Group A *Streptococcus* strains causing meningitis without distinct invasive phenotype

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Abstract

Group A streptococcal (GAS, aka *Streptococcus pyogenes*) meningitis is a fulminant

disease associated with high morbidity and mortality. To elucidate the mechanisms

underlying the invasiveness of GAS in meningitis, we compared GAS isolates derived from

five cases of meningitis, to otitis and colonizing isolates. We did not observe differences in

adherence to and invasion of human brain microvascular endothelial cells, virulence factors

activity or barrier disruption. Whole genome sequencing did not reveal particular

invasiveness traits. Most patients previously suffered from otitis media suggesting that

meningitis likely resulted from a continuous spread of the infection rather than being

attributable to changes in pathogen's virulence.

Introduction

Streptococcus pyogenes or group A *Streptococcus* (GAS) asymptomatically colonizes the human throat and skin. In addition to asymptomatic colonization, GAS causes mild to life-threatening infections, such as necrotizing fasciitis and meningitis. GAS meningitis cases are rare and represent only 1-4% of all invasive GAS cases detected across Europe and the United States [1-3]. GAS meningitis is a fulminant disease associated with high morbidity, including neurologic sequelae, and mortality. It occurs mainly in patients with predisposing factors such as otitis, sinusitis, recent head injury or neurosurgery, presence of a neurosurgical device, cerebrospinal fluid leak, or alcoholism [4, 5]. This suggests extension *per continuitatem* from a neighboring infection such as otitis media, resulting in otogenic meningitis. However, in a third of cases there are no predisposing factors present. The prevalence of the predisposing factors for GAS meningitis in children are acute otitis media and acute mastoiditis (2-3% [6, 7] and 11% [8], respectively). GAS is also responsible for 41% of acute mastoiditis and 14% of acute otitis cases that require hospitalization in children [9, 10] and for 19% of acute otitis and mastoiditis cases requiring hospitalization in adults [11]. The most common host-associated risk factors for invasive GAS infections in adults are crowding, diabetes, cardiac disease, cancer and corticosteroid use [12]. Other risk factors are breaching of the physical barriers skin and mucosa by trauma or preceding viral infections such as varicella [13] and influenza [14, 15].

On the pathogen's side, GAS components such as the streptococcal virulence factors Streptolysin O (SLO), streptococcal DNases, the IL-8 protease SpyCEP and the M-protein, play key roles in the invasiveness and ability of the pathogen to successfully invade and survive within the host environment [16]. SLO is a pore forming toxin that can lead to inhibition of phagocytosis and cell lysis of target cells [17], the DNase sda1 was shown to degrade

neutrophils extracellular traps and to prevent TLR9-dependent recognition, leading to

enhanced pathogen spread in the host [18-20], while SpyCEP is involved in degradation of

the cytokine IL-8, impairing neutrophils recruitment to the site of infection [21].

The M-protein is a GAS surface protein promoting GAS survival in the host by increasing

adherence to host cells [22], escape from opsonophagocytosis [23] and neutrophil

activation, leading to inflammation [24]. To date, more than 200 different M-protein types

(*emm*-types) have been described, based on the sequence of the variable region of the M-

protein. Genotyping methods used for the classification of GAS strains include serotyping of

the M-protein as well as Multilocus Sequence Typing (MLST), based on the sequences of

seven housekeeping genes [25]. MLST is normally used to study the epidemiology and

population genetics of a specific set of isolates, proving particularly useful for the detection

and classification of outbreaks. Strains of *emm* type 1 (*emm*-1) and MLST-28 (ST28) or emm-

28 ST52 are highly prevalent among GAS meningitis isolates [26].

The aim of this study focused on assessing whether specific pathogen traits distinguish

GAS meningitis isolates from colonizing GAS isolates or GAS causing otitis and whether these

specific traits could be associated with disease severity. We investigated five patients

suffering from GAS meningitis, analyzing both host and pathogen characteristics. On the host

side, various clinical parameters were analyzed. On the pathogen side, we phenotypically

and genetically compared five GAS isolates derived from the abovementioned meningitis

cases with ten GAS isolates derived either from otitis patients or asymptomatic colonization.

To our knowledge, this is the first study comparing the virulence traits of meningitis isolates

with otitis or colonizing ones.

Material and methods

Patients and ethics

Five patients diagnosed between 2013 and 2017 with clinically confirmed community-

acquired GAS meningitis in Switzerland were retrospectively included in this study (Tab.1).

No specific surveillance program for GAS meningitis exists in Switzerland and the cases were

selected using a convenience sampling strategy. Otitis isolates were also selected using a

convenience sampling strategy while GAS colonizing isolates were selected to reflect the

population present in the meningitis and otitis groups (*emm*-1 and *emm*-28 types). This

study was approved by the regional committee for medical research ethics (BASEC-ID 2016-

00145 and 2017-02225).

Bacterial strains and cell lines

All meningitis, otitis and colonizing GAS clinical isolates are listed in Tab.S1 and were

cultured in THY [Todd-Hewitt broth (BD) plus 2% yeast extract (Oxoid)] at 37°C. Immortalized

human brain microvascular endothelial cells (HBMECs) [27] were cultured in Roswell Park

Memorial Institute (RPMI) medium supplemented with 10% FBS and L-glutamine, unless

specified otherwise.

Adherence and invasion assays

Adherence and invasion assays were performed as previously described, with minor

modifications [28]. HBMECs were seeded in 24 well plates to $2.2*10^4$ cells/well in RPMI

medium supplemented with 10% FBS and L-glutamine and incubated at 37ºC in the presence

of 5% CO_2 for 48h. Overnight (ON) cultures of the GAS strains were grown at 37ºC in a static

incubator in 5ml THY. Bacterial cultures were diluted 1:10 in fresh THY medium and grown

to an optical density at 600nm (OD_{600}) of 0.4, washed once with PBS and resuspended in

infection medium (RPMI+L-glutamine+0.4% bovine serum albumin). Prior to infection,

HBMECs were washed once with PBS and bacteria, resuspended in 500 ul of infection

medium, were added to reach a multiplicity of infection (MOI) of 1 or 10 for adherence and

invasion assays respectively. The plates were spun down at 1200g for 5 minutes to and

subsequently incubated 37ºC in the presence of 5% CO_2. To assess adherence, infected

HBMECs were incubated for 30 minutes, washed six times with PBS to remove unbound

bacteria and subsequently lysed using 500 ul of deionized sterile water (dH2O). To assess

invasion, infected HBMECs were incubated for 2 hours and washed three times with PBS to

remove excess bacteria. 500 ul of infection medium supplemented with penicillin and

gentamycin, at a final concentration of 10 ug/ml and 100 ug/ml respectively, were added to

each well to kill extracellular bacteria. The plates were then incubated for 2 hours at 37ºC in

the presence of 5% CO_2 and the cells were subsequently lysed with 500 ul of dH2O after

three washes with PBS. Ten-fold serial dilution of the cell lysates were plated on THY plates

to enumerate bacterial colonies after ON incubation at 37ºC. The percentage of adherence

and invasion were calculated relatively to the initial inoculum.

Viability assay

Cell viability was assessed as follows. HBMECs and bacteria were prepared for adherence

and invasion assays, as described above. After 30 minutes infection followed by six washes

with PBS or after the two hours incubation with antibiotics followed by three washes with

PBS, HBMECs were harvested using 100 µl of trypsin (Gibco), washed and resuspended in

500 µl PBS. Three wells/condition were pooled and the experiment was carried out in

duplicate.

250 µL of the cell suspension were transferred to the wells of a conical 96well plate, spun

down at 470g for 5min, washed with Annexin V binding buffer (ThermoFisher) and spun

down at 470g for 5min. Dead cells were stained using 40 µl/well of a cocktail of Annexin V –

FITC (#640906; Biolegend, 1:50 dilution) and 7 AAD (#420404; Biolegend 1:25 dilution)

diluted in Annexin V binding buffer, for the detection of apoptotic and late apoptotic cells.

The cells were incubated for 20min at room temperature, 160 µL of Annexin V binding buffer

were added and data were acquired using the AttuneNxT flow cytometer (ThermoFisher).

Virulence factors activity

Activity of the virulence factors SLO, streptococcal DNases and SpyCEP was assessed in

the supernatants of exponentially growing GAS as previously described [29]. ON cultures of

the various strains were statically grown at 37ºC in THY medium. Bacterial cultures were

diluted 1:10 in fresh THY medium and grown to an optical density at 600nm (OD_{600}) of OD

0.4. After centrifugation, the supernatants were removed and filter sterilized using 0.22 µm

filtration membranes.

SLO activity was assessed as follows. 4 mM DTT and 0.0004% Trypan-blue were added to

the supernatants, the samples were incubated for ten minutes at room temperature (RT)

and two-fold dilutions were made in PBS in a 96 well plate. PBS and dH_2O alone were used

respectively as negative and positive controls. Erythrocytes were diluted to 2% v/v in PBS

and added to the diluted supernatants (1 part erythrocytes, 4 parts supernatant) or PBS or

water for no lysis and full lysis controls, respectively. Samples were incubated at 37ºC for 30

minutes and spun down at 3000g for five minutes. The OD_{541} of the supernatants, directly

proportional to the activity of SLO, was used as a readout.

Streptococcal DNase activity was assessed as follows. 5 ul of bacterial supernatant or 5 ul

of THY (negative control) were mixed with 5 ul of 10X DNase buffer (50 mM CaCl2, 500 mM

Tris, pH 7.9) and 5ul of calf thymus DNA (1mg/ml, Sigma) in a total volume of 50 ul and

incubated at 37ºC for five minutes. The reaction was stopped by adding 8 ul of 0.5M EDTA

and the samples were loaded on a 0.8% agarose gel to assess DNA degradation.

Streptococcal DNase activity was evaluated with a score corresponding to no activity (0=no

degradation of calf thymus DNA), intermediate activity (1=partial degradation of calf thymus

DNA) and high activity (2=complete degradation of calf thymus DNA).

SpyCEP activity was assessed by measuring the degradation of the cytokine IL-8. Bacterial

supernatants were incubated with 1ng/ul IL-8 for 16 hours at 37ºC after which an IL-8 ELISA

was carried out according to the manufacturer's instruction (R&D). THY was used as a

negative control. SpyCEP activity is depicted as the percentage of IL-8 cleavage compared to

the negative control (medium only).

Electric cell-substrate impedance sensing (ECIS®) assay

ECIS® was performed with the ECIS® Z-Theta instrument (Applied Biophysics) to monitor

the effects of the GAS strains on human brain microvascular endothelial cells (HBMECs) [27]

barrier integrity. Before seeding, the wells of the arrays (ECIS 8W10E+, IBIDI) were washed

twice with sterile, nuclease-free dH_2O, coated with 5 µg/ml collagen IV for 1h at room

temperature (RT), washed three times with dH_2O and incubated with 10 mM L-cysteine for

2h at RT. The wells were washed two times with dH_2O before 200 µl of culture medium were

added and the assay started (37ºC, 5% CO2). After 30 min, 200µl culture medium containing

10^5 HBMECs were seeded per well. After 22 h incubation 50 µl of medium were removed

and 2h later HBMECs were infected with log phase GAS strains resuspended in

DMEM+10%FCS at an MOI of 10. DMEM+10%FCS alone was used as a control. After 1.5h

penicillin (10 µg/ml) and gentamycin (100 µg/ml) were added. Barrier integrity was

monitored at 4000 Hz from seeding to 70h post infection. Resistance data were normalized

to the time of infection.

Bacterial genomics

Whole genome sequencing and genomic data analysis were performed as previously

described [30]. Briefly, from 150 bp paired end reads, *de novo* assemblies were generated

with SPAdes v.3.10 [30] and annotated with Prokka v.1.13 [31]. A pangenome was

constructed with Roary v.3.12 [32] and the resulting core-gene alignment was used to build

a maximum likelihood phylogenetic tree with Fasttree v.2.1.10 [33]. *Emm*-typing was

performed *in-silico* by querying assemblies against the curated CDC database [34, 35] with

blastn v.2.9.0. Virulence genes were identified by querying the assemblies against the full

VFDB database, i.e. VFDB_setB_nt.fasta, downloaded on the 04.08.23 [36], using ABRIcate

v.0.5 (minimum identity and coverage: 85%). ENA accession number: PRJEB57816.

Statistics

Three biological replicates were carried out for each strain, except for HBMECs viability and

ECIS assessment where two biological replicates were carried out for each strain. The

average of the biological replicates for each strain is represented on the graphs as a single

data point. The average of these values represents the final value depicted on the graphs.

Error bars represent standard deviations. The Mann-Whitney test was used to assess

statistical significance for all assays apart from HBMECs viability data that were analyzed

using the Kruskal-Wallis test.

Results

Cases description

Five GAS meningitis cases, occurring between 2013 and 2017 in Switzerland, were

included in this study and the corresponding GAS isolates were retrospectively retrieved for

analysis. The median age of the patients was 64 years (42 to 79 years) and 60% (n=3) were

female. The mortality rate was 20% (n=1) while 2 out of 6 patients (33%) experienced

neurological sequelae (patients 3 and 4), which included slight hearing loss, vertigo and

slight paresis of the limbs. The median duration of the hospital stay was 17 days (3 to 29

days) and 4 patients needed intensive care treatment with endotracheal intubation.

Relevant clinical and laboratory findings are depicted in table 1. Mastoiditis was the

underlying diagnosis in three cases. During the course of the disease one patient had septic

thrombosis and one developed a brain abscess (Fig.1A-1B). Surgical treatment such as

antrotomy, mastoidectomy, or burr-hole trepanation was required in three patients. In

patient 4, histological analysis from the surgically-removed mastoid tissue showed necrotic

tissue and individual mucosal fragments covered by simple cuboid epithelium with severe

acute and chronic inflammation and showing extracellular accumulation of bacterial cells

(Fig.1C-1D).

Bacterial virulence

The assessment of adherence to and invasion of HBMECs, the activity of the streptococcal

virulence factors DNases, SLO and SpyCEP as well as barrier integrity of HBMECs upon GAS

challenge was carried out on meningitis, otitis and colonizing clinical isolates. No significant

difference between adherence and invasion behavior to HBMECs upon infection with

meningitis, otitis or colonizing GAS isolates was found (Fig.2A-2B) and the toxicity of the

isolates toward HBMECs during infection did not vary according to their sampling location

(Fig.S1A-S1B). Furthermore, no differences in virulence factors activity or the isolates

potential to disrupt the barrier formed by HBMECs among the three groups were observed

(Fig.2C-2F). A trend, albeit not significant, towards reduced SpyCEP activity was observed in

the meningitis group (Fig.2D).

Genotyping

Typing of meningitis and otitis isolates showed a high prevalence of emm-1/ST28 (n=4/5 and

n=5/5, respectively, Tab.S1). Since one meningitis clinical isolate belonged to the emm-

28/ST52, we chose to include four emm-1/ST28 and one emm-28/ST52 in the colonizing

isolates group. Only one meningitis clinical isolate belonged to the *emm*-28/ST52 and one

colonizing isolate to the *emm*-28/ST458. A core genome phylogeny revealed that, within the

M1 clade, the meningitis isolates did not form a monophyletic group. (Fig.S2A-S2B). The

presence of genes encoding virulence factors was analyzed and this showed no specific

pattern for meningitis isolates when compared to otitis or colonizing ones (Fig. S2C). We

additionally checked for the presence of mutations in the GAS CovRS regulatory genes that

may lead to upregulated virulence factors activity (Tab.S2) [37]. No mutations were found in

CovR, while we found CovS mutations in two of the meningitis isolates (CI1224, *emm*-1 and

CI407, *emm*-28) and in one of the colonizing ones (CI571, *emm*-28). An in-frame deletion of

the proline in position 16 was found in CI1224. CI407 presented a deletion of the first 46

amino acids as well as non-synonymous mutations in position 47 (Leu-Met), 226 (Glu-Gly)

and 332 (Val-Glu). Isolate CI571 had the same non-synonymous mutations as CI407 in

position 226 (Glu-Gly) and 332 (Val-Glu), common to the emm-28 lineage [38]. However,

these mutations did not result in a reduced CovS function or increased virulence factors

activity.

Discussion

We found no particular phenotypic or genotypic characteristics in the five GAS isolates causing meningitis as compared to otitis or colonizing isolates. The only common denominator was preceding otitis and mastoiditis on the host's side. GAS meningitis is an extremely rare condition, representing only the 0.2%-0.9% of cases of bacterial meningitis in the United States and Brazil [39-41] and 1-4% of invasive GAS cases in Europe and the United States [1-3]. A report from Switzerland described nine cases of GAS meningitis registered between 1983 and 1999 across all major Swiss hospitals, again emphasizing the rarity of this disease [42]. A recent report showed an increase in the number of GAS meningitis case in the Netherlands in 2022-2023, compared with the previous 40 years, with emm-1 being the most represented M-protein variant [43].

Meningitis occurs either by hematogenous spread or by direct spread of bacteria from a neighboring infection. Hematogenous spread normally occurs when colonizing bacteria breach the nasopharyngeal mucosa, reach the blood [44] and subsequently cross the blood-brain barrier. It was suggested that spread from the upper respiratory tract's mucosa may be due to an altered balance in the host microbe population, possibly favoring GAS number increase and subsequent spread [5]. This interesting hypothesis was however never tested and remains a speculation. On the other hand, direct spread to the brain can also occur *per continuitatem* from infected adjacent anatomic structures, such as the ear or the mastoid bone [44].

We searched for common traits in the clinical presentation of our patients and characteristics shared by the meningitis clinical isolates, as opposed to otitis and colonizing ones, to find possible explanations for the spreading of GAS to the brain. In our patients' cohort, the presence of mastoiditis in four patients and thrombosis in one supports the idea

of direct spread through bone erosion from neighboring oto-mastoiditis and osteo-

thrombophlebitis. Moreover otitis, a condition often associated with GAS meningitis [5], was

present in the majority of cases suggesting continuous spread as a likely cause of infection.

In contrast, hematogenous spread was less common and a plausible cause of infection for

patient 3 only. There is no way to determine whether the presence of bacteria in the blood

in patients 1, 2 and 5 occurred before or after spread to the central nervous system.

However, given the infection focus and the evidence pointing to continuous spread from the

ear and mastoid bone as one of the main causes of GAS meningitis, we postulate that

hematogenous spread, In the case of patients 1, 2 and 5, was not the cause of meningitis.

The presence of a local focus of infection may explain the high surgery rate in our GAS

meningitis cohort, which has been reported previously. In a nationwide cohort study in the

Netherlands, 46% of GAS meningitis patients required surgical intervention, while this

remains an absolute rarity only described in case reports for the more common meningitis

pathogens (Pneumococcus, Meningococcus) [4, 45].

On the pathogen side, no common features explaining bacterial spread to the brain were

found among the meningitis isolates, as compared to otitis and colonizing ones. This points

to host rather than pathogen factors playing a role in facilitating bacterial invasion, as

previously described for *S. aureus* [30]. Adherence and invasion assays towards HBMECs

confirmed the absence of a particular tropism towards brain microvascular tissue for the

meningitis isolates, excluding an advantage in crossing or damaging the blood brain barrier,

as compared to otitis and colonizing ones, in the tested settings. Moreover, virulence factors

activity did not significantly vary among the three groups of isolates. The *emm*-type of the

strains did not generally seem to influence adherence, invasion or virulence factors activity,

although the number of strains and *emm*-types tested is too little to be able to make a

conclusive statement. A trend towards decreased SpyCEp activity was observed in the

meningitis group with the M28 strain CI407 acting as an outlier and presenting a level of

activity remarkably higher than the other meningitis isolates. CI407 was characterized by a

pronounced deletion of the first portion of the CovS protein, a key regulator of virulence

factors activity in GAS, as well as by the presence of three non-synonymous mutations

(Tab.S2), as compared to the sequence of the CovS protein of GAS M1T1 5448 [37]. Normally

a complete loss of CovS function, as for the GAS M1T1 5448 AP strain, would lead to a

complete IL-8 degradation leading to 100% cleavage [29], which we did not observe. SLO and

DNase activities were also identical to the non-mutated strains. A partial loss of CovS

functionality could be postulated in this case [46], although the mutations in CI407 mainly

affect the N-terminal portion of the CovS protein where no functional domains are present

[18, 47]. Further investigations involving a larger group of clinical isolates, including different

emm-types, is needed to address the question whether a decreased IL-8 degradation

potential might facilitate invasion of the blood brain barrier. Isolates from the *emm*-1 and

emm-28 lineages represented both in the meningitis and colonizing groups, behaved very

similarly in terms of virulence factors activity.

In accordance with previous studies stating an association of *emm*-1/ST28 with GAS

meningitis [26], 4/5 meningitis isolates belonged to the *emm*-1/ST28 lineage. The fifth

isolate belonged to the *emm*-28 lineage, which is another prevalent *emm*-type previously

described in GAS meningitis [26] and, together with *emm*-1, among the most representative

emm-types for invasive GAS infections across Europe and nord America [48]. All otitis and

colonizing isolates belonged either to the *emm*-1 or the *emm*-28 lineages as well. A

prevalence of the emm-1 linage in acute otitis media in children (around 30%) was

previously reported [49]. The serotype of colonizing isolates was however specifically

selected to reflect the serotypes present in the meningitis and otitis groups. It cannot be

excluded that the overrepresentation of *emm*-1 serotypes in meningitis and otitis might

reflect the global success and dominance of *emm*-1 strains causing GAS invasive infections

[48]. The presence or absence of GAS virulence determinants did not reveal a specific

pattern for meningitis isolates when compared to otitis and colonizing ones.

GAS meningitis occurs very rarely, leading to the availability of a small number of cases

that could be included in this study. Nevertheless, despite the small sample size, our data

suggest that the most common pathophysiological cause in this patients' cohort was local

spread from a site of bacterial infection in neighboring tissues, with hematogenous spread a

possible explanation in one case.

In the absence of distinct virulence determinants characterizing the meningitis isolates,

we conclude that, in this particular cohort, host factors such as breach of the mucosa or local

infections play a preponderant role in the transition from colonization to invasion. These

results call for future studies including a higher number of cases and clinical isolates and a

more comprehensive phenotypic and genotypic investigation to support our conclusion.

Author contributions

LM: Experimental design, acquisition, analysis and interpretation of data, writing of the

manuscript.

FA: Experimental design, acquisition, analysis and interpretation of data, writing of the

manuscript.

MB: Acquisition, analysis and interpretation of data, writing of the manuscript

TAS: Experimental design, data acquisition, data analysis and critical reading of the

manuscript

DMH: Experimental design, data acquisition, data analysis

EP: Experimental design, data acquisition, data analysis

SH: Experimental design, data acquisition, data analysis

JE: Data acquisition, critical reading of the manuscript

DM: Data acquisition

AKR: Data acquisition

EMM: Histological analysis

RAS: Funding

BH: Funding

SMS: Experimental design, funding, critical reading of the manuscript

SDB: Conceptualization, funding, critical reading of the manuscript

ASZ: Conceptualization, funding, critical reading of the manuscript

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374   Conflict of interest

375   The   authors   have   no   conflict   of   interest   to   declare.

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 Figure legend

Figure 1 - Patients scans and histology results

A) CT-Scan: septic thrombosis with mastoiditis and otitis media on the right side. The red

arrow indicates thrombosis **B)** MRI-Scan: T2 sequence showing penetration of lower

temporal lobe with abscess formation. The red arrow indicates an abscess **C)** HE Staining of

mastoid tissue: the black arrow indicates necrotic mastoid tissue, the dashed black arrow

indicates mixed neutrocytic and lymphocytic infiltration, foam cells and cholesterol crystals

(not depicted) indicating chronic inflammation. **D)** Brown-Brenn staining of mastoid tissue:

The arrows indicates accumulation of extracellular bacteria in chains.

Figure 2 - Bacterial strains virulence determinants

A) Adherence to and **B)** invasion of human brain vascular endothelial cells (HBMECs) were

assessed after 30 minutes (MOI 1) or two hours (MOI 10) of infection, respectively.

Adherence and invasion percentages were calculated based on the initial bacterial inoculum

used to infect HBMECs. Bacterial adherence above 100% occurred in one case and could be

explained by increased initial bacterial growth or by disruption of long bacterial chains or

clumps due to sheer pipetting forces during the eukaryotic cells lysis step. Bacterial virulence

factors activity of **C)** Streptococcal DNases, **D)** the IL-8 protease SpyCEP and **E)** the pore-

forming toxin SLO were assessed in supernatants of exponentially growing GAS. **F)** HBMECs

barrier function disruption was carried with the ECIS® Z-Theta instrument (Applied

Biophysics), measuring the impedance generated by the cell barrier in the presence or

absence of bacteria. Control experiments were carried out using medium only. **A, B, C, D)**

The average of the biological replicates for each strain is depicted as a single data point on

the graphs, so that each data point represents a single strain. The two strains indicated in

the figure legend with a black symbol (CI407 and CI571) are the only strains of *emm*-type 28

(M28), as opposed to all other strains that are *emm*-type 1 (M1). **A, B, C, D, E)** Three

biological replicates were carried out for each strain. The average values for strains

belonging to a single group of isolates (meningitis, otitis or colonizing) is depicted in the

graphs and the error bars represent the standard deviation. **F)** Two biological replicates were

carried out for each strain and the average of the strains belonging to the same group is

depicted on the graph. Control=medium only. No significant differences in adherence,

invasion, virulence factors activity or barrier disruption was observed among the different

groups of isolates. A significant difference was found between the respective controls and

meningitis, otitis or colonizing isolates for DNase (p=0.0159, 0.0159 and 0.0159, respectively)

and SpyCEP (p=0.0179, 0.0357 and 0.0179, respectively) activity. Statistical significance was

assessed using the Mann-Whitney test. Men=meningitis, Col=colonizing, Ctrl=control.

A
CT scan

B
MRI

C
H&E stain

D
Brown-Brenn stain

Patient ID Sex Age(y) Isolate Nr.	Associated health conditions	Clinical presentation	Complications	Site of GAS identification	Antibiotic treatment regimen	Death
1 F 73 CI1224	• Perforated otitis media • Diabetes • Hypothyreosis • Depression	• Altered vigilance and mental state • Vertigo • Hypothermia • Hypotension • Acute abdomen	• Partial septic sinus transversus thrombosis • Coma • Irreversible hypoxic brain damage • ICU involvement	CSF Blood Ear swab	3d ceftriaxon 3d amoxicillin 3d clindamycin	yes
2 M 55 CI1271	• Otitis media • Depression	• Altered vigilance • Meningism • Slight one-sided arm paralysis • Otalgia • Diarrhea • Nausea	• Coma • Mastoiditis • Cerebral abscess • ICU involvement	Blood Mastoid tissue	2d ceftriaxon 1d amoxicillin Escalation to: 18d penicillin 5d metronidazole 5d clindamycin + Surgical intervention	no
3 F 64 CI1293	• Bicuspid aortic valve	• Acute reduction of general condition • Meningism • Right sided paresis of upper and lower limbs • Right sided tongue deviation • Anisocoria • Petechial bleeds	• Thrombocytopenia • Acute kidney failure • Aortitis • Nosocomial Influenza infection	CSF Blood	Initially ceftriaxon then change to penicillin due to elevated transaminases, total 18d 3d gentamycin 15d steroids 10d vancomycin 10d clindamycin	no
4 F 79 CI407	• Otitis media (perforated) • Coronary heart disease • Pre-diabetic metabolism	• High temperatures • Vertigo • Altered mental state and vigilance	• Tegmen tympani penetration • Mastoiditis • Labyrinthitis • Coma • Long-term one sided deafness • ICU involvement	Mastoid tissue	Initially 1d ceftriaxon, then escalation to meropeneme for 2d, then back to ceftriaxone for 20 days Initially 2d acyclovir 1d amoxicillin 4d mephamesone + Surgical intervention	no
5 M 42 CI543	• Otitis media • Bullous • Myringitis with insertion of tympanostomy tube	• One day after insertion of tympanostomy tube acute reduction of vigilance • Altered mental state and meningism	• Mastoiditis • Subdural empyema • Cerebritis • Ventrikulitis • Coma • ICU involvement	Blood CSF Mastoid tissue	9d ceftriaxon, then escalation to 18d ceftazidime and de-escalation back to ceftriaxone for another 35d 2d amoxicillin + clavulanic acid 2d acyclovir 2d metronidazol 2d, pause 3d, then another 2d 9d intrathecal gentamycin 9d intrathecal vancomycin + Surgical intervention	no

Table 1 – Patients description. Clinical findings, complications and treatment regimen of the patients included in this study. Abbreviations: CSF=cerebrospinal fluid; d=day; ICU=intensive care unit.