The complete genome sequence of a reassortant field strain (IND2014/01) of Bluetongue virus (BTV) serotype 16, isolated from sheep from southern India in 2014, was sequenced. The total genome size was 19,186 bp. Sequence comparisons of all genome segments, except segment 5 (Seg-5), showed that IND2014/01 belonged to the major eastern topotype of BTV.


