

| Gene Name | | Probes (10 μM) | Primers (90 μM) |
|---------------|--------------|---|--|
| MLV Env | | EcoP 5'-CAGACCAACAGCCACT-3' | EcoF 5'-GGCAAACCCCGTTCTG-3' EcoR 5'-ACTTAACAGGTTTGGGCTTGGA-3' |
| Synaptophysin | | Roche Universal ProbeLibrary Set, Human #72 | F 5'-ACAACAAGGGCCAATGAT-3' R 5'-TAGCCACATGAAAGCGAACA-3' |
| Vif | EcoHIV/NL4-3 | NLVFQP 5'-AGTAGTAATACAAGATAATAGTGA-3' | NLVFQ5 5'-CGCGCACGGCAAGAG-3' NLVFQ3 5'-CTTTGCTTTTCTTCTTGGCACTACT-3' |
| | EcoHIV/NDK | QVFP 5'-AGTAGTAATACAAGACAATAGTG-3' | QVF52 5'-AAGAGGCGAGGGGCAGCGA-3' QVF3 5'-TCTTTACTTTTCTTCTTGGTACTACCTTTATG-3' |
| Tat | EcoHIV/NL4-3 | NLTatP 5'-AAAGCCTTAGGCATCTC-3' | NLTatF 5'-CCTAAAAGTCTTGTACCAATTGC-3' NLTatR2 5'-GTCGGGTCCCCTCGGGATTGGGAG-3' |
| | EcoHIV/NDK | NDTatP 5'-AAAGGCTTAGGCATCTC-3' | NDTatF 5'-CCTAGGACTGCTTGTAAATAAGTGT-3' NDTatR2 5'-GTCGGGTCCCCTCGGGACTGGGAG-3' |
| Mouse GAPDH | | 20x mix (ABI 4352932 E) | |

Supplementary Table S1. Gene sequences of probes and primers for QPCR

| | 129x1/SV | | C57BL/6 | |
|----------------------------------|------------|-------------|-------------|----------------|
| Microglia size (μm) | Uninfected | Infected | Uninfected | Infected |
| Average | 35.373 | 144.571 | 28.8 | 163.357 |
| STD error | 6.282 | 23.409 | 5.304 | 21.289 |
| <i>t</i> -test | | | | |
| 129x1/SV Uninfected vs. | | 0.000000213 | 0.159374 | 0.000000000103 |
| 129x1/SV Infected vs. | | | 0.000000215 | 0.299825 |
| C57BL/6 Unintected vs. | | | | 0.000000000141 |

Supplementary Table S2. (for Figure 4)

Microglia cell size measurement of EcoHIV/NDK-EGFP-infected cell percentage to compare infectivity of EcoHIV/NDK-EGFP in difference mice strains, C57BL/6 and 129x/SV mice in brain glial cell culture.