

WRNA2-057**Comparison of the intracellular distribution of lyssavirus matrix proteins***Pollin Reiko¹, Finke Stefan²*¹Friedrich-Loeffler-Institut, Greifswald - Insel Riems, Germany²Friedrich-Loeffler Institut, Greifswald - Insel Riems, Germany

Lyssaviruses are neurotropic rhabdoviruses with bats as the main reservoir. Of the different genotypes, only the one comprising classical Rabies Virus (RABV) is able to sustain in non-bat host population, suggesting that lyssaviruses differ in virus-host interactions required for efficient virus replication. Based on the recent finding, that matrix (M) proteins of RABV and bat-adapted European bat lyssavirus type 1 (EBLV-1) may differ in the targeting of cellular membranes, we compared the intracellular distribution of RABV and EBLV-1 M proteins. According to the essential role of M in virus budding, both M were detected at the cytoplasm and intracellular membranes. For the latter, a remarkable difference was observed: in contrast to RABV M, EBLV-1 M perfectly colocalized with Golgi-markers, confirming the postulated differences in membrane targeting. The use of a novel, highly cross-reactive antibody also allowed the detection of both M not only in the cytoplasm but also in the nucleus of infected cells, suggesting that lyssavirus M proteins are also involved in the regulation of nuclear processes. Furthermore, colocalization of both M proteins with cytoplasmic inclusion bodies for the first time showed the presence of lyssavirus M proteins at the suspected sites of virus replication, suggesting a direct influence of lyssavirus M on viral RNA synthesis in or nucleocapsid export from inclusion bodies. The identification of similar and distinct features of M proteins of bat and non-bat adapted lyssaviruses may allow the further identification of restrictive virus-host interactions and their contributions to host barriers and virus tropism.