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1	The emergence of enterovirus D68 in England in autumn 2014 and the necessity
2	for reinforcing enterovirus respiratory screening.
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27 SUMMARY

28 In autumn 2014, enterovirus D68 (EV-D68) cases presenting with severe respiratory or neurological disease were described in countries worldwide. To describe the 29 epidemiology and virological characteristics of EV-D68 in England, we collected 30 clinical information on laboratory-confirmed EV-D68 cases detected in secondary 31 care (hospitals), between September 2014 and January 2015. In primary care (general 32 33 practitioners), respiratory swabs collected (September 2013-January 2015) from patients presenting with influenza-like-illness were tested for EV-D68. In secondary 34 care 55 EV-D68 cases were detected. Among those, 45 cases had clinical information 35 36 available and 89% (40/45) presented with severe respiratory symptoms. Detection of EV-D68 among patients in primary care increased from 0.4% (4/1074; 95%CI: 0.1-37 1.0) (September 2013-January 2014) to 0.8% (11/1359; 95%CI: 0.4-1.5) (September-38 39 2014 to January-2015). Characterization of EV-D68 strains circulating in England since 2012 and up to winter 2014/2015 indicated that those strains were genetically 40 similar to those detected in 2014 in USA. We recommend reinforcing enterovirus 41 surveillance through screening respiratory samples of suspected cases. 42

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50 BACKGROUND

The Enterovirus genus (family Picornaviridae) includes twelve species (A-J) of 51 which four (A-D) are associated with disease in humans; each of these species 52 encompasses multiple types. Enterovirus infection can be associated with a broad 53 range of clinical presentations, including meningitis, encephalitis, hand, foot and 54 mouth disease (HFMD), respiratory illness, and more rarely, neurological symptoms. 55 Multiple enterovirus types co-circulate, and some types are known to emerge 56 occasionally in a population to cause outbreaks or clusters of cases of acute 57 58 neurological disease [1, 2]. Increasing recognition of the disease burden associated with enterovirus infections, and the description of 're-emerging' enteroviruses such as 59 EV-C99, EV-C104, EV-D70, EV-A71 [3, 4], provide evidence that enteroviruses have 60 61 rapidly evolving genomes, are ubiquitous worldwide, spread rapidly via faecal-oral and environmental routes, but also via respiratory transmission, and have the ability to 62 cause severe disease. Enterovirus D68 (EV-D68) is one of five enterovirus types 63 assigned to *Enterovirus* species D. The virus was first described in 1962, isolated from 64 children with pneumonia and bronchitis [5]; in 2002 a human rhinovirus 87 was 65 reclassified as EV-D68, based on phylogenetic analysis [6]. In countries with 66 enterovirus established surveillance EV-D68 has been detected sporadically in cases 67 68 of acute respiratory illness, particularly in children, and occasionally was associated 69 with acute neurological disease and other severe outcomes [7, 8]. In autumn 2014, the United States of America (USA) experienced a nationwide 70

ni autanin 2014, the officed states of America (OSA) experienced a nation wide
 outbreak of EV-D68 associated with severe acute respiratory illness [9]. This
 outbreak coincided with an apparent increase in incidence of reported cases of
 neurological disease characterized as acute flaccid myelitis (AFM), suggesting that
 AFM is a rare yet severe clinical manifestation of EV-D68 infection [10-12].

75 In 2014, following the USA outbreak, EV-D68 was detected in European countries [13]. In Norway, EV-D68 was detected in 11% of 303 paediatric nasopharyngeal 76 samples collected from children hospitalized with acute respiratory infection. EV-D68 77 78 was associated with acute flaccid paralysis in one child [14]. In the Netherlands, 1% (18/1896) of respiratory samples taken from patients with respiratory symptoms tested 79 positive for EV-D68 [15]. Phylogenetic analysis showed that strains detected in the 80 Netherlands were genetically similar to those circulating in the USA in 2014 [15]. 81 Public Health England (PHE) has an established national Enterovirus Surveillance 82 System (ESS), with emphasis on further investigation of enterovirus-positive cases 83 presenting with acute neurological symptoms as part of enhanced poliovirus 84 surveillance [16]. Following reports of the emergence of EV-D68 in USA, PHE issued 85 guidance on October 2014 that EV-D68 should be considered as a possible cause of 86 87 disease, particularly in children presenting with severe acute respiratory infections and/or unexplained neurological symptoms, and where standard respiratory virus 88 89 screens were negative, or if a combined rhinovirus/enterovirus positive result was initially detected. 90 This study aimed to quantify the circulation of EV-D68 in England in primary and 91 secondary care before and after the identification of EV-D68 in North America in 92

autumn 2014. Our objectives were i) to determine whether there was an increase in
number of cases between autumn/winter 2013-2014 and autumn/winter 2014-2015,
and ii) to describe the clinical presentation and phylogenetic characteristics of this
virus, including severity in England in the autumn/winter 2014-2015.

97 METHODS

A confirmed case was any patient presenting to primary or secondary care with a
specimen positive for EV-D68 detected initially by one-step real-time reverse
transcription PCR (rRT-PCR) [15]. Cases investigated in this study were identified
from two surveillance sources.

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1. Secondary care (clinical presentations)

103 As part of the ESS, samples (including stool, cerebrospinal fluid (CSF) and respiratory samples) from individuals who presented to hospitals with acute 104 neurological syndromes and in whom an enterovirus is detected are sent for 105 enterovirus typing. This typing is carried out at the Virus Reference Department, PHE, 106 Colindale. From November 2014, referrals to the ESS were enhanced, to include 107 108 enterovirus-positive paediatric cases with acute respiratory symptoms. The Respiratory Diseases Department of the PHE Centre for Infectious Disease 109 Surveillance and Control (CIDSC) coordinated the epidemiological investigation of 110 each confirmed case. 111

Using a standardized questionnaire the treating clinicians collected information on all confirmed EV-D68 cases from October 2014 to February 2015. The information gathered included: patient characteristics, clinical presentation (including details on the neurological and respiratory presentations), testing performed (including all pathogens detected and type of samples) and disease outcome.

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2. Primary care (EV-D68 circulation)

118To estimate the rate of circulation of EV-D68 in the community, we tested respiratory119samples from the general practitioner (GP) influenza sentinel surveillance system.120Patients presenting with acute febrile respiratory illness considered by their GP to be121influenza-like-illness (ILI) had nasopharyngeal virology specimens sent to PHE, as

part of the Royal College of General Practitioners Research and Surveillance Centre 122 (RCGP RSC) influenza sentinel surveillance weekly returns service. These specimens 123 were tested for EV-D68. This service, run by RCGP RSC since 1957, provides 124 125 clinical surveillance data from a sentinel network of over 100 general practices (GP) (primary health care centres) together with virological respiratory specimens, from a 126 subset of patients presenting with ILI across England [17]. Each GP sends around 127 128 five combined nose and throat swabs per week from patients consulting with ILI, particularly during the winter season, from around week 40 of one year to around 129 130 week 20 of the next year. Among the samples received from the RCGP RSC from week 39-2013 to week 15-2015 (September 2013 to April 2015) only those testing 131 negative for influenza virus were tested for EV-D68. 132

133 Data analysis

Changes in circulation of EV-D68 were estimated by calculating EV-D68 percent 134 135 positivity in two periods. Percent positivity was calculated as number of EV-D68 positive samples divided by total of samples tested for EV-D68 in each period. 136 Percent positivity in the period week 39-2013 to week 05-2014 (September 2013 to 137 138 January 2014) (in 2013 RCGP RSC influenza testing only started in week 39) was compared to percent positivity in the period week 36-2014 to week 05-2015 139 (September 2014 to February 2015), using Chi-square test. For primary care samples, 140 we only looked at percent positivity and demographic characteristics (age and sex) of 141 142 cases.

143 Laboratory detection and characterization

144Detection of EV-D68 was performed by rRT-PCR as previously described [15].145Further characterization of EV-D68 strains was performed by amplification and

146	partial sequencing of the VP1 coding region using a previously developed in-house
147	assay (M. Iturriza-Gómara, University of Liverpool, UK). Briefly, primers Ent68F
148	[5'-GAAGCCATACAAACTCGCAC-3'] and Ent68R [5'-
149	ATTWGCAATGCTCATGTATGG-3'] were used to amplify a ~670nt fragment of
150	the VP1 coding region. A high fidelity PCR system was used (Expand High Fidelity
151	PCR Kit, Roche, UK) according to manufacturer's instruction, with the following
152	specifications (per reaction): magnesium chloride final concentration 2.5mM, 40pmol
153	each primer, and 1.75U DNA polymerase. Amplification was performed under the
154	following conditions: initial denaturation: 95°C, 240 seconds; thermal cycling 95°C,
155	30 seconds – 42°C, 60 seconds – 72°C, 60 seconds for 40 cycles; final extension:
156	72°C, 240 seconds.
157	Sequence analysis was performed using Bionumerics v6.1 (Applied Maths,
158	Kortijk,Belgium) and phylogenetic and molecular evolutionary analyses were
159	conducted using MEGA version 6 [18].
160	For phylogenetic analysis, we included representatives of EV-D68 strains detected in
160	
161	the USA in 2014 [19] and 2015 [20], EV-D68 strains detected in England in 2010-
162	2013 as part of national enterovirus surveillance and enhanced poliovirus surveillance
163	[21] and representatives of other EV-D68 strains reported globally [22-24].
164	
165	RESULTS
166	Clinical presentation of EV-D68 in England
167	From week 42-2014 to week 05-2015 (October 2014 to February 2015), 433
168	respiratory samples were collected from patients presenting to secondary care in

169	England. During this period, 13% (55/433) of these respiratory samples were positive
170	for EV-D68, compared with 11% (3/26) of EV-D68 positive samples detected in the
171	same period in the previous year (Figure 1). Sixty-nine percent (38/55) of the EV-D68
172	cases (October 2014 to February 2015) were in children under 12 years of age; the
173	median age of cases was 4 years (range 0-71), and there was no difference in the
174	gender distribution (46% male) (Figure 2). The majority of these EV-D68 positive
175	specimens were nasopharyngeal aspirates, 44% (24/55), and nose/throat swabs,
176	31%(17/55). The remaining 14 were sputum, mouth swabs and tracheal/bronco
177	alveolar aspirates. EV-D68 was not detected in any CSF specimen tested (n=562).
178	Detailed clinical information was available for 82% (45/ 55) EV-D68 cases (Table 1).
179	None of these cases reported having recently travelled outside of the United Kingdom
180	(UK). Ninety-one percent (41/45) of the cases presented with acute respiratory
181	symptoms, one case presented with HFMD, one case had leukaemia (respiratory
182	symptoms not reported) and two cases presented with acute neurological symptoms.
183	These cases showing neurological presentations were in children <5 years of age: one
184	case presented with meningoencephalitis and paucity of limb movements [22]; the
185	second presented with afebrile seizures and respiratory arrest. These cases were
186	previously healthy children who required intensive care unit (ICU) treatment and
187	subsequently recovered; in both cases, EV-D68 was detected in nasopharyngeal
188	aspirate specimens. Thirty-three percent $(13/39)$ of the cases presented with asthma or
189	wheezing, 84% (11/13) of them were children under 15 years of age (median age: 2
190	years) (information for asthma was only collected for 39 patients). Forty-seven
191	percent (21/45) of the cases were immunocompromised or had an underlying chronic
192	condition. Eighty-two percent (37/45) of the cases were admitted to hospital, with
193	27% (12/45) requiring treatment in ICU, all of them children (median age: <1 year)

(Table 1). Two patients in whom EV-D68 was detected died: both were infants with
underlying neurological problems. Information on cause of death was only accessible
for one of them: it was likely due to complications in the underlying neurological
problems. Two other infant cases had never left the hospital since birth.

198 Estimate of prevalence of EV-D68 in England

A total of 3575 nose/throat swab samples were collected from week 39-2013 to week 15-2015 (September 2013 to April 2015) as part of the RCGP RSC influenza sentinel surveillance weekly returns service. The median age of the patients providing these samples was 33 years (range 0-99), 17% of them were children <12 years of age. Sixty-one percent were female.

EV-D68 was detected in 0.4% (15/3575) of all the community respiratory samples 204 205 analysed. Among the cases, the median age was 40 years, (range 5-84 years); nine (60%) of them were female (Figure 2). Among those 3575 samples, 2433 206 (1074+1359) had been collected during the two periods of comparison and were 207 included in the percent positivity analysis. Positivity in the autumn/winter 2013/14 208 (week 39-2013 to week 05-2014) was 0.4% (4/1074; 95%CI: 0.1-1.0). Positivity in 209 the autumn/winter 2014/15 (week 36-2014 to week 05-2015) was 0.8% (11/1359) 210 (95%CI: 0.4-1.5) (p value=0.17). All primary care (RCGP RSC) EV-D68 positive 211 samples in 2013 were collected between weeks 40 to 47, while all the community EV-212 D68 positive samples from 2014 were collected between weeks 49 to 52 (Figure 1). 213

214 **Phylogenetics**

215 Partial VP1 sequence was obtained from 88% (57/70) of the EV-D68 cases

216 identified from primary and secondary care in this study (four from community cases recruited via the RCGP network and 53 from cases referred to the ESS). 217 Of the 57 EV-D68 sequences detected in this study, 56 clustered with EV-D68 genetic 218 clades (A and B as described elsewhere [23]): 49 (85%) in clade B and 8 (14%) in 219 clade A. The remaining 2014 strain clustered with the prototype US/Fermon/1962 220 strain (Figure 3). 221 EV-D68 strains detected in England in 2014 and 2015 clustered both with strains 222 detected in the USA during the same period, and with strains circulating in England 223 224 prior to 2014. Among the clade B viruses there were the two sequences found in cases with neurological presentation. Both of these sequences were highly similar to other 225 co-circulating viruses from the same period. 226 227 Amino acid sequences were highly similar within clades, with most amino acid differences occurring in the BC and DE loops (data not shown). There were no 228 specific amino acid changes in either the BC or DE loop, or elsewhere in the regions 229 230 of VP1 sequenced that distinguished strains associated with neurovirulence from other EV-D68 strains detected. 231 DISCUSSION 232 We used two established surveillance systems for this study. From the national 233 enterovirus surveillance system data we described the burden of illness due to EV-234 235 D68 and the severity of clinical presentation in England. We used the influenza sentinel GP surveillance system to look at the changes in EV-D68 circulation in 236 primary care. Clinical information from the cases identified in the ESS (secondary 237 238 care) found that cases with EV-D68 positive samples were mainly presenting with severe respiratory symptoms in the autumn/early winter 2014-2015. In primary care, 239 retrospective testing of samples revealed a non-significant increase in circulation of 240

EV-D68 in autumn/early winter 2014-2015 compared to the same period the previous year.

Our findings from secondary care suggest that, when detected, EV-D68 is most 243 frequently associated with respiratory disease, usually in children, and more rarely 244 with neurological disease. Previous studies presented comparable results [9, 24-27]. 245 Further, in our study the majority of hospitalized cases presented with severe 246 respiratory disease and one third of those were admitted to ICU (all of them children) 247 and required respiratory support. Our results on cases with severe respiratory 248 249 presentations, mostly children, associated with EV-D68 infection were comparable to similar findings in the Netherlands and the USA [15, 27]. We found that 33% of 250 cases presented with asthma/wheezing compared to the 70% of cases presenting with 251 252 these symptoms in the 2014 USA outbreak [27], this difference could be attributable to the small number of cases for whom we had information on asthma/wheezing. In 253 our study, similarly to other studies, almost half of the patients admitted with 254 respiratory disease were immunosuppressed or had severe underlying conditions [15, 255 28]. Our findings reinforce the importance of clinicians considering EV-D68 (and 256 other enteroviruses) as one of the differential diagnoses of severe acute respiratory 257 illness, particularly in children. 258

In the only two acute neurological presentations in our study, EV-D68 was not detected in the CSF of these children and the detection of EV-D68 in the respiratory samples may have been coincidental. However, both cases were previously healthy and no other infectious cause was identified for the disease. Therefore we cannot exclude the possibility of an association between EV-D68 and their severe acute presentations. Both children made a complete recovery. Severe neurological presentations associated with EV-D68 in previously healthy children with a preceding 266 respiratory infection have been reported in different European countries: in Norway, two cases (Autumn 2014) [29]; a case in France (September 2014) [30] and two cases 267 in Wales (UK) (January 2016) [31], EV-D68 was not identified in the CSF in any of 268 269 these cases, but rather from upper respiratory tract samples. Similarly, we found that EV-D68 RNA was detectable in respiratory specimens, but that other sample types 270 (e.g. CSF or stool) were not sensitive for detection of EV-D68. We were able to 271 recover infectious EV-D68 virus by inoculation of permissive cell lines with RT-PCR-272 positive respiratory specimens, but not other specimen types. This highlights the 273 274 importance of collecting all clinically relevant samples for differential diagnosis in such acute presentations and indicates that in EV-D68 suspected cases respiratory 275 specimens are the most appropriate for diagnostic testing and virological 276 277 characterization.

278 The rate of positivity in cases of acute respiratory disease presenting to primary care in England doubled from autumn/winter 2013-2014 to autumn/winter 2014-2015. 279 280 Nonetheless this increase was not statistically significant and the number of cases and 281 positivity rates were low. This suggests that the high number of severe cases observed in secondary care in the period post autumn 2014 is presumably explained by the 282 change in case definition and enhanced case ascertainment. Similarly, in the 283 Netherlands, a non-significant increase in the percentage of EV-D68 positives was 284 observed from 2013 (0.3%) to 2014 (0.6%) among patients presenting to the GP [32]. 285 In another similar study in Germany from 2014, a higher percentage (7.7%) of EV-286 D68 positive samples from outpatients presenting with ILI, and/or acute respiratory 287 infection was reported [28]. Our primary care cases had similar distribution across all 288 age groups, but in those in secondary care, age was skewed toward infants and young 289 290 children (Figure 2). This could be related to the PHE guidance, suggesting considering 291 EV-D68 as a possible cause of disease in children with severe respiratory symptoms. Or, taking into account similar results from other studies, it may demonstrate a 292 different age profile of the EV-D68 cases presenting with severe symptoms in 293 secondary care compared to the profile of the mild presentations in primary care. 294 We characterized EV-D68 strains circulating in England since 2012 and up to winter 295 2014-2015, among which we identified strains genetically similar to those detected in 296 2014 in USA. These findings are consistent with similar results from Netherlands and 297 298 Germany [15, 28]. These studies describe emergence of distinct genetic clusters, which reflects the continuous evolution of EV-D68 strains in circulation. 299 With extensive genetic variation in the EV-D68 genome, continuous collection of 300 301 genomic data by representative surveillance systems is required to ensure that PCRbased detection methods are capable of detecting contemporary circulating viruses. 302 Genotyping assays are generally based upon sequence analysis of the VP1 coding 303 304 region, whilst primary detection assays usually target the 5'-untranslated region of the 305 genome. With recognized sequence variation in both regions, complete genome sequences representing contemporary circulating viruses is essential in assuring 306 307 performance of molecular diagnostic and characterization assays. However, to fully understand the clinical significance of detection of EV-D68 in 308 patients, alongside genomic characterization, isolation of the virus, characterization by 309 neutralization and measurement of virus-specific antibody titres is required. 310 311 Integrating 'classical' virology approaches and molecular techniques will better establish links between infection and disease in patients, as well as provide valuable 312 313 technical information for improving laboratory diagnostics, monitoring genetic drift in the virus genome which may result in degradation of PCR primer/probe-binding sites. 314

Some molecular assays have been found to exhibit cross-reactivity between EV-D68 and rhinovirus which can lead to false-positive results. Furthermore, with reports of other emerging and novel enterovirus types [1, 2, 33], it is important to continually enhance the virological, genomic and epidemiological understanding of these viruses by maintaining up-to-date diagnostic and reference methods, and monitoring the factors determining local and temporal differences in atypical disease outbreaks due to enteroviruses.

The detection of EV-D68 worldwide in 2014-2015 may be the 'tip-of-the-iceberg' of circulation of EVD-68. EV-D68 is rarely associated with severe disease, which is probably why fewer cases have been detected by current surveillance systems that are designed to capture more severe (neurological) infections. In our study, detection of the virus in community and in hospitalized patients with respiratory disease suggest EV-D68, and perhaps other enteroviruses, are a more important cause of infection and potentially disease than captured by current surveillance systems.

329 STRENGTHS AND LIMITATIONS

Our study had some strengths and limitations. In terms of strengths, we were able to 330 look at the EV-D68 epidemiology and virology in primary care before and after the 331 signal in the US, using an identical surveillance system, which allowed the 332 comparison between the two periods with no ascertainment bias. In terms of 333 weaknesses, first, we could not compare the EV-D68 positivity rate in ESS with 334 335 previous years given the change in case definitions in autumn 2014 following the PHE guidance mentioned before. Second, the RCGP RSC influenza sentinel surveillance 336 337 samples are predominantly restricted to the winter months when influenza is circulating, considering the previously described EV-D68 seasonal pattern, majority 338

339 of cases occurring in late summer and autumn [34, 35], we may have missed cases if they had occurred in summer or late spring. Furthermore, the representativeness of the 340 RCGP RSC influenza sentinel surveillance is not ideal for an estimate of the real 341 circulation of this virus in patients presenting to primary care, as the patients tested 342 present with acute febrile respiratory illness, and during the 2014 outbreak in the US 343 only 48% of the hospitalized cases presented with fever; additionally EV-D68 may be 344 a more common cause of mild respiratory illness both in primary care and in the 345 community among cases who self-manage without seeking healthcare intervention. 346 347 Nonetheless, the RCGP RSC is a representative network, with only small differences with the national population and therefore the results of this study provide a useful 348 baseline for future monitoring of the circulation of EV-D68 and its severity in 349 350 England[17]. Finally, we did not sequence the whole genome which might contain 351 virulence markers elsewhere. For example, changes in the 5'UTR and IRES have been proposed to associate with changes in translation initiation [36] and virulence [37, 38]. 352

353 CONCLUSIONS

In this study we detected EV-D68 in cases of severe acute respiratory illness and, less commonly, in neurological illness, particularly in children and those with underlying disease from autumn 2014 when enterovirus surveillance system was enhanced and respiratory sampling was included. We also provided evidence of EV-D68 in cases of ILI presenting in primary care before and after autumn 2014, with strains genetically similar to those detected in 2014 in the USA.

360 EV-D68 should be considered as a possible cause of disease in patients presenting
 361 with severe acute respiratory infections and/or with unexplained neurological
 362 symptoms. Our findings emphasize the necessity of reinforcing enterovirus

363	surveillance in England. We recommend the screening of respiratory samples for
364	enterovirus, particularly EV-D68, from patients presenting with severe acute
365	respiratory infections and/or with unexplained neurological symptoms, when all other
366	respiratory virus screen are negative or if an indeterminate rhinovirus/enterovirus
367	positive result was initially detected. The surveillance of enterovirus, specifically EV-
368	D68, in respiratory samples will help us to better understand the epidemiology of EV-
369	D68 and to inform surveillance and laboratory-testing guidance.
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384	Conflict of interest

385	None
386	Ethical approval
387	No ethical approval was required for this study.

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TABLES AND FIGURES

Table 1. Clinical presentation and severity	y of EV-D68 positive cases in secondar	y care in England, September 2014 to January 201	15.

Clinical Presentations-Severity (N=45)		%
Respiratory symptoms (including one case with		89
neurological symptoms) Asthma or wheezing		33 (N=39)
Immunocompromised or severe underlying diseases		
(all)		46
Admitted to hospital	37	82
Respiratory presentations	34	75
Intensive Care	12	27
Neurological presentations	2	4
Deaths	2	4

Figure 1. Distribution of EV-D68 cases detected in the Royal College of General Practitioners (RCGP) influenza sentinel surveillance system and in the Enterovirus Surveillance System (ESS) (including positives from before September 2014 and other later cases for whom clinical information is not available) by week of sample, England, September 2013 to March 2015.

Figure 2. Number of EV-D68 cases by sex, age and surveillance system where detected, England September 2013- January 2015. ((RCGP: Royal College of General Practitioners influenza sentinel surveillance system, (primary care.) EES: Enterovirus Surveillance System (secondary care))

Figure 3. Phylogenetic analysis of partial VP1 sequences of EV-D68 covering the BC and DE loops of the VP1 region. The evolutionary history was inferred using the Neighbour-Joining method and the optimal tree with the sum of branch length = 0.95104738 is shown. The percentages of replicate trees in which the associated taxa clustered together in the bootstrap test (1000 replicates) are shown next to the branches (where >90%). The tree is drawn to scale. Analyses were conducted in MEGA6 [18]. \blacksquare (green)=ESS specimens; \blacksquare (blue)=RCGP specimens; \blacksquare =Cases associated with neurological illness; \blacktriangle =Reference sequences







