Trichodysplasia spinulosa-associated polyomavirus (TSV) is responsible for a rare skin cancer. Using metagenomic approaches, we determined the complete genome sequence of a TSV first detected in Brazil in spicules of an immunocompromised patient suspected to have trichodysplasia spinulosa.


ACKNOWLEDGMENTS
This study was supported by FAPESP #2012/15381-7.
We thank Roberto Ribeiro for assistance with Ion Torrent sequencing.

REFERENCES

