

Supporting Information

Bergthaler et al. 10.1073/pnas.1011998107

Table S1. Comparison of the glycoprotein 260 and long-segment 1,079 position in widely used lymphocytic choriomeningitis virus strains

Gene and amino acid position*	GP ₂₆₀	L ₁₀₇₉
LCMV strain		
Armstrong [†]	F	K
Clone 13 [‡]	L	Q
WE [§]	L	K
Aggressive [¶]	L	K
Docile	L	K
Pasteur**	L	K
Traub ^{††}	L	K

GP, glycoprotein; L, long segment; LCMV, lymphocytic choriomeningitis virus.

*Counted from the first amino acid of the respective ORF.

[†]GenBank AY847350 and AY847351.

[‡]GenBank DQ361065 and DQ361066.

[§]GenBank M22138 and AF004519.

[¶]GenBank EU480450 and EU480451.

^{||}GenBank EU480452 and EU480453.

**GenBank DQ868485 and DQ868486.

^{††}GenBank DQ868487 and DQ868488.