Here, we announce the first complete genome sequence of a field isolate of a peste des petits ruminants virus (PPRV) from northern Africa. This isolate is derived from an Alpine goat that suffered from severe clinical disease during the 2008 outbreak in Morocco. The full genome sequence of this isolate clusters phylogenetically with the lineage IV isolates of PPRV, sharing high levels of sequence identity with other lineage IV isolates.
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REFERENCES