

**Pathogens Penetrating the Central Nervous System: Infection Pathways and the Cellular and Molecular Mechanisms of Invasion**

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Published

2014

Journal Title

Clinical Microbiology Reviews

DOI

[10.1128/CMR.00118-13](https://doi.org/10.1128/CMR.00118-13)

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**Pathogens penetrating the central nervous system: infection pathways and the cellular and molecular mechanisms of invasion**

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Running title: Bacterial mechanisms of brain invasion

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## **Table of Contents**

<b>SUMMARY</b>	<b>5</b>
<b>INTRODUCTION</b>	<b>6</b>
<b>Invasion of the CNS by Bacteria: a Clinical Perspective</b>	<b>6</b>
<b>Meningitis</b>	<b>6</b>
i. <b>Pneumococcal meningitis</b>	<b>8</b>
ii. <b>Meningococcal meningitis</b>	<b>8</b>
iii. <b><i>H. influenzae</i>, <i>L. monocytogenes</i> and other bacterial causes of meningitis</b>	<b>9</b>
<b>Encephalitis</b>	<b>10</b>
<b>Focal infections</b>	<b>12</b>
<b>BARRIERS OF THE CNS</b>	<b>13</b>
<b>The Blood-Brain Barrier</b>	<b>14</b>
<b>The Blood-Cerebrospinal Fluid Barrier</b>	<b>15</b>
<b>Immunosurveillance of the CNS</b>	<b>15</b>
<b>MICROBIAL ENTRY TO THE CNS</b>	<b>17</b>
<b><i>In vivo</i> and <i>in vitro</i> models of bacterial CNS invasion</b>	<b>18</b>
<b>BACTERIAL INVASION THROUGH THE BLOOD-BRAIN AND BLOOD-CEREBROSPINAL FLUID BARRIERS</b>	<b>19</b>
<b>Sites of Entry</b>	<b>20</b>

<b>Mechanisms of Entry</b>	<b>21</b>
<b>Transcellular Penetration of Brain Microvascular Endothelial Cells</b>	<b>22</b>
<i>E. coli</i>	23
<i>N. meningitidis</i>	25
<i>S. pneumoniae</i>	27
<b>Group B streptococcus</b>	<b>28</b>
<i>S. suis</i>	31
<i>L. monocytogenes</i> and <i>B. pseudomallei</i>	32
<i>M. tuberculosis</i>	35
<b>Paracellular Penetration of Brain Microvascular Endothelial Cells</b>	<b>36</b>
<i>E. coli</i>	37
<i>H. influenzae</i>	39
<i>N. meningitidis</i>	40
<i>Streptococcus</i> spp.	41
<b>Trojan Horse Penetration of Brain Microvascular Endothelial Cells</b>	<b>43</b>
<b>THE OLFACTORY NERVE AS A PORTAL TO THE BRAIN</b>	<b>45</b>
<b>THE TRIGEMINAL NERVE AS A PORTAL TO THE BRAIN</b>	<b>46</b>
<b>PROTECTING THE CNS FROM MICROBIAL INVASION VIA THE NASAL CAVITY</b>	<b>47</b>

<b>PATHOGENS THAT ENTER THE BRAIN THROUGH THE NOSE</b>	<b>49</b>
<b>Bacteria</b>	<b>49</b>
<i>S. pneumoniae</i>	<b>50</b>
<i>N. meningitidis</i>	<b>51</b>
<i>B. pseudomallei</i>	<b>53</b>
<i>L. monocytogenes</i>	<b>55</b>
<b>Viruses</b>	<b>56</b>
<b>Protozoa</b>	<b>62</b>
<b>Yeasts and fungi</b>	<b>63</b>
<b>CONCLUDING REMARKS</b>	<b>65</b>
<b>REFERENCES</b>	<b>67</b>
<b>AUTHOR BIOS</b>	<b>123</b>

## SUMMARY

The brain is well protected against microbial invasion by cellular barriers, such as the blood-brain barrier (BBB) and the blood-cerebrospinal fluid barrier (BCSFB). In addition, cells within the central nervous system (CNS) are capable of producing an immune response against invading pathogens. Nonetheless, a range of pathogenic microbes make their way to the CNS and the resulting infections can cause significant morbidity and mortality. Bacteria, amoebae, fungi and viruses are capable of CNS invasion, with the latter using axonal transport as a common route of infection. In this review we compare the mechanisms by which bacterial pathogens reach the CNS and infect the brain. In particular, we focus on recent data regarding mechanisms of bacterial translocation from the nasal mucosa to the brain, which represents a little explored pathway of bacterial invasion but has been proposed as being particularly important in explaining how infection with *Burkholderia pseudomallei* can result in melioidosis encephalomyelitis.

# 1 INTRODUCTION

## 2 Invasion of the CNS by bacteria: a clinical perspective

3 Microbial infection involving the CNS is an important and relatively common presentation.  
4 CNS infections are frequently caused by viruses, such as the enteroviruses, which cause the  
5 majority of cases of aseptic meningitis and meningoencephalitis (1-3). Other neurotropic  
6 viruses such as human cytomegalovirus, herpes simplex viruses, varicella zoster virus and the  
7 emerging viruses West Nile virus, henipaviruses, Japanese encephalitis virus, chikungunya  
8 virus, Ebola virus and rabies virus may also cause CNS infections (4-7). There are many  
9 bacterial pathogens that are associated with CNS invasion. Rapid detection of a bacterial  
10 agent in such presentations and the initiation of appropriate antibiotic therapy influences  
11 morbidity and mortality (8-10). The clinical presentations of bacterial CNS infection range  
12 from meningitis and meningoencephalitis to focal CNS syndromes.

## 13 Meningitis

14 Meningitis, inflammation of the meninges, is usually acute but can also be subacute and most  
15 frequently presents with headache, fever and neck stiffness (11). Meningitis can be pyogenic  
16 (pus-forming) - associated with common bacterial etiologies (described below), or aseptic, in  
17 which pyogenic bacteria are not isolated from the cerebrospinal fluid (CSF) and the causative  
18 agent may be viral or mycobacterial or there is a non-infective etiology. Features of CSF,  
19 which would generally exclude a bacterial etiology for meningitis, include a protein  
20 concentration of  $< 600$  mg/L and a white cell count of  $< 90 \times 10^6$ /L (2). The classic triad of  
21 fever, neck stiffness and a change in mental state is observed in 44% of patients with  
22 bacterial meningitis (11).

23 A prospective, laboratory-based surveillance study in 1986 defined the epidemiology of

24 bacterial meningitis in a population of 34 million in the United States (12). This study  
25 reported an average of 6.7 cases of bacterial meningitis per 100, 000 population across six  
26 states, and demonstrated that *Haemophilus influenzae*, *Neisseria meningitidis* and  
27 *Streptococcus pneumoniae* caused 77% of cases, which was consistent with a previous report  
28 by the National Surveillance of Bacterial Meningitis (13). More recently, it was reported that  
29 the incidence of bacterial meningitis in the United States declined by 31% during the  
30 surveillance period of 1998 to 2007, most likely due to the success of vaccine programs (14).  
31 *S. pneumoniae* was the most common cause of bacterial meningitis (58.0%), followed by  
32 *Streptococcus agalactiae* (group B streptococcus, 18.1%), *N. meningitidis* (13.9%), *H.*  
33 *influenzae* (6.7%) and *Listeria monocytogenes* (3.4%). *Staphylococcus aureus* and  
34 *Escherichia coli* are also important causes of bacterial meningitis (15) and *Streptococcus suis*  
35 is emerging as a zoonotic etiology of meningitis (16); however, it must be noted that  
36 meningitis can be caused by a plethora of different bacteria (17). The most common  
37 alternative bacterial cause of meningitis is *Mycobacterium tuberculosis*, which is recognized  
38 particularly in pediatric populations of the developing world (18, 19). These infections  
39 involve a clinical onset that is usually insidious with hydrocephalus being a prominent  
40 feature.

41 Bacterial meningitis is usually preceded by nasopharyngeal or middle ear colonization,  
42 followed by: (i) microbial invasion of the tissue and intravascular space, and (ii) bacteremia.  
43 Meningeal invasion occurs following penetration of the cellular barriers of the CNS. High-  
44 level bacteremia was shown to be necessary for the development of meningitis in  
45 experimental animal models (20, 21), which is consistent with clinical observations in  
46 humans (22-24). Bacteria that commonly cause meningitis express anti-phagocytic capsular  
47 polysaccharide, which enables survival/multiplication within the blood. Several species-  
48 specific factors that promote bacterial survival in blood, resistance to complement and



49 survival within phagocytes have been identified (25-30). Bacterial meningitis resulting from  
50 non-hematogenous spread is less common and is a consequence of infection pre-existing in  
51 an adjacent intracranial site. This would include mastoiditis, frontal sinusitis or rupture of an  
52 intracerebral abscess into the ventricles of the brain. Such infections may occur following  
53 head trauma or neurosurgical procedures.

#### 54 **Pneumococcal meningitis**

55 There are no specific distinguishing clinical features of pneumococcal meningitis. While a  
56 rash is uncommon, it is occasionally seen in splenectomised patients with overwhelming  
57 sepsis. There is a close correlation between bacteremic and meningitic serotypes of *S.*  
58 *pneumoniae* implicating hematogenous spread as the commonest mode of acquisition (31). It  
59 is the most frequent cause of bacterial meningitis in the United States and has a case fatality  
60 rate of 14.7% (14). Since the introduction of the 7-valent pneumococcal conjugate vaccine  
61 (which contains serotype antigens 4, 6B, 9V, 14, 18C, 19F and 23F) in the United States, the  
62 incidence of pneumococcal meningitis declined by 30.1% from 1.13 cases per 100, 000  
63 population in 1998 – 1999 to 0.79 cases per 100, 000 population in 2004 – 2005 (32). In  
64 patients < 2 years and > 65 years of age, the incidence of pneumococcal meningitis  
65 throughout the surveillance period declined by 64.0% and 54.0% respectively. Across all age  
66 groups, the number of meningitis cases caused by *S. pneumoniae* serotypes covered by the 7-  
67 valent vaccine dropped by 73.3% (32). Similar reductions in invasive pneumococcal disease  
68 caused by 7-valent serotypes have also been reported in Australia, England and Wales (33-  
69 35); however, this has led to a replacement phenomenon in which the rates of disease caused  
70 by non-7-valent *S. pneumoniae* serotypes have significantly increased (32, 34, 35).

#### 71 **Meningococcal meningitis**

72 *N. meningitidis* is most likely to cause meningitis in children and adolescents, and has a

73 mortality rate of 10.1% in the United States (14). The serotypes most commonly implicated  
74 are A, B, C, W135 and Y (14, 36-38) and meningitis without shock is the most common  
75 presentation (38). The predominant clinical feature, which can distinguish *N. meningitidis*  
76 from other causes of bacterial meningitis, is the presence of a petechial rash. It can rapidly  
77 become purpuric, which in the presence of meningitis or sepsis, would strongly implicate *N.*  
78 *meningitidis* as the etiologic agent. These lesions are a consequence of meningococci  
79 adhering to the endothelial cells of the capillaries and small veins in the skin, thereby altering  
80 the antithrombotic surface of the endothelium. This results in the formation of clots and the  
81 extravasion of erythrocytes, which appears as skin hemorrhages. Lesions > 1.0 cm usually  
82 occur in patients developing shock, with high levels of circulating endotoxin (38).

83 Immunization against *N. meningitides* serotypes A and C has resulted in a significant  
84 decrease in meningococcal disease. The United Kingdom was the first country to introduce  
85 routine immunization with a monovalent serotype C conjugate vaccine in 1999. During the  
86 first follow-up surveillance period (2000 - 2001), the incidence of serotype C meningococcal  
87 disease fell by 80% in target age groups (39). Meningococcal serotype C vaccination  
88 programs in Australia have also led to a marked decrease in disease due to *N. meningitidis*  
89 serotype C. Of the 90% of meningococcal cases in Australia in 2010 that had serotype data  
90 available, *N. meningitidis* serotype B caused 84.0% of cases (40). In 2013, a recombinant  
91 meningococcus serotype B vaccine (Bexsero®) was added to the Australian Register of  
92 Therapeutic Goods for use in individuals > 2 months of age, and was also approved for use in  
93 Europe. In New Zealand, the introduction of an outer membrane vesicle vaccine against an  
94 epidemic *N. meningitidis* serotype B clone in New Zealand was reported to be 80.0%  
95 effective in fully immunized children aged 6 months to < 5 years (41).

96 ***H. influenzae*, *L. monocytogenes* and other bacterial causes of meningitis**

97 *H. influenzae* as a cause of meningitis is now uncommon in most western countries due to  
98 widespread immunization against *H. influenzae* capsular type B, which was previously  
99 responsible for the majority of cases of *H. influenzae* meningitis. As a cause of meningitis, *H.*  
100 *influenzae* has fallen from 45.0% in 1986 (12) to 6.7% in 2003 – 2007 (14). *L.*  
101 *monocytogenes* is also an infrequent cause of meningitis; however, it is associated with a high  
102 mortality rate (17.0% - 28.0%) (14, 42, 43). *L. monocytogenes* meningitis occurs primarily in  
103 children < 2 months of age and in immunocompromised and elderly individuals (14, 42, 43).  
104 The signs and symptoms of *L. monocytogenes* meningitis are not different from those  
105 associated with other causes of community-acquired meningitis, but CSF findings may be  
106 atypical (43, 44). Listeriosis is associated with the consumption of contaminated food,  
107 implicating the gastrointestinal tract as the portal of entry into the blood stream.

108 Group B streptococcus and *E. coli* are common causes of meningitis in neonates (45, 46),  
109 with group B streptococcus accounting for 86.1% of bacterial meningitis cases in patients < 2  
110 months of age (14). Gram negative aerobic bacteria such as *Klebsiella pneumoniae*, *Serratia*  
111 *marcescens*, *Salmonella* spp. and *Pseudomonas aeruginosa* can occasionally be responsible  
112 for meningitis in certain patient groups, including infants and patients with head injuries, or  
113 those undergoing post-neurosurgical procedures or immunosuppression (31, 47, 48). *M.*  
114 *tuberculosis* is a common cause of chronic or subacute meningitis (49) and is associated with  
115 non-specific symptoms and an absence of neck stiffness during early disease (50).  
116 Tuberculosis meningitis occurs following hematogenous spread of bacilli from the lungs and  
117 the establishment of a caseating Rich focus (tuberculosis granuloma) in the brain cortex,  
118 meninges or choroid plexus. Bacteria are subsequently released from the foci and enter the  
119 subarachnoid space (51).

## 120 **Encephalitis**

121 Acute meningitis can, in certain circumstances, be clinically indistinguishable from acute  
122 encephalitis, which refers to inflammation of the brain parenchyma in association with  
123 neurologic dysfunction (52). Encephalitis is typically characterized by headache with an  
124 acute confusional state, with or without seizures. The California Encephalitis Program  
125 identified the clinical profiles and etiologies of encephalitis in a large cohort of  
126 immunocompetent patients from 1998 – 2005 (7). A confirmed or probable etiology was  
127 identified only in 15.8% of cases; of these, viruses were the causative agent in 68.5% of  
128 cases. Herpes simplex virus is the most common cause of encephalitis in western countries  
129 (6, 53, 54). Enteroviruses, varicella zoster virus, Epstein-Barr virus, measles virus and  
130 arboviruses such as Japanese encephalitis virus, West Nile virus and Murray Valley  
131 encephalitis virus have also been implicated as causes of encephalitis (6, 7, 54, 55). Beckham  
132 and Tyler provide a comprehensive list of other important and emerging viral causes of  
133 encephalitis (52). *L. monocytogenes* may cause meningoencephalitis, a syndrome where  
134 features of both meningitis and encephalitis are present.

135 *Burkholderia pseudomallei*, the causal agent of melioidosis, has a relatively uncommon  
136 association with CNS involvement, accounting for around 4% of cases in northern Australia  
137 (56, 57), with distinctive clinical features being brain stem encephalitis, often with cranial  
138 nerve palsies, together with peripheral motor weakness, or occasionally just flaccid  
139 paraparesis alone (myelitis) and presenting with urinary retention (56-58). Fevers are usually  
140 a prominent feature, and headaches are common but neck stiffness is usually mild or absent.  
141 The CT scan is often normal, but dramatic changes are seen on magnetic resonance imaging,  
142 most notably a diffusely increased T<sub>2</sub>-weighted signal in the midbrain, brain stem, and spinal  
143 cord, with variable symmetry and enhancement with gadolinium. CSF shows high protein,  
144 glucose normal or slightly decreased and most notably a pleocytosis with usually  
145 lymphocytes predominant. Bulbar palsy and decreased respiratory drive often necessitate

146 ventilation, which may be required for many weeks (56, 59-62). In northern Australia patients  
147 infected with the flavivirus Murray Valley encephalitis virus can present with clinical  
148 features indistinguishable from those seen with melioidosis encephalomyelitis.

149 Neurologic melioidosis is occasionally seen outside Australia, although it appears to be less  
150 common in melioidosis patients in Southeast Asia than in Australia. Rather than the  
151 distinctive brainstem encephalitis and/or myelitis seen in Australia, patients with neurological  
152 melioidosis in Southeast Asia usually present with macroscopic cortical brain abscesses (63),  
153 suggesting that in these cases there has been bacteremic spread to the brain. Frank meningitis  
154 is rarely seen in melioidosis. A recent report described the first case of neurological  
155 melioidosis in the United States, in a 58 year old male who presented with low-grade fever,  
156 nausea, vomiting, headaches, diplopia, left-sided ptosis and transient episodes of marked  
157 somnolence, after recent travel to Cambodia (64). CSF demonstrated increased levels of  
158 glucose, protein and lymphocytes; however, microbiological investigation consistently  
159 showed no organisms by Gram stain and failed to detect any organisms on culture medium.  
160 *B. pseudomallei* was only diagnosed following the culture of biopsy material collected during  
161 a suboccipital craniectomy. The mortality of neurological melioidosis is up to 30%, with  
162 ongoing residual neurological deficit in up to 50% overall (56-58).

### 163 **Focal infections**

164 The range of focal CNS infections includes brain abscesses, subdural empyema and epidural  
165 abscesses. Brain abscess refers to an intracerebral infection that usually begins as localized  
166 cerebritis, which develops into a collection of pus that is surrounded by a well-vascularized  
167 capsule (65). The clinical presentations of brain abscesses are variable and depend on the  
168 anatomical area of the brain that is compressed or affected. Brain abscesses commonly  
169 originate from a contiguous focus of infection, such as the middle ear, mastoid cells or

170 paranasal sinuses. Brain abscesses may also occur following: (i) hematogenous spread of  
171 bacteria from a distant focus of infection, or (ii) brain trauma or neurosurgery (66). A recent  
172 systematic review and meta-analysis of 123 brain abscess studies reported between 1935 and  
173 2012 demonstrated that *Streptococcus* and *Staphylococcus* spp. were the most frequently  
174 isolated microorganisms, causing 34.9% and 18.9% of infections respectively (67). Gram-  
175 negative enteric bacteria were responsible for 15.0% of cases of brain abscess and were  
176 frequently associated with polymicrobial infections. Other less common causative agents  
177 include anaerobic bacteria (*Bacteroides* spp., *Peptostreptococcus* spp., *Fusobacterium* spp.),  
178 Actinomycetales, *Haemophilus* spp. and *Pseudomonas* spp. (66-68).

179 Subdural empyema refers to a collection of pus in the subdural space, which is located  
180 between the dura mater and arachnoid mater layers of the meninges (see ‘barriers of the  
181 CNS’ below). In contrast, epidural abscesses are defined as a collection of pus between the  
182 dura mater and the skull or vertebral column (69). The etiologies of subdural empyema and  
183 epidural abscesses are similar to those of brain abscesses; however, due to the anatomical  
184 location of these infections bacteria do not need to penetrate the BBB to establish disease.

## 185 **BARRIERS OF THE CNS**

186 The CNS is physically protected from injury by bony structures such as vertebrae and the  
187 cranium, and is also enveloped by the meninges and surrounded by CSF. The meninges  
188 comprise three layers: the dura mater, arachnoid mater and pia mater. The arachnoid mater  
189 and pia mater are separated by the subarachnoid space, which is filled with CSF that is  
190 produced by the choroid plexus in the ventricles of the brain (Figure 1) (70). The CSF  
191 provides mechanical support, regulates ion composition and maintains chemical stability.  
192 Once it has circulated through the ventricles of the brain, the CSF enters the subarachnoid

193 space and drains into the blood through arachnoid villi and granulations. CSF also drains  
194 through the cribriform plate into the lymphatics within the nasal mucosa (71-73).

195 The complex functions of the brain are dependent on homeostatic mechanisms that regulate  
196 the ionic composition of the interstitial fluid that surrounds neurons in the parenchyma (74).  
197 This homeostasis is maintained by two cellular barriers, which separate the CNS from the  
198 systemic circulation: the BBB and the BCSFB (Figure 1). These barriers protect the brain  
199 from abrupt changes in blood biochemistry while at the same time allowing delivery of  
200 nutrients and removal of metabolites. Importantly, these barriers also protect the CNS from  
201 invasion by microbes that may be present within the blood.

## 202 **The blood-brain barrier**

203 The BBB (Figure 1) exists at the level of cerebral microvessels and represents the largest  
204 interface for blood-brain exchange (75). The BBB is composed of endothelial cells that line  
205 cerebral microvessels in association with pericytes, astrocytes and the basement membrane  
206 (collectively referred to as a neurovascular unit in association with neurons, microglia and  
207 peripheral immune cells). Pericytes are attached to the abluminal surface of the endothelium  
208 and together with the astrocytic endfeet form a contiguous membrane barrier. A physical  
209 barrier is formed at the interendothelial cleft by complex junctions that regulate the flux of  
210 ions, polar molecules and macromolecules from the systemic circulation (74, 75).

211 At the interendothelial cleft, two types of junctions comprise the BBB: adherens junctions  
212 and tight junctions. These junctions block the transport of a wide range of molecules,  
213 restricting paracellular permeability across the BBB (75). In addition, endothelial cells  
214 forming the BBB lack fenestrations in their plasma membranes and have a reduced number of  
215 pinocytotic vesicles compared to endothelial cells in other tissues, which restricts  
216 transcellular flux (76). Despite this, certain solutes and macromolecules may be transported

217 across the BBB *via* specific and non-specific mechanisms, such as passive diffusion, receptor  
218 and/or adsorptive-mediated transcytosis, or *via* ATP-binding cassette transporters and solute  
219 carrier transporters, which are expressed on cerebral microvessels (75, 77-79). The  
220 transcellular and paracellular routes of entry into the CNS are highly relevant for microbial  
221 pathogens.

## 222 **The blood-cerebrospinal fluid barrier**

223 The BCSFB (Figure 1) is formed by tight junctions between: (i) epithelial cells located at the  
224 choroid plexus in the lateral, third and fourth ventricles of the brain and (ii) endothelial cells  
225 of the veins and venules within the subarachnoid space. Tight junctions between the epithelial  
226 cells of the arachnoid mater also contribute to the BCSFB, by forming a barrier between the  
227 subarachnoid space and the leaky fenestrated blood vessels of the dura mater. Of these  
228 BCSFB locations, the choroid plexus has a large vascular surface area and represents the  
229 major interface between fenestrated blood vessels and CSF. Similar to the BBB, the BCSFB  
230 functions to protect the CNS and helps to maintain homeostasis; however, the choroidal  
231 epithelial cells of the BCSFB also secrete CSF from the choroid plexus into the brain  
232 ventricular system, and the tight junctions between the epithelial cells have a lower electrical  
233 resistance compared to BBB endothelial cells (80). Thus, the choroid plexus BCSFB may be  
234 more vulnerable to microbial penetration *via* paracellular mechanisms. The BCSFB at the  
235 arachnoid mater has a smaller total surface area compared to other barriers of the CNS, and is  
236 not thought to be an important route for the entry of solutes into the brain (74). However, the  
237 arachnoid BCSFB may represent an important site of microbial entry.

## 238 **Immunosurveillance of the CNS**

239 The brain parenchyma and spinal cord are populated throughout by resident immune cells,  
240 the microglia, which are highly specialized tissue macrophages that are maintained through *in*



241 *situ* self-renewal without reconstitution from the bone marrow (81-83). Microglia  
242 continuously survey the brain parenchyma by sampling their microenvironment with highly  
243 motile processes and protrusions (84). Recent data has also revealed cells expressing CD11c  
244 are localized within the juxtavascular parenchyma of the healthy mouse brain, and extend  
245 processes to the glia limitans (astrocytic foot processes at the parenchymal basement  
246 membrane, surrounding the brain under the pia mater) (85). In agreement with a previous  
247 study (86), this raises the possibility that dendritic cells, which are characterized by high-  
248 level CD11c expression, may reside within the brain parenchyma. However, a range of cells  
249 of the myeloid lineage (including activated microglia) also express CD11c, albeit at lower  
250 levels. Thus the CD11c-expressing cells described within the brain parenchyma may also  
251 represent a subpopulation of microglia (85).

252 Other resident immune cells of the CNS include perivascular macrophages in addition to  
253 small populations of blood-derived dendritic cells and macrophages within the choroid plexus  
254 and meninges. Dendritic cells constitute approximately 1% of immune cells within the  
255 choroid plexus and meninges (87, 88), but unlike the putative dendritic cells of the brain  
256 parenchyma, choroid plexus and meningeal dendritic cells are distinct from microglia and  
257 closely resemble classical dendritic cells that require FMS-like receptor tyrosine kinase 3  
258 ligand for development (88). The CSF also contains a trafficking population of mononuclear  
259 cells, comprising T cells (90%), B cells (5%), monocytes (5%) and dendritic cells (less than  
260 1%) (89).

261 Normal CNS immune surveillance occurs when central memory T cells migrate from the  
262 blood across the choroid plexus epithelium, enter the CSF and scan the subarachnoid space.  
263 In the absence of recognizing ligands presented by local antigen presenting cells, T cells exit  
264 the subarachnoid space and drain to the secondary lymphoid organs *via* the nasal mucosa (90,  
265 91). In contrast, recognition of presented antigens results in the formation of immunological

266 synapses and is accompanied by cellular signalling that results in the recruitment of  
267 peripheral inflammatory cells to the CNS *via* paracellular or transcellular diapedesis (92).

268 The production of cytokines, free radicals (such as reactive oxygen species) and matrix  
269 metalloproteinases by inflammatory cells within the CSF spaces can subsequently result in  
270 degradation of the glia limitans, BBB/BCSFB dysfunction and cellular death (93-95). In  
271 bacterial meningitis, neuronal injury also occurs due to the direct effects of bacterial toxins  
272 and virulence factors (96-98). A recent study of human bacterial meningitis cases  
273 demonstrated that the profile of pro- and anti-inflammatory cytokines and matrix  
274 metalloproteinases within the CSF is dependent upon the causative pathogen (99). The  
275 sequelae of bacterial meningitis and the associated neuroinflammation include vasculitis and  
276 cerebral venous thrombosis, which can lead to cerebral ischemia. An increase in CSF outflow  
277 resistance in combination with interstitial edema results in raised intracranial pressure, which  
278 causes hypoxic brain damage and may lead to death (100).

## 279 **MICROBIAL ENTRY TO THE CNS**

280 Microbial pathogens may access the CNS (i) by penetrating the BBB or the BCSFB; or (ii)  
281 *via* the olfactory and/or trigeminal nerves (101). Peripheral nerves other than the olfactory  
282 and trigeminal cranial nerves have also been implicated as routes of entry for some viruses,  
283 but these will not be reviewed here. Penetration of the BBB and BCSFB represent common  
284 routes of CNS invasion, resulting in encephalitis or brain abscesses, and meningitis,  
285 respectively. In contrast to a diverse group of neurotropic viruses for which mechanisms are  
286 described for invasion of the brain *via* the olfactory sensory nerves, bacterial infection *via* the  
287 olfactory epithelium and the olfactory nerve bundles has been infrequently described. Even  
288 less well understood is that, in addition to the olfactory nerve, the trigeminal nerve may also  
289 present a potential pathway to the brain.

290 ***In vivo* and *in vitro* models of CNS invasion**

291 Much of our understanding of the mechanisms of microbial CNS invasion has been obtained  
292 from *in vivo* and *in vitro* studies. Mice and rats are the most common animals used to model  
293 CNS infections. These animal models have been used to investigate the sites of microbial  
294 entry (20, 102-106) and the roles of virulence determinants in CNS invasion and pathogenesis  
295 (107-111). Despite clear differences in the size and complexity of mouse and human brains,  
296 comparative transcriptomic studies between mice and humans have demonstrated that gene  
297 expression profiles of the healthy brain and choroid epithelium are highly conserved between  
298 the species (112-114).

299 There are, however, several important differences between the mouse and human CNS that  
300 are relevant to microbial invasion studies. In mice, the leptomeninges (comprising the  
301 arachnoid mater and pia mater) are significantly thinner and less vascular than in humans  
302 (115). Additionally, the human choroid plexus epithelium has more abundant fibrovascular  
303 stroma along the vascular channels and has a relatively larger surface area compared to  
304 mouse choroid epithelium. Mice also possess very large olfactory bulbs compared to humans  
305 (115), which may result in a higher incidence of CNS invasion *via* the olfactory route  
306 compared to humans. Furthermore, mice lack several anatomically defined subclasses of  
307 astrocytes that are present in humans (116). These combined anatomical differences may  
308 influence the initial site of microbial attachment and subsequent CNS invasion.

309 *In vitro* cell culture systems have been used to model the BBB and elucidate the cellular and  
310 molecular mechanisms of microbial CNS invasion. Primary human, rodent and bovine brain  
311 microvascular endothelial cells have been used in some studies (97, 117-119); however, the  
312 isolation of these cells is technically demanding and *in vitro* culture dramatically alters the  
313 transcriptome leading to downregulation of genes involved in BBB function (120, 121).

314 Human umbilical vein endothelial cells (HUVECs) (122-124), caco-2 cells and Mading-  
315 Darby canine kidney cells (125) have been used as surrogate models of the BBB, although  
316 the non-cerebral origin of these cells may limit their relevance. The availability of  
317 immortalized brain endothelial cell lines (for example: human brain microvascular  
318 endothelial cells [HBMECs], hCMEC/D3, TY10 and BB19) has enabled the human BBB to  
319 be studied. When grown on Transwell membranes, HBMECs: (i) form tight junctions, as  
320 measured by transendothelial electrical resistance; (ii) express the adherens junction and tight  
321 junction proteins zonula occludens-1, junctional adhesion molecule A, claudin-5, occludin  
322 and vascular endothelial-cadherin; and (iii) demonstrate negligible paracellular permeability  
323 (126-129).

324 It must be noted that brain microvascular endothelial cells grown in static monocultures  
325 represent a surrogate model for the human BBB, as they lack other components of the  
326 neurovascular unit that are required for the development of true BBB properties. Studies  
327 using pericyte deficient mice have shown that pericytes (see Figure 1) are required for BBB  
328 integrity. Compared to wild type mice, pericyte deficient mice demonstrated an accumulation  
329 of water and tracer dyes within the brain, structural abnormalities in cerebral endothelial tight  
330 junctions and increased expression of genes known to increase vascular permeability,  
331 endothelial vesicle trafficking and immune cell recruitment (130-132). Astrocytes also play a  
332 key role in regulating the BBB (133-136), and as such, a range of co-culture systems have  
333 been developed to more closely model the BBB properties of the neurovascular unit (137-  
334 139). Choroid plexus epithelial cell lines that display the characteristic properties of the  
335 BCSFB have been described (140, 141); however, the majority of *in vitro* microbial invasion  
336 studies have been performed using monocultures of brain microvascular endothelial cells.

## 337 **BACTERIAL INVASION THROUGH THE BLOOD-BRAIN AND BLOOD-** 338 **CEREBROSPINAL FLUID BARRIERS**

339 **Sites of entry**

340 Classical bacterial meningeal pathogens may potentially enter the CSF by penetrating the  
341 BBB of cerebral microvessels and entering the extracellular fluid of the brain, which is  
342 continuous with the CSF. However, a more likely route involves direct entry into the CSF *via*  
343 penetration of the BCSFB at either the arachnoid, or the choroid plexus (142). The highly  
344 vascularized choroid plexus has been favored as a major site of bacterial entry to the CSF due  
345 to lower electrical resistances between choroid epithelial cells. Autopsies of neonates who  
346 died from bacterial meningitis demonstrated that meningitis was frequently (84.0%)  
347 associated with an accumulation of a purulent exudate within the ventricles (ventriculitis) and  
348 the choroid plexus (plexitis) (143). Ventriculitis preceding meningitis would support the role  
349 of the choroid plexus as the initial site of bacterial invasion; however, ventriculitis frequently  
350 occurs as a complication of meningitis, and the site of bacterial CNS invasion remains  
351 incompletely understood.

352 In experimentally-infected rhesus macaques, the choroid plexus was the site at which the  
353 earliest histopathologic lesions were observed in *H. influenzae* meningitis (144). Following  
354 intra-nasal inoculation, *H. influenzae* was isolated from the CSF within the lateral cerebral  
355 ventricle and this was associated with intravascular inflammatory infiltration and mild  
356 choroid plexitis (145). In 5 – 7 week old pigs, *S. suis* was localized within the choroid plexus  
357 and caused disruption of the brush border of the choroid epithelium and loss of the apical  
358 cytoplasm (146). Interestingly, autopsy of a 2 month old infant who died of fulminant  
359 meningococemia prior to the onset of meningitis revealed adherent bacteria within the  
360 capillaries of the choroid plexus and, to a much lesser extent, the capillaries within the  
361 meninges (147). This would appear to support the choroid plexus as the preferential site of *N.*  
362 *meningitidis* entry to the CSF; however, no bacteria were found within or between choroid  
363 epithelial cells. In contrast, a study of autopsy material from symptomatic patients with

364 meningococcal disease revealed the presence of *N. meningitidis* within the choroid plexus  
365 epithelium, interstitium and blood vessels (148).

366 Neonatal rat models of *E. coli* meningitis have been used to identify the potential site of entry  
367 to the CSF. Parkkinen *et al.* demonstrated that *E. coli* binds to the luminal surface of choroid  
368 epithelial cells, vascular endothelial cells and ependymal cells lining the brain ventricles  
369 (102). Similar findings were reported by Zelmer *et al.*, who suggested that *E. coli* K1 enters  
370 the CSF *via* the choroid plexus, circulates to the subarachnoid space and then adheres to the  
371 meninges (103). In contrast, Kim *et al.* reported that in the brains of 5 day old rats, *E. coli*  
372 was localized within the perivascular areas of the subarachnoid space, and not within the  
373 choroid plexus (20). Recently, the spatiotemporal events of CNS invasion were investigated  
374 in a BALB/c mouse model of pneumococcal meningitis (104). *S. pneumoniae* preferentially  
375 adhered to vessels within the subarachnoid space at 1 hour post-infection and formed clusters  
376 around the endothelium at this location at later time points. Remarkably, pneumococci could  
377 only be detected in the choroid plexus at 8 hours post-infection, suggesting that this is not the  
378 site of initial entry to the CSF.

### 379 **Mechanisms of entry**

380 Bacteria may cross the BBB and BCSFB to access the CNS by transcellular penetration,  
381 paracellular entry or *via* infected leukocytes from the peripheral circulation (the “Trojan  
382 horse” mechanism; Figure 2). Transcellular penetration occurs following bacterial adhesion  
383 to endothelial or epithelial cells. Bacteria are subsequently translocated across these barriers  
384 by pinocytosis or receptor-mediated mechanisms. Alternatively, bacteria may invade the CNS  
385 paracellularly following disruption of the tight junctions between cells forming the BBB  
386 and/or BCSFB, resulting in increased permeability. Microorganisms may exploit more than  
387 one of these mechanisms to access the CNS.

388 Bacterial attachment to HBMECs is a prerequisite for transcellular and paracellular  
389 penetration. There has been significant research into the host-pathogen interactions that occur  
390 prior to bacterial invasion of brain microvascular endothelial cells (Table 1), including the  
391 identification of host receptors that mediate these interactions. *E. coli*, *H. influenzae*, *N.*  
392 *meningitidis* and *S. pneumoniae* bind to HBMECs via the 37/67 kDa laminin receptor (149-  
393 152). The 37 kDa laminin receptor is a precursor for the mature 67 kDa laminin receptor,  
394 which binds laminin-1 within the basement membrane on the abluminal surface of eukaryotic  
395 cells (153). Kim *et al.* (150) demonstrated that the 67 kDa laminin receptor is expressed on  
396 the basolateral side of HBMECs (i.e. the abluminal surface); however, the cytotoxic necrotizing  
397 factor-1 (CNF-1) of *E. coli* triggered clustering of the 67 kDa laminin receptor on the apical  
398 side (i.e. the luminal surface), where it could potentially interact with bacteria within the  
399 cerebral microvasculature. In an *in vivo* model, Orihuela *et al.* reported that fluorescent  
400 microspheres coated with laminin receptor-binding adhesins (such as the choline binding  
401 protein A (CbpA) of *S. pneumoniae*) adhered to the cerebral endothelium of mice via the  
402 37/67 kDa laminin receptor following intravascular injection, as demonstrated by cranial  
403 window imaging (151). Several bacterial meningeal pathogens have been shown to bind host  
404 glycoproteins such as fibrinogen, vitronectin, fibronectin, laminin and collagen, which may  
405 act as bridging molecules between the bacterium and HBMECs (109, 154-159). In addition,  
406 the heat shock protein gp96 and a gp96 homologue act as receptors for the surface-expressed  
407 virulence protein (Vip) of *L. monocytogenes* (160) and the outer membrane protein A  
408 (OmpA) of *E. coli* (161, 162) respectively.

#### 409 **Transcellular penetration of brain microvascular endothelial cells**

410 Transcellular penetration of brain microvascular endothelial cells, mainly via receptor-  
411 mediated mechanisms (Figure 2), has been demonstrated for several bacteria, including *E.*

412 *coli*, *N. meningitidis*, *S. pneumoniae*, group B streptococcus, *S. suis*, *L. monocytogenes* and  
413 *M. tuberculosis*.

414 ***E. coli***

415 *E. coli* translocates from the upper compartment to the lower compartment in a HBMEC  
416 Transwell model without increasing cellular permeability, thus supporting a transcellular  
417 mechanism of penetration (163). Electron microscopy studies have demonstrated that *E. coli*  
418 invades HBMECs through a zipper-like, receptor-mediated endocytosis mechanism (164).  
419 Intracellular *E. coli* E44 was observed in membrane-bound vesicles within HBMECs 30  
420 minutes post-infection; by 45 minutes the bacteria had translocated to the basolateral side  
421 (164). Both early (early endosome antigen 1 and transferrin receptor) and late (ras-related  
422 protein-7 and lysosomal-associated membrane protein-1) endosomal markers were recruited  
423 to vacuoles containing *E. coli* K1. However, the lysosomal hydrolytic enzyme cathepsin D  
424 did not accumulate within the vacuoles, demonstrating that lysosomal fusion was prevented.  
425 This was found to occur in a capsule-dependent manner (165).

426 Several *E. coli* proteins contribute to the adhesion to, and invasion of, HBMECs (Table 1).  
427 Attachment of *E. coli* to HBMECs induces host cytoskeletal rearrangements and actin  
428 condensation below adherent bacteria (164). This is associated with a CNF-1-mediated  
429 recruitment of focal adhesion kinase and the cytoskeletal protein paxillin to the 67 kDa  
430 laminin receptor, which forms clusters that co-localize with adherent *E. coli* (150).  
431 Interactions between *E. coli* and HBMECs result in tyrosine phosphorylation of focal  
432 adhesion kinase and paxillin. The activity of focal adhesion kinase and its  
433 autophosphorylation site tyrosine 397 was shown to be essential for *E. coli* invasion of  
434 HBMECs (166). Furthermore, the Src kinase-dependent activation of phosphatidylinositol 3-  
435 kinase and its interaction with focal adhesion kinase were required for *E. coli* K1 invasion



436 and host cytoskeletal rearrangement (167, 168). Sukumaran *et al.* demonstrated that the  
437 association of phosphatidylinositol 3-kinase with focal adhesion kinase resulted in the  
438 downstream activation of protein kinase C $\alpha$  in an Omp-A-dependent manner (169). Activated  
439 protein kinase C $\alpha$  was recruited to the *E. coli* entry site, where it interacted with its substrate  
440 (myristoylated alanine-rich C kinase substrate), leading to an accumulation of actin (169).  
441 The activated protein kinase C $\alpha$  also interacted with caveolin-1 co-localized with condensed  
442 actin underneath the bacterial entry site to form caveolae, plasma membrane invaginations  
443 that are involved in endocytosis and signal transduction (170). CNF-1 (*via* the 37/67 kDa  
444 laminin receptor) and type 1 fimbrial adhesin (FimH, *via* CD48) have also been shown to  
445 induce cytoskeletal rearrangements through the activation of the GTPase RhoA (149, 171,  
446 172).

447 In an independent mechanism, *E. coli* outer membrane protein A (OmpA) and new  
448 lipoprotein-1 (Nlp1) promote the activation of host cytosolic phospholipase A<sub>2</sub> (173, 174).  
449 The activation of cytosolic phospholipase A<sub>2</sub> generated arachidonic acid metabolites and  
450 induced host actin cytoskeletal rearrangements, and was essential for *E. coli* K1 invasion of  
451 HBMECs (173, 175-177). OmpA and invasion of brain endothelial cell protein A (IbeA) also  
452 trigger the phosphorylation of STAT3, which results in the activation of Rac1, a Rho family  
453 GTPase that regulates host cytoskeletal rearrangements (178). Notably, OmpA is highly  
454 conserved between Gram negative bacteria and may play a role in CNS invasion by those in  
455 which the mechanisms of entry are unknown. Similarly, a *B. pseudomallei* toxin,  
456 *Burkholderia* lethal factor-1 (BLF-1), is structurally similar to *E. coli* CNF-1 and possesses a  
457 number of conserved residues that, in CNF-1, promote deamidation of glutamine and result in  
458 the activation of RhoA and subsequent host cytoskeletal rearrangements (179). Thus, OmpA  
459 and BLF-1 may be attractive targets for future studies into the mechanisms of *B.*  
460 *pseudomallei* CNS invasion.

461 *N. meningitidis*

462 The attachment of *N. meningitidis* to HBMECs is mediated by type IV pili. This initial  
463 attachment is inhibited by high cerebral microcirculation shear stress levels; however, once  
464 attached, *N. meningitidis* proliferates and is able to resist high blood velocities by forming  
465 microcolonies with strong bacteria-bacteria interactions (124) and bacteria-host interactions  
466 (122). CD46 was previously identified as the receptor for type IV pili (180); however, this  
467 was not corroborated by additional studies and it is likely that the type IV pilus attaches to  
468 host cells independently of CD46 (181). The type IV pilus protein PilQ, in addition to the  
469 outer membrane porin PorA, have more recently been demonstrated to bind to the 37/67 kDa  
470 laminin receptor on HBMECs (151). Furthermore, studies with human bronchial epithelial  
471 cells show that the meningococcal type IV pilus binds to the platelet activating factor (PAF)  
472 receptor, and synergy between a pilin-linked glycan and phosphorylcholine decorating  
473 moieties is required for pili to efficiently engage the receptor (182).

474 *N. meningitidis* is thought to primarily invade HBMECs by paracellular mechanisms (see  
475 below); however, *in vitro* studies have demonstrated that small numbers of both encapsulated  
476 and non-encapsulated meningococci are internalized by HUVECs and HBMECs (183-185).  
477 Following initial bacterial attachment, the meningococcal minor pilin protein, PilV, triggers a  
478 cellular response in which endothelial cells form protrusions around *N. meningitidis* (122).  
479 These cellular projections further protect the bacterial microcolony from the mechanical  
480 stresses associated with high velocity blood flow (122), and also lead to the engulfment and  
481 internalization of *N. meningitidis* (186). The host signalling events that contribute to these  
482 interactions have been elucidated. Following type IV pilus-mediated attachment, *N.*  
483 *meningitidis* activates the host cell  $\beta$ 2-adrenoceptor, leading to the translocation of  $\beta$ -arrestin  
484 to the site of bacterial attachment at the plasma membrane (187). Several host proteins,  
485 including ezrin and moesin, which regulate the cortical cytoskeleton through F-actin binding

486 sites, are then recruited to the site of bacterial attachment and accumulate in honeycombed  
487 shaped molecular complexes, referred to as cortical plaques, underneath the meningococcal  
488 microcolonies (186, 188, 189). The translocation of  $\beta$ -arrestin also results in the docking and  
489 activation of the protein tyrosine kinase Src, which phosphorylates cortactin (187, 188).  
490 Cortical actin is then polymerized in a Rho GTPase- and Cdc42-dependent manner, which  
491 leads to the formation of cell membrane protrusions (186). In human bone marrow  
492 endothelial cells, the recruitment of cortactin and formation of membrane protrusions was  
493 also shown to be dependent on the activation of a phosphoinositide-3-kinase/Rac1 signalling  
494 pathway by lipooligosaccharide (190).

495 Once internalized, the membrane-bound vesicles containing meningococci associate with  
496 transferrin receptor and lysosomal-associated membrane protein-1 endosomal markers (185).  
497 It is likely that lysosomal fusion is subverted, as live meningococci have been shown to  
498 translocate from the basolateral side to the apical side in a Transwell BCSFB invasion model  
499 (140). *In vitro* studies have demonstrated that capsular polysaccharide inhibits the invasion of  
500 *N. meningitidis* into HBMECs (191) and human choroid plexus epithelial cells (140), but is  
501 essential for survival and replication within the intracellular niche (185).

502 In non-encapsulated meningococcal strains, the invasion of HBMECs is dependent upon  
503 outer membrane protein C (Opc). Opc binds to the human serum factor fibronectin, which  
504 acts as a bridging molecule and anchors the bacterium to the  $\alpha_5\beta_1$  integrin receptor of  
505 HBMECs (191). This interaction results in the activation of mitogen-activated protein kinase  
506 (c-Jun N-terminal kinase (JNK)1, JNK2 and p38) and protein tyrosine kinase signal  
507 transduction pathways (192). Pre-treatment of cells with specific JNK1, JNK2 and protein  
508 tyrosine kinase inhibitors significantly reduced the internalization of *N. meningitidis* by  
509 HBMECs, without affecting bacterial adherence. Blocking of the  $\alpha_5\beta_1$  integrin receptor of  
510 HBMECs also decreased JNK activation in the presence of *N. meningitidis*. Furthermore, the

511 use of Opc-deficient mutant strains demonstrated that JNK signalling, but not p38 signalling,  
512 was mediated by Opc expression (192). A more recent study identified that Opc preferentially  
513 binds to activated vitronectin within human serum and this interaction promotes *N.*  
514 *meningitidis* invasion of HBMECs (155). *In vitro*, the enhanced invasion of non-encapsulated  
515 *N. meningitidis* strains is thought to be due to the unmasking of Opc; however, the role of  
516 Opc *in vivo* is yet to be determined.

### 517 *S. pneumoniae*

518 *S. pneumoniae* is internalized by human and rat brain microvascular endothelial cells *via*  
519 receptor-mediated endocytosis. This uptake does not involve the formation of cellular  
520 membrane protrusions, as discussed for *N. meningitidis* (193). The adhesion of pneumococci  
521 to HBMECs is mediated by interactions between CbpA and the 37/67 kDa laminin receptor  
522 (151), and neuraminidase A (NanA) and an unknown cellular receptor(s) (194). Uchiyama *et*  
523 *al.* demonstrated that NanA, a surface-anchored sialidase, was necessary and sufficient to  
524 promote pneumococcal adhesion to, and invasion of, HBMECs (195). Following initial  
525 attachment, the laminin G-like domain of NanA initiates chemokine signalling and  
526 inflammatory activation of endothelial cells (194). The activation of endothelial cells results  
527 in the increased expression of PAF receptor - the receptor for pneumococcal  
528 phosphorylcholine, which is a component of the cell wall and acts as a molecular mimic of  
529 the chemokine PAF (196).

530 In HBMECs, *S. pneumoniae* within vacuoles may: (i) transit through the cell to the  
531 basolateral surface; (ii) recycle back to the apical surface; or (iii) be killed within the vacuole  
532 (193). Opaque *S. pneumoniae* variants, which express more capsular polysaccharide and less  
533 phosphorylcholine than transparent variants, were efficiently killed, suggesting that vacuoles  
534 containing these strains were targeted for lysosomal fusion. In contrast, transparent

535 pneumococci may undergo some recycling back to the apical surface; however, by 5 hours  
536 post-infection in a Transwell system, the majority of bacteria had translocated to the  
537 basolateral chamber (193). The invasion of HBMECs by transparent *S. pneumoniae* depends  
538 on phosphorylcholine, the PAF receptor and  $\beta$ -arrestin, which targets G-protein coupled  
539 receptors (such as PAF receptor) to clathrin-coated vesicles (193, 197). *S. pneumoniae*  
540 induces the translocation of  $\beta$ -arrestin from the cytosol to the plasma membrane, where it co-  
541 localizes with the PAF receptor. This interaction also stimulates mitogen-activated protein  
542 kinase signalling, which is required for pneumococcal uptake (197). Vacuoles containing *S.*  
543 *pneumoniae* co-localized with early and late endosomal markers; however increased  
544 expression of  $\beta$ -arrestin subverted these vacuoles from lysosomal trafficking and promoted  
545 the transcytosis of viable bacteria. The small number of pneumococci-containing vacuoles  
546 that underwent recycling to the apical surface demonstrated co-localisation with ras-related  
547 protein-11, which regulates endosome recycling. These findings suggest that the  
548 pneumococcal-induced interactions between PAF receptor and  $\beta$ -arrestin contribute to the  
549 transcellular penetration of HBMECs by *S. pneumoniae* (197).

#### 550 **Group B streptococcus**

551 Several group B streptococcal proteins promote the adhesion to, and invasion of, HBMECs  
552 (Table 1). The hypervirulent group B streptococcus adhesin (HvgA) was identified as a  
553 sequence type-17 (ST-17)-specific virulence factor, which is anchored to the group B  
554 streptococcal cell wall by sortase A (198). ST-17 is a clonal complex belonging to the group  
555 B streptococcus capsular serotype III, and is responsible for > 80% of neonatal meningitis  
556 cases. Therefore, it was proposed that HvgA expression may contribute to the hypervirulence  
557 of ST-17. *In vitro* studies using an isogenic *hvgA* mutant demonstrated that HvgA was  
558 required for efficient group B streptococcal adhesion to HUVECs, primary rodent choroid  
559 plexus epithelial and brain microvascular endothelial cells, and immortalized HBMECs.

560 Furthermore, in a murine model of hematogenous meningitis, HvgA was required for CNS  
561 invasion, although had no effect on the levels of bacteremia (198).

562 A common characteristic of several other group B streptococcal adhesins/invasins is the  
563 ability to bind components of the HBMEC extracellular matrix, such as laminin, collagen and  
564 fibronectin (109, 157, 159, 199). The group B streptococcal pilus tip adhesin (PilA) and the  
565 recently identified streptococcal fibronectin binding protein A (SfbA) bind immobilised  
566 collagen and fibronectin respectively. The collagen and fibronectin then associate with  
567 HBMECs *via* integrins, and these interactions facilitate the entry of group B streptococcus  
568 (109, 199). Similarly, the group B streptococcal serine-rich repeat (Srr-1) glycoprotein binds  
569 the fibrinogen A $\alpha$  chain within human blood through a dock, lock and latch mechanism (200,  
570 201). During this interaction, fibronectin docks between two IgG-like folds (N2 and N3  
571 domains) of the binding region of the adhesin, which initiates a conformational change  
572 whereby a flexible region of the N3 domain latches to form a  $\beta$ -strand and completes a  $\beta$ -  
573 sheet within the N2 domain, effectively locking the ligand in place (200). Seo *et al.*  
574 demonstrated that deletion of the latch-like domain of the C-terminal end of the Srr-1  
575 fibrinogen binding region significantly reduced group B streptococcal adhesion to HBMECs  
576 (201). The fibrinogen-binding protein (FbsA) also mediates group B streptococcal adherence  
577 to HBMECs by binding immobilised fibrinogen (156), although it is unknown if this  
578 interaction also occurs *via* a dock, latch and lock mechanism.

579 Electron microscopy studies have demonstrated that *in vitro*, group B streptococcus  
580 associates closely with the cell membrane of HBMECs, and is enveloped by microvillus  
581 structures (202). Group B streptococci are internalized within membrane-bound vesicles and  
582 translocate from the apical side to the basolateral side of HBMECs without a marked change  
583 in transendothelial electrical resistance. Serotype III group B streptococci were shown to  
584 invade HBMECs more efficiently than representative strains of serotypes Ia, Ib, II and V;

585 however, the serotype III capsule itself did not facilitate group B streptococcal invasion  
586 (202). Internalized group B streptococci did not undergo significant replication and survived  
587 within HBMECs for up to 20 hours (202). Intracellular survival of group B streptococcus  
588 may be promoted by the CiaR/CiaH two-component regulatory system, which regulates  
589 several genes associated with stress tolerance and the subversion of host defences (203). A  
590 serotype III strain that was deficient in the CiaR response regulator demonstrated similar  
591 adhesion and invasion levels to wild type group B streptococcus in HBMECs, but was  
592 associated with a significant decrease in intracellular survival. Broth inhibition assays  
593 demonstrated that CiaR conferred resistance to antimicrobial peptides, lysozyme and reactive  
594 oxygen species (203).

595 The host signal transduction pathways that are involved in the uptake of group B  
596 streptococcus by HBMECs are similar to those discussed for *E. coli*. *In vitro*, the tri-  
597 molecular interactions between PilA, collagen and  $\alpha_2\beta_1$  integrins on HBMECs stimulate the  
598 phosphorylation of focal adhesion kinase at tyrosine 397, which then induces host  
599 cytoskeletal rearrangements and the uptake of group B streptococci *via* the recruitment and  
600 activation of phosphatidylinositol 3-kinase and paxillin (109, 204). It has also been  
601 demonstrated that group B streptococcus invasion occurs *via* cytoskeletal rearrangements that  
602 are induced following the activation of RhoA and Rac1 (205). Furthermore, group B  
603 streptococcus induces serine 505 phosphorylation of host cytosolic phospholipase A<sub>2</sub>, which  
604 leads to the release of arachidonic acid metabolites including cysteinyl leukotrienes (206). *In*  
605 *vitro* pharmacological inhibition studies demonstrated that the activation of cytosolic  
606 phospholipase A<sub>2</sub> was required for efficient group B streptococcal invasion of HBMECs. In  
607 addition, brain colonization by group B streptococcus was significantly reduced in mice  
608 deficient in cytosolic phospholipase A<sub>2</sub> compared to wild type animals in a model of  
609 hematogenous meningitis. Following cytosolic phospholipase A<sub>2</sub> activation and the release of

610 cysteinyl leukotrienes, downstream signalling activates protein kinase C $\alpha$ , which is involved in  
611 the regulation of actin cytoskeletal rearrangements (206).

## 612 *S. suis*

613 The zoonotic pathogen *S. suis* adheres to, but does not invade, HBMECs *in vitro* (207). In  
614 contrast, invasion of porcine brain microvascular endothelial cells has been observed for this  
615 swine pathogen. Inhibition studies demonstrated that *S. suis* invasion of porcine brain  
616 microvascular endothelial cells required actin filaments, but not microtubular cytoskeletal  
617 elements, or active bacterial RNA or protein synthesis (208). Electron microscopy studies  
618 have shown that *S. suis* adheres to porcine brain microvascular endothelial cells 5 minutes  
619 post-infection, and at 2 hours post-infection the streptococci were in close contact with the  
620 cells, within invaginated cell membrane structures and underneath the cell surface behind the  
621 cell membrane. Unlike group B streptococci, following internalization within membrane-  
622 bound vacuoles, the number of viable intracellular *S. suis* steadily decreased over time, and  
623 viable cocci were not detected after 7 hours (208). However, similar to *S. pneumoniae* and  
624 group B streptococcus, the capsular polysaccharide of *S. suis* *partially* inhibited interactions  
625 with brain microvascular endothelial cells (208).

626 The *S. suis* proteins and virulence factors that mediate invasion of porcine brain  
627 microvascular endothelial cells remain largely unknown. Pre-treatment of porcine brain  
628 microvascular endothelial cells with *S. suis* lipoteichoic acid reduced, but did not abolish, the  
629 adhesion and invasion of *S. suis* in a dose-dependent manner, suggesting that lipoteichoic  
630 acid may play a role in *S. suis* interactions with porcine brain microvascular endothelial cells  
631 (209). Vanier *et al.* screened a transposon mutagenesis library of *S. suis* and identified several  
632 genes that may contribute to the invasion of porcine brain microvascular endothelial cells  
633 (210). These genes were characterized as belonging to the following groups: surface proteins,



634 transport/binding proteins, regulatory functions, metabolism, amino acid synthesis, protein  
635 synthesis and hypothetical proteins. However, the role of these genes in *S. suis* invasion of  
636 porcine brain microvascular endothelial cells has not been confirmed using isogenic mutants.

637 As discussed above, the choroid plexus may be the site of entry into the CNS for *S. suis*. In  
638 order to study the interactions between *S. suis* and choroid plexus epithelial cells, Tenenbaum  
639 *et al.* developed an inverted Transwell model, in which the translocation of *S. suis* from the  
640 basolateral ‘blood’ side to the apical ‘CSF’ side could be investigated (211). Following *S.*  
641 *suis* infection of the basolateral side, the transepithelial electrical resistance and paracellular  
642 permeability remained constant for up to 4 hours. *S. suis* invaded porcine choroid plexus  
643 epithelial cells from the basolateral side in the inverted Transwell system; however, invasion  
644 was rare when *S. suis* was inoculated into the apical chamber in the standard Transwell  
645 model. These findings were also replicated in a human choroid plexus epithelial cell inverted  
646 Transwell model (140). This suggested that *S. suis* may interact with components of the  
647 extracellular matrix that are accessible on the basolateral side. The translocation of *S. suis*  
648 through porcine brain microvascular endothelial cells was inhibited when cells were treated  
649 with a phosphatidylinositol 3-kinase inhibitor, demonstrating that phosphatidylinositol 3-  
650 kinase may be involved in the uptake of *S. suis* (211).

#### 651 ***L. monocytogenes* and *B. pseudomallei***

652 *L. monocytogenes* can enter the CNS by the Trojan horse mechanism (see below) or by  
653 transcellular penetration of HBMECs. Using HUVECs and HBMECs, it was demonstrated  
654 that *L. monocytogenes* can invade endothelial cells directly, or *via* cell-to-cell spread from  
655 adherent, infected mononuclear phagocytes (212, 213). In HUVECs, *L. monocytogenes*  
656 attaches to the cellular surface and induces membrane ruffling, which leads to internalization  
657 (214). In HBMECs, *L. monocytogenes* invasion is preceded by intimate interactions between

658 the bacterium and microvilli on the cell surface (215). The invasion of *L. monocytogenes* into  
659 HBMECs requires actin microfilaments (213), but unlike other meningeal pathogens, does  
660 not involve the activation of phosphatidylinositol 3-kinase (213) or cytosolic phospholipase  
661 A<sub>2</sub> (175).

662 Once internalized, *L. monocytogenes* degrades the phagosome *via* the activity of a pore-  
663 forming toxin (listeriolysin) and two phospholipases (PlcA and PlcB), and escapes into the  
664 cytoplasm using actin-based motility, which is driven by the actin assembly-inducing protein,  
665 ActA. Following escape into the cytoplasm, *L. monocytogenes* can replicate within HUVECs  
666 and HBMECs and use actin-based motility to spread to adjacent cells (212, 213), thus  
667 avoiding an extracellular lifestyle. However, *L. monocytogenes* can be isolated from the CSF  
668 (216), suggesting that translocation through the cellular barriers of the CNS also occurs.

669 *B. pseudomallei* has a similar intracellular lifestyle to *L. monocytogenes*: the type III  
670 secretion system of *B. pseudomallei* (particularly the BopE and BopA effector proteins, and  
671 the Bsa translocation apparatus) is required for escape from the vacuole and the subversion of  
672 autophagy (217-222). In an epithelial cell line or a murine macrophage-like cell line, cell-to-  
673 cell spread then occurs by cell fusion, with the formation of multinucleated giant cells, in a  
674 type VI secretion-dependent process; BimA, necessary for actin-mediated motility, or Fla2,  
675 which is thought to mediate intracellular motility are also required and are considered to  
676 mediate prior cell-to-cell contact prior to cell fusion (223-227). To our knowledge, the ability  
677 of *B. pseudomallei* to directly invade HBMECs has not been investigated, but it would be  
678 reasonable to hypothesize that the intracellular lifestyle of *B. pseudomallei* may contribute to  
679 the pathogenesis of CNS invasion. Interestingly, a recent study investigated the variable  
680 virulence factors of *B. pseudomallei* associated with melioidosis in Australia, and reported  
681 that *B. pseudomallei* strains harboring a *bimA* allele that shares 95% homology with *B. mallei*  
682 *BimA* (*bimA*<sub>Bm</sub>) (228) were significantly associated with neurological melioidosis (229).

683 Patients that were infected with *B. pseudomallei* harboring the *bimA*<sub>Bm</sub> allele were found to be  
684 14 times more likely to present with neurological involvement compared to patients infected  
685 with strains harboring the *bimA*<sub>Bp</sub> variant (229), indicating a role for actin-mediated motility  
686 in either transgression of the BBB/BCSFB or transit to the brain *via* the olfactory or  
687 trigeminal nerve pathways (see below).

688 Several studies have shown that internalin B (InlB) is required for *L. monocytogenes* invasion  
689 of HUVECs (212-214) and HBMECs (213, 215). Greiffenberg *et al.* demonstrated that  
690 although *L. monocytogenes* deficient in InlB could adhere to HBMECs at comparable levels  
691 to the wild type strain, the invasive capability of the InlB mutant was reduced by > 100-fold  
692 (215). In contrast, one study reported that neither InlA nor InlB were required for *L.*  
693 *monocytogenes* invasion into HUVECs (230). It was suggested that these inconsistent  
694 findings may be attributable to differences in experimental conditions, especially involving  
695 the addition of normal human serum, which markedly affects InlB-mediated invasion due to  
696 the presence of anti-*Listeria* antibodies (231). In an inverted Transwell model of human  
697 choroid plexus epithelial cells, the cellular receptors for InlA and InlB (E-cadherin and met  
698 receptor tyrosine kinase respectively) were expressed on the basolateral ‘blood’ side, but not  
699 the apical ‘CSF’ side (232). In this model, *L. monocytogenes* invaded human choroid plexus  
700 epithelial cells exclusively from the basolateral side, and both InlA and InlB were required  
701 for efficient invasion.

702 Other *L. monocytogenes* virulence factors that may play a role in the attachment to, and  
703 invasion of, HBMECs include Vip and the autolysin IspC. Vip is a surface-expressed protein  
704 that is absent from non-pathogenic *Listeria* and binds to gp96 on host cells (160). A *vip*  
705 allelic replacement mutant was shown to be significantly less invasive than wild type *L.*  
706 *monocytogenes* in Caco-2 cells and L2071 mouse fibroblasts, but was not investigated in  
707 HBMECs. However, in intravenously inoculated mice,  $\Delta vip$  was attenuated for virulence and

708 associated with a significant decrease in bacterial loads within the brain (160). Similarly,  
709 IspC was also shown to contribute to *L. monocytogenes* invasion of the brain in a mouse  
710 model of hematogenous meningitis (233). Interestingly, the deletion of IspC did not affect the  
711 ability of *L. monocytogenes* to attach to, and invade HBMECs *in vitro*. In contrast,  $\Delta$ *ispC*  
712 demonstrated a significant reduction in attachment and invasion in sheep choroid plexus  
713 epithelial cells (compared to wild type *L. monocytogenes*), and purified IspC was capable of  
714 binding to these cells. Proteomic analyses revealed that IspC regulates the expression of other  
715 virulence factors, including ActA, InlC2 and a flagellin homologue (FlaA); therefore the  
716 phenotype observed by the  $\Delta$ *ispC* strain may be a result of the deletion of IspC in  
717 combination with a reduction in the expression of these factors (233).

#### 718 ***M. tuberculosis***

719 *M. tuberculosis* translocates from the apical chamber to the basolateral chamber in a  
720 Transwell model of HBMECs (118). *In vitro*, *M. tuberculosis* interacts with microvillus-like  
721 protrusions on HBMECs and is observed intracellularly, but not paracellularly. Antibiotic  
722 protection assays demonstrated that *M. tuberculosis* invaded HBMECs. Furthermore, *M.*  
723 *tuberculosis* co-localized with actin on the cell surface. Treatment of HBMECs with  
724 cytochalasin D significantly decreased *M. tuberculosis* invasion, suggesting that actin  
725 polymerization is required for internalization (118).

726 The mechanisms and host-pathogen interactions involved in CNS invasion by *M. tuberculosis*  
727 remain incompletely understood. In an *in vitro* BBB model consisting of bovine brain  
728 microvascular endothelial cells (BBMECs) co-cultured with rat astrocytes, the addition of  
729 purified recombinant heparin-binding haemagglutinin adhesin (rHBHA) induced actin  
730 filament rearrangement (234). This was shown to be dependent on the ability of rHBHA to  
731 bind to heparan sulfate glycosaminoglycans on the surface of BBMECs. The role of HBHA

732 in brain microvascular endothelial cell invasion by *M. tuberculosis* has not been investigated  
733 in the context of live bacteria. Furthermore, gene expression profiling of *M. tuberculosis*  
734 associated with HBMECs *in vitro* (versus non-adherent bacteria), did not identify HBHA as a  
735 gene that was significantly upregulated during infection (118). Jain *et al.* demonstrated that  
736 33 genes were upregulated > 8 fold in *M. tuberculosis* associated with HBMECs. Transposon  
737 mutants were created to determine if these upregulated genes were associated with HBMEC  
738 invasion; of the 33 tested mutants, 4 genes (*Rv0980c*, *Rv0987*, *Rv0989c* and *Rv1801*) were  
739 required for efficient invasion and/or intracellular survival (118).

740 *In vivo* screening of transposon mutant libraries has also been employed to identify potential  
741 *M. tuberculosis* virulence factors that are associated with CNS invasion. Be *et al.* inoculated  
742 pools of *M. tuberculosis* mutants intravenously into BALB/c mice and determined that  
743 mutants deficient in the expression of 5 genes (*Rv0311*, *Rv0805*, *Rv0931c*, *Rv0986* and  
744 *MT3280*) were significantly attenuated in the brain, but not the lungs. *In vitro* studies  
745 confirmed that these mutants invaded HBMECs at significantly reduced levels compared to a  
746 negative control transposon mutant (235). *Rv0931c* (also known as *pknD*) encodes a serine-  
747 threonine protein kinase, and was also identified as a key microbial factor required for brain  
748 infection in a guinea pig model of hematogenous meningitis, from a library of 398 mutants  
749 (158). When studied independently in a mouse model of infection, the *pknD* mutant was  
750 associated with a significant reduction in bacterial loads within the brain, but not the lungs. *In*  
751 *vitro*, *pknD* was required for efficient invasion and intracellular survival within HBMECs,  
752 this phenotype was not observed in HUVECs, A549 lung epithelial cells or J447  
753 macrophages. Microspheres coated with *pknD* adhered to HBMECs at significantly higher  
754 levels than microspheres coated with bovine serum albumin, and it was proposed that *pknD*  
755 binds to HBMECs *via* interactions with laminin within the extracellular matrix (158).

## 756 **Paracellular penetration of brain microvascular endothelial cells**

757 The breakdown of tight junctions between cells of the BBB and/or BCSFB may occur due to  
758 the activity of microbial virulence factors/toxins, and/or inflammatory-mediated processes.  
759 Early studies in intracisternally infected rats demonstrated that *E. coli* K1, *S. pneumoniae* and  
760 *H. influenzae* type B caused an increase in pinocytotic vesicle formation in the cerebral  
761 capillary endothelium, in addition to the complete separation of intercellular junctions (236).  
762 Further experiments with *H. influenzae* showed that these cerebral capillary morphological  
763 changes were associated with a significant increase in BBB permeability to circulating <sup>125</sup>I-  
764 albumin (236). This suggests that bacterial meningeal pathogens may open a paracellular  
765 route of CNS entry, although these data should be interpreted cautiously as intracisternal  
766 inoculation cannot be used to model the physiological events that occur during hematogenous  
767 meningitis, as the inoculum is injected directly into the cisternal spaces between the  
768 arachnoid and pia mater. The development of *in vitro* models of the human BBB/BCSFB and  
769 *in vivo* models of hematogenous meningitis has enabled microbial invasion *via* the  
770 paracellular route to be studied in some detail. From these studies (reviewed below for  
771 specific pathogens), it appears that microbes may initially attach to and transcellularly invade  
772 brain microvascular endothelial cells, and that these host-pathogen interactions may lead to a  
773 subsequent increase in barrier permeability and a loss of tight junctions. Thus it is likely that  
774 both transcellular and paracellular mechanisms of CNS entry are exploited by bacterial  
775 pathogens. It is of considerable clinical importance that the breakdown of the BBB/BCSFB in  
776 bacterial meningitis also results in generally higher levels of antibiotics in the CSF.

#### 777 *E. coli*

778 Several *E. coli* virulence factors promote the destruction of tight junctions between brain  
779 microvascular endothelial cells. In addition to triggering the internalization of *E. coli* K1 into  
780 HBMECs, the binding of OmpA to its gp96 homologue receptor stimulates inducible nitric  
781 oxide synthase and nitric oxide production, which enhances the generation of cyclic GMP

782 (237). This increase in cyclic GMP activates protein kinase C $\alpha$  (237), which leads to a ras  
783 GTPase-activating-like protein (IQGAP1)-mediated dissociation of  $\beta$ -catenin from vascular  
784 endothelial-cadherin at adherens junctions (238). Immunofluorescence experiments  
785 demonstrated that vascular endothelial-cadherin was redistributed from the intercellular  
786 junctions to the sites of bacterial attachment (239). This resulted in a significant increase in  
787 HBMEC permeability to horseradish peroxidase and a parallel decrease in transendothelial  
788 electrical resistance (239). Furthermore, the activation of host cytosolic phospholipase A<sub>2</sub> by  
789 *E. coli* OmpA and Nlp1 may play a role in reducing the integrity of the BBB. Using co-  
790 cultures of primary BBMECs and primary bovine retinal pericytes seeded onto Transwell  
791 membranes, Salmeri *et al.* demonstrated that the arachidonic acid released following  
792 cytosolic phospholipase A<sub>2</sub> activation acts as a substrate for cyclooxygenase to produce  
793 prostaglandins, which trigger the synthesis of vascular endothelial growth factor by brain  
794 endothelial cells (119). This was associated with a significant decrease in transendothelial  
795 electrical resistance and a dramatic detachment of pericytes from the Transwell membrane.  
796 These effects were not observed when the pericyte vascular endothelial growth factor  
797 receptor-1 was blocked using a polyclonal antibody. These findings suggest that the vascular  
798 endothelial growth factor released by brain endothelial cells may bind to the vascular  
799 endothelial growth factor receptor-1 on adjacent pericytes, and trigger ablation of the  
800 pericytes from the basal membrane, potentially opening a paracellular route (119).

801 *In vitro* studies using the mouse brain microvascular endothelial cell line, Bend.3, have  
802 demonstrated a role for lipopolysaccharide (LPS, *E. coli* 055:B5) in inducing BBB  
803 permeability. Similar to OmpA, LPS was shown to activate protein kinase C isoforms, which  
804 led to the activation of RhoA, and the stimulation of downstream nuclear factor- $\kappa$ B signalling  
805 and myosin light chain phosphorylation (240, 241). These signalling events were associated  
806 with a decrease in the expression of the tight junction proteins claudin-5 and zonula

807 occludens-1, and a reduction in transendothelial electrical resistance. LPS also induced  
808 changes in F actin organization, leading to paracellular gaps and stress fiber formation (240).  
809 In Shiga toxin-producing *E. coli* associated with hemolytic uremic syndrome and CNS  
810 involvement, both Shiga toxin-1 and Shiga toxin-2 induce cytotoxic effects that lead to a  
811 decrease in BBB integrity and cellular injury *in vivo* and *in vitro* (242-245). LPS may mediate  
812 these effects (242, 245).

### 813 *H. influenzae*

814 Quagliariello *et al.* first demonstrated that *H. influenzae* caused an increase in BBB  
815 permeability and tight junction disruption in adult Wistar rats following intracisternal  
816 inoculation of approximately  $10^6$  colony forming units (236). Using this intracisternal  
817 inoculation model, the role of *H. influenzae* type B lipooligosaccharide (LOS) was  
818 investigated. Purified LOS and outer membrane vesicles containing non-cell associated LOS  
819 induced a dose-dependent increase in BBB permeability to  $^{125}$ I-bovine serum albumin and  
820 CSF pleocytosis (246, 247). In leukopenic rats, no increases in BBB permeability and CSF  
821 leukocyte counts were observed following LOS or outer membrane vesicle challenge,  
822 suggesting that the decrease in BBB integrity is mediated by the intact host leukocyte  
823 response (246, 247). In contrast, Tunkel *et al.* demonstrated that *H. influenzae* type B LOS  
824 significantly increased the permeability of rat brain microvascular endothelial cells to  $^{125}$ I-  
825 albumin in the absence of host inflammatory cells *in vitro* (248). Furthermore, in the absence  
826 of monocytic cells, purified LOS was cytotoxic to BBMECs. However, *in vivo* studies have  
827 shown that nitric oxide production within the CSF is required for LOS-mediated pleocytosis  
828 and increased BBB permeability (249), and that host PAF synergizes with *H. influenzae* LOS  
829 to augment these events (250).



830 Peptidoglycan within the *H. influenzae* cell wall may also contribute to BBB/BCSFB  
831 dysfunction *in vivo*. Intracisternally inoculated peptidoglycan induced an increase in BBB  
832 permeability to <sup>125</sup>I-albumin (251) and CSF leukocytosis (252), although the CSF  
833 leukocytosis was attenuated compared to animals challenged with LOS (252). It was  
834 subsequently hypothesized that *H. influenzae* cell wall components, such as peptidoglycan,  
835 activate brain microvascular endothelial cells and induce the separation of intercellular  
836 junctions. LOS may then augment this response by activating the recruited leukocytes to  
837 produce pro-inflammatory cytokines that further contribute to BBB breakdown (252);  
838 however, the mechanisms by which this may occur have not been elucidated. The *H.*  
839 *influenzae* type B porin may also contribute to these events. The inoculation of purified porin  
840 into the fourth cerebral ventricle of rats induced the expression of interleukin (IL)-1 $\alpha$ , tumor  
841 necrosis factor- $\alpha$  (TNF- $\alpha$ ) and macrophage inflammatory protein-2, which preceded an  
842 increase in BBB permeability to serum proteins and leukocytes, and increased brain water  
843 content (108).

#### 844 *N. meningitidis*

845 As discussed above, the *N. meningitidis*-induced  $\beta$ 2-adrenoceptor/ $\beta$ -arrestin signalling in  
846 HBMECs results in the formation of cortical plaques and membrane protrusions. In parallel,  
847 the type IV pilus-mediated translocation of  $\beta$ -arrestin to the cortical plaque also acts as a  
848 molecular scaffold for additional *N. meningitidis*-induced signalling events that open the  
849 paracellular route (187). These events include the recruitment of the Par3/Par6/PKC $\zeta$  polarity  
850 complex to the cortical plaque in a Cdc42-dependent manner, and the accumulation of  
851 adherens junction (vascular endothelial-cadherin, p120-catenin,  $\beta$ -catenin) and tight junction  
852 (zonula occludens-1, zonula occludens-2, claudin-5) proteins underneath the site of bacterial  
853 adhesion (253). Labelling of vascular endothelial-cadherin demonstrated that *N. meningitidis*  
854 delocalized this protein from the adherens junctions between HBMECs and caused it to be

855 redistributed to the cortical plaque. This depletion of junctional proteins was associated with  
856 an increase in barrier permeability to Lucifer yellow and the formation of gaps between cells,  
857 thus opening a paracellular route of entry (253). Interestingly, the recruitment of junctional  
858 components by *N. meningitidis* occurs only in endothelial cells, as it was demonstrated that  
859 meningococci do not activate the  $\beta$ 2-adrenoceptor/ $\beta$ -arrestin signalling pathway in epithelial  
860 cells (254). Therefore, different host-pathogen interactions may occur at the BBB and  
861 BCSFB.

862 In an inflammatory-mediated process, *N. meningitidis* stimulates HBMECs to produce matrix  
863 metalloproteinase-8, which was shown to cleave occludin and cause it to dissociate from tight  
864 junctions *in vitro* (255). The cleavage of occludin was prevented in the presence of matrix  
865 metalloproteinase-8 inhibitors and in cells transfected with matrix metalloproteinase-8  
866 siRNA. Inhibition of matrix metalloproteinase-8 activity also significantly decreased the *N.*  
867 *meningitidis*-induced permeability of HBMECs to 40 kDa FITC-dextran at 24 hours.  
868 Furthermore, the inhibition of matrix metalloproteinase-8 prevented the caspase-independent  
869 detachment of HBMECs from a matrigel matrix on a solid support following *N. meningitidis*  
870 infection (255).

#### 871 ***Streptococcus spp.***

872 Group B streptococcus, *S. pneumoniae* and *S. suis* invade brain microvascular endothelial  
873 cells *via* transcellular mechanisms; however, the associated production of pro-inflammatory  
874 cytokines, the activity of pore-forming toxins and the induction of plasmin activity may also  
875 lead to a concurrent weakening of the BBB, potentially opening a paracellular route of entry.  
876 *In vivo* and *in vitro* studies have shown that group B streptococci, pneumococci and *S. suis*  
877 induce the production of a range of cytokines and chemokines, including IL-1 $\beta$ , IL-6, IL-8,  
878 IL-10, TNF- $\alpha$ , MCP-1, GM-CSF, Gro- $\alpha$  and Gro- $\beta$  in brain microvascular endothelial cells

879 (109, 256-258) and brain tissue (94, 259, 260). The stimulation of HBMECs with cytokines  
880 (including TNF- $\alpha$ , IL-1 $\beta$ , IL-6 and IL-17F) leads to cytoskeletal rearrangements and a  
881 redistribution of tight junction and adherens junction proteins, resulting in a decrease in  
882 barrier integrity (261-267). Therefore, the host inflammatory response to group B  
883 streptococcus, *S. pneumoniae* and *S. suis* meningitis may contribute to the disruption of tight  
884 junctions. In HBMECs, the group B streptococcal Pila-mediated activation of focal adhesion  
885 kinase induces bacterial uptake, but also triggers a parallel mitogen-activated protein  
886 kinase/extracellular signal-regulated kinase pathway, which leads to the secretion of IL-8  
887 (109). Furthermore, the production of the IL-8 homologue (KC) in mice intravenously  
888 inoculated with group B streptococci resulted in the recruitment of neutrophils and an  
889 increase in BBB permeability to Evans blue and FITC-albumin (109). It is likely that  
890 disruption of the BBB/BCSFB by host inflammatory factors represents a common  
891 physiological event that occurs during bacterial CNS invasion.

892 Using a separate mechanism, group B streptococcus, *S. pneumoniae* and *S. suis* may hijack  
893 the host plasminogen system to cause cellular injury and disrupt tight junctions between  
894 HBMECs. Plasminogen is found in plasma and extracellular fluids, and upon activation is  
895 converted to plasmin, which degrades the extracellular matrix and may upregulate matrix  
896 metalloproteinases that degrade tight junctions (268). *In vitro*, plasminogen binds to  
897 streptococcal glyceraldehyde-3-phosphate dehydrogenase (269-271). Enolase and choline-  
898 binding protein E have also been identified as additional pneumococcal plasminogen binding  
899 proteins (272, 273). Once plasminogen is bound to its binding protein, it becomes activated  
900 and is subsequently converted to plasmin by exogenous tissue plasminogen activator and  
901 urokinase (269, 270, 274). Plasmin bound to the surface of group B streptococci enhanced the  
902 ability of bacteria to adhere to, and invade HBMECs, and induced a significant decrease in  
903 transendothelial electrical resistance (275). Furthermore, streptococcal-bound plasmin

904 contributed to cellular injury, characterized by human brain microvascular cell detachment  
905 and lactate dehydrogenase release (275). In the human vascular endothelial cell line EaHy,  
906 plasmin bound to the pneumococcal cell surface cleaved vascular endothelial-cadherin from  
907 adherens junctions and increased bacterial translocation across endothelial barriers (276).  
908 Taken together, group B streptococcus, *S. pneumoniae* and *S. suis* may bind host  
909 plasminogen and hijack the proteolytic activity of plasmin to potentially open a paracellular  
910 route between cells. Plasminogen has also been shown to bind to *N. meningitidis* (277, 278),  
911 suggesting that this mechanism may be exploited by several common meningeal pathogens.

912 Group B streptococcus, *S. pneumoniae* and *S. suis* produce pore-forming hemolysins named  
913  $\beta$ -hemolysin/cytolysin, pneumolysin and suilysin respectively. In HBMECs, these  
914 hemolysins are cytotoxic and promote lactate dehydrogenase release, the loss of cytoplasmic  
915 density, discontinuity of cytoplasmic membranes, clumping of nuclear chromatin, dislocation of  
916 the endoplasmic reticulum, cell rounding and detachment (97, 202, 207). These  
917 morphological changes most likely result in the formation of paracellular gaps. The effects of  
918 pneumolysin have also been investigated in rat astrocytes; the addition of pneumolysin to  
919 monolayers of primary astrocytes resulted in cell shrinkage and the subsequent separation of  
920 cells from each other (107). In rat brain slices, exposure to pneumolysin led to astrocytic  
921 process retraction and reorganization of astrocytes within the glia limitans. These  
922 pneumolysin-induced changes were associated with increased interstitial fluid retention and  
923 BBB permeability, which facilitated the penetration of macromolecules and bacteria into  
924 brain slices (107). These data suggest that in addition to mediating cytotoxic effects in brain  
925 microvascular endothelial cells, bacterial hemolysins may also induce morphological changes  
926 in other components of the neurovascular unit.

## 927 **Trojan horse penetration of brain microvascular endothelial cells**

928 Bacteria that are capable of surviving within host peripheral immune cells have the ability to  
929 invade the CNS *via* the Trojan horse route. This route of entry has been suggested for *L.*  
930 *monocytogenes* (279), *B. pseudomallei* (280), *S. suis* (146) and *M. tuberculosis* (281). In  
931 C57Bl/6 x DBA/2 mice, *L. monocytogenes*-infected peripheral blood leukocytes disseminated  
932 to the CNS and were shown to induce brain colonization more efficiently than extracellular *L.*  
933 *monocytogenes* (279). Furthermore, CNS infection was not reduced following the elimination  
934 of extracellular *L. monocytogenes* with gentamicin delivered by surgically implanted osmotic  
935 pumps (282), suggesting that intracellular bacteria were responsible for neuroinvasion.  
936 Drevets *et al.* demonstrated that in C57Bl/6 mice, *L. monocytogenes* associated with CD11b+  
937 monocytes, and further analysis revealed that most of these infected monocytes belonged to  
938 the Ly-6C<sup>high</sup> subset (283). Following systemic *L. monocytogenes* infection, 90% of CD11b+  
939 leukocytes isolated from the brain were Ly-6C<sup>high</sup> and the influx of these cells correlated with  
940 an increase in brain colonization, suggesting that Ly-6C<sup>high</sup> monocytes may act as Trojan  
941 horses *in vivo* (283). In BALB/c mice, bone marrow myelomonocytic cells (CD31+: Ly-6C+:  
942 CD11b+: Ly-6G<sup>low</sup>) may also transport *L. monocytogenes* to the CNS (284).

943 In a hematogenous model of melioidosis meningitis, *B. pseudomallei* established a primary  
944 focus of infection in the spleen, followed by migration to the bone marrow and subsequent  
945 spread to the brain (280). *B. pseudomallei* associated with CD11b+ cells within the bone  
946 marrow and spleen, and it was hypothesized that these cells may serve as Trojan horses that  
947 facilitate bacterial spread to the brain (280). In adoptive transfer experiments, splenic  
948 CD11b+ cells and bone marrow CD11b+ cells harboring *B. pseudomallei* induced brain  
949 colonization. In contrast, bacterial burdens within the brain were attenuated following the  
950 adoptive transfer of *B. pseudomallei*-infected CD11b- cells, and equivalent numbers of  
951 extracellular *B. pseudomallei* were unable to colonize the brain. These data suggest that *B.*  
952 *pseudomallei*-infected CD11b+ cells within the spleen and bone marrow may disseminate to

953 the CNS and contribute to the pathogenesis of neurological melioidosis with meningitis.  
954 Further experiments demonstrated that the expression of selectin on *B. pseudomallei*-infected  
955 CD11b+ cells was required for the development of meningitis in recipient mice (280).

## 956 **THE OLFACTORY NERVE AS A PORTAL TO THE BRAIN**

957 The olfactory system (Figure 3A) comprises the odor detecting sensory system that exists in  
958 the peripheral nervous system within the nasal cavity and, within the CNS, the olfactory bulb  
959 and higher processing centers of the brain. Within the nasal cavity, the olfactory epithelium  
960 lines the more dorsal and caudal regions of the cavity. Olfactory sensory neurons reside  
961 within the olfactory epithelium and project a dendritic knob-like swelling with 20-30 cilia  
962 into the mucous layer lining the nasal cavity. The olfactory sensory neuron cilia possess  
963 odorant receptors (285), which bind a large spectrum of ligands (286). Binding of an inhaled  
964 odorant to its receptor results in signal amplification and the activation of a signal  
965 transduction pathway, leading to the generation of an action potential (287). The olfactory  
966 epithelium also comprises sustentacular cells, which are glia-like cells whose apical surfaces  
967 form the epithelial surface lining the nasal cavity. These supporting cells form tight junctions  
968 with each other and with the dendrites of the olfactory sensory neurons, forming the primary  
969 barrier from the environment (288). The olfactory epithelium also contains basal cells, which  
970 are stem and multipotent progenitors from which new olfactory sensory neurons are  
971 generated; and developing neurons, which are newly generated from the basal cells (289,  
972 290). The axons of olfactory sensory neurons penetrate the basement membrane beneath the  
973 epithelium entering the lamina propria where they are met by specialized glial cells, olfactory  
974 ensheathing cells. Olfactory ensheathing cells surround multiple axons and bundle them  
975 together in larger fascicles that comprise the olfactory nerve (291, 292). Within the lamina  
976 propria are characteristic Bowman's glands of the olfactory mucosa, whose ducts penetrate  
977 the epithelium above to supply specialized mucus to the epithelium.

978 The axons of the olfactory nerve course through the lamina propria towards the brain,  
979 penetrating the skull through the cribriform plate and enter the brain at the olfactory bulb  
980 (Figure 3A-C). Within the olfactory bulb the sensory axons form specialized structures,  
981 glomeruli, where they synapse with mitral cells, which carry the sensory signal to higher  
982 brain structures (293, 294). Olfactory sensory neurons are directly exposed to the external  
983 environment *via* the nasal cavity; therefore microbes within the nasal cavity may potentially  
984 exploit the olfactory pathway and access the subarachnoid space and the olfactory bulb. From  
985 the olfactory bulb, viruses have been shown to migrate to higher brain regions including the  
986 basal nuclei, thalamus, hypothalamus, cerebrum and cerebellum in animal models of  
987 infection (295-297).

#### 988 **THE TRIGEMINAL NERVE AS A PORTAL TO THE BRAIN**

989 The trigeminal nerve is the largest cranial nerve whose afferent branches carry touch, pain  
990 and noxious stimuli from the face, the corneas of the eyes, and the oral and nasal cavities.  
991 The trigeminal nerve innervates the olfactory and respiratory epithelia of the nasal cavity *via*  
992 branches of the ophthalmic (V1) and maxillary (V2) nerves (Figure 4A) (298). The trigeminal  
993 nerve endings are separated from the lumen of the nasal cavity *via* an apical tight junction  
994 complex at the epithelial surface (299, 300). The same trigeminal nerves that innervate the  
995 olfactory epithelium also branch to innervate the olfactory bulb, providing an alternative  
996 route for entry of pathogens (301). Lipid soluble trigeminal irritants within the nasal cavity  
997 may reach their receptors located on trigeminal nerve endings by diffusing across tight  
998 junctions, potentially using paracellular mechanisms (302), or by interacting with specialized  
999 solitary chemosensory cells within the nasal and respiratory epithelium, which form synaptic  
1000 contacts with trigeminal nerve fibers (303). Solitary chemosensory cells isolated from mice  
1001 were shown to detect bacterial signals such as acyl-homoserine lactones and trigger  
1002 downstream signalling pathways associated with bitter irritant transduction (304).

1003 The trigeminal nerve fibers course to the brain forming the trigeminal ganglion outside the  
1004 brainstem but beneath the dura mater (298). From the trigeminal ganglion the sensory axons  
1005 innervate the trigeminal nucleus in the brainstem. Therefore the trigeminal nerve, like the  
1006 olfactory nerve, could provide a direct pathway by which bacteria within the nasal mucosa  
1007 may access the brain independently of the blood. It should be noted that bacterial entry to the  
1008 brain *via* the trigeminal nerve might not be limited to intranasal infection. For example,  
1009 viruses such as herpes simplex virus type 1 access the trigeminal nerve through infection of  
1010 the cornea (305) and potentially any epithelial infection of the facial skin or oral cavity could  
1011 allow access to the trigeminal nerve.

## 1012 **PROTECTING THE CNS FROM MICROBIAL INVASION VIA THE NASAL** 1013 **CAVITY**

1014 The nasal cavity is constantly exposed to inhaled microbes, allergens and particulate material.  
1015 The epithelium lining the nasal nares (306, 307) and the nasal cavity (308) harbor normal  
1016 bacterial flora; and CNS pathogens such as *S. aureus*, *S. pneumoniae*, *N. meningitidis* and *H.*  
1017 *influenzae* may asymptotically colonize the nasopharynx in some individuals (306, 309,  
1018 310). Therefore, the nasal cavity features innate defense mechanisms to filter inhaled air and  
1019 prevent microbes from invading deeper tissues.

1020 Goblet cells and submucosal glands secrete mucins (primarily MUC5AC and MUC5B) into  
1021 the lumen of the airway and nasal cavity. MUC5AC and MUC5B cross-link to form the  
1022 structural framework of a mucous barrier, which floats on top of a periciliary layer of  
1023 membrane-bound mucins (including MUC1 and MUC4) and glycoproteins (311). Through a  
1024 mucociliary clearance process, inhaled bacteria and particulate matter become trapped in the  
1025 mucous layer, and are swept towards the pharynx by the coordinated beating of cilia located  
1026 on the surface of epithelial cells. A recent study demonstrated that MUC5B, but not



1027 MUC5AC, is essential for mucociliary clearance activity and the prevention of bacterial  
1028 spread from the nasal cavity to the lower respiratory tract (312). Mucous secretion and  
1029 mucociliary clearance are enhanced in the presence of inflammatory mediators (313) and  
1030 microbial pathogens (314-316), and following exposure to cigarette smoke (317).  
1031 Antimicrobial substances, including enzymes (lysozyme), protease inhibitors (secretory  
1032 leukoprotease inhibitor, elastase inhibitor,  $\alpha$ 1-antiprotease, antichymotrypsin), antimicrobial  
1033 peptides ( $\beta$ -defensins, L37) and oxidants (nitric oxide and hydrogen peroxide) are secreted by  
1034 epithelial cells into the airway surface liquid, and provide a further line of defense against  
1035 microbes (318). Finally, tight junctions and adherens junctions between the epithelial cells  
1036 lining the nasal cavity and airway create a cellular barrier that prevents microbial spread.  
1037 Dendritic cells localized beyond the epithelium extend processes through these tight junctions  
1038 to interact with ligands and collaborate with resident macrophages to remove foreign antigens  
1039 (319).

1040 Despite the host defences that exist within the nasal cavity and upper airways, bacteria  
1041 colonizing the epithelium may invade deeper tissues and cause disease. In the case of  
1042 asymptomatic colonizers, the transition from a commensal to a pathogenic phenotype may be  
1043 due to within-host evolution that results in genetic changes to the regulation of virulence  
1044 factors (320, 321). Alterations in susceptibility to disease in immunocompromised individuals  
1045 may also promote the spread of bacteria from the nasal cavity (318). Respiratory pathogens  
1046 such as *Pseudomonas aeruginosa* and *Burkholderia cenocepacia* have been demonstrated to  
1047 trigger the disassembly of tight junctions between epithelial cells to migrate to the lower  
1048 respiratory tract (322, 323), whereas *N. meningitidis* redistributes junctional proteins at the  
1049 olfactory epithelium (324). Harris *et al.* (325) demonstrated that the intact olfactory  
1050 epithelium usually represents an effective barrier against infiltration of *S. aureus* into the  
1051 nasal lamina propria. However, following mild detergent damage to the nasal epithelium of

1052 C57Bl/6 mice, intranasally delivered *S. aureus* was detected in the olfactory epithelium,  
1053 olfactory nerve and the olfactory bulb as early as 6 hours post infection (325). Indeed,  
1054 damage to the olfactory epithelium within the nasal cavity appears to be an important and  
1055 common event in bacterial spread to the CNS *via* the olfactory nerve (324-327).

## 1056 **PATHOGENS THAT ENTER THE BRAIN THROUGH THE NOSE**

### 1057 **Bacteria**

1058 Bacteria within the nasal cavity could potentially penetrate the olfactory epithelium by  
1059 directly infecting olfactory sensory neurons; they could enter *via* the ducts of the Bowman's  
1060 glands; or they could enter after reactive inflammatory processes compromise the epithelial  
1061 tissue leading to extravasion and subsequent tissue damage after exposure to bacterial  
1062 virulence factors. Bacterial signalling molecules, such as acyl-homoserine lactones, are  
1063 detected by solitary chemosensory cells within the nasal mucosa that synapse with the  
1064 trigeminal nerve, triggering trigeminally-mediated reflex reactions (304). Mouse solitary  
1065 chemosensory cells exposed to *P. aeruginosa* acyl-homoserine lactones responded with an  
1066 increase in intracellular Ca<sup>2+</sup>. Furthermore, *in vivo* studies demonstrated that mice exposed to  
1067 acyl-homoserine lactones *via* the retronasal stream experienced significant reductions in  
1068 respiratory rate, suggesting that interactions between bacterial acyl-homoserine lactones and  
1069 chemosensory cells are capable of inducing responses indicative of nasal trigeminal irritants  
1070 (304). The activation of nasal solitary chemosensory cells by acyl-homoserine lactones  
1071 triggers a pro-inflammatory response (328); this response could damage the integrity of the  
1072 epithelium and enable bacteria to access the trigeminal nerve endings. Once bacteria have  
1073 penetrated the epithelium, they could potentially travel along the olfactory and/or trigeminal  
1074 nerves to the brain within axons, between the axons, within the surrounding glia (olfactory

1075 ensheathing cells or Schwann cells), or between the glia and the connective tissue sheath  
1076 (perineurium) that contains the nerve bundle.

1077 Meningeal pathogens such as *S. pneumoniae* and *N. meningitidis* unequivocally enter the CSF  
1078 *via* the hematogenous route. Animal models of infection have also highlighted an alternative  
1079 route of CNS entry, where these bacteria may directly invade the olfactory bulb within the  
1080 brain from the nasal mucosa. *B. pseudomallei* has also been shown to invade the brain *via* the  
1081 olfactory and trigeminal nerves, whereas *L. monocytogenes* demonstrates a predilection for  
1082 the trigeminal nerve and brainstem. The following section reviews the evidence for CNS  
1083 invasion from the nasal cavity for these pathogens.

#### 1084 *S. pneumoniae*

1085 Early studies by Rake demonstrated that *S. pneumoniae* rapidly enters the olfactory bulb in  
1086 Swiss mice following intranasal inoculation (329). Remarkably, pneumococci were isolated  
1087 from the olfactory bulbs of the brain 1 minute post-infection. At this time point, bacteria were  
1088 not isolated from the blood, effectively excluding a hematogenous route of CNS invasion. At  
1089 2 minutes post-infection, pneumococci were observed between the sustentacular cells of the  
1090 olfactory epithelium, within the perineural space of the olfactory nerve and within the  
1091 subarachnoid space (329). Similar findings were reported by van Ginkel *et al.*, who  
1092 demonstrated that intranasally delivered *S. pneumoniae* (strain EF3030) could be recovered  
1093 from nasal washes, the olfactory nerve and epithelium, and the olfactory bulb 24 hours post-  
1094 infection in CBA/CAHN/*xid* mice (330). These sites remained colonized throughout the  
1095 duration of the experiment, until day 39 post-infection; however, this did not lead to  
1096 extensive brain infection. In contrast, non-encapsulated pneumococci were unable to persist  
1097 within the nasal cavity and olfactory epithelium, and did not invade the olfactory bulb within  
1098 the brain. Briles *et al.* confirmed that the pneumococcal capsular polysaccharide is an

1099 important mediator of olfactory bulb infection in CBA/CAHN/*xid* mice, as only opaque  
1100 variants were isolated from the olfactory bulb following intranasal inoculation (331). In these  
1101 studies, intranasally delivered *S. pneumoniae* did not lead to bacteremia, suggesting a role for  
1102 direct brain invasion from the nasal cavity. *S. pneumoniae* was also recovered from the  
1103 trigeminal ganglia, further suggesting that the trigeminal nerve bundles may also be a  
1104 potential route to the brain for this bacterial pathogen (330). However, it must be noted that  
1105 these studies utilised CBA/CAHN/*xid* mice, which possess a mutation in the Bruton's  
1106 tyrosine kinase gene and thus do not respond to thymus-independent type II antigens (332).  
1107 These mice also fail to respond to capsular polysaccharide; therefore pneumococcal infection  
1108 in this mouse strain is unlikely to accurately model human disease.

1109 Incubation of *S. pneumoniae* with gangliosides prior to intranasal infection resulted in  
1110 reduced colonization of the nasal mucosa, olfactory system and brain (330).  
1111 Phosphorylcholine residues on the pneumococcal cell wall have been shown to bind to  
1112 gangliosides (333), suggesting that gangliosides could be an important target for *S.*  
1113 *pneumoniae* attachment in the neuroepithelium. *In vitro* studies have also demonstrated that  
1114 *S. pneumoniae* can invade olfactory ensheathing cells *via* mannose receptor-mediated  
1115 endocytosis (334); however, the role of olfactory ensheathing cells in pneumococcal infection  
1116 and the exact mode of transport of *S. pneumoniae* along the olfactory and/or trigeminal  
1117 nerves has not yet been studied.

### 1118 *N. meningitidis*

1119 In meningococcal meningitis, >60% of patients develop meningitis without septic shock  
1120 (335), suggesting the possibility that *N. meningitidis* may also penetrate the CNS without  
1121 prior bacteremia. In a mouse model of intranasal *N. meningitidis* infection, 20% of mice  
1122 developed lethal meningitis; however, no bacteria were isolated from the blood (324). At 3

1123 days post-infection, lesions and polymorphonuclear cells were observed within the olfactory  
1124 epithelium. Furthermore, *N. meningitidis* infection was associated with significant damage to  
1125 the olfactory epithelium, and reduced its thickness in discrete, non-continuous sections of  
1126 epithelium by more than 50% compared to controls. Immunofluorescence studies  
1127 demonstrated *N. meningitidis* within the olfactory epithelium, basement membrane, lamina  
1128 propria, along the olfactory nerves in the cribriform plate and within the nerve fiber layer of  
1129 the olfactory bulb. Meningococci were also isolated from the CSF. Due to the absence of  
1130 bacteremia, it was suggested that *N. meningitidis* may travel from the nasal cavity to the  
1131 meninges and subarachnoid space *via* the olfactory nerves (324). In rhesus macaques,  
1132 commensal neisseriae (RM *Neisseria*) were shown to be transmitted between animals, and  
1133 naturally colonized the epithelium covering the cribriform plate (336), suggesting that  
1134 migration of neisseriae along the olfactory pathway is not a phenomenon that is observed  
1135 only in animals experimentally-inoculated with pathogenic *N. meningitides*. In the African  
1136 “meningitis belt” the onset of meningococcal meningitis epidemics coincides with harsh, dry  
1137 harmattan winds. It has been hypothesized that these harsh environmental conditions may  
1138 irritate the mucous membranes (such as the olfactory mucosa), and enable *N. meningitidis* to  
1139 penetrate the epithelium and invade the CNS (337), although this has not yet been  
1140 investigated.

1141 Sjölander and Jonsson demonstrated that the expression of the junctional protein N-cadherin  
1142 in the olfactory epithelium and lamina propria was significantly reduced in mice intranasally  
1143 infected with *N. meningitidis* (324). These data suggest that *N. meningitidis* may invade the  
1144 olfactory epithelium *via* the destruction or rearrangement of cellular junctions between  
1145 sustentacular cells and olfactory sensory neurons, although additional studies are required to  
1146 confirm these findings. The mechanism by which *N. meningitidis* may travel along the  
1147 olfactory nerve pathway to invade the nerve fiber layer of the olfactory bulb, the meninges

1148 and CSF are also unknown. The meninges and the subarachnoid space extend over the  
1149 cribriform plate and further into the olfactory foramen where the olfactory nerves pass  
1150 through the cribriform plate (338, 339). Bacteria travelling along the olfactory pathway could  
1151 potentially invade the meninges after transversing the cribriform plate, and subsequently  
1152 enter the CSF. However, there is a positive pressure from the brain to the nasal lymphatics  
1153 due to the drainage of CSF through the cribriform plate, and it is unknown how *N.*  
1154 *meningitidis* could travel against this pressure gradient.

### 1155 ***B. pseudomallei***

1156 The olfactory pathway has been identified as a route of CNS invasion by *B. pseudomallei*.  
1157 Owen *et al.* used a capsule-deficient strain, which fails to survive within the blood, to  
1158 demonstrate that bacteria were present in the olfactory epithelium and olfactory bulb of  
1159 BALB/c mice 24 hours post-infection, in the absence of bacteremia. It was also shown that  
1160 when colonization occurred only in one side of the nasal cavity, wild-type *B. pseudomallei*  
1161 was detected in the olfactory epithelium and olfactory bulb of the infected side only.  
1162 Combined, these data suggest that *B. pseudomallei* is able to directly invade the brain *via* the  
1163 olfactory pathway without blood-borne infection (105). These findings also demonstrate that  
1164 the presence of the polysaccharide capsule is not a prerequisite for CNS invasion *via* the  
1165 olfactory pathway, presumably in contrast to systemic invasion *via* the BBB/BCSFB.  
1166 However, it might be the case that an absence of capsule does diminish translocation *via* the  
1167 olfactory or trigeminal nerves, since it is not possible to definitively compare with a presence  
1168 of capsule due to hematogenous spread in the latter case. Figure 3 (D-E) illustrates *B.*  
1169 *pseudomallei* invasion of the nasal cavity, olfactory epithelium and ventral nerve fiber layer  
1170 of the olfactory bulb in a mouse model of acute melioidosis. It may be noted that mice, like  
1171 most mammals, are naturally susceptible to melioidosis.

1172 The mechanisms by which *B. pseudomallei* migrates along the olfactory nerves have recently  
1173 been investigated (326). In BALB/c mice, we demonstrated that intranasally-delivered *B.*  
1174 *pseudomallei* was associated with widespread crenellation of the olfactory epithelium and  
1175 loss of the neuron cell bodies. This neuronal loss led to the degeneration of olfactory axons  
1176 within 24 – 48 hours, and the axon-devoid, hollow olfactory nerve fascicles that are  
1177 surrounded by olfactory ensheathing cells provided an open conduit for bacterial passage  
1178 from the nasal cavity through the cribriform plate and into the nerve fiber layer of the  
1179 olfactory bulb. No bacteria were observed within the supporting sustentacular cells of the  
1180 olfactory epithelium, nor within the Bowman’s glands (326). By migrating within the nerve  
1181 sheath, *B. pseudomallei* would be shielded from the flow of CSF out of the brain. This may  
1182 also explain why *B. pseudomallei* is not often isolated from the CSF in neurological  
1183 melioidosis (56). During our initial studies in mice, we also demonstrated that *B.*  
1184 *pseudomallei* was present in the brain stem, increasing in number to 48 hours post infection,  
1185 indicating, though not definitively demonstrating, that bacteria may enter *via* the trigeminal  
1186 nerve after intranasal inoculation (105). We have now confirmed that *B. pseudomallei*  
1187 penetrates the intact respiratory epithelium in mice and migrates along Schwann cell-encased  
1188 trigeminal nerve bundles to the cranial cavity (326). Figure 4 (B-D) demonstrates *B.*  
1189 *pseudomallei* localized within the branches of the trigeminal nerve. As noted above, clinical  
1190 evidence points to a role for BimA, and hence actin-mediated motility, in neurological  
1191 melioidosis (229); assuming inhalational acquisition to be associated with such cases, it  
1192 remains to be determined where in the olfactory/trigeminal pathways such motility is utilized.

1193 Although there is currently no direct evidence of *B. pseudomallei* penetration of the brain *via*  
1194 the olfactory and/or trigeminal pathways in humans, two important considerations provide  
1195 support for CNS invasion from the nasal mucosa. First, *B. pseudomallei* can colonize the  
1196 nasal cavity in humans (340), and this may represent a potential reservoir of infection. Our

1197 unpublished data also demonstrate that following the inhalation of aerosolized *B.*  
1198 *pseudomallei*, considered a major route of infection in humans, bacteria can be recovered in  
1199 high numbers from the mouse olfactory epithelium. Second, in human cases of primary  
1200 neurological melioidosis, Currie *et al.* reported that bacteremia was only observed in 3 out of  
1201 14 patients (21%) (57). In addition to our animal studies, these data provide evidence that *B.*  
1202 *pseudomallei* can invade the CNS by a non-hematogenous route. Furthermore, the trigeminal  
1203 route of CNS invasion may explain the brainstem-related neurological presentations of  
1204 melioidosis patients, without requiring frank encephalitis.

### 1205 *L. monocytogenes*

1206 In ruminants naturally infected with *L. monocytogenes*, bacteria migrate along several cranial  
1207 nerves (including the trigeminal nerve) to the brain stem and cause encephalitis (341). Similar  
1208 findings are observed in human cases of *L. monocytogenes* brain stem encephalitis, where  
1209 inflammatory lesions may be observed in the nuclei and tracts of the V<sup>th</sup>, VII<sup>th</sup>, IX<sup>th</sup>, X<sup>th</sup> and  
1210 XII<sup>th</sup> cranial nerves innervating the oropharynx (342). Of the cranial nerves, the trigeminal  
1211 nerve pathway is thought to represent a primary route by which *L. monocytogenes* may  
1212 invade the brain stem from the oropharynx (343, 344). It has been proposed that *L.*  
1213 *monocytogenes* may bind to E-cadherin on Schwann cells surrounding trigeminal nerves and  
1214 subsequently invade the trigeminal axons by receptor-independent cell-to-cell spread (345).  
1215 However, binding of *L. monocytogenes* to E-cadherin-expressing Schwann cells has not yet  
1216 been demonstrated.

1217 In animals spontaneously infected and experimentally infected with *L. monocytogenes*,  
1218 bacteria were observed within the axons of the trigeminal nerve, suggesting that these  
1219 bacteria may use intra-axonal spread to migrate to the CNS (343, 344, 346). In a mouse  
1220 model of brain stem encephalitis, *L. monocytogenes* was inoculated into the snout of



1221 immunodeficient mice and was observed within the cytoplasm of trigeminal nerve cell bodies  
1222 at 7 days post-infection (344). Following this, bacteria were then observed within the brain  
1223 stem, but not other regions of the brain. In this model, IFN- $\gamma$  release by natural killer cells  
1224 was required for efficient control of *L. monocytogenes* neuroinvasion *via* the trigeminal route  
1225 (344). *In vivo* and *in vitro* studies have demonstrated that colony-stimulating factor 1-  
1226 dependent cells (including macrophages and dendritic cells) facilitate the neuronal spread of  
1227 *L. monocytogenes* to the CNS (347, 348). Following replication and escape from the  
1228 phagosome in colony-stimulating factor 1-dependent cells, *L. monocytogenes* may use actin-  
1229 based motility to propel itself into adjacent non-phagocytic cells such as axon terminals.  
1230 Indeed, the presence of *L. monocytogenes* with actin tails has been demonstrated *in vivo* in  
1231 the axons of sheep trigeminal nerves (346), and *in vitro* in mouse hippocampal neurons and  
1232 hypothalamic neurons (349). Dramsi *et al.* showed that *L. monocytogenes* spreads from J774  
1233 macrophages to neurons *in vitro*, in a process that was dependent on actin polymerization by  
1234 the bacterium (348), suggesting that the formation of actin tails is critical for neuronal spread  
1235 in listeriosis.

## 1236 **Viruses**

1237 Data from animal studies and human cases have demonstrated that the olfactory and/or  
1238 trigeminal nerve pathway represents a major route of CNS entry for several groups of viruses.  
1239 *In vivo*, herpes viruses demonstrate a tropism for the olfactory epithelium, but not the  
1240 respiratory epithelium (350-353). Expression of the herpes virus receptors heparan sulfate  
1241 and nectin-1 on the apical side of the olfactory epithelium may facilitate binding to the  
1242 neuroepithelium (350, 351). In the respiratory epithelium, these receptors are either expressed  
1243 on the basal side of the epithelium (and are thus inaccessible), or are not highly expressed  
1244 (350, 351). Herpes simplex virus type 1 (354), bovine herpesvirus-5 (355) and equine  
1245 herpesvirus-9 (352, 356) spread from the nasal mucosa to the CNS *via* the olfactory nerves in

1246 animal models of infection. In bovine herpesvirus-5 CNS invasion, the viral protein Us9 and  
1247 the glycine-rich epitope-coding region of the glycoprotein E are required for transport from  
1248 the olfactory sensory neurons to the olfactory bulb (353, 357). In suckling hamsters, equine  
1249 herpesvirus-9 antigen was detected within olfactory sensory neurons 12 hours after intranasal  
1250 infection (352). At 48 hours post-infection, viral antigen was detected within the olfactory  
1251 nerve and olfactory bulb, and at 60 hours post-infection virus was observed within the frontal  
1252 and temporal lobes of the cerebral cortex. Some positive staining occurred within the  
1253 trigeminal nerve, trigeminal ganglia and the region where the trigeminal sensory nerve root  
1254 connects to the brain stem, although this was observed at the later time points, suggesting that  
1255 the olfactory nerve is likely to be the primary route of infection (352). Interestingly,  
1256 Shivkumar *et al.* demonstrated that intranasally delivered herpes simplex virus type 1  
1257 targeted the olfactory epithelium of mice and subsequently travelled along the trigeminal  
1258 nerve branches and re-emerged peripherally within the facial skin after 5 days (350). In this  
1259 model, the virus rarely reached the olfactory bulbs within the brain. However, in another  
1260 study, herpes simplex virus type 1 was isolated from the olfactory bulbs and higher brain  
1261 regions of mice 3 days after intranasal inoculation (354).

1262 In a study of human autopsy material, Harberts *et al.* reported that herpesvirus-6 was detected  
1263 at high frequencies within the olfactory bulb and tract (358). Herpesvirus-6B was the variant  
1264 present in most positive samples, and quantitative real-time TaqMan PCR of selected samples  
1265 detected viral loads of  $1 \times 10^3$  and  $1 \times 10^4$  copies per million cells. In the same study, the  
1266 prevalence of herpesvirus-6 within the nasal mucosa was determined in 3 cohorts of patients:  
1267 (i) healthy controls; (ii) multiple sclerosis patients; and (iii) patients with loss of smell.  
1268 Overall, herpesvirus-6 DNA was detected in 41.3% of nasal mucous samples, and the  
1269 prevalence did not differ between the patient cohorts. These findings suggest that the nasal  
1270 cavity may be a reservoir for herpesvirus-6, and that virus within the nasal cavity may travel

1271 to the olfactory bulbs and tract *via* the olfactory pathway (358). Immunohistological evidence  
1272 from fatal cases of herpes simplex encephalitis demonstrated that herpes simplex virus type 1  
1273 antigen was detected within the olfactory tract, the olfactory cortex, and regions of the limbic  
1274 system that are connected by the olfactory pathway. In contrast, viral antigen was not  
1275 detected within the trigeminal pathway (359). Combined, these studies suggest that the  
1276 olfactory route of CNS entry is highly relevant in human cases of symptomatic and  
1277 asymptomatic herpes virus infection. The role of the trigeminal nerve as a portal of entry for  
1278 herpes viruses in humans is less clear, although the sensory neurons of the trigeminal ganglia  
1279 are the principal site of herpes simplex virus type 1 latent infection in humans (360, 361).

1280 Several subtypes of influenza virus A, including H5N1 (362), WSN/33 (363) and H1N1  
1281 (364), invade the CNS *via* the olfactory nerve pathway in animal models of infection. In  
1282 intranasally infected C57Bl/6 mice, H1N1 (strain PR8) mRNA was detected within the  
1283 olfactory bulb as early as 4 hours post-infection, and viral antigen was visualized within the  
1284 olfactory nerves and the glomerular layer of the olfactory bulb at 15 hours (364). Iwasaki *et*  
1285 *al.* demonstrated that hematogenous dissemination of H5N1 from the lungs to the CNS may  
1286 not be an important entry mechanism, as viral antigen was detected within the brain prior to  
1287 the lungs in a murine model (362). In ferrets intranasally-infected with different H5N1  
1288 strains, 3D imaging demonstrated that brain lesions were distributed: (i) along the olfactory  
1289 pathway; (ii) along the olfactory pathway and within the brain stem; or (iii) surrounding the  
1290 brain vasculature (365). These data suggest that there may be different routes of entry used by  
1291 H5N1 strains; however, the olfactory pathway was identified as the most common route used  
1292 by the small number of strains that were investigated (365). Further studies in ferrets have  
1293 demonstrated that the Vietnam/1203/2004 H5N1 strain spread along the olfactory nerve  
1294 filaments passing through the cribriform plate and into the olfactory bulb (366). From the  
1295 olfactory bulb, the virus migrated through the olfactory tract and into the anterior olfactory

1296 nucleus and anterior commissure, and subsequently to the pyriform lobe, cerebral cortex and  
1297 Ammon's horn (366). Schauwen *et al.* demonstrated that the multibasic cleavage site in the  
1298 haemagglutinin of H5N1 was required for virus spread from the nasal cavity to the olfactory  
1299 bulb and the rest of the CNS (296).

1300 The neurovirulence of influenza A subtypes may be influenced by the ability of the virus to  
1301 disseminate from the olfactory bulb into other regions of the brain (367), which in turn may  
1302 be controlled by the host immune response. *In vivo*, mouse olfactory sensory neurons infected  
1303 with influenza A R404BP virus displayed apoptotic neurodegradation and were subsequently  
1304 phagocytosed by Iba1-expressing microglia/macrophages (368). The infection was therefore  
1305 restricted to the neuroepithelium, and did not spread to the olfactory bulb. This suggested that  
1306 apoptosis of olfactory sensory neurons might be a mechanism by which the host is protected  
1307 from microbial invasion from the nasal cavity (368). Influenza A virus also stimulates a host  
1308 pro-inflammatory cytokine response within the olfactory bulb (369), which may also act to  
1309 protect the host from further CNS invasion (370). Autopsy of a severely  
1310 immunocompromised 11 month old infant revealed influenza virus A antigen within the  
1311 olfactory bulb, olfactory tract and gyrus rectus, which is located inferolaterally to the  
1312 olfactory bulb (371). Viral antigen was not detected within any other regions of the CNS,  
1313 respiratory tract or any other organs. Viral RNA was also not detected within plasma,  
1314 suggesting that viremia was not present. These findings provide evidence for influenza A  
1315 entry into the CNS *via* the olfactory route in a severely immunocompromised infant (371).

1316 Paramyxoviruses including Nipah virus, Hendra virus and parainfluenza virus may enter the  
1317 CNS directly from the nasal mucosa. *In vivo*, the Sendai strain of parainfluenza virus infected  
1318 mouse olfactory sensory neurons, but not sustentacular cells, and travelled to the glomeruli of  
1319 the olfactory bulb (372, 373). Infection of second order neurons and virus spread to the rest of  
1320 the brain did not occur (372-374). The Sendai virus nucleoprotein gene was consistently

1321 detected within the olfactory bulb up to 168 days post-infection, indicating that persistence  
1322 may occur within the olfactory bulb (373). In hamsters, Nipah virus was detected in olfactory  
1323 sensory neurons as they passed through the cribriform plate into the olfactory bulb, providing  
1324 evidence of direct brain infection following intranasal infection (297). Similar results were  
1325 reported in a porcine model of Nipah virus infection, in which Nipah virus antigen was  
1326 detected within a cross section of the olfactory nerve (375). Temporal analysis demonstrated  
1327 that Nipah virus entered the olfactory bulb within 4 days in mice (297), whereas the virus  
1328 spread from the olfactory nerve to the granular cells of the olfactory bulb within 7 days in  
1329 pigs (375). The related Hendra virus was also shown to target the olfactory pathway and  
1330 invade the brain directly from the nasal cavity (in the absence of viremia) in a mouse model  
1331 of encephalitis (376). Weingartl *et al.* demonstrated that Nipah virus antigen was present  
1332 within the endothelial cells and smooth muscle cells of the meningeal veins at days 5-7 days  
1333 post-infection (375), which is consistent with findings from autopsies of fatal human Nipah  
1334 virus cases (377). Thus, it is likely that Nipah virus exploits both the hematogenous and  
1335 olfactory routes of invasion.

1336 Eastern, western and Venezuelan equine encephalitis viruses can cause encephalitis in horses  
1337 and humans, and are transmitted by mosquitos or following aerosol exposure. Using a  
1338 bioluminescent western equine encephalitis virus, Phillips *et al.* demonstrated the progression  
1339 of CNS invasion in intranasally infected mice (295). The bioluminescent signal was initially  
1340 detected in the nasal turbinates and olfactory bulb, and was amplified in the basal nuclei,  
1341 thalamus and hypothalamus. The distribution of lesions within the brain and the detection of  
1342 viral antigen by immunohistochemistry supported the olfactory pathway as the route of  
1343 infection, and suggested that the trigeminal nerve may provide a secondary conduit to the  
1344 brain (295). Venezuelan equine encephalitis virus also targeted both the olfactory (primary  
1345 route) and trigeminal (secondary route) nerve pathways for CNS entry (378), whereas eastern

1346 equine encephalitis virus appeared to infect only the olfactory nerve (379). In CD-1 mice,  
1347 ablation of the olfactory epithelium and the main olfactory bulb prevented invasion of  
1348 Venezuelan equine encephalitis virus into the brain *via* the olfactory nerve; however, the  
1349 virus was still able to spread to the CNS along the trigeminal nerve (378). Interestingly,  
1350 replication of Venezuelan equine encephalitis virus within the nasal mucosa induced the  
1351 expression of pro-inflammatory cytokines, matrix metalloproteinase-9 and intracellular  
1352 adhesion molecule-1 within the olfactory bulb, which led to subsequent breakdown of the  
1353 BBB (380). These events enabled circulating virus to penetrate the brain, suggesting that in  
1354 addition to the olfactory and trigeminal routes of entry, Venezuelan equine encephalitis virus  
1355 may also enter the CNS by a hematogenous route.

1356 Within the *Rhabdoviridae* family, rabies virus and vesicular stomatitis virus within the nasal  
1357 cavity directly invade the olfactory bulbs within the brain. In a fatal human case of airborne  
1358 rabies encephalitis, rabies virions were observed only within the nerve fibers of the olfactory  
1359 bulb, and not any other regions of the brain (381). Data from animal studies have also  
1360 demonstrated that intranasally-delivered rabies virus selectively targets the olfactory  
1361 epithelium and migrates to the olfactory bulb, including the glomeruli, mitral cells and tufted  
1362 cells (382). Rabies virus antigen was also detected in the mouse trigeminal nerve. The  
1363 tropism of rabies virus to the olfactory epithelium may be due to the expression of neural cell  
1364 adhesion molecule by olfactory sensory neurons (383), which was identified as a receptor for  
1365 rabies virus *in vitro* (384). Similar to rabies virus, intranasal vesicular stomatitis virus  
1366 infected the olfactory epithelium, but not the respiratory epithelium in a mouse model (385).  
1367 By 6 hours post-infection viral antigen was observed within the olfactory sensory neurons; at  
1368 12 hours sustentacular cells and basal cells within the neuroepithelium were also infected, as  
1369 were olfactory nerve bundles within the lamina propria; and at 24 hours the Bowman's glands

1370 and olfactory bulb were infected. In contrast to rabies virus, the trigeminal nerve was not  
1371 implicated as a portal of CNS entry for vesicular stomatitis virus (385).

## 1372 **Protozoa**

1373 *Naegleria fowleri* is a free-living amoeba that causes primary amoebic meningoencephalitis,  
1374 a rare but almost always fatal disease in humans. *N. fowleri* is found in warm fresh water and  
1375 human infection occurs following the inspiration of contaminated water, and is usually  
1376 associated with swimming. Contaminated tap water used to reconstitute saline for nasal  
1377 irrigation, or for ablution of the nasal cavity has also been implicated as a source of infection  
1378 (386, 387). Pathological investigations of fatal human cases revealed hemolytic, necrotic  
1379 encephalitis of the olfactory area, the contiguous forebrain and cerebellum (388, 389). The  
1380 suspected route of CNS entry was the olfactory route, due to the presence of amoebae and  
1381 acute inflammation within the nasal mucosa and the olfactory nerve bundles (388, 389).

1382 In murine models of the early stages of primary amoebic meningoencephalitis, intranasal *N.*  
1383 *fowleri* invaded the cribriform plate and olfactory bulb and was observed within the olfactory  
1384 nerves (390, 391). *In vivo* studies have shown that *N. fowleri* induces mucous production  
1385 within the nasal cavity as early as 1 hour post-infection, and that by 6 hours the amoebae  
1386 were covered by mucous and surrounded by neutrophils (392). In the mucin-producing cell  
1387 line, NCI-H292, *N. fowleri* induced the release of reactive oxygen species, which led to the  
1388 activation of the epidermal growth factor receptor, which in turn stimulated production of the  
1389 mucin MUC5A and IL-8 secretion (314). Despite this host response, by 12 hours *N. fowleri*  
1390 was observed adhering to and invading the olfactory epithelium in mice, suggesting that the  
1391 amoebae may have pathogenic mechanisms to efficiently penetrate the mucous layer (392).  
1392 *In vitro*, it was shown that both live trophozoites and crude total *N. fowleri* extracts  
1393 demonstrated mucinolytic activity. A 37 kDa cysteine protease of *N. fowleri* was

1394 subsequently identified as being responsible for this mucinolytic activity (392); this most  
1395 likely enabled the amoebae to penetrate the nasal mucosa and invade the neuroepithelium.  
1396 Penetration of the olfactory epithelium in mice occurred without cellular disruption or  
1397 damage (390), and microscopy studies have shown that *N. fowleri* may be ingested by  
1398 sustentacular cells within the epithelium, or may migrate between the sustentacular cells  
1399 (393). A recent study demonstrated that *N. fowleri* degraded the epithelial tight junction  
1400 proteins claudin-1 and zonula occludens-1 *in vitro*, highlighting a potential paracellular  
1401 mechanism of penetration (394).

1402 *Balamuthia mandrillaris*, an opportunistic free-living amoeba that can cause granulomatous  
1403 amoebic encephalitis, was also shown to enter the CNS directly from the nasal cavity after  
1404 penetrating the olfactory epithelium and cribriform plate in immunodeficient mice following  
1405 intranasal infection (395). However, the implications of these findings in humans are unclear  
1406 as *B. mandrillaris* is primarily thought to cause CNS infections due to hematogenous spread  
1407 from the lungs, or through cuts or skin abrasions (396).

#### 1408 **Yeasts and fungi**

1409 The encapsulated yeast *Cryptococcus neoformans* is an important cause of fungal  
1410 meningoencephalitis worldwide and can enter the CNS by penetrating the BBB using  
1411 transcellular, paracellular and Trojan horse mechanisms, following blood-borne  
1412 dissemination from the lungs (as reviewed elsewhere (397)). Although the hematogenous  
1413 route of CNS is well accepted for cryptococcal meningoencephalitis, some strains of *C.*  
1414 *neoformans* isolated from the CSF of human patients were shown to be rhinotropic in mice  
1415 (398, 399), and persistent *C. neoformans* colonization of the nasal cavity (90 days) was  
1416 reported in one study (400). These findings prompted investigations into the possibility of an  
1417 alternative direct route of CNS entry from the nose. Gomes *et al.* followed the progression of



1418 intranasal *C. neoformans* infection in immunodeficient mice, and demonstrated colonization  
1419 of the olfactory mucosa, invasion along the olfactory nerve and meningeal involvement  
1420 (401). In a study of the pathological features of three patients with acquired immune  
1421 deficiency syndrome and cryptococcal meningitis, *C. neoformans* was observed in the  
1422 subarachnoid space around the olfactory tracts and bulbs and in the olfactory nerve fascicles  
1423 (402). No cryptococci were located within the lamina propria or the olfactory epithelium, and  
1424 thus the authors suggested that *C. neoformans* within the CSF may have been transported to  
1425 the olfactory regions due to the drainage of CSF from the subarachnoid space through the  
1426 cribriform plate (402). Using a guinea pig model, Lima and Vital provided further evidence  
1427 that the olfactory route is unlikely to represent a portal of *C. neoformans* CNS entry, as  
1428 intranasally instilled cryptococci were unable to penetrate the olfactory epithelium and were  
1429 cleared from the olfactory mucosa (316).

1430 In contrast, the fungal infection rhinocerebral mucormycosis has been demonstrated to spread  
1431 to the brain *via* the trigeminal nerve. Rhinocerebral mucormycosis refers to infection caused  
1432 by fungi within the order Mucorales, and usually affects individuals with poorly controlled  
1433 diabetes mellitus or the immunocompromised. These organisms display a predilection for the  
1434 nasal cavity and paranasal sinuses; from these sites the organisms typically invade blood  
1435 vessel walls and then spread to the cavernous sinus, internal carotid artery and the brain (403-  
1436 406). However, fungal hyphae and lesions have been demonstrated within the trigeminal  
1437 nerve and the pons within the brain stem in the absence of leptomeningitis, suggesting that  
1438 direct invasion occurred from the sinuses to the brain along the trigeminal nerve (407, 408).  
1439 The perineural route of CNS entry was thought to be atypical; however, a study of the  
1440 histologic features of patients with mucormycosis demonstrated that perineural invasion,  
1441 characterized by fungal hyphae within the perineurium that surrounds the nerves, was a

1442 common feature that occurred concurrently with angioinvasion (409). The mechanisms of  
1443 rhinocerebral mucormycosis CNS infection have not been investigated.

#### 1444 **CONCLUDING REMARKS**

1445 A wide range of microbes can invade the CNS, and any organisms that can enter the CSF  
1446 have the potential to cause meningitis. Many bacteria and other pathogens have developed  
1447 sophisticated mechanisms to penetrate the BBB and/or BCSFB by transcellular and  
1448 paracellular transport, or the Trojan horse route. The olfactory and trigeminal nerves  
1449 represent pathways by which microbes enter the brain, without encountering the BBB or the  
1450 BCSFB. These nerves are well recognized as portals of entry for many viruses, as well as  
1451 protozoa and fungi, and there is now evidence from animal models that some bacteria can  
1452 infect the brain *via* the olfactory and trigeminal nerves. The purpose of this review is to  
1453 highlight an alternative route of entry that has thus far received little attention and may  
1454 explain some of the pathological features observed in human disease. These routes of  
1455 bacterial invasion require investigation in humans, especially in cases where the etiological  
1456 agent is known to colonize the nasal mucosa. In such cases, clinical assessment of olfactory  
1457 function and the nasal mucosa should be considered. Furthermore, several questions remain  
1458 unanswered; first, what is the route for bacteria to travel along olfactory nerves – within the  
1459 axons/nerve bundles, within the glia or within the perineural space? We have demonstrated  
1460 that *B. pseudomallei* caused destruction of the olfactory epithelium and axonal death, which  
1461 provided an open conduit for migration to the olfactory bulb within the nerve fascicles (326).  
1462 It is unknown if similar mechanisms of transport are used by other bacterial pathogens such  
1463 as *S. pneumoniae* or *N. meningitidis*. Remarkably, *L. monocytogenes* appears to travel intra-  
1464 axonally. Second, which bacterial virulence factors are required for penetration of the  
1465 olfactory epithelium and invasion of the brain *via* the olfactory and trigeminal nerves? This  
1466 review has highlighted the requirement for additional research to characterize the role of the

1467 olfactory and trigeminal nerves in bacterial penetration of the brain; and determine the  
1468 molecular and cellular mechanisms by which bacterial pathogens may exploit these  
1469 pathways.

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2832 of Technology, Australia, where she studied the pathogenic mechanisms of the human  
2833 *Ureaplasma* species during intrauterine infection. She was subsequently appointed as a  
2834 postdoctoral research fellow at the Institute for Glycomics, Griffith University, and undertook  
2835 research to investigate the virulence of *Burkholderia pseudomallei* in a murine model of  
2836 acute melioidosis. She has recently accepted a postdoctoral research position at Monash  
2837 University, to research dendritic cell biology in the central nervous system during  
2838 autoimmune disease.

2839 **Alan Mackay-Sim**

2840 Professor Alan Mackay-Sim obtained his PhD in 1980 at Macquarie University, Australia. He  
2841 undertook a postdoctoral position at the University of Pennsylvania and had faculty positions  
2842 at the University of Sydney, the University of Wyoming and the University of Adelaide  
2843 before moving to Griffith University in 1987 where he has held positions as Deputy Director  
2844 of the Eskitis Institute for Cell and Molecular Therapies and Director of the National Centre  
2845 for Adult Stem Cell Research. He is currently Professor of Neuroscience, Griffith University.  
2846 With over 30 years of research in the field, Professor Mackay-Sim is a leading expert on the  
2847 human sense of smell and the biology and development of the olfactory mucosa. In the last  
2848 16 years he has concentrated on the clinical applications of olfactory cells and their use for  
2849 neural regeneration therapies, as well as the involvement of the olfactory nerve pathway in  
2850 the development of disease.

2851 **Robert Norton**

2852 Dr Robert Norton graduated in Medicine in 1980 and has worked in a variety of clinical  
2853 positions including five years in Australian indigenous communities. He trained in  
2854 Microbiology at the Institute of Medical and Veterinary Science in Adelaide between 1991  
2855 and 1995. He gained an MD from the University of Adelaide in 1998. In his current capacity  
2856 as Director of Microbiology at Townsville Hospital, Queensland, he has collaborated with  
2857 researchers locally and nationally on projects relating to melioidosis, rheumatic fever,  
2858 invasive group A streptococcal disease and Q-fever. Dr Norton is part of the Infectious  
2859 Diseases and Immunopathogenesis Research Group which includes clinicians and academic  
2860 staff of James Cook University. He is an Associate Professor at James Cook University and is  
2861 the Chief Examiner in Microbiology for the Royal College of Pathologists in Australasia. He  
2862 has over 100 peer-reviewed publications and has been successful in obtaining local and  
2863 national collaborative grants.

2864 **Bart J. Currie**

2865 Bart Currie is Head of Infectious Diseases at Royal Darwin Hospital and Professor in  
2866 Medicine at the Northern Territory Medical Program, Flinders and Charles Darwin  
2867 Universities. He is also Program Leader for Tropical and Emerging Infectious Diseases in the  
2868 Global and Tropical Health Division of the Menzies School of Health Research and Director  
2869 of RHD Australia. Areas of interest include clinical and epidemiological aspects of tropical  
2870 and emerging infections, development of treatment guidelines and clinical toxinology. He  
2871 initiated the Darwin Prospective Melioidosis Study in 1989 and this remains the basis for  
2872 ongoing multi-disciplinary collaborations on melioidosis.

2873 **James A. St John**

2874 Dr James St John obtained his PhD in Agricultural Science at the University of Melbourne,  
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2876 Melbourne and The University of Queensland. He is currently a Senior Research Fellow at  
2877 the Eskitis Institute for Drug Discovery, Griffith University. Since his PhD he has worked in  
2878 the field of mammalian olfactory nervous system development and regeneration and has  
2879 published over 40 papers on olfactory axon guidance and the role of olfactory glia. He  
2880 performs detailed microscopic anatomical studies of the olfactory system and has identified  
2881 subpopulations of olfactory glia using live cell imaging of in vitro cultures. He is particularly  
2882 interested in the role of olfactory glia in protecting the brain from bacterial infections.  
2883 Together with Dr Jenny Ekberg, he recently performed the majority of the work which  
2884 identified the intranasal route of infection via the olfactory nerve by *Burkholderia*  
2885 *pseudomallei*.

2886 **Jenny A. K. Ekberg**

2887 Dr Jenny Ekberg obtained her PhD in neuroscience/biomedical science from The University  
2888 of Queensland in 2005. During her PhD and first postdoctoral appointment, also at UQ, she  
2889 studied novel modulatory mechanisms of voltage-gated ion channels in neurons using  
2890 electrophysiological techniques. In 2008, she moved to Griffith University as a Research  
2891 Fellow, now focussing on neuron-glia interactions and neural regeneration in the olfactory  
2892 nervous system. In 2012, she took up a position as Senior Lecturer at Queensland University  
2893 of Technology. She continues her work on neural repair, and has expanded into the field of  
2894 bacterial infections of the central nervous system. Together with Dr St John and Prof.  
2895 Beacham, she recently investigated how *B. pseudomallei* can invade the brain via the  
2896 olfactory nerve. The focus of Dr. Ekberg's research is to investigate how cells in the olfactory  
2897 nerve prevent microorganisms from entering the brain, whilst simultaneously promoting  
2898 nerve regeneration.

2899 **Michael Batzloff**

2900 Dr Michael Batzloff completed his PhD from Griffith University in 2001 after which he  
2901 accepted a postdoctoral position at the Queensland Institute of Medical Research, Australia.  
2902 He has been a recipient of two fellowships from the National Heart Foundation of Australia  
2903 for his research into vaccine development for *Streptococcus pyogenes*. He was subsequently  
2904 appointed the inaugural head of the Bacterial Vaccines Laboratory at the Queensland Institute  
2905 of Medical Research and recently accepted a position at the Institute for Glycomics at Griffith  
2906 University. His research interests include neglected tropical diseases focusing on  
2907 pathogenesis and vaccine development for the bacterial pathogens *Streptococcus pyogenes*  
2908 and *Burkholderia pseudomallei*.

2909 **Glen C. Ulett**

2910 Associate Professor Glen C. Ulett received his PhD from James Cook University (Australia)  
2911 in 2001, and completed post-doctoral research at St Jude Children's Research Hospital, the  
2912 University of Queensland, and the University of Alabama at Birmingham. He is currently an  
2913 Australian Research Council (ARC) Future Fellow in Microbiology at Griffith University  
2914 where he leads a research team studying bacterial pathogenesis and mechanisms of host  
2915 defence against infection. His laboratory focuses on infections related to the urogenital tract,  
2916 programmed cell death, and mechanisms of virulence and disease associated with  
2917 *Escherichia coli*, *Streptococcus agalactiae* and *Burkholderia pseudomallei*. He has been a  
2918 microbiology researcher in the field of bacterial pathogenesis for seventeen years and is a  
2919 two-time-recipient of the George McCracken Infectious Disease Fellow Award from the  
2920 American Society for Microbiology. His research program in microbiology and infectious  
2921 diseases is supported by funding from the National Health and Medical Research Council of  
2922 Australia.

2923 **Ifor R. Beacham**

2924 Emeritus Professor Ifor Beacham undertook undergraduate studies in biochemistry at the  
2925 University of Otago, New Zealand, and obtained his PhD in microbial genetics from the  
2926 University of Leicester, UK. He has undertaken postdoctoral research at the University of  
2927 Leicester, Washington University, St Louis, and the University of California, Santa Barbara,  
2928 and has held teaching positions at the University of Wales, Aberystwyth, UK, and Griffith  
2929 University, Australia. He is currently a Research Leader at the Institute for Glycomics,  
2930 Griffith University. He has worked in the general area of molecular microbiology on a variety  
2931 of bacteria before undertaking work with *Burkholderia pseudomallei* 16 years ago. These  
2932 latter studies were motivated by the endemicity of melioidosis in northern Australia,  
2933 difficulties with genetic manipulation and the enigmatic status of *B. pseudomallei* as a  
2934 pathogen. He hopes his continuing work will contribute to a greater understanding of the  
2935 molecular nature of virulence of *B. pseudomallei*.



Bacterial pathogen	Ligand	HBMEC receptor	Function	References
<i>Escherichia coli</i> (and other Gram negative enterics)	FimH	CD48	Adhesion to HBMECs; FimH also promotes invasion by activation of RhoA, which induces host cytoskeletal rearrangements	(154, 410)
	Nlp1	Unknown	Adhesion to and invasion of HBMECs; Nlp1 activates host cytosolic phospholipase A <sub>2</sub>	(174)
	OmpA	Gp96 homologue	Loops 1, 2 and 3 of OmpA function in adhesion to, and invasion of, HBMECs; ompA-mediated invasion is due to Rac1 activation and activation of host cytosolic phospholipase A <sub>2</sub>	(161, 162, 173, 178)
	CNF-1	37/67 kDa laminin receptor	Invasion of HBMECs; CNF-1 activates RhoA and recruits focal adhesin kinase to the site of bacterial entry	(149, 150, 171, 172)
	IbeA	Vimentin, PSF	Invasion of HBMECs; activates Rac1	(178, 411-413)
	Ibe10	45 kDa Ibe10 receptor	Invasion of HBMECs	(414, 415)
	AslA	Unknown	Invasion of HBMECs	(416)
	TraJ	Unknown	Invasion of HBMECs	(417)
	<i>yijP</i>	Unknown	Invasion of HBMECs	(418)
	Flagella	Unknown	Adhesion to and invasion of HBMECs	(419)
	<i>Haemophilus influenzae</i>	Phosphorylcholine	PAF receptor	Potentially involved in invasion of HBMECs
OmpP2		37/67 kDa laminin receptor	Adhesion to HBMECs	(151, 152)
<i>Listeria monocytogenes</i>	InlA	E-cadherin	Invasion of polarized human choroid plexus epithelial cells	(232)
	InlB	gC1q-R, met receptor tyrosine kinase	Invasion of HBMECs and	(213, 232,

			polarized human choroid plexus epithelial cells	421, 422)
	Vip	Gp96	Invasion of the brain <i>in vivo</i>	(160)
	IspC	Unknown	Adhesion to and invasion of sheep choroid plexus epithelial cells, but not HBMECs	(233)
<i>Mycobacterium tuberculosis</i>	<i>Rv0980c, Rv0987, Rv0989c, Rv1801</i>	Unknown	Invasion of HBMECs	(118)
	HBHA	Heparan sulfate glycosaminoglycans	Adhesion to and invasion of BBMECs	(234)
	<i>Rv0311, Rv0805, Rv0931c, Rv0986, MT3280</i>	Unknown	Invasion of HBMECs	(235)
	PknD	Laminin	Attachment to, invasion of and survival within HBMECs	(158)
<i>Neisseria meningitidis</i>	Type IV pili	Unknown	Adhesion to HBMECs; type IV pilus-mediated adhesion leads to the formation of cortical plaques and host membrane protrusions	(122, 124)
	PilQ and PorA	37/67 kDa laminin receptor	Adhesion to HBMECs	(151, 152)
	Opc	$\alpha_5\beta_1$ integrin (mediated by binding to the bridging molecules vitronectin and fibronectin)	Adhesion to and invasion of HBMECs; activates JNK1, JNK2 and p38, and protein tyrosine kinase signalling	(155, 191, 192)
<i>Staphylococcus aureus</i>	<i>ypfP</i> and LTA	Unknown	Invasion of HBMECs	(423)
<i>Streptococcus agalactiae</i>	Lmb	Laminin	Invasion of HBMECs	(159)
	<i>iagA</i> and LTA	Unknown	Invasion of HBMECs; <i>iagA</i> anchors LTA to the streptococcal cell wall and is required for invasion	(424)
	FbsA	Fibrinogen	Adhesion to HBMECs	(156)
	SfbA	Integrins (via <i>S. agalactiae</i> binding to	Invasion of HBMECs	(199)

		immobilised fibronectin)		
	PilA	$\alpha_2\beta_1$ integrin (mediated by <i>S. agalactiae</i> binding to immobilised collagen)	Adhesion to and invasion of HBMECs; pilA binding stimulates focal adhesion kinase phosphorylation and downstream phosphatidylinositol-3 kinase activation	(109, 425)
	PilB	Unknown	Invasion of HBMECs	(425)
	Srr-1	Fibrinogen (srr-1 binds amino acids 283 – 410 of the fibrinogen $\alpha$ chain via a dock, lock and latch mechanism)	Adhesion to and invasion of HBMECs	(200, 201, 426)
	HvgA	Unknown	Adhesion to HBMECs	(198)
	ACP	Glycosaminoglycans	Adhesion to and invasion of HBMECs; invasion potentially occurs following Rho GTPase-dependent actin rearrangements	(157, 427)
<i>Streptococcus pneumoniae</i>	CbpA	37/67 kDa laminin receptor	Adhesion to HBMECs	(151)
	Phosphorylcholine	PAF receptor	Invasion of HBMECs; binding to platelet-activating factor receptor induces the co-localisation of $\beta$ -arrestin	(193, 197)
	NanA	Unknown	Adhesion to and invasion of HBMECs	(194, 195)

**Table 1:** Known bacterial ligands and their host receptors for adhesion to and invasion of the blood-brain barrier

ACP = alpha C protein; AsIA = aryl-sulfatesulfohydrolase; BBMEC = bovine brain microvascular endothelial cells; CbpA = choline binding protein A; CNF-1 = cytotoxic necrotizing factor-1; FbsA = fibrinogen-binding protein; FimH = type 1 fimbrial adhesin; HBHA = heparin-

binding haemagglutinin adhesin; HBMECs = human brain microvascular endothelial cells; HvgA = hypervirulent GBS adhesion; IbeA = invasion of brain endothelial cell protein A; iagA = invasion associated gene; InlA = internalin A; InlB = internalin B; LTA = lipoteichoic acid; Lmb = laminin-binding protein; NanA = neuraminidase A; Nlp1 = new lipoprotein 1; OmpP2 = outer membrane protein P2; Opc = outer membrane protein C; PAF = platelet activating factor; PilQ = pilus secretin protein; PorA = major outer membrane protein; PSF = polypyrimidine-tract-binding protein (PTB)-associated splicing factor; SfbA = streptococcal fibrinogen binding protein A; Srr-1 = serine-rich repeat 1; Vip = virulence protein.

## Figure legends

**Figure 1:** Anatomical location and organization of the blood-brain and blood-cerebrospinal fluid barriers, and the olfactory portal.

The skull and the meninges, which consist of the dura mater, arachnoid mater and pia mater, protect the brain. The arachnoid mater and pia mater are connected by strands of connective tissue called arachnoid trabeculae, which course through the CSF-containing subarachnoid space. At the cellular level, the BBB and BCSFB maintain homeostasis of the brain. The BBB is formed by tight junctions between endothelial cells lining cerebral microvessels, in addition to pericytes and astrocytes. The BCSFB is formed by tight junctions between: (i) epithelial cells at the choroid plexus, (ii) endothelial cells of the veins and venules within the subarachnoid space, and (iii) the epithelial cells of the arachnoid mater. The olfactory system bypasses the cellular barriers of the CNS and provides a direct portal from the nasal cavity to the olfactory bulb within the brain.

**Figure 2:** Mechanisms of blood-cerebrospinal fluid penetration by bacterial pathogens.

Bacteria (purple) may invade the CNS *via* the BCSFB by (A) transcellular penetration involving either pinocytosis or receptor-mediated mechanisms; (B) by paracellular entry following the disruption of junctions (comprising tight junctions and adherens junctions) between choroidal epithelial cells, endothelial cells of veins/venules within the subarachnoid space, or the cells of the arachnoid membrane; and (C) by the “Trojan horse” mechanism, where microbes may transmigrate with infected leukocytes (such as macrophages, as shown).

**Figure 3:** The olfactory system is a direct portal for bacterial pathogens to the brain.

(A) The cilia of olfactory sensory neurons penetrate the nasal mucosa and provide a direct pathway from the external environment to the CNS. Olfactory sensory neurons in the

olfactory epithelium are supported by sustentacular cells and replaced by proliferation and differentiation of basal stem cells, and their axons pass through the lamina propria and cribriform plate of the skull to synapse with mitral cells in the glomeruli of the olfactory bulb. Microbial pathogens can potentially access the brain through the olfactory epithelium *via* axonal transport, by travel within olfactory ensheathing cells that surround the axons, or external to these cells within the perineural space and passage through holes in the cribriform plate to access the subarachnoid space. Adapted from (101, 428). (B-C) A coronal section through the olfactory system of transgenic reporter mice OMP-ZsGreen x S100 $\beta$ -DsRed (429, 430). (B) Primary olfactory neurons (green) reside in the olfactory epithelium, which lines the nasal cavity (NC); septum is indicated as S. The neurons project axons to the olfactory bulb (OB) within the CNS. Boxed region is shown in C. (C) Bundles of olfactory axons project from the olfactory epithelium (OE) through the cribriform plate (CP; chondrocytes are bright red) and enter the nerve fiber layer (NFL), which forms the outer layer of the olfactory bulb. Olfactory ensheathing cells (dull red, arrows) surround the axon bundles. (D) A sagittal section through the olfactory bulb and nasal cavity of a mouse that was intranasally inoculated with *B. pseudomallei* (green) with invasion of the nerve fiber layer of the olfactory bulb in the region indicated by the arrow. (E) A higher power view of the ventral nerve fiber layer shows bacterial infestation within the nasal cavity (NC), the olfactory epithelium (OE) and invasion of the NFL by the bacteria. Inset box shows higher power view of *B. pseudomallei* (green) within the nerve fiber layer. Scale bar is 500  $\mu$ m in B; 65  $\mu$ m in C; 750  $\mu$ m in D; 300  $\mu$ m in E; 300  $\mu$ m in E; 70  $\mu$ m in inset in E.

**Figure 4:** The trigeminal nerve route of entry.

(A) Schematic showing the three branches of the trigeminal nerve, V1, V2 and V3. The branches V1 and V2 innervate the nasal cavity and project to the brain stem (BS). (B-D) In coronal sections of the mouse nasal cavity, *B. pseudomallei* (green) is localized within

branches of the trigeminal nerve that innervates the septum (S) within the nasal cavity (NC). Boxed area in B is shown in C. The bacteria (arrows) are contained within the trigeminal nerve while adjacent olfactory nerve (arrows with tail) bundles do not contain bacteria. (D) A higher magnification view of the trigeminal nerve in a nearby section. Scale bar is 650  $\mu\text{m}$  in B; 40  $\mu\text{m}$  in C; 15  $\mu\text{m}$  in D. OE = olfactory epithelium.