

Methanobrevibacter smithii tonsillar phlegmon: a case report

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1	Methanobrevibacter smithii tonsillar phlegmon: A case report.
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20	ABSTRACT

- 21 Untreated tonsillar phlegmon is a life-threatening condition commonly caused by
- 22 Streptococcus pyogenes and Fusobacterium necrophorum among other pathogens. Here using
- 23 specific laboratory tools, we detected *Methanobrevibacter smithii* in addition to *S. pyogenes*.
- 24 This unprecedented observation questions the role of methanogens in phlegmon, and the
- 25 optimal treatment of these mixed infection.
- 26
- Keywords: *Methanobrevibacter smithii*, methanogens, tonsillar phlegmon, *Streptococcus pyogenes*.
- 29

30 INTRODUCTION

Tonsillar phlegmon, a retropharyngeal abscess resulting from the infection of the peritonsillar space and the pharyngeal constrictor muscle through the connective tissue of the amygdala [1], is a life-threatening situation in the case of cellulitis with extension to the mediastinum and potential extensive venous thrombosis up to the cavernous sinus [2]. Tonsillar phlegmon is affecting particularly adolescent and young adults [3], complicating acute pharyngitis, recurrent pharyngitis, and chronic tonsillitis [4].

Current microbiological documentation, mostly based on direct microscopic
observation and culture of pus specimen, yields a polymicrobial infection of aerobic and
anaerobic bacteria, dominated by *Streptococcus pyogenes* (Group A *Streptococcus*), the only
established pathogen in tonsillar abscess [5-7]. Also, Streptococci (S. *pyogenes*, Streptococcus *milleri*, Streptococcus *viridans*), *Haemophilus spp* and anaerobes including *Fusobacterium necrophorum*, *Fusobacterium nucleatum*, *Prevotella melaninogenica*, *Prevotella intermedia*and *Peptostreptococcus* spp are isolated from most patients (82% to 90%) [2, 8, 9].

- In this situation of bacterial tonsillar phlegmon, detection of methanogenic archaea,
 mainly represented by *Methanobrevibacter smithii* (*M. smithii*), the most abundant
 methanogenic archaea species in the human gut [10] and recently emerging as co-pathogen in
 various disease situations [11-14] would requires specific laboratory methods.
- 48 Accordingly, Here, we have used such specific methods we are mastering, to explore49 the presence of methanogens in one case of tonsillar phlegmon.

50 **Case presentation.** A 22-year-old man was admitted to the emergency unit for a two-day

51 progressively worsening dysphagia with odynophagia and bilateral pharyngeal pain,

- 52 associated with sweating without fever. The patient self-treated with paracetamol without
- 53 antibiotic nor non-steroidal anti-inflammatory drugs. Clinical examination revealed bilateral

inflammatory oropharynx and trismus, no sign of cellulitis, nor palpated cervical 54 lymphadenopathy. Nasofibroscopy disclosed no lesion in the oropharynx with mobile and free 55 vocal cords. Remarkable laboratory test values included leukocytosis at 17 G/L with 56 neutrophilic cells 12G/L and C-reactive protein at 132 mg/L. A cervical CT scan showed 57 bilateral tonsillar hypertrophy with a left tonsillar phlegmon without extension to the 58 prevertebral space and without anomalies on the internal jugular venous, and bilateral cervical 59 adenopathies (Figure 1). Intravenous amoxicillin and clavulanic acid 1gr three times a day 60 and metronidazole 500 mg three times a day was started 24 hours before the patient benefited 61 from a surgical drainage of the left phlegmon. Per-operative papillomatosis was observed on 62 the left tonsil, as confirmed by pathology which disclosed inflammatory papilloma without 63 dysplasia. The pus collected from the phlegmon was immediately inoculated into one aerobic 64 and one anaerobic blood culture bottles (Virtuo; bioMérieux, La Balme les Grottes, France), 65 66 incubated in the BACT/ALERT® VIRTUO® automaton (bioMérieux). Growth was detected in aerobic and anaerobic bottles after 3.6 and 4.3-hour incubation respectively, and direct 67 microscopic examination yielded Gram-positive cocci. Broth culture was inoculated on 68 Chocolate agar PolyViteX medium (bioMérieux) and COLUMBIA ANC medium 69 (bioMérieux), incubated at 37°C under 5% CO₂ for 5 days to grow aerobic bacteria; and 5% 70 sheep blood Columbia agar medium (bioMérieux) incubated under strict anaerobiosis at 37°C 71 72 for 10 days to grow anaerobic bacteria. After 24-hour incubation, colonies grown on Columbia and Chocolat PolyVitex media, and then identified as *Streptococcus pyogenes* 73 using matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (MALDI-74 TOF-MS) as previously described [15]. Antibiotic susceptibility testing was done on the 75 isolated strain as previously described [16] and indicated a wild-type S. pyogenes susceptible 76 77 to amoxicillin, erythromycin, vancomycin, teicoplanin, clindamycin and pristinamycin; and exhibiting a low-level susceptibility to gentamicin. The final antibiotherapy was amoxicillin 78

- and clavulanic acid 1gr three times a day and metronidazole 500 mg three times a day for 10
- 80 days. The patient was cured after 15 days of follow up.

81 For molecular detection of methanogens, DNA was extracted from the pus sample using the

82 automat extractor EZ1 advanced XL with EZ1 DNA Tissue kit (Qiagen, Courtaboeuf, France)

after 20-minute sonication. Amplification of the archaeal 16S rRNA gene was performed as

previously described [14]. Sequencing the amplicon yielded 100 % sequence similarity with

85 the reference *M. smithii* ATCC 35061 homologous sequence (GenBank accession number

86 NC_009515).

87 Further, to determine whether the pus sample was monomicrobial or polymicrobial, we

completed the routine bacteriological investigations by using the rapid culturomic protocol as

89 previously described [17]. Seven cultured bacterial species including *Morganella morganii*,

90 Klebsiella pneumoniae, Klebsiella oxytoca, Proteus mirabilis, Escherichia coli, Enterococcus

91 *faecalis* and *Bacteroides vulgatus*, were finaly identified by MALDI-TOF-MS [15]. However,

92 tentative isolation of *M. smithii* failed.

93 **DISCUSSION**

We are reporting the unprecedented detection of the methanogen *M. smithii* in the pus
sample collected from a tonsillar phlegmon. While this detection was ascertained by the
negativity of the negative controls and the fact that amplicon was sequenced, the very same
pus also yielded a total of eight different bacterial species, in agreement with previous
knowledge regarding the bacterial species reported to be associated with tonsillar phlegmon
[18-20].

Until now, the presence of methanogens has never been reported in tonsil infections,
this is due to the fact that methanogens are not systematically searched, and their culture
remains fastidious. Here, the reason for no growth of *M. smithii* may be attributed to the fact

that methanogens are extremely sensitive to oxygen and to the fact that phlegmon pus was
collected after the patient received metronidazole, one of few antibiotics shown to be *in vitro*effective against methanogens [21].

Although previously unreported, here detection of *M. smithii* in a tonsillar pus was not surprising as *M.smithii* is known to colonize the oral fluid microbiota [22], being further implicated in oral cavity dysbiosis such as periodontitis and peri implantitis [23, 24]. The presence of methanogens in saliva reinforces the hypothesis of a possible inflammation of the salivary glands that would cause tonsillar phlegmon [2, 25].

In this patient, *M. smithii* was detected in a pus sample also containing eight different 111 bacterial species and such an association of *M. smithii* with bacteria, chiefly anaerobes, is 112 constant in all the previous descriptions of *M. smithii* coinfections [11-14]. These bacteria 113 essentially belong to the enterobacterial and Bacteroidal orders comprising of hydrogen 114 115 producers [26, 27]. Here, we suggest the possible involvement of *M. smithii* in this mixed infection by promoting the growth of aerobic and anaerobic bacteria via syntrophic 116 117 interactions which are the real pathogens. The exact mechanism is still poorly understood, but 118 it would seem that methanogens control the production of essential growth factors for bacteria, such as short-chain fatty acids, by continuously maintaining a low partial pressure of 119 H₂ [28, 29] and may also play occasional roles in the development of tonsillar phlegmon as 120 121 well as spread of infection.

In conclusion, this case report indicates that tonsillar phlegmon should no longer be regarded as a bacterial infection, but rather an infection mixing anaerobes with methanogens. When the relative role of these different microorganisms should be reevaluated as methanogens are producing methane which is a tissue and cell damage [30, 31], potentially responsible for gas radiologically visible in some cases of tonsillar phlegmon [32, 33], it

- 127 further questions the importance of prescribing antibiotics active against methanogens, such
- as in this patient which medical evolution was favorable.

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156 Conflict of interest:

- 157 The authors declare no conflicts of interest.
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Author Contributions: D.K; performed the experiments; data interpretation; manuscript

167 drafting; G.F; study design and manuscript drafting; M.J and R.T; sample collection and data

168 interpretation; D.M; study design and implementation, data analysis, results interpretation,

- and manuscript writing; G.G; study design and implementation, data analysis, and manuscript
- 170 writing. All authors have read and agreed to the published version of the manuscript.

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- Figure 1. head and neck CT-scan sagittal plan showing the tonsillar phlegmon
- 279 (arrow) in which PCR detected *Methanobrevibacter smithii*, in addition to cultured
- 280 bacteria.



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