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Abstract:	<p>Picornaviruses are the most commonly encountered infectious agents in mankind. They typically cause mild infections of the gastrointestinal or respiratory tract, but sometimes also invade the central nervous system. There, they can cause severe diseases with long-term sequelae and even be lethal. The most famous picornavirus is polio that was a huge burden for mankind for a long time. A successful vaccination campaign brought polio close to eradication, but neurological diseases caused by other picornaviruses have been increasingly reported since the late 1990s. In this review we focus on enterovirus 71, coxsackievirus A16, enterovirus 68 and human parechovirus 3 that have recently drawn attention because of their links to severe neurological diseases. We discuss the clinical relevance of these viruses, the primary role of humoral immunity to control them and summarize current knowledge on neutralization of such viruses by antibodies.</p>

1 Human Picornaviruses Associated with Neurological Diseases and Their
2 Neutralization by Antibodies

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18

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20 **ABSTRACT**

21 Picornaviruses are the most commonly encountered infectious agents in mankind.
22 They typically cause mild infections of the gastrointestinal or respiratory tract, but
23 sometimes also invade the central nervous system. There, they can cause severe
24 diseases with long-term sequelae and even be lethal. The most famous picornavirus is
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26 campaign brought polio close to eradication, but neurological diseases caused by
27 other picornaviruses have been increasingly reported since the late 1990s. In this
28 review we focus on enterovirus 71, coxsackievirus A16, enterovirus 68 and human
29 parechovirus 3 that have recently drawn attention because of their links to severe
30 neurological diseases. We discuss the clinical relevance of these viruses, the primary
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32 neutralization of such viruses by antibodies.

33 INTRODUCTION

34 *Picornaviridae* is one of the largest viral families. According to the International
35 Committee on Taxonomy of Viruses (ICTV) it contains 31 genera that together
36 enclose 54 viral species (Adams *et al.*, 2015). They infect diverse hosts, from lower
37 vertebrates to mammals. The genera *Kobuvirus*, *Salivirus*, *Cosavirus*, *Cardiovirus*,
38 *Hepatovirus*, *Parechovirus*, and *Enterovirus* infect humans (Fig. 1) (Tapparel *et al.*,
39 2013).

40 *Hepatovirus A*, *Parechovirus A* and multiple enterovirus genera can cause
41 symptomatic infections in humans. They typically result in mild disease of
42 gastrointestinal or respiratory tract, but sometimes are associated with severe
43 conditions. For instance, coxsackievirus B (CVB) type 3 has an established role in
44 viral myocarditis (Fairweather *et al.*, 2012), and CVB4 is well-documented as a viral
45 trigger of type 1 diabetes onset (Yeung *et al.*, 2011).

46 Several human picornaviruses can target the central nervous system (CNS)
47 and cause severe neurological diseases. The most well known of them is polio (PV). It
48 caused outbreaks of flaccid paralytic disease in children and was a health care burden
49 for a long time, until development of vaccines and a worldwide vaccination campaign
50 brought it close to eradication (Morales *et al.*, 2016). However, other neurotropic
51 picornaviruses still have potential to cause outbreaks of neurological diseases, such as
52 severe and life-threatening meningoencephalitis, encephalitis or acute flaccid
53 paralysis (AFP). A recent metagenomic study identified members of *Cosavirus*,
54 *Cardiovirus*, *Kobuvirus*, *Enterovirus* and *Parechovirus* genera in clinical samples
55 from AFP children (Victoria *et al.*, 2009). This study corroborated epidemiological
56 and experimental work that has already established firm connections between

57 *Enterovirus* and *Parechovirus* and neurological diseases in humans and is described
58 below in detail.

59

60 **PICORNAVIRUS CNS TARGETING**

61 Picornaviruses spread via the fecal-oral or respiratory routes, and the primary sites of
62 their replication are the gastrointestinal or respiratory tracts. Nevertheless, at least
63 some enteroviruses (EV) and human parechoviruses (HPeV) are routinely neurotropic
64 (Rhoades *et al.*, 2011; Wiley *et al.*, 2015).

65 Picornaviruses utilize a variety of widely expressed molecules as their entry
66 receptors (Evans & Almond, 1998). Such receptors are often present on the surface of
67 cells within the CNS. For example, a receptor for PV—CD155—is expressed in the
68 motor neurons of the spinal cord anterior horns, which are affected during
69 poliomyelitis (Gromeier *et al.*, 2000). Human scavenger receptor class B member 2
70 (hSCARB2) that is utilized by EV71 and CVA16, is expressed on a variety of cells,
71 including neurons and glial cells (Jiao *et al.*, 2014). Thus, the CNS cells are
72 susceptible for infection. In addition, the nervous tissue has reduced immune
73 surveillance and weaker interferon (IFN) responses, and is a plausible site for
74 replication of IFN-sensitive picornaviruses (Ida-hosonuma *et al.*, 2005). Hence, the
75 CNS cells are also permissive for viral replication.

76 There is molecular evidence suggesting that picornaviruses can invade the
77 CNS by three possible mechanisms: peripheral nerve infection, blood-brain barrier
78 crossing and “Trojan horse” invasion.

79 The first mechanism is peripheral nerve infection followed by retrograde
80 axonal transport and trans-synaptic spread in nervous tissue (Fig. 2 (a) and (b)). The
81 evidence for this came from tissue culture studies and *in vivo* experimental models for

82 PV and also from EV71 patient material (Chen *et al.*, 2007; Daley *et al.*, 2005; Ren &
83 Racaniello, 1992; Wong *et al.*, 2008).

84 The second mechanism proposes that during viremia viruses cross the blood-
85 brain barrier (BBB) and infect neural cells. Indeed, high levels of viremia and
86 inflammation can decrease tight junction protein expression, disrupt BBB integrity
87 and facilitate viral invasion (Fig. 2 (c)) (Chai *et al.*, 2014; Daniels *et al.*, 2014).

88 Although the inflammation-induced BBB breakdown has not been directly shown for
89 picornaviruses, their prolonged viremia correlates with severe CNS infections and
90 supports this possibility (Cheng *et al.*, 2014). Picornaviruses can also cross the BBB
91 in an active manner: PV can move through the BBB at a rate comparable to a BBB-
92 crossing antibody (Fig. 2 (d)) (Yang *et al.*, 1997). Such trafficking happens
93 independently of the PvR and appears to rely on transferrin receptor 1 (Mizutani *et*
94 *al.*, 2016).

95 The third mechanism of neurotropism involves migration of infected cells,
96 such as dendritic cells, monocytes, macrophages, T- and B-cells and nestin⁺ myeloid
97 cells to the CNS, and is called a “Trojan horse” invasion (Fig. 2 (e)) (Tabor-Godwin
98 *et al.*, 2010; Vuorinen *et al.*, 1996; Wahid *et al.*, 2005).

99 Neurotropic picornaviruses often target different regions of the CNS, and
100 hence vary in their clinical manifestations. Infection of meningeal cells or cells of the
101 ventricular lining results in aseptic meningitis—a non-bacterial inflammation of
102 tissues lining the brain (Irani, 2008). Infection of neurons with subsequent
103 inflammation of brain parenchyma results in encephalitis that can have long-term
104 sequelae or be fatal (Verboon-Maciolek *et al.*, 2008). Inflammation of the spinal cord
105 grey matter results in myelitis and can lead to limb paralysis (Irani, 2008). All these

106 conditions can be caused by different picornaviruses and their incidences are highest
107 in children (Nicolosi *et al.*, 1986).

108 **NEUROTROPIC PICORNAVIRUSES IN FOCUS**

109 Confirmed neurotropic picornaviruses are members of *Enterovirus* and *Parechovirus*
110 genera. The genus *Enterovirus* includes many recognized pathogens, such as PV,
111 CVA, coxsackieviruses B (CVB), rhinoviruses and EV, whereas genus *Parechovirus*
112 is smaller and includes one human pathogenic species—Parechovirus A. The
113 infections are common, and in the US alone over 10 million symptomatic EV cases
114 are reported annually (Strikas *et al.*, 1986). Human EV and HPeV can be responsible
115 for about 80% of aseptic meningitis cases (Esposito *et al.*, 2014) and 11% of reported
116 encephalitis cases (Koskiniemi *et al.*, 2001). Several types of EV can trigger myelitis
117 with limb paralysis (Kincaid & Lipton, 2006).

118 Not all serotypes of EV and HPeV cause CNS diseases. Enteroviruses
119 associated with CNS infections include PV types 1, 2 and 3, echovirus types 9, 11, 30
120 and 33, CVA type 16, CVB types 3 and 5 (Mistchenko *et al.*, 2006), EV types 68
121 (Messacar *et al.*, 2015) and 71 (McMinn *et al.*, 2001). Parechovirus CNS infections
122 are almost exclusively caused by HPeV3 (Piralla *et al.*, 2014). In this review we will
123 discuss EV71, CVA16, EV68 and HPeV3 that have gained attention due to their
124 recent emergence and connection with CNS infections.

125

126 **Enterovirus 71 and Coxsackievirus A16**

127 EV71 was initially discovered as a CNS-targeting picornavirus: the first isolates came
128 from two children with neurological symptoms in 1969 in California (Schmidt *et al.*,
129 1974). In 1973 it was identified as an etiological agent for hand-foot-and-mouth

130 disease (HFMD), a childhood exanthema characterized by rashes on the palms and
131 soles, oral ulcers and brief febrile illness, but cases of aseptic meningitis were also
132 observed (Hagiwara *et al.*, 1978). In the middle of 1970s it caused a few small
133 outbreaks of aseptic meningitis in the USA, Europe and Australia (Alexander *et al.*,
134 1994; Blomberg *et al.*, 1974; Ishimaru *et al.*, 1980; Kennett *et al.*, 1974) and two
135 rather large outbreaks of polio-like disease in Bulgaria and Hungary that affected
136 predominantly infants with up to 21% of paralytic manifestations of which over a
137 quarter were lethal (Chumakov *et al.*, 1979; Nagy *et al.*, 1982; Shindarov *et al.*,
138 1979).

139 EV71 became a major health care threat in the late 1990s after a series of large
140 outbreaks across the Asia-Pacific region. Most of the affected individuals were
141 children; they often developed HFMD or herpangina, but upper respiratory tract
142 infections (URTI) and non-specific rashes were occasionally observed. A fraction of
143 EV71 infections had neurological and systemic manifestations, but, unlike earlier
144 outbreaks when aseptic meningitis was the most frequent neurological manifestation,
145 more recent outbreaks were characterized with the increased incidence of much more
146 severe brainstem encephalitis and high mortality rate (Chan *et al.*, 2000; Huang *et al.*,
147 1999). The first and largest outbreak in this series occurred in 1998 in Taiwan.
148 Sentinel physicians reported almost 130,000 cases of HFMD, and over 400 cases of
149 severe neurological involvement with almost 20% mortality rate (Ho *et al.*, 1999b)
150 and in total that outbreak affected about 1.5 million people (Solomon *et al.*, 2010). In
151 1999 an outbreak in Western Australia resulted in 6000 cases of HFMD and 29 cases
152 of CNS disease, at least nine of which were severe and four developed long-term
153 neurological sequelae (McMinn *et al.*, 2001). EV71 outbreaks, most of which had
154 lethal cases, continued in Korea, Singapore, Japan, Malaysia, Vietnam and Thailand

155 (Solomon *et al.*, 2010). In 2008 there was a large EV71 outbreak in China with almost
156 half a million reported HFMD cases and 122 fatal cases (Yang *et al.*, 2009). The virus
157 kept on circulating in China and contributed to HFMD cases with CNS complications
158 and fatalities at least until 2014 (Liu *et al.*, 2015a). By then, over 7.5 million cases of
159 HFMD were reported in China alone, of which over 80,000 had neurological
160 involvement and Chinese government declared the development of measures to
161 control EV71 spread as a national priority (Liu *et al.*, 2015b).

162 Enterovirus 71 infections have been lately detected in Europe, including
163 Denmark, France, Spain and the Netherlands, and although the incidence of EV71
164 infections there is low, occasional lethal cases have already been reported (Cabrerizo
165 *et al.*, 2014; Fischer *et al.*, 2014; van der Sanden *et al.*, 2011; Schuffenecker *et al.*,
166 2011). Interestingly, German researchers have recently described a case of pediatric
167 encephalitis caused by a novel EV71 genotype that likely arose from a recombination
168 event (Karrasch *et al.*, 2016). Although no epidemic activity of EV71 has so far been
169 reported in Europe, the European Centre for Disease Prevention and Control (ECDC)
170 risk assessment reported increased detections of EV71 in the first half of 2016 as
171 compared to previous years, necessitating preparedness to control EV71 spread in
172 Europe (ECDC, 2016).

173 Importantly, EV71 often alternates or co-circulates with other *Enterovirus A*
174 genotypes, mostly with CVA16, another recognized HFMD agent. Coxsackievirus
175 A16 also predominantly infects children but, unlike EV71, typically causes milder
176 symptoms and has much lower morbidity and mortality rate. In a comparative study
177 of 177 EV71 and 64 CVA16 patients in Taiwan, only 6.3% of CVA16 infections
178 developed aseptic meningitis, whereas 32% of EV71 cases resulted in aseptic
179 meningitis, encephalitis, polio-like syndrome, encephalomyelitis and fatal pulmonary

180 edema of which 7.9% were lethal (Chang *et al.*, 1999). However, occasional severe
181 and fatal CVA16 cases have been reported in USA (Wright *et al.*, 1963), Taiwan
182 (Wang *et al.*, 2004), France (Legay *et al.*, 2007), Japan (Goto *et al.*, 2009), and China
183 (Chen *et al.*, 2015).

184 The outbreaks of CVA16 were documented in Canada (1957) (Robinson *et al.*,
185 1958), Australia (1991) (Ferson & Bell, 1991), England and Wales (1994) (Bendig &
186 Fleming, 1996) and India (2009) (Kar *et al.*, 2013) and were linked to HFMD. It co-
187 circulated with EV71 in 1998 during the unprecedented HFMD epidemic in Taiwan
188 (Ho *et al.*, 1999a) and continued circulating in Taiwan becoming dominant in years
189 2002 and 2003 (Chang, 2008). The co-circulation of CVA16 and EV71 was
190 documented during HFMD outbreaks in Vietnam in 2005 (Tu *et al.*, 2007) and in
191 China in 2008 onwards (Liu, 2014). In Singapore CVA16 was the major cause of
192 HFMD epidemics in years 2002, 2005 and 2007, whereas in 2006 it was EV71 (Ang
193 *et al.*, 2009).

194 Co-circulation of EV71 and CVA16 allows viral co-infections (He *et al.*, 2013)
195 and recombination, which happened during the HFMD outbreak in China (Zhang *et*
196 *al.*, 2010a). Recombination between EV71 and CVA16 and accumulation of point
197 mutations can give rise to viruses with altered antigenicity, thus limiting population
198 protection and complicating control of viral spread. Both EV71 and CVA16 require
199 attention as clinically important pathogens.

200

201 **Enterovirus 68**

202 Enterovirus 68 belongs to the D species of the *Enterovirus* genus. Although
203 genetically it is an enterovirus, EV68 shares properties of both entero- and
204 rhinoviruses and in most cases causes respiratory infections (Oberste *et al.*, 2004). It

205 was first isolated in 1962, but only 26 cases were reported until 2005 and EV68
206 received little attention from clinical and scientific communities (Khetsuriani *et al.*,
207 2006). This changed in 2008–2010 when several outbreaks of acute respiratory illness
208 caused by EV68 were reported by the Centers for Disease Control and Prevention
209 (CDC) in the Philippines, Japan, the Netherlands and the USA (Imamura *et al.*, 2011).
210 The affected individuals typically presented with URTI symptoms—cough, fever,
211 rhinorea, difficulties in breathing and hypoxia—although severe lower respiratory
212 tract infections (LRTI) were also detected (Khetsuriani *et al.*, 2006). Individuals
213 infected with EV68 often required hospitalization: for example, prospective study in
214 the Netherlands reported that out of 24 EV68-positive subjects, 23 were hospitalized
215 (Imamura *et al.*, 2011). Deaths were reported in the Philippines and in Japan, but not
216 in the US and the Netherlands (Imamura *et al.*, 2011). In all studies EV68 was
217 reported as a paediatric pathogen, except for the work done by Meijer *et al.* who
218 reported a significant number of patients over 50 years old (Meijer *et al.*, 2014).

219 Following the initial outbreaks, EV68 continued its seasonal circulation, and
220 was occasionally detected in respiratory samples from paediatric patients with URTI
221 and severe LRTI in different countries, further supporting its clinical relevance and
222 place as a concern for medical society (Esposito *et al.*, 2015; Gimferrer *et al.*, 2015;
223 Imamura *et al.*, 2013; Lu *et al.*, 2014). The concern has been raised further after the
224 EV68 outbreak in the USA in 2014 (Khan, 2015) when over one thousand patients
225 from 47 states have been diagnosed with acute respiratory illness (ARI) caused by
226 EV68. The outbreak resulted in a significant increase in hospital admissions: in
227 Kansas City alone over 300 patients were hospitalized, of which 15% were admitted
228 to ICU and 15 cases were fatal. Simultaneously, an increased incidence of EV68 ARI
229 was reported in Canada, where over 200 cases have been identified, resulting in 140

230 hospitalizations and one death (Khan, 2015). In 2016 EV68 was detected in patients
231 with neurological manifestations in the Netherlands, France, UK, Italy, Portugal and
232 Germany (ECDC, 2016).

233 Intriguingly, the 2014 EV68 outbreak in North America overlapped with an
234 outbreak of a polio-like disease with the brain stem and the spinal cord grey matter
235 lesions and AFP. Infection with EV68 was confirmed in 5 out of 11 (45%) of the
236 American AFP patients (Messacar *et al.*, 2015). Furthermore, EV68 has been also
237 detected in four cases of AFP in Canada, two in Norway and one in France (Khan,
238 2015; Lang *et al.*, 2014; Pfeiffer *et al.*, 2015). A recent retrospective study identified
239 EV68 in respiratory secretions from 12 of 25 (48%) patients with sporadic paralysis,
240 strengthening the EV68 link to CNS disease (Greninger *et al.*, 2015). Interestingly, all
241 the EV68 strains identified in association with paralytic disease formed a distinct
242 genetic cluster suggesting ongoing emergence and adaptation of this virus (Du *et al.*,
243 2015). Direct linkage of EV68 to neurological disease has been complicated by
244 difficulties to detect EV68 or its RNA in patients' CSF and so far only two studies
245 succeeded to detect EV68 in patients' CSF (Khetsuriani *et al.*, 2006; Kreuter *et al.*,
246 2011). However, we should note that other neurological picornaviruses—PV and
247 EV71—are also rarely recovered from the CSF, and therefore neurological
248 involvement of EV68 cannot be ruled out on the basis of negative CSF samples.

249 The incidence of EV68 has clearly increased over the last decade. Although
250 this observation could be related to significant improvements in the detection
251 techniques, the accumulating clinical data suggests that EV68 should be considered
252 an emerging pathogen. The concerns raised by respiratory EV68 infections and
253 especially by their possible link to AFP necessitate careful surveillance of the virus

254 spread, detailed studies of its pathogenesis and evolution and development of
255 preventative and/or treatment options.

256

257 **Human parechovirus 3**

258 Human parechovirus 3 belongs to species A within genus *Parechovirus* and is
259 the second most common human parechovirus (Wolthers *et al.*, 2008). HPeV3 was
260 isolated in 1999 from an infant with severe CNS disease (Ito *et al.*, 2004), and since
261 then it has been recognized as the most or second most prevalent virus causing CNS
262 infections in infants under 3 months old (Harvala *et al.*, 2009, 2011; Piralla *et al.*,
263 2014; van der Sanden *et al.*, 2008). Outbreaks of HPeV3 usually occur in summer-
264 autumn seasons and have a distinct biennial pattern (Harvala *et al.*, 2011; Wolthers *et*
265 *al.*, 2008). They have been documented in Europe (Benschop *et al.*, 2006), North
266 America (Boivin *et al.*, 2005), Asia (Yamamoto *et al.*, 2009), and Australia
267 (Cumming *et al.*, 2015) and are regularly associated with a variety of clinical
268 presentations, from mild gastrointestinal or respiratory illness to life-threatening
269 conditions in neonates (Esposito *et al.*, 2014; Harvala *et al.*, 2011; Tapia *et al.*, 2008).
270 It can cause systemic infections with possible neurological involvement in infants that
271 are collectively described as “sepsis-like illnesses” (Benschop *et al.*, 2006;
272 Selvarangan *et al.*, 2011; Wolthers *et al.*, 2008). Such illnesses typically present with
273 fever, seizures, irritability, respiratory and gastrointestinal problems and occasional
274 rash being indistinguishable from severe EV infections (Shoji *et al.*, 2013; Verboon-
275 Maciolek *et al.*, 2008). The fraction of symptomatic HPeV3-infected infants that
276 develop sepsis-like illness can exceed 80%; most of such patients require
277 hospitalization and up to one third of them are admitted to the ICU (Schuffenecker *et*
278 *al.*, 2012; Selvarangan *et al.*, 2011; Shoji *et al.*, 2013). The CNS symptoms of

279 HPeV3 infection can include meningitis, meningoencephalitis, encephalitis or
280 cerebral hemorrhage with occasional white matter alterations (Khatami *et al.*, 2015;
281 Kurz *et al.*, 2015). Whereas HPeV3 meningitis typically has good prognosis,
282 meningoencephalitis entailing white matter alterations may have long-term sequelae
283 such as cerebral palsy, learning disabilities, epilepsy or visual impairment (Verboon-
284 Maciolek *et al.*, 2008). In addition, HPeV3 is occasionally associated with
285 hemophagocytic lymphohistiocytosis (Aviner *et al.*, 2014) and sudden death
286 syndrome in infants (Schuffenecker *et al.*, 2012). Furthermore, in Japan it was linked
287 to myositis in children and epidemic myalgia in adults (Mizuta *et al.*, 2013;
288 Yamamoto *et al.*, 2015). Fatal cases of HPeV3 infections are known: they resulted
289 from encephalitis in the absence of an immune response and sometimes also involved
290 white matter necrosis (Bissel *et al.*, 2015; van Zwol *et al.*, 2009).

291 Overall, HPeV3 represents a significant threat to neonatal health care. The
292 incidence of HPeV3 infections and number of CNS disease cases increases (Harvala
293 *et al.*, 2011) with no treatment options available necessitating search for antivirals and
294 understanding of HPeV neutralization by antibodies (Wildenbeest *et al.*, 2010).

295

296 **NEUTRALIZING ANTIBODIES IN PICORNAVIRUS INFECTIONS**

297 Like with other viruses, the severity and outcome of picornavirus infections depends
298 both on the viral and host factors. Host immune status is a key regulator of infection,
299 and failure to mount an appropriate response inevitably leads to severe disease. The
300 viral infection is detected by the specific pattern recognition receptors of the innate
301 immunity that establish a complex signalling network, triggering the expression of
302 antiviral genes in infected cells and the activation of specific adaptive responses
303 (Dotzauer & Kraemer, 2012). The adaptive responses rely on the specific populations

304 of T-cells and antibody-producing B-cells. Although both innate and cellular adaptive
305 immunity are essential, a large body of evidence indicates that efficient production of
306 specific antibodies by the B-cells is primary for the control of picornaviral infections.

307 Picornaviruses typically infect children, likely due to their naive immune
308 system. Severe picornavirus infections in healthy adults are uncommon. However,
309 immunocompromised adults, in particular those with impaired B-cell responses, are
310 susceptible to prolonged and/or severe picornavirus infections. For example, patients
311 with X-linked agammaglobulinemia (XLA) have markedly reduced levels of B-cells
312 and serum antibodies and are susceptible to enterovirus infections (Halliday *et al.*,
313 2003) with severe neurological manifestations (Quartier *et al.*, 2000). Moreover,
314 patients with a- or hypogammaglobulinemia can also develop chronic HPeV1
315 infection (van de Ven *et al.*, 2011), as well as HPeV3 myocarditis and encephalitis
316 that are extremely uncommon in adults (Mardekian *et al.*, 2015). Individuals
317 undergoing immunosuppressive therapy further support the critical role of antibody
318 responses in the control of picornaviruses. For instance, cancer treatment with
319 rituximab leads to prolonged B-cell deficiency and hypogammaglobulinemia and
320 patients receiving such therapy are susceptible to severe and even lethal enterovirus
321 infections (Servais *et al.*, 2010).

322 Direct confirmation for the role of antibodies in picornavirus infections comes
323 from controlled infections in animal models. Experimental infections in mice with
324 different immunodeficiencies proved the significance of adaptive responses: whereas
325 up to 70% of mice deficient in innate immune responses survived EV71 infection,
326 mice with severe combined immunodeficiency developed limb paralysis and died in
327 almost 100% of cases (Liao *et al.*, 2014). In Theiler's encephalomyelitis virus
328 (TMEV)-infected mice—a common model for neurotropic picornaviruses—

329 immunosuppression with anti-IgM antibodies led to virus-induced demyelination
330 (Rodriguez *et al.*, 1990).

331 At the moment there are no antivirals for treatment of severe picornavirus
332 infections (Linden *et al.*, 2015; Wildenbeest *et al.*, 2010) and the only therapeutic
333 option is intravenous immunoglobulin (IVIG). However, because of the low BBB
334 permeability to antibodies, IVIG is rarely effective in CNS infections although
335 intraventricular immunoglobulin administration may be beneficial (Quartier *et al.*,
336 2000). Antibodies can also be effective at mucosal sites and prevent picornavirus
337 viremia and CNS invasion (Nathanson & Bodian, 1962). Successful management of
338 severe picornavirus infections using IVIG (Wildenbeest *et al.*, 2013) indicates the
339 potential efficacy of passive immunization to control picornavirus infections.
340 However, the presence of specific neutralizing antibodies and their titres cannot be
341 controlled in IVIG preparation, and the reliable options of passive immunization
342 against picornaviruses should be based on the production of specific neutralizing or
343 broadly neutralizing antibodies that target known viral epitopes. Production of
344 specific antibodies could also contribute to rapid and specific serology-based
345 diagnostics of picornavirus infections that are beneficial at time-critical point-of-care
346 setups. In addition to passive immunization, the success of the polio vaccine
347 encourages development of vaccines against other picornaviruses. Controlling
348 picornavirus infections with vaccines is not a feasible approach for the entire
349 *Picornaviridae* family, but can be realistic for some virus types. Both vaccine and
350 antibody development require thorough understanding of viral neutralization, and
351 below we summarize the current knowledge of the neutralization of CNS-invading
352 picornaviruses.

353 NEUTRALIZATION OF PICORNAVIRUSES

354 Neutralization of EV71 and CVA16

355 EV71 is genetically diverse and contains 14 genotypes: A, B1–B5, C1–C5, D, E, and
356 F (Bessaud *et al.*, 2014); C4 is further classified into two lineages C4a and C4b
357 (Zhang *et al.*, 2010a). Genotypes B3, B4, B5, C1, C2, C4 and C5 contributed to recent
358 outbreaks (Chong *et al.*, 2015). CVA16 is less genetically diverse showing relatively
359 slower evolutionary rate and has 3 genotypes: A, B1 (B1a, B1b, B1c) and B2 (Zhang
360 *et al.*, 2010b). EV71 can undergo intra- and intergenotype shifts that occur due to
361 recombination events during co-circulation with CVA16 or different EV71 genotypes
362 and correlate with most outbreaks (Bible *et al.*, 2007). An effective vaccine should
363 neutralize multiple serotypes of EV71 and also CVA16. This necessity underpins
364 difficulties in EV71 vaccine development.

365 The first live-attenuated vaccine strain was reported by Arita *et al.* in 2005. It
366 induced broadly-neutralizing responses in immunized monkeys, but was neurotropic
367 when inoculated intravenously and its further development was halted (Arita *et al.*,
368 2007). Development of inactivated vaccines was more successful: five such vaccines
369 developed by different organizations entered clinical trials and three of them have
370 already completed Phase III showing 80.4–97.4% efficacy against EV71-induced
371 HFMD in humans (Liu *et al.*, 2015a). Two C4-based vaccines developed by the
372 Chinese Academy for Medical Sciences (CAMS) and Sinovac Biotech Co Ltd were
373 approved by the Chinese Food and Drug Administration (CFDA) as of January 2016
374 (Mao *et al.*, 2016). In addition to live-attenuated and inactivated vaccines, virus-like
375 particle (VLP) vaccine candidates were generated in baculovirus or *Saccharomyces*
376 *cerevisiae* systems using co-expression of viral protein precursor P1 with viral
377 protease 3CD (Chung *et al.*, 2008; Li *et al.*, 2013). The VLP vaccine candidate

378 produced in baculovirus system showed promise in *in vivo* studies: the survival rate of
379 VLP-immunized mouse pups after lethal EV71 challenge was superior to those
380 immunized with inactivated virus (Chung *et al.*, 2008) and it also induced protective
381 responses in monkeys (Lin *et al.*, 2012). The approved inactivated vaccines cross-
382 protected against B1, B5 and C4a (CAMS vaccine) (Chou *et al.*, 2013) and B4, B5,
383 C2 and C5 (Sinovac vaccine) (Mao *et al.*, 2013); VLP vaccines protected monkeys
384 against B4, B5, C3, C4 and C5 (Lin *et al.*, 2012), but none of them neutralized
385 CVA16.

386 Multivalent vaccines may offer protection against EV71 and co-circulating
387 CVA. Bivalent EV71/CVA16 vaccines based on inactivated viruses or VLP can elicit
388 high titres of neutralizing antibodies in immunized mice and protect from EV71 and
389 CVA16 infections (Ku *et al.*, 2014). Yet broader protection is desired to mitigate
390 other HFMD contributors, such as coxsackievirus A6 (CVA6) (Liu *et al.*, 2014), and
391 one trivalent EV71/CVA16/CVA6 inactivated vaccine candidate protecting mice
392 from lethal challenge with these viruses was reported (Caine *et al.*, 2015).

393 Peptide vaccines consisting of well-defined neutralizing epitopes represent a
394 promising approach to target against several heterologous viruses (Li *et al.*, 2014a).
395 They are easier to produce compared to inactivated virus vaccines, do not require
396 handling live virus and allow immunization with lower protein load. Neutralizing
397 epitopes on EV71 are localized on viral structural proteins VP1, VP2 and VP3 (Fig. 3
398 (a) and (b)). Three continuous neutralizing epitopes have been localized to VP1
399 residues 163–177 (known as SP55) and 215–219 (part of SP70 which localizes to
400 residues 208–222) and to region 240–260 (Chang *et al.*, 2010; Foo *et al.*, 2007; Lim *et*
401 *al.*, 2012). SP55 has 85–100% sequence identity within A–C4 groups (Foo *et al.*,
402 2007) and SP70 is 100% conserved across EV71 genotypes A–C4 and thus is

403 universal for them. In addition, VP1 encompasses a strain-specific discontinuous
404 neutralization epitope at the 5-fold symmetry axis with residue 145 critically
405 contributing to antibody interaction (Lee *et al.*, 2013). Two other neutralizing
406 epitopes were localized to residues 136–150 of VP2 (known as VP2-28) (Liu *et al.*,
407 2011), and to VP3 residues 55–69 that form a “knob” and are 100% conserved across
408 EV71 subgenogroups A-C4 (Kiener *et al.*, 2014). Much less is known about
409 neutralizing epitopes in CVA16. Most of the experimentally proven neutralizing
410 epitopes of CVA16 were localized to VP1 (GH, EF, C-terminal loops and B and C β -
411 sheets) (Ren *et al.*, 2015; Shi *et al.*, 2013); one more continuous epitope was found in
412 GH loop of VP3 (Chong *et al.*, 2012) (Fig 3 (b)). Additional antigenic sites of CVA16
413 were predicted *in silico* in EF and HI loops of VP2 (Ren *et al.*, 2015). Although
414 peptide vaccines are usually less immunogenic compared to the viral particles, this
415 can be mitigated using adjuvants or fusing viral epitopes with highly immunogenic
416 antigens. Using such an approach, a tandem of three well described EV71 epitopes—
417 SP55, SP70 and VP2-28—separated by Gly-Ser linker and fused to thioredoxin was
418 expressed in *E. coli*. The recombinant protein induced EV71-specific neutralizing
419 responses in immunized mice, serving as a proof-of-concept for peptide vaccines
420 against HFMD (Li *et al.*, 2014b). In another study, a hepatitis B virus-like particle
421 displaying EV71 SP55 and VP2-28 epitopes induced neutralizing responses in
422 immunized mice, and could also be cross-protective against CVA16 (Xu *et al.*,
423 2015b). In terms of immunogenicity EV71 is one of the best-studied picornaviruses
424 with two EV71 vaccines approved by CFDA and further work will be driven by the
425 necessity of multivalent HFMD vaccines.
426

427 **Neutralization of enterovirus 68**

428 EV68 is an emerging virus and little is known about its antigenicity. Imamura *et al.*
429 studied the immunogenic properties of twelve EV68 genotypes belonging to all three
430 genetic lineages of EV68—A, B and C (Imamura *et al.*, 2013). Immunized guinea
431 pigs generated high titres of neutralizing antibodies to the original virus and to viruses
432 of the same lineage, but very little cross-neutralization between genetic lineages was
433 found (Imamura *et al.*, 2014). Importantly, the majority of sequence variation between
434 EV68 lineages is localized to the VP1 BC and DE surface loops (Imamura *et al.*,
435 2013, 2014; Liu *et al.*, 2015b), which are the most variable and likely are epitope-
436 containing in enteroviruses. Indeed, VP1 likely contains antigenic determinants as the
437 increase in its gene diversity correlates with an increase in the number of EV68
438 detections (Meijer *et al.*, 2012), reflecting the appearance of antigenically new viruses
439 in the population. Substitution dynamics within the viral genome also suggests the
440 localization of antigenic epitopes to VP1: several positions in VP1 BC and DE loops
441 (Fig. 3 (b)) are undergoing positive selection and might be associated with antigenic
442 differences between EV68 genetic lineages (Imamura *et al.*, 2014). Nevertheless, to
443 date the exact immunogenic epitopes of EV68 are only predictive and have not been
444 mapped.

445 Not much is also known about the distribution of EV68 neutralizing antibodies
446 in human population. One study done in Finland addressed this question reporting
447 high titres of neutralizing responses to EV68 in 80% of the studied individuals (Smura
448 *et al.*, 2010). However, neutralization responses in this study were addressed against
449 the prototype EV68 Fermon strain, which is antigenically very different from the
450 currently circulating EV68 strains (Greninger *et al.*, 2015; Meijer *et al.*, 2012).

451 Therefore, neither EV68 antigenicity, nor the population protection levels are
452 understood well at the moment.

453

454 **Neutralization of HPeV3**

455

456 Almost nothing is known about HPeV3 antigenicity. Of the parechoviruses, only
457 antigenicity of HPeV1 has been studied (Shakeel *et al.*, 2015) and so far three
458 neutralizing epitopes on HPeV1 structural proteins VP0, VP1 and VP3 are described.
459 One is localized to residues 83-97 of VP0 (Joki-Korpela *et al.*, 2000), another is found
460 on VP1 and encompasses receptor recognizing arginine-glycine-aspartic acid (RGD)
461 motif (Alho *et al.*, 2003) and the third one is formed by VP0 and VP3 (Shakeel *et al.*,
462 2015; Westerhuis *et al.*, 2015). Whereas antisera generated against HPeV1 VP0
463 peptide was not tested for HPeV3 neutralization, two other monoclonal antibodies did
464 not cross-react with HPeV3 (Shakeel *et al.*, 2015; Westerhuis *et al.*, 2015). For
465 HPeV3 only non-neutralizing epitope has been described (Shakeel *et al.*, 2016)

466 The data on HPeV3 seroprevalence in population is also sparse. A study of
467 sera from populations in Finland and Netherlands revealed neutralizing responses to
468 HPeV3 only in about 10% of the samples and very low titres of neutralizing
469 antibodies post infection (Westerhuis *et al.*, 2013). On the contrary, researchers in
470 Japan detected neutralizing responses to HPeV3 in 67% of individuals between 7
471 months and 40 years old (Ito *et al.*, 2004), and reported high titres of neutralizing
472 antibodies after 3 months post infection in all studied individuals (Aizawa *et al.*,
473 2015). The virus strains used in European and Japanese studies were different,
474 suggesting that some strains of HPeV3 are strongly immunogenic, whereas others are
475 not. The determinants of immunogenicity within this virus are currently unknown and

476 investigation of the Japanese A308/99 strain immunogenicity may shed light on the
477 neutralization of HPeV3.

478 The treatment option for human parechovirus infections could be IVIG
479 (Wildenbeest *et al.*, 2013), but titres of neutralizing antibodies against HPeV3 in
480 European IVIG preparations are very low (Westerhuis *et al.*, 2012). Developing
481 vaccines does not seem feasible because subjects of severe HPeV3 infections are
482 infants. Passive immunization protects against HPeV3 (Aizawa *et al.*, 2015) thus
483 developing therapeutic antibodies is a necessity for which studies of HPeV3
484 antigenicity are required.

485

486 **CONCLUSIONS AND FUTURE PERSPECTIVES**

487 Our understanding of antigenicity of picornaviruses that target CNS is very poor and
488 is largely limited to studies of EV71. Although studies of EV71 have already resulted
489 in two CFDA-approved HFMD vaccines, multivalent HFMD vaccines to control
490 different EV71 genotypes and also co-circulating CVA16 are the next goal. Proof-of-
491 concept studies of such vaccines are promising (Caine *et al.*, 2015; Sun *et al.*, 2014),
492 but further work is needed to identify the optimal combination of antigens for
493 balanced, broadly protective immunity. Targeting multiple viruses with a single shot
494 also requires delivery systems for effective presentation of multiple epitopes. In this
495 regard, VLP and peptide vaccines may be preferable over inactivated vaccines,
496 offering tailored solutions in terms of presented antigens together with comparable
497 immunogenicity, high safety and less tedious production, and economical feasibility
498 (Li *et al.*, 2014a).

499 We know almost nothing about EV68 and HPeV3 antigenicity. Classical
500 approach to study virus antigenicity and develop vaccines relies on animal models,

501 which is just being developed for EV68 and not available for HPeV3, hampering
502 investigation of these viruses. Therefore, their antigenicity should be studied directly
503 from human sera using novel methods, such as peptide arrays (Hansen *et al.*, 2013),
504 classical phage display or phage display enhanced with next generation sequencing
505 (NGS) (Christiansen *et al.*, 2015). Successful use of custom phage display library and
506 NGS was reported by Xu *et al.* who analysed antibody-peptide interactions in sera of
507 over 550 donors and identified numerous previously undescribed viral epitopes,
508 proving the utility of NGS-enhanced phage display for epitope identification (Xu *et*
509 *al.*, 2015a).

510 Another future direction is the search for EV68 and HPeV3 neutralizing
511 antibodies. In the absence of antivirals (Linden *et al.*, 2015; Wildenbeest *et al.*, 2010),
512 such antibodies could be valuable therapeutics, especially for HPeV3 that infects
513 infants for whom vaccination is not a suitable option. A useful approach for that is
514 identification of individual's immune response to a given virus followed by respective
515 B-cell cloning (Kwakkenbos *et al.*, 2014). This approach was utilized to generate two
516 broadly neutralizing antibodies against HPeV (Shakeel *et al.*, 2015; Westerhuis *et al.*,
517 2015). Unfortunately, these antibodies did not neutralize HPeV3. High throughput
518 approaches, such as sequencing of antibody repertoire (Georgiou *et al.*, 2014) and
519 screening of large antibody fragments libraries using ribosomal, bacterial, yeast or
520 phage display (Bradbury *et al.*, 2011) are also utilized for antibody discovery. For
521 instance, phage display technology has already yielded monoclonal neutralizing
522 antibodies with therapeutic potential for EV71 (Zhang *et al.*, 2015). Over 50 phage-
523 display derived antibodies have been approved by the U.S. Food and Drug
524 Administration (FDA) or European Medicines Agency (EMA) as of May, 2016.
525 About 500 antibodies are undergoing clinical trials. These include antibodies targeting

526 infectious agents, for example Rabies virus, which is now in Phase II (Frenzel *et al.*,
527 2016).

528 Overall, picornaviruses antigenicity and neutralization studies have already
529 brought encouraging results, however more challenges are ahead. The detailed
530 understanding of viral immunogenicity is clearly an important task to focus on, but it
531 should be carried out in parallel with broader studies of viral biology and spread.
532 Although many research groups in Europe, US and Asia are very active in the field of
533 picornavirus research, we are still far from a thorough understanding of viral
534 epidemiology, pathogenesis, evolution and inhibition, which are necessary for
535 effective virus control. Apart from scientific challenges, development protective
536 measures against picornaviruses may face additional financial and regulatory
537 difficulties due to the endemic nature of diseases that they cause. Hence, drawing the
538 public attention to the health care threats that picornaviruses impose is another
539 important activity area for the medical and scientific communities.

540

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550

551 **CONFLICTS OF INTEREST**

552 The authors declare no conflicts of interest.

553

554 **ETHICAL STATEMENT**

555 The authors declare that there are no ethical issues.

556

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FIGURE LEGENDS

Fig. 1. Classification of picornaviruses. The scheme shows picornavirus genera and species that infect humans. Clinically important genotypes are also shown. The genotypes associated with neurological infections are highlighted with heavy font.

Fig. 2. Routes of picornavirus entry to the CNS. (a) Picornaviruses can infect peripheral nerve and invade CNS via retrograde axonal transport. (b) They can and spread further in CNS in a trans-synaptic manner. (c) During viremia picornaviruses can enter the CNS via hematogenous route through a disintegrated blood-brain barrier. (d) They can also cross the blood-brain-barrier in an active manner, possibly relying on cellular transferrin receptor 1. (e) Picornaviruses can infect migrating cells and invade CNS in a “Trojan horse” manner.

Fig. 3. Enterovirus structure and localization of immunogenic epitopes on viral surface. (a) Icosahedral picornavirus capsid, shown for EV71 (PDB ID: 3VBS), consists of 60 identical structural units (asymmetric units). Each asymmetric unit is composed of four viral structural proteins: VP1 (dark green), VP2 (grey), VP3 (light green) and VP4 (attached to the inner surface of the capsid and not seen in the cartoon). (b) Localization of known immunogenic epitopes on EV71 (PDB ID: 3VBS) (upper panel), CVA16 (PDB ID: 5C4W) (middle panel) and predicted epitope-containing loops on EV68 (PDB ID: 4WM8) (lower panel). For simplicity, only one structural unit is mapped on each virus. Structural proteins VP1 (dark green), VP2 (grey) and VP3 (light green) are shown. The epitopes are marked in orange and are pointed with arrows on a zoomed image. Overlapping epitopes are marked in red.

Figure 1

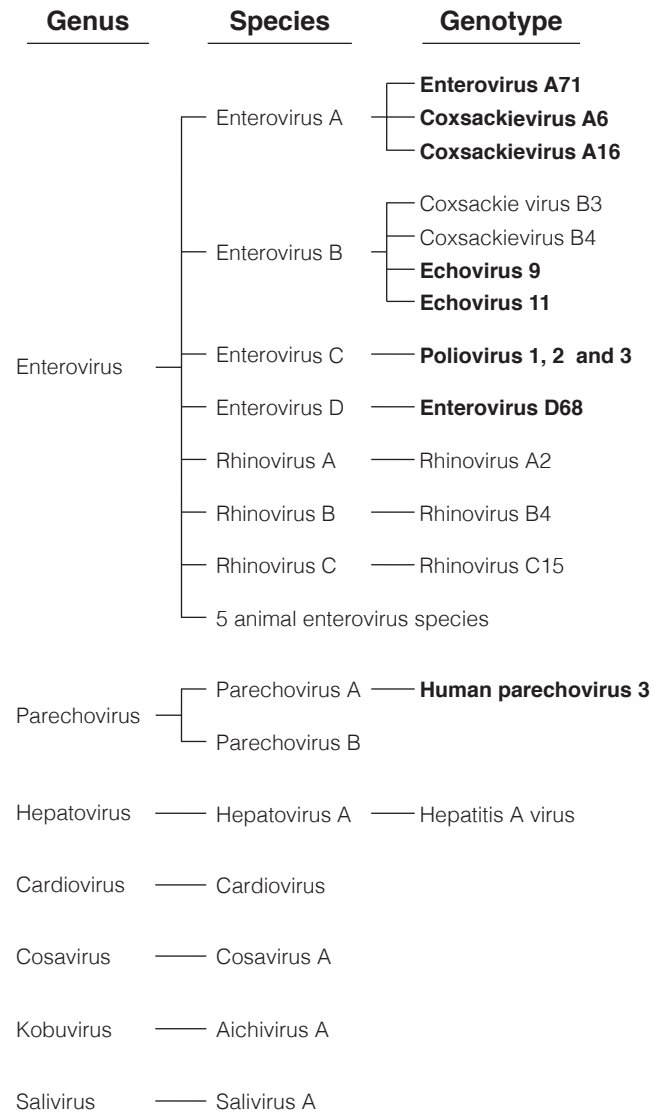


Figure 2

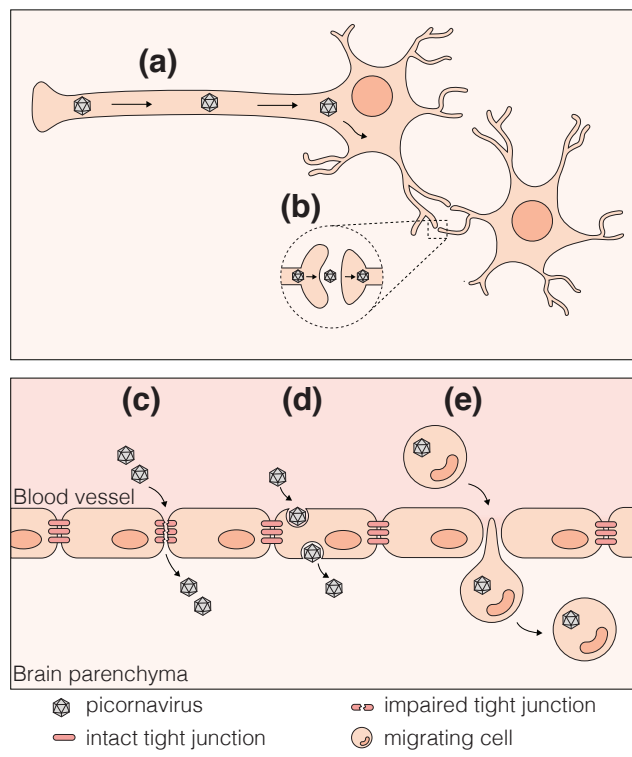
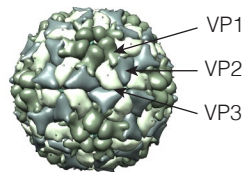
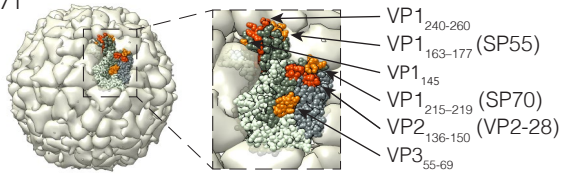


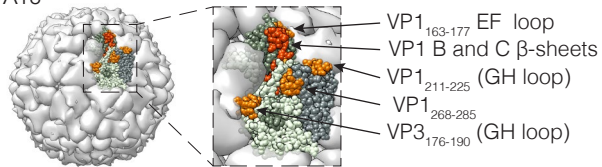
Figure 3

(a)**(b)**

EV71



CVA16



EV68

