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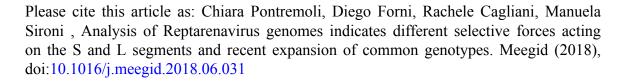
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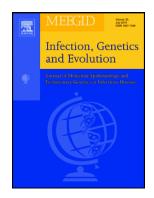
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Analysis of *Reptarenavirus* genomes indicates different selective forces acting on the S and L segments and recent expansion of common genotypes

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