

**Mutational analysis of the Potyviridae transcriptional slippage site utilized for expression of the P3N-PIPO and P1N-PISPO proteins**

Olspert, Allan; Carr, John P.; Firth, Andrew E. Nucleic acids research 2016 / p. 7618-7629 : ill <http://dx.doi.org/10.1093/nar/gkw441>

**Potyvirus helper component-proteinase and coat protein (CP) have coordinated functions in virus-host interactions and the same CP motif affects virus transmission and accumulation**

Andrejeva, Jelena; Puurand, Ülo; **Merits, Andres**; Rabenstein, Frank; **Järvekülg, Lilian**; Valkonen, J.P.T. Journal of general virology 1999 / p. 1133-1139 [https://www.researchgate.net/publication/12942693\\_Potyvirus\\_helper\\_component-proteinase\\_and\\_coat\\_protein\\_CP\\_have\\_coordinated\\_functions\\_in\\_virus-host\\_interactions\\_and\\_the\\_same\\_CP\\_motif\\_affects\\_virus\\_transmission\\_and\\_accumulation](https://www.researchgate.net/publication/12942693_Potyvirus_helper_component-proteinase_and_coat_protein_CP_have_coordinated_functions_in_virus-host_interactions_and_the_same_CP_motif_affects_virus_transmission_and_accumulation)

**The complete nucleotide sequence of the ryegrass mosaic potyvirus indicates that it is a recombinant between members of two different genera in the family Potyviridae**

Schubert, J.; Fauquet, C.; **Merits, Andres**; Rabenstein, Frank Zeitschrift für Pflanzenkrankheiten und Pflanzenschutz 1999 / p. 392-404 <https://www.jstor.org/stable/43216179?seq=1>

**3' terminal sequence analysis and homologies for two members of genus Rymovirus (Potyviridae)**

Merits, Andres; **Andrejeva, Jelena**; Kibe, Kristiina; Rabenstein, Frank; Proll, Eckhard; **Järvekülg, Lilian** International Symposium 75 Years of Phytopathological and Resistance Research at Aschersleben, 12-16 June, 1995, Aschersleben, Germany : programme, abstracts, list of participants 1995 / p. P18

**Turnip Mosaic Virus Transcriptional Slippage Dynamics and Distribution in RNA Subpopulations**

Kärblane, Kairi; Olspert, Allan; Firth, Andrew E. Molecular plant-microbe interactions 2022 / p. 835-844 : ill

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