Response from Ralf Dietzgen to “Comment on The complete nucleotide sequence and genome organization of pea streak virus (genus *Carlavirus*)”

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Apparently not being aware of the recently published ALV genome sequence when claiming priority in reporting the PeSV genome sequence as a ‘first’ was likely an oversight by the authors that was unfortunately not picked up by two expert reviewers and the handling editor. There is however no basis for Dr Nemchinov’s accusation of malicious misrepresentation by the authors, and intentional exclusion of his data during the review process.

Both publications report sequences of two different ‘strains’ of PeSV and represent valuable contributions to the field. Furthermore, the scientific conclusions of the *Archives of Virology* article are not affected by the availability of KP784454 sequence. However, clearly the earlier report should have been credited. Our editors and reviewers will need to pay closer attention to accessible sequences in order to avoid such conflicts in the future.

Yours sincerely,
Ralf Dietzgen
Editor

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