plotted against spleen LCMV titers (Figure 1h). (p<0.001 indicates slope significantly non-zero).





Supplementary Figure 3: cDC and pDC gating strategy of Figure 2c



Supplementary Figure 4: LCMV-Docile infection led to defective IFN-I production.

C57BL/6 mice were infected with $2x10^4$ pfu (**a**+**c**+**d**) or $2x10^6$ pfu (**b**+**e**+**f**) LCMV-WE, LCMV-Docile, co-infected WE and Docile or Clone 13. (**a**-**b**) At indicated days post-infection, serum IFN- α (left panel) or IFN- β (right panel) were quantified (n=4). (**c**+**e**) At day 8 or day 20 post-infection, blood cells were re-stimulated with LCMV-specific CD8⁺ T cell epitopes as indicated or left untreated (negative control: n.c.) followed by staining for IFN- γ and TNF- α (n=3-4). (**d**+**f**) At day 20 post-infection, virus titers were determined in the spleen, liver, lung, and kidney tissue (n=3-4).



Supplementary Figure 5: LCMV-WE infection promotes enhanced dendritic cell activation.

(a) BMDC gating strategy of Figure 3a. (b) GM-CSF induced BMDCs were infected with LCMV-WE or LCMV-Docile, or co-infected at the indicated MOI's. IFN- α concentration was determined in the supernatant of infected BMDCs 48h and 72h post-infection (n=9). (c-e) GM-CSF induced BMDCs were infected with LCMV WE or Docile, or co-infected at the indicated MOI's, 48h, and 72h post-infection and (c) IL-6, (d) TNF- α , (e) IL-1 β levels were determined in the supernatant of infected BMDCs (n=7). (f) GM-CSF induced BMDCs were infected with LCMV-WE, LCMV-Docile, or co-infected at MOI 1. At the indicated time points, IL-1 β was determined by immunoblot analysis (A representative of n=4 immunoblots is shown). (g) uncrop scan of Western Blots in (f), (h) GM-CSF induced BMDCs were infected with LCMV-WE or LCMV-Docile, or co-infected at a MOI =10. 24 h post-infection, the oxygen consumption rate (OCR) was determined in real time after the addition oligomycin, FCCP, and antimycin-A/rotenone (n=9-11).



Supplementary Figure 6: Uncropped scans of Western blots in Fig. 4a



Supplementary Figure 7: Both LCMV-WE and LCMV-Docile infections induced NF-κB and MAP kinase activation.

(a) GM-CSF induced BMDCs were infected with LCMV-WE, LCMV-Docile, or co-infected at a MOI=1. p-IKB α , total IKB α , pERK1/2, ERK1/2, p-JNK, total JNK, and their loading control Tubulin were assessed by immunoblot analysis at the indicated time points (one of n=4 representative blot was shown). (b) Uncropped scans of Western blots in (a).



Supplementary Figure 8: Uncropped scans of Western blots in Fig. 4d

| | 10 | 20 | 30 | 40 | 50 | 60 | 70 | 80 | 90 | 100 |
|-----------|-----------------|------------|-----------|------------|------------|--------------|------------|------------|------------|-------|
| WE NP | MSLSKEVKSFOWTO | LRRELOSFTS | VKAAVIKDA | TSLLNGLDFS | EVSNVORIME | KERRDDKDL | RLRSLNOTVH | SLVDLKSTSK | KNVLKVGRLS | AEELM |
| Docile NP | | T | | | | | | E | | |
| | | | | | | | | | | |
| | 110 | 120 | 130 | 140 | 150 | 160 | 170 | 180 | 190 | 200 |
| WE NP | TLAADLEKLKAKIM | TERPOASGVY | GNLTAQOLD | ORSOILOMVG | MRRPOOGASG | VVRVWDVKDS | SLLNNQFGTM | PSLTMACMAR | SOTPLNDVM | ALTD |
| Docile NP | s | ST | | | RS.N. | | | | | |
| | | | | | | | | | | |
| | 210 | 220 | 230 | 240 | 250 | 260 | 270 | 280 | 290 | 300 |
| WE NP | LGLLYTVKYPNLSDI | ERLKDKHPVL | VITEQUSSI | NISGYNFSLG | AAVKAGAALL | DGGNMLESI | IKPSNSEDLL | KAVLGAKKKL | MEVSDOVGD | RNPYE |
| Docile NP | N. | R | | | | | | R | | |
| | | | | | | | | | | |
| | 310 | 320 | 330 | 340 | 350 | 360 | 370 | 380 | 390 | 400 |
| WE NP | NILYRVCLSGEGWPY | IACRTSWGR | WENTTIDLT | NEKLVANSSR | PVPGAAGPPC | VGLSYSQTMI | LKDLMGGIDP | NAPTWIDIEG | RENDEVEIAI | OPON |
| Docile NP | | I | | TPP. | | | | | | |
| | | | | | | | | | | |
| | 410 | 420 | 430 | 440 | 450 | 460 | 470 | 480 | 490 | 500 |
| WE NP | GOFIHFYREPTDOK | FKDDSKYSHG | DLADLFNAC | PGLTSSVIGA | LPQGMVLSC | GSDDIRKLLI | SONRRDIKLI | DVEMTREASR | EYEDKVWDKY | WLCK |
| Docile NP | V | | ••••• | | | | | R | | |
| | | | | | | | | | | |
| | ···· ···· ···· | 320 | 580 | 540 | | | | | | |
| WE NP | MHTGVVRDKKKKEIT | PHCALMDCII | ESASKARLF | DLKTVHNILP | HDLIFRGPNV | ∕ ∕TL | | | | |
| Docile NP | I.K | | | I | | | | | | |

Supplementary Figure 9: LCMV-WE and LCMV-Docile NP share the same DIEG motif NP Amino acids sequence were between WE and Docile Strain. The DIEG motif is highlighted in red box.



Supplementary Figure 10: LCMV-WE and LCMV-Docile did not block IFNAR signalling.

GM-CSF induced BMDCs were infected with LCMV-WE, LCMV-Docile, or co-infected at MOI 1. 12 h post-infection, BMDCs were treated with recombinant IFN $\alpha 4$ at 100U/ml and 30 minutes later, (a) p-STAT1, total STAT1 were measured by immunoblotting (A representative blot of n=3 is shown). (b) Uncropped scans of Western blots in (a). (c) Quantification of (a) is shown (n=3). (Error bars show SEM, *p<0.05, **p <0.01 between the indicated groups).



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Supplementary Figure 11: Uncropped scans of Northern blots in Fig. 5h

LCMV NP staining on BHK21- cells



Supplementary Figure 12: BHK-21 LCMV NP intracellular staining gate strategy of Figure 5k



Supplementary Figure 13: LCMV-WE infection increases IFN-I transcripts.

C57BL/6 mice were infected with LCMV-WE, LCMV-Docile, or co-infected for 1 day. Various types of IFN-I subtype mRNA transcripts were determined in spleen tissue (n=6).



Supplementary Figure 14: LCMV Docile RNA production was inhibited in the presence of LCMV WE.

(a-b) BHK-21 cells were infected with LCMV-WE, LCMV-Docile, or both at the indicted time points (MOI=1). 24h post-infection, BHK-21 cellular RNA was isolated and (a) WE-GP RNA (left panel), Docile-GP RNA (right panel) (b) WE-NP RNA (left panel), Docile-NP RNA (right panel) were assessed by RT-PCR (n=3). (Error bars show SEM, ***p < 0.001, and ns indicates statistically not significant between the indicated groups).



Supplementary Figure 15: Chimeric LCMV virus is attenuated in WT animals.

C57BL/6 mice were infected with $2x10^5$ pfu of chimeric virus S(WE)/L(Clone 13) or S (Docile)/L(Clone 13). At day 12 post-infection, virus titers were determined in spleen, liver, lung, and kidney tissue (n=7-8).