

# The Role of *Ureaplasma* spp. in the Development of Nongonococcal Urethritis and Infertility among Men

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<b>SUMMARY</b> .....	1
<b>INTRODUCTION</b> .....	1
Background Biology of <i>Ureaplasma</i> spp. ....	2
<b>THE PROINFLAMMATORY POTENTIAL OF UREAPLASMA SPP.</b> .....	3
Human Volunteer Experiments with <i>Ureaplasma</i> Species Infection of the Urethra .....	3
Animal Models of <i>Ureaplasma</i> Species-Induced Urethritis .....	3
<i>In Vitro</i> Cell Line Models of <i>Ureaplasma</i> Species-Induced Inflammation .....	4
<b>RISK FACTORS LINKED WITH DEVELOPMENT OF UREAPLASMA SPECIES-ASSOCIATED NGU</b> .....	4
Risk Factor 1: the Species of <i>Ureaplasma</i> Present within the Urethra .....	4
Risk Factor 2: the Sexual History of the Patient .....	8
Risk Factor 3: the Bacterial Load of <i>Ureaplasma</i> spp. within the Male Urethra .....	9
<b>IMPACT OF UREAPLASMA SPP. ON MALE FERTILITY</b> .....	9
Clinical Studies Associating <i>Ureaplasma</i> spp. with Infertility .....	9
Proposed Mechanisms of <i>Ureaplasma</i> Species-Associated Infertility .....	11
<b>TREATMENT OF GENITAL TRACT INFECTIONS IN MEN CAUSED BY UREAPLASMA SPP.</b> .....	12
<b>CONCLUSIONS</b> .....	13
<b>ACKNOWLEDGMENT</b> .....	14
<b>REFERENCES</b> .....	14
<b>AUTHOR BIOS</b> .....	16

**SUMMARY** *Ureaplasma* spp. are a genus of bacteria for which two human-associated species exist: *Ureaplasma urealyticum* and *Ureaplasma parvum*. Their definition as a pathogen in the context of nongonococcal urethritis (NGU) and infertility among males remains highly controversial, largely due to historically high rates of isolation of these bacteria from the urethra of seemingly healthy men. This review summarizes the emerging evidence suggesting a true pathogenic role of these bacteria under specific conditions, which we term risk factors. We examine the historical, clinical, and experimental studies which support a causal role for *Ureaplasma* spp. in the development of NGU as well as some of the proposed mechanisms behind the association of *Ureaplasma* spp. and the development of infertility. Finally, we discuss the potential for developing a case-by-case risk-based approach toward the management of men who present with seemingly idiopathic NGU but who are positive for *Ureaplasma* spp.

**KEYWORDS** *Ureaplasma parvum*, *Ureaplasma urealyticum*, infertility, nongonococcal urethritis

## INTRODUCTION

Nongonococcal urethritis (NGU) is a leading sexually acquired condition among men. It is defined by inflammation of the urethra in the absence of *Neisseria gonorrhoeae* and includes signs and symptoms such as penile discharge, dysuria, as well as irritation inside and around the urethra. *Chlamydia trachomatis* has long been regarded as the predominant infectious agent among patients suffering from NGU,

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with 20 to 50% of individuals being positive for the pathogen, whereas more recently, *Mycoplasma genitalium* has achieved recognition as a pathogen and may be isolated from 10 to 30% of NGU patients (1). Although *C. trachomatis* and *M. genitalium* account for many cases of NGU, of concern is the high prevalence of up to 45% of idiopathic NGU cases in which classic pathogens are not identified (2).

Bacteria from the genus *Ureaplasma* are leading candidates to fill the void presented by this idiopathic condition among NGU patients. The first documented isolation of these bacteria was from male patients experiencing NGU (3). Many reports have followed up this observation with a view to gather evidence to support the idea of *Ureaplasma* spp. being an etiological agent of NGU, but the combination of inconsistencies in reporting and study design and the high prevalence of between 5 and 15% among healthy males aged 16 to 44 years has prevented the acknowledgment of these organisms as true pathogens in the context of genitourinary medicine (GUM) (4). For these reasons, the idea of *Ureaplasma* spp. as GUM pathogens remains controversial among GUM practitioners. Additionally, the potential role of *Ureaplasma* spp. as agents with a causal role in male infertility has been debated. Many of the recognized GUM pathogens, such as *N. gonorrhoeae* and *C. trachomatis*, have been implicated in complications, such as male infertility, but more work is required to gain a clear understanding of the implications associated with a failure to clear *Ureaplasma* spp. from the urethra (5).

In this review, we present an update from the current literature to discuss the potential of *Ureaplasma* spp. to be a risk factor for male genital tract infections, with specific reference to NGU and infertility. In the context of NGU, we present the arguments for and against a role for these bacteria in disease development with a focus on some of the unique risk factors which have been overlooked historically. Increasing interest has focused on *Ureaplasma* spp. and their potential role in the development of male infertility. We discuss the clinical evidence as well as the proposed mechanisms which have been neglected when taking into account markers for infertility. Finally, the potential therapeutic considerations are evaluated, and we discuss the potential for risk-based screening approaches as an effective means to manage patients with seemingly idiopathic NGU in the face of growing concerns over antimicrobial resistance among GUM pathogens.

### **Background Biology of *Ureaplasma* spp.**

*Ureaplasma* spp. are recognized as some of the smallest self-replicating, free-living microorganisms. They are a unique genus of bacteria due to their essential requirement for urea in the synthesis of ATP, with further defining characteristics being shared with the closely related mycoplasmas, including a low G+C genomic content, the lack of a peptidoglycan-containing cell wall, and a requirement of cholesterol for growth (6, 7).

*Ureaplasmas* were first isolated from male NGU patients in 1954, and due to the tiny colony size upon agar plates, these bacteria were originally referred to as "T-strain" or "tiny" mycoplasmas (3). Following the establishment of the essential requirement for urea, the genus name *Ureaplasma* was adopted (8). A single species of human-associated *Ureaplasma*, *Ureaplasma urealyticum*, was initially recognized, and was further subdivided into two biovars. The nomenclature of *U. urealyticum* for describing all human-associated isolates was embedded until the work by Robertson et al. in 2002, which made a substantial contribution to redefining these bacteria into two antigenically distinct human-associated species (9). These were defined as *U. urealyticum* and *Ureaplasma parvum*. The two species are divided into 14 serovars, with serovars 1, 3, 6 and 14 being assigned to *U. parvum* and the remaining serovars (serovars 2, 4, 5, and 7 to 13) being defined as *U. urealyticum*.

Numerous clinical manifestations have been associated with *Ureaplasma* spp. Among the most notable is the role of *Ureaplasma* spp. in adverse pregnancy outcomes, such as chorioamnionitis and preterm premature rupture of membranes leading to preterm birth (10, 11). The subsequent colonization of *Ureaplasma* spp. within the lungs of premature neonates has been associated with a 7.9-fold increased risk of

bronchopulmonary dysplasia, a 3.3-fold increased risk of intraventricular hemorrhage, and a 2.5-fold increased risk of necrotizing enterocolitis (12). In adults, attention has been drawn to the development of an atypical hyperammonemia, in which lung transplant patients and, potentially, kidney transplant patients have increased serum ammonia levels as a result of systemic *Ureaplasma* species infection (13–15). If left untreated with antibiotics, such increased serum ammonia levels can lead to delirium, cerebral edema, and, eventually, death.

Historically, the link between *Ureaplasma* spp. and the development of NGU, as well as infertility, prostatitis, and epididymitis, among men has been inconsistent. The reason for this, however, is certainly not from a lack of studies examining potential associations between *Ureaplasma* spp. and NGU (2, 16–21). Rather, the lack of conclusive evidence may reflect the complex interaction between host and pathogen, as discussed later, combined with the high prevalence of *Ureaplasma* spp. among control groups, which suggests that they are innocent bystanders present at the time of screening. Although *Ureaplasma* spp. were isolated approximately 30 years prior to the isolation of *Mycoplasma genitalium*, the latter has risen to prominent pathogen status more rapidly, and new guidelines for its management are now in place in the United Kingdom (3, 22, 23).

### THE PROINFLAMMATORY POTENTIAL OF UREAPLASMA SPP.

#### Human Volunteer Experiments with *Ureaplasma* Species Infection of the Urethra

To demonstrate the pathogenicity of *Ureaplasma* spp., several investigators have undertaken human participant experiments (24, 25). The first such experiment, by Jänsch in 1972, identified a polymorphonuclear leukocyte (PMN) response following inoculation with an unknown and poorly defined *Ureaplasma* sp. (24). Although the experiment was poorly designed and controlled for, this gave an initial insight into the inflammatory nature of a human infection with a *Ureaplasma* sp. A second, more defined experiment was conducted with two human participants. The first participant received an intraurethral inoculation of a clinically relevant titer of a low-passage clinical isolate of *U. urealyticum* serovar 5 isolated from a patient experiencing NGU in which no other organisms were present (25). The participant subsequently developed symptoms of dysuria and signs of urethritis in the form of a PMN response. The serum recovered from the volunteer demonstrated seroconversion with high titers of specific antibodies. Upon administration of tetracycline, both signs and symptoms resolved. The second participant received an alternative isolate of *U. urealyticum* serovar 5 isolated from a second patient presenting with NGU. Again, signs and symptoms ensued, but upon administration of tetracycline, signs, such as urinary threads, persisted in the absence of viable cultures. Seminal samples collected after antibiotic treatment indicated that the *U. urealyticum* isolate had disseminated, suggesting the potential involvement of the prostate and highlighting the potential adverse sequelae associated with exposure to *Ureaplasma* spp. Although such experiments are ethically questionable by today's standards, these studies provided initial evidence that exposure of the male urethra to clinically relevant titers of *U. urealyticum* has the capacity to elicit a PMN immune response in the presence of symptoms which reflect those seen among NGU patients. It should be noted, however, that what a clinically relevant titer of *U. urealyticum* is was not defined in this study and is a notable limitation for interpretation of these data.

#### Animal Models of *Ureaplasma* Species-Induced Urethritis

Due to the substantial ethical implications of human volunteer studies, investigators turned to model urethral infection caused by *Ureaplasma* spp. utilizing animal models. Like *Neisseria gonorrhoeae*, *Ureaplasma* spp. are host specific, resulting in an early reliance upon chimpanzee models due to the close ancestry with humans. Initial experiments with intraurethral inoculation saw the rapid multiplication of the bacteria within the urethra, but in the absence of a PMN response (26). A possible explanation for this lack of an immune response was suggested to be a loss of virulence from *in vitro*

passage. To examine this hypothesis, a second study was conducted with a larger number of chimpanzees (27). The inoculum for this study consisted of *Ureaplasma* spp. from men with NGU resuspended in a transport medium which was directly inoculated into the chimpanzees via intraurethral inoculation. Unlike the first study, a substantial PMN response was noted in conjunction with an increase in the *Ureaplasma* species titer. For reasons which are unknown, the species of *Ureaplasma* which was inoculated during this study was not determined.

Due to the lack of availability of chimpanzee models, investigators have moved to murine models to investigate colonization of the genital tract (28–30). Although many of these models have relied upon female mice and vaginal colonization, due to the physiology of the male mouse urethra, an inflammatory response characterized by increased tumor necrosis factor alpha (TNF- $\alpha$ ) and PMN recruitment has been described. A key confounding variable has been the essential requirement to pretreat the mice with estradiol to allow colonization to establish. This requirement for estradiol is likely due to suppression of the innate immune system, but the presence of estradiol binding proteins, as seen in other pathogens, is yet to be ruled out (31, 32).

### **In Vitro Cell Line Models of *Ureaplasma* Species-Induced Inflammation**

The difficulty in assessing *Ureaplasma* species infection of the urethra has resulted in a reductionist approach utilizing specific cell lines in isolation *in vitro* to look at cytokine responses. Some studies have focused on immune cells, such as THP-1 monocytes, phorbol myristate acetate (PMA)-differentiated macrophages, and primary human macrophages derived from lung fluid, which were then stimulated with *U. urealyticum* serovar 8 (33). In all cell types examined, stimulation with *U. urealyticum* resulted in a dose-dependent increase in the levels of interleukin-6 (IL-6) and TNF- $\alpha$  at both the mRNA and the protein levels. However, it should be noted that studies examining cytokine expression in relation to stimulation by *Ureaplasma* spp. tend to examine a single bacterial isolate and therefore do not give a true representation of the diversity of stimulating properties of *Ureaplasma* spp. It has been suggested that the predominant antigen found on the surface of *Ureaplasma* spp., known as the multiple-banded antigen (MBA), may account for differences in the inflammatory response (34). Sweeney et al. noted that the size and number of MBA repeats had an effect upon the levels of IL-8, which is a primary chemoattractant of PMNs, such as neutrophils (34). Although many of these studies were generalized for the immunogenic properties of *Ureaplasma* spp., they provide evidence for the inflammatory potential for these bacteria. An obvious limitation of these studies is the lack of consideration of the complexities of a full biological system, such as an adaptive immune response, the presence of other microorganisms which may permit infection, as well as the response to chronic exposure over time.

## **RISK FACTORS LINKED WITH DEVELOPMENT OF UREAPLASMA SPECIES-ASSOCIATED NGU**

*Ureaplasma* spp. can be detected in genital samples from men with NGU as well as healthy controls, which has fueled much of the controversy surrounding the role of *Ureaplasma* spp. in NGU (Table 1). Much of this historic evidence may now be questioned due to developments in the reclassification of *Ureaplasma* spp., better techniques for species differentiation, fully quantitative reporting of sample titer, as well as a better understanding of patient sexual histories. A proposed overview of the natural history of *Ureaplasma* spp. taking into account these risk factors is presented in Fig. 1.

### **Risk Factor 1: the Species of *Ureaplasma* Present within the Urethra**

Until 2002, human ureaplasmas were recognized as a single species subdivided into two biovars. The result of phenotypic and genotypic analysis later saw the official recognition of two independent species, *U. parvum* and *U. urealyticum* (9). This absence of species differentiation meant that studies prior to 2002 solely reported results as *U.*

**TABLE 1** Published studies examining the relationship between *Ureaplasma* spp. and NGU<sup>a</sup>

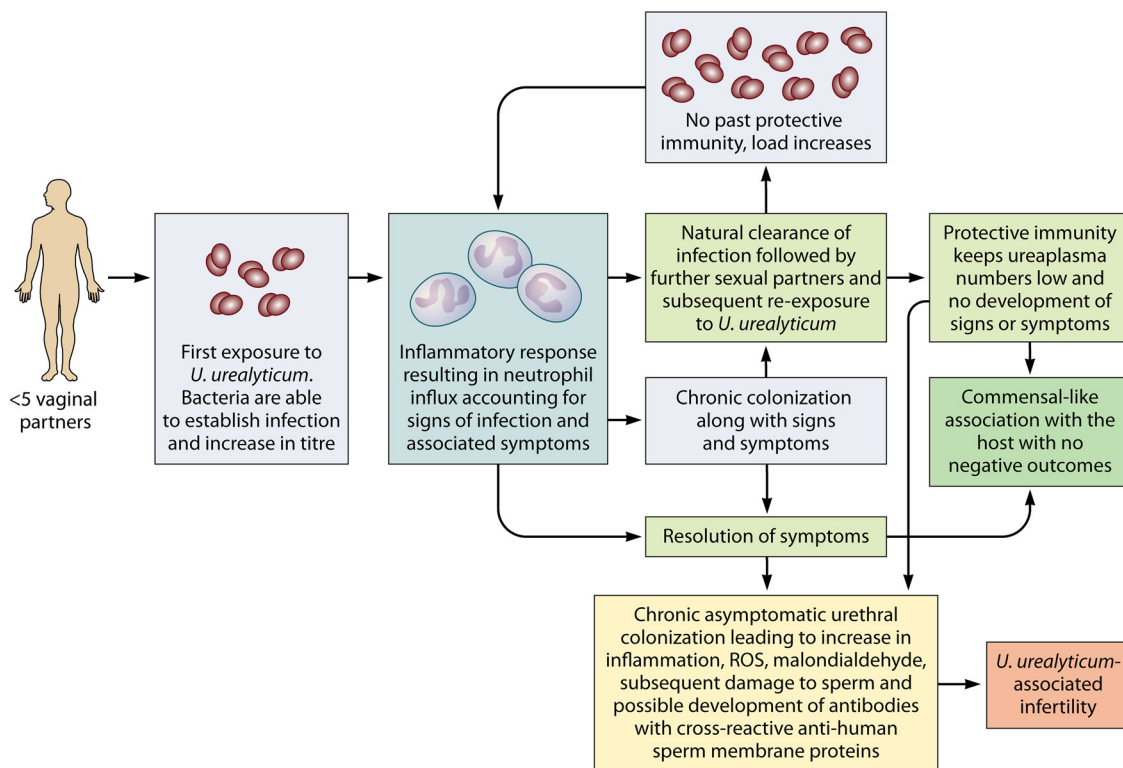
Authors (yr)	Reference	Country of study	Patient group	Sample type	No. of participants	Method of identification	Differentiation of <i>Ureaplasma</i> spp.	Key findings relating to <i>Ureaplasma</i> spp.
Frolund et al. (2016)	45	Sweden	Male patients attending STD clinic	First-void urine	187 men with acute NGU, 24 men with chronic NGU, and 73 men without NGU	Species-specific qPCR	Yes	The number of lifetime sexual partners was negatively associated with the <i>U. urealyticum</i> load. Urine containing <i>U. urealyticum</i> with >1.3 × 10 <sup>3</sup> genome equivalents/ml was associated with NGU. The prevalence of <i>U. parvum</i> was significantly higher in the NCNGU group.
Cox et al. (2016)	41	UK	Male patients attending a GUM clinic	Urine	75 men with NCNGU and 90 men without NCNGU	Species-specific real-time PCR	Yes	There was no association between <i>U. urealyticum</i> and NCNGU. Only four patients were positive for <i>U. urealyticum</i> .
Khatib et al. (2015)	19	UK	Males attending an urban sexual health clinic	Urine	83 men with urethritis	Multiplex PCR	Yes	A <i>U. parvum</i> load of >5 × 10 <sup>3</sup> cells/ml was significantly associated with >12.5 leucocytes/μl of urine. 83% of subjects had <5 × 10 <sup>3</sup> cells/ml, suggesting a low bacterial load and a lack of signs of inflammation.
Deguchi et al. (2015)	48	Japan	Retrospective study of men attending a urology clinic	First-void urine	15 symptomatic men and 38 asymptomatic men	qPCR	Yes	No significant difference in the positive rate was found between the undifferentiated <i>Ureaplasma</i> species-NGU and control groups. When the species was differentiated, <i>U. urealyticum</i> was significantly associated with the NGU group, whereas <i>U. parvum</i> was significantly associated with the control group.
Zhang et al. (2014)	40	Multiple countries	Meta-analysis	NA	1,507 men with NGU and 1,223 men without NGU	NA	Yes	The bacterial load of <i>U. urealyticum</i> was positively correlated with NGU and the number of leukocytes in urine. <i>U. urealyticum</i> was associated with NGU. The association was significantly stronger when analyzing men with <10 vaginal partners. The association was further strengthened when analyzing men with <5 vaginal partners. <i>U. parvum</i> was not associated with NGU.
Shimada et al. (2014)	47	Japan	Retrospective study of men attending a urology clinic	First-void urine	25 symptomatic men and 26 asymptomatic men	Species-specific qPCR	Yes	
Wetmore et al. (2011)	17	USA	Men attending an STD clinic	Urine	329 men with NGU, control group 1 consisting of 191 males without NGU attending an STD clinic, and control group 2 consisting of 193 males without NGU attending an emergency room	Culture	Yes	

(Continued on next page)

**TABLE 1** (Continued)

Authors (yr)	Reference	Country of study	Patient group	Sample type	No. of participants	Method of identification	Differentiation of <i>Ureaplasma</i> spp.	Key findings relating to <i>Ureaplasma</i> spp.
Couldwell et al. (2010)	38	Australia	Men attending a sexual health clinic	First-void urine	237 men with NGU and 268 controls	PCR	Yes	<i>U. urealyticum</i> was significantly associated with NGU in the absence of another urethral pathogen.
Ondondo et al. (2010)	21	USA	Archived samples from heterosexual males attending an STD clinic	Urine	119 men with NGU and 117 controls	PCR	Yes	<i>U. urealyticum</i> was strongly associated with NGU. This association was the strongest in men <28 yr of age.
Yu et al. (2008)	18	Hong Kong	Males attending a government STD clinic	Urine	98 men with NGU and 235 controls	Real-time PCR targeting the urease gene	No	<i>U. parvum</i> was not associated with NGU. Neither <i>Ureaplasma</i> nor <i>M. genitalium</i> was associated with symptomatic NGU.
Bradshaw et al. (2006)	39	Australia	Men attending a sexual health clinic	First-stream urine	329 men with NGU and 307 controls	PCR	Yes	Neither <i>U. urealyticum</i> nor <i>U. parvum</i> was associated with NGU.
Povlsen et al. (2002)	36	Sweden	Men attending a sexual health clinic	Urethral swab	125 men with NGU and 205 without NGU	PCR	Yes	No difference was found between the NGU and non-NGU group if ureaplasmas were not differentiated to the species level. When differentiated, significantly more <i>U. urealyticum</i> bacteria were associated with males with NGU than those without.
Horner et al. (2001)	16	UK	Heterosexual men with NGU and a control group	First-pass urine	114 men with NGU and 64 without NGU	Culture	No	Ureaplasmas were not associated with acute NGU. Ureaplasmas were associated with NGU during follow-up. Ureaplasmas were associated with chronic NGU.

<sup>a</sup>qPCR, quantitative PCR; NCNGU, nonchlamydial nongonococcal urethritis; NA, not applicable.



**FIG 1** Proposed natural history of *U. urealyticum* urethral infection in men following initial exposure. A hypothetical scenario in an immunologically naive male when exposed to *U. urealyticum* for the first time is described. The lack of prior exposure to *U. urealyticum* results in an increased bacterial titer and subsequent polymorphonuclear neutrophil influx (signs of infection) with accompanying symptoms. Depending on the adaptive immunological response to *U. urealyticum*, the infection may clear without intervention or result in persistent urethral colonization. With an increase in the number of sexual contacts, the presence of an adaptive immune response is able to keep the titer of any newly acquired *U. urealyticum* bacteria below the threshold which results in inflammation. In the absence of an adaptive immune response, signs and symptoms may be present again. Persistent urethral colonization may result in a commensal-like association with the host, accounting for the high prevalence among healthy individuals, or, alternatively, may result in the factors which are associated with the development of infertility.

*urealyticum* and therefore may have overrepresented this species among clinical samples from both cases and control groups. The legacy of the original nomenclature is still evident today, with publications still referring to *U. urealyticum* or just *Ureaplasma* spp., and may be partially attributed to the use of culture-based rapid diagnostic kits which are commercially available (35). Over time, studies have begun to differentiate the *Ureaplasma* spp. found into the respective species, with some studies identifying the presence of *U. urealyticum* more often among NGU patients than among controls, whereas the inverse of this was found for *U. parvum* (17, 21, 36–39). In many of these studies, patient numbers in both NGU and control groups were seen to be low and therefore lacked the power to confidently associate *U. urealyticum* with NGU. For this reason, Zhang et al. performed a meta-analysis which included seven eligible case-control studies encompassing 1,507 NGU patients and 1,223 controls from four separate continents (40). The findings identified that *U. urealyticum* was more prevalent among NGU patients than among controls and that *U. parvum* was significantly more associated with the control group than with those with NGU. This analysis gave significant weighting toward the idea that *U. urealyticum* is the most commonly associated species of *Ureaplasma* among NGU patients and presents as a substantial risk factor for the development of disease, although some studies have found the inverse result (41).

Due to the link between species and disease, it is essential that future studies differentiate *Ureaplasma* spp. from clinical samples to the species level using molecular methods to aid in epidemiological studies, which will either support or refute the role

of these bacteria in the development of NGU. One of the limiting factors inhibiting this is the use of culture-based rapid diagnostic tests available commercially, which are able to yield semiquantitative data with regard to the titers within a sample, as well as an indication of antibiotic susceptibility, but which are of limited diagnostic use in the instance of NGU due to the failure to differentiate between species (35). If reference or research facilities are accessible, molecular-based techniques are available which can differentiate the two species based on the size of the amplicons generated following a PCR that targets the 5' region of the *mba* gene or that uses real time-based molecular probes (42–44). Additionally, multiplex molecular assays are also commercially available and may play an important role in the species-level identification of *Ureaplasma* spp., alongside more traditional pathogens responsible for sexually transmitted infections (STI). As discussed below, the presence of *Ureaplasma* spp. alone may not be sufficient to warrant clinical intervention and the results of such tests being reported back to the clinician may result in inappropriate use of antibiotics, and therefore caution should be taken in reporting and/or interpretation of these results.

### **Risk Factor 2: the Sexual History of the Patient**

It would be very simplistic to assume that the species of *Ureaplasma* was the sole differential which accounts for NGU among men and the inconsistency in reporting in previous cases. A fascinating insight most likely relating to the immune response of the host and the transition from pathogen to commensal has instead been indicated. For many STIs, the risk of symptomatic disease is proportional to the number of sexual partners (36). In contrast to this, the relationship between *Ureaplasma* spp. and the number of sexual partners is inversely correlated (17, 45). Some of the pioneering work in this area was identified by Wetmore et al., who examined 329 patients with NGU, defined as  $\geq 5$  PMNs per high-powered field and/or a visible discharge (17). In this study, two control groups, consisting of 191 attendees to a sexually transmitted disease (STD) clinic and 193 patients who were attending the emergency department who did not have NGU, were also assessed. Upon initial analysis, *U. urealyticum* was only marginally associated with NGU compared with the association found for the STD control group (adjusted odds ratio, 1.6) or the emergency department group (adjusted odds ratio, 1.7), but when the analysis considered the number of sexual partners, the adjusted odds ratio rose to 2.9 for the STD group and 3.2 for the emergency department group when focusing on less than 10 vaginal partners. This association was even greater when the number of vaginal partners was restricted to less than five, with the adjusted odds ratios increasing to 6.2 and 5.2, respectively. When the same analysis was performed on patients positive for *U. parvum*, there was no association in any group, adding further weighting to the argument to differentiate the species of *Ureaplasma*. A similar finding was noted by Frolund et al., who examined a Danish cohort of 211 NGU patients and 73 asymptomatic controls (45). Again, a similar finding was observed, with the increase in the number of sexual partners being associated with a reduced likelihood of disease.

These studies suggest that *Ureaplasma* species infections resulting in NGU are associated with patients with fewer sexual contacts. At first, this may seem counterintuitive, considering that other sexually transmitted infections are positively associated with the number of sexual contacts, which therefore represents a significant risk factor. The scenario with *Ureaplasma* spp. suggests a significant role for the adaptive immune system in the presentation of disease. Early work by Brown et al. examined the serological response to *Ureaplasma* spp. among NGU patients in acute- and convalescent-phase serum (46). They noted that a change in antibody titer was identified in 68% of patients, and greater than 80% of these patients saw a change in IgM titer, suggesting an active infection. When examining the titers of IgG and IgA, the immunoglobulins responsible for protective immunity, only 10% of patients had an increase in titer. When the data were stratified by prior NGU or not, there was no significant difference in IgG levels in either acute- or convalescent-phase serum, whereas prior NGU accounted for a greater IgA response. These data suggest that some



patients gained protective immunity following a previous NGU, whereas others did not. This ability to develop protective immunity may account for why some individuals do not develop NGU on future reexposure, whereas others may.

### **Risk Factor 3: the Bacterial Load of *Ureaplasma* spp. within the Male Urethra**

The third significant risk factor linking *U. urealyticum* and, in some cases, *U. parvum* to the development of NGU is the bacterial load within the urethra. As mentioned previously, *in vitro* stimulation of monocytic cell lines with *U. urealyticum* resulted in a dose-dependent response between the *U. urealyticum* titer added and mRNA and cytokine production for the proinflammatory cytokines IL-6 and TNF- $\alpha$  (33). This *in vitro* evidence is supported by clinical findings from a number of studies (45, 47, 48). Frolund et al. reported that in the presence of *U. urealyticum* at concentrations of  $\geq 1.3 \times 10^3$  genome equivalents/ml of urine, corresponding to approximately  $1 \times 10^3$  bacteria/ml, there was a significant association with the development of NGU (45). This figure was similar to that in earlier papers which looked at cutoff points for the bacterial load for both *U. urealyticum* and *U. parvum* (47, 48). It is important to note that the study by Deguchi et al. reported that 83% of subjects which were positive for *U. parvum* had less than  $5 \times 10^3$  bacteria/ml urine, and 80% of these subjects had less than 12.5 leukocytes/ml (48). This gives further weighting to the idea that *U. parvum* is less proinflammatory, but in situations in which high titers of *U. parvum* are present, it has the capacity to generate an inflammatory response. An observation by Frolund et al. ties in risk factor 2 (sexual history of the patient) with risk factor 3 (bacterial load) (45). Analysis of their cohort identified that as the number of vaginal sexual partners increased, the load of *Ureaplasma* isolated decreased, with a predicted drop by 2.2% with each additional sexual partner. It is conceivable that the host immune response which develops due to multiple exposures to *Ureaplasma* spp. may have a direct impact on keeping the titer of *Ureaplasma* spp. low and therefore under the threshold to mount a significant proinflammatory response like that which would result in a PMN response, but these findings need to be expanded by future studies.

The data from the studies presented here suggest that a simple qualitative result would not be enough to predict a causal relationship between the presence of *Ureaplasma* spp. and NGU, a factor which has been overlooked by previous studies. From these data it may be possible to develop an objectively determined titer of *Ureaplasma* spp. which clinical laboratories could use to differentiate between causality and association.

### **IMPACT OF UREAPLASMA SPP. ON MALE FERTILITY**

The link between sexually transmitted pathogens and a negative impact on fertility in females is well established. In males, however, such a link is less well defined, yet it has been controversially suggested by some that *Ureaplasma* spp. may be associated with male infertility (Table 2). The chronic and often asymptomatic carriage of *Ureaplasma* spp. may therefore have important implications on the development and progression of infertility among men. In this section, we discuss the studies which have contributed to this argument and examine the clinical and mechanistic studies which have contributed to the argument for a causal role of *Ureaplasma* spp. in impaired male fertility.

#### **Clinical Studies Associating *Ureaplasma* spp. with Infertility**

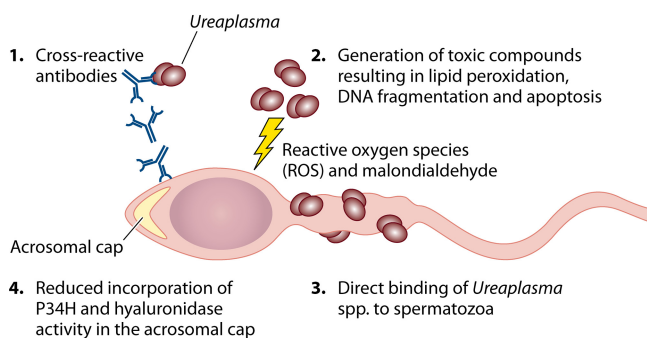
As discussed above, *Ureaplasma* spp. can be found in the urethra of seemingly healthy males, but they have also been isolated from expressed prostatic secretions, urine following prostatic massage, and prostate tissue (49–53). This colonization therefore permits a source to contaminate semen during ejaculation and serves as a means to impact male fertility.

Many studies have examined the clinical association between the presence of *Ureaplasma* spp. in men who are infertile and that in a control group of men without signs of infertility (53–57). In one relatively small study of 100 infertile and 100 control

**TABLE 2** Published studies examining the relationship between *Ureaplasma* spp. and male infertility

Authors (yr)	Reference	Country of study	Patient group	Sample type	No. of participants	Method of identification	Species differentiated	Key findings relating to <i>Ureaplasma</i> spp.
Huang et al. (2016)	57	China	Men attending a reproductive center	Semen	19,098 infertile men and 3,368 fertile men	Culture	No	<i>Ureaplasma</i> spp. were significantly associated with infertility. <i>Ureaplasma</i> spp. were significantly associated with reduced motility and normal forms in infertile men compared with fertile controls. <i>U. urealyticum</i> was significantly associated with infertility. <i>U. parvum</i> was not associated with infertility.
Huang et al. (2015)	56	Multiple countries	Meta-analysis	NA <sup>a</sup>	611 infertile men and 506 fertile men	NA	Yes	<i>U. urealyticum</i> was significantly associated with infertility.
Zhang et al. (2014)	54	China	Men attending an infertility clinic	Semen	223 infertile men and 146 fertile men	Culture	Yes	<i>U. urealyticum</i> was significantly associated with infertility compared with <i>U. parvum</i> . Semen positive for <i>U. urealyticum</i> showed a decreased concentration of spermatozoa and decreased motility. Ureaplasmas were found more frequently among samples from infertile men (10.8%) than among those from fertile men (5.7%). Ureaplasmas were detected significantly more often in semen from infertile men than in semen from controls. <i>U. urealyticum</i> was detected in 9% of infertile men vs 1% of control men. <i>U. parvum</i> was detected in 3% of infertile men vs 2% of control men.
Abusarah et al. (2013)	55	Jordan	Men attending a urology clinic	Semen and first-void urine	93 infertile men and 70 fertile men	PCR	Yes	
Zeighami et al. (2009)	53	Iran	Men attending an infertility center	Semen	100 infertile men and 100 fertile controls	PCR	Yes	

<sup>a</sup>NA, not applicable.



**FIG 2** Mechanisms associated with *Ureaplasma* species-induced infertility in men. A number of mechanisms have been proposed to account for the clinical observational studies showing decreased fertility among men who experience urethral colonization with *Ureaplasma* spp. These include (i) cross-reactivity of host-generated antibodies against the UreG protein of *Ureaplasma* spp. to the autoantigenic sperm protein; (ii) the generation of toxic compounds, such as reactive oxygen species (ROS), which contributes to lipid peroxidation, DNA fragmentation, and subsequent apoptosis; (iii) direct binding to spermatozoa, which may result in reduced motility; and (iv) reduced incorporation of P34H and hyaluronidase activity in the acrosomal cap, which may reduce the capacity of spermatozoa to penetrate the egg.

individuals, the authors identified 12% of infertile men but only 3% of fertile men to be *Ureaplasma* species positive by PCR (53). The individuals which were *Ureaplasma* species positive and infertile had significantly lower volumes of seminal fluid, lower concentrations of sperm cells, and higher levels of sperm cells with an abnormal morphology than the *Ureaplasma* species-negative infertile patients. Of significance was the finding that the frequency of *U. urealyticum* in the semen of infertile men was higher (9%) than that in the healthy controls (1%), whereas the frequency of the presence of *U. parvum* was 3% in the infertile group and 2% in healthy men, suggesting that *U. parvum* may not have a causal role in infertility in this patient group. In a fate similar to that of previous NGU studies, many studies have neglected to differentiate between species (and in one instance, in excess of 19,000 samples were examined), and although a significant negative impact of *Ureaplasma* spp. on semen quality was identified, further power may have been afforded if species-level discrimination had been conducted (57, 58). To further investigate the species-specific association with infertility, a meta-analysis which examined 14 studies comprising 611 cases and 506 controls was conducted (56). These studies suggested an association between *U. urealyticum* and a negative impact on fertility, whereas there was little evidence for a role of *U. parvum*, which draws a parallel to the findings for NGU patients discussed above.

### Proposed Mechanisms of *Ureaplasma* Species-Associated Infertility

It is important to discuss the proposed mechanisms behind these observations that *Ureaplasma* spp. contribute to infertility among men (Fig. 2). Sexually transmitted pathogens are known to affect sperm quality by reducing motility, negatively affecting sperm morphology, and inducing apoptosis (5). Several mechanisms have been proposed to account for infertility in men because of *Ureaplasma* species colonization. These include the direct binding of *Ureaplasma* spp. to spermatozoa, which may impede swimming motility (59–62); the production of toxic metabolites, which can damage spermatozoon membranes and result in DNA fragmentation (54, 63); as well as the host generation of cross-reactive antibodies between *Ureaplasma* spp. and sperm surface proteins (64–66).

Work by Potts et al. identified 17 out of 50 chronic prostatitis patients to be positive for *Ureaplasma* spp., and the levels of reactive oxygen species (ROS) were significantly higher among the *Ureaplasma*-positive infertile patients than among the *Ureaplasma*-negative infertile patients and the control group (63). ROS have the potential to induce lipid peroxidation, therefore compromising the integrity of sperm membranes and leading to impaired fertilization capabilities. Of interest was the finding that only 1 out

of 17 of the positive samples in which ROS levels were elevated showed signs of leukocytospermia, suggesting that in some cases traditional signs of prostatitis, such as leukocytospermia, may not be indicative of *Ureaplasma* species infection. The potential for *Ureaplasma* spp. to contribute to lipid peroxidation through the generation of ROS, as well as malondialdehyde formation, by either *U. urealyticum* or *U. parvum* was further developed and stratified (54). The levels of ROS, malondialdehyde formation, and DNA fragmentation were all significantly higher in *U. urealyticum*-positive samples than in *U. parvum*-positive samples. The high levels of ROS could therefore result in DNA fragmentation and subsequent apoptosis (67).

Some studies have suggested that the presence of *Ureaplasma* spp. in seminal fluid has no real impact on semen quality (68, 69). One possibility is that the mechanism associated with infertility is one which cannot be identified by classic markers of infertility but is one which may impact the interaction between the sperm and the egg. P34H is a key membrane-bound protein which is essential for sperm-zona pellucida interactions. P34H is incorporated into membranes covering the acrosomal cap as it transits across the epididymis and therefore can serve as a marker for epididymal function (67). The levels of P34H were significantly lower in the *Ureaplasma* species-positive group than in the control groups, as determined by Western blotting and immunofluorescence imaging, which identified that 38% of the sperm in the *Ureaplasma* species-positive group and 73% of the sperm in the control group had P34H. These data suggest a potential impact of chronic asymptomatic infection of the epididymis. The acrosomal cap also contains the enzyme hyaluronidase (HYD), which is essential for the sperm to penetrate the egg. In the *Ureaplasma* species-positive group, the levels of HYD activity were significantly different from those of the infertile *Ureaplasma* species-negative group and the fertile controls (67). By reducing the activity of HYD, the likelihood of successful sperm penetration into the egg is therefore reduced.

An alternative mechanism to *Ureaplasma*-related infertility is the development of cross-reactive antibodies to human sperm membrane proteins following exposure to *Ureaplasma* spp. (64–66). Shi et al. demonstrated that antibodies raised against the UreG protein of *Ureaplasma* spp. were able to cross-react with human nuclear autoantigenic sperm protein (NASP) (66). A higher titer of anti-UreG antibody was found in the serum of the infertile men than in that of the fertile controls. In an *in vitro* fertility assay, sperm which had been pretreated with anti-UreG antibodies had significantly lower binding and fusion to eggs than nontreated control sperm.

The evidence presented here suggests that the impact of *Ureaplasma* spp. on male infertility is similar to that described for NGU, although unlike in the context of NGU, the effect of the bacterial titer has yet to be investigated. The lack of species differentiation has hindered studies, but association as well as mechanistic studies are pointing toward a potential for *U. urealyticum* to be the primary *Ureaplasma* spp. associated with infertility in men. Furthermore, some of the traditional markers for infertility may not indicate a causal role for *Ureaplasma* spp. in male infertility.

### TREATMENT OF GENITAL TRACT INFECTIONS IN MEN CAUSED BY *UREAPLASMA* SPP.

A position statement from the European STI Guidelines Editorial Board states that routine testing of asymptomatic or symptomatic men for the presence of *Ureaplasma* spp. is not recommended; however, one of the key messages that the authors made states that *Ureaplasma urealyticum* at high bacterial loads can cause a small proportion of cases of male NGU (3 to 11% of NGU cases) (4). The authors also noted that NGU caused by *U. urealyticum* was more likely to develop in younger men and men with fewer lifetime sexual partners. They highlight that there is a paucity of well-designed large controlled studies which investigate the role of *Ureaplasma* spp. in STI syndromes and NGU.

In the light of mounting evidence of a causal role for *Ureaplasma* spp. in the development of NGU and male infertility, the question remains whether we should treat individuals who are *Ureaplasma* spp. positive with symptoms. Currently, a position statement from the European STI Guidelines Editorial Board does not recommend

routine testing or treatment of either asymptomatic or symptomatic men for any *Ureaplasma* spp.; however, this position statement also suggests that *U. urealyticum* is causal in up to 11% of NGU cases, which contradicts this recommendation (4). The evidence presented in this review suggests that in a subset of men with symptoms of NGU and the absence of other etiological factors, a risk-based approach could be used to guide the treatment of these patients. For example, symptomatic NGU patients with the absence of other sexually transmitted infections, younger age, a low number of partners, and high titers of *U. urealyticum* may benefit from treatment. Currently, it is clinically difficult to implement a risk-based approach in countries such as the United Kingdom, as sexual health care settings do not widely test for *Ureaplasma* spp., and when testing is done, it almost never involves the differentiation of *U. urealyticum* and *U. parvum* or determination of the bacterial load.

Currently, guidelines set out by the British Association for Sexual Health and HIV (BASHH) suggest treatment of a first episode of NGU with a 7-day course of doxycycline at 100 mg twice daily or, if contraindicated, azithromycin at 1 g *stat.*, followed by 500 mg once daily for 2 days, or ofloxacin at 200 mg twice daily or 400 mg once daily for 7 days (70). With recurrent episodes of NGU where the possibility of reinfection has been excluded, the recommended first-line regimen is azithromycin at 1 g *stat.* and then at 500 mg once daily for the next 2 days plus metronidazole at 400 mg twice daily for 5 days. If symptoms still persist, treatment with moxifloxacin at 400 mg once daily for 10 to 14 days plus metronidazole at 400 mg twice daily for 5 days is the recommended regimen. It is considered reasonable to provide epidemiological treatment to the partners of men with NGU using the same antimicrobial regimen that resulted in cure in the index case. In practice, if *Ureaplasma* spp. were present and responsible for the symptoms of NGU, the first-line treatment recommended (a short course of doxycycline at 100 mg twice a day for 7 days) would be adequate to treat it in the United Kingdom at the moment, in light of the low levels of *tet(M)*-mediated doxycycline resistance among these organisms (71). Any decision to treat would need to be carefully weighed with the risk of inappropriate prescribing. Antibiotic resistance, in particular, azithromycin resistance, among recognized GUM pathogens, such as *M. genitalium* and *N. gonorrhoeae*, is of growing concern (72, 73).

## CONCLUSIONS

The role that *Ureaplasma* spp. play in the development of genitourinary medicine-related infections is still a controversial area for many, but there is mounting evidence that these bacteria, especially *U. urealyticum*, have a causative role in infection under very specific conditions. *Ureaplasma* spp. are by no means a leading cause of NGU, but this is not to say that they do not contribute to cases which are currently classified as idiopathic; as such, these patients are no less deserving of attention or correct management. Furthermore, the role of *Ureaplasma* spp. in the development of infertility among men is beginning to be recognized, but further work exploring the mechanism, as well as appropriate criteria for identifying patients with *Ureaplasma* species-induced infertility, is required.

*Ureaplasma* spp. have a proven proinflammatory capacity in cell lines and in animal and human models of disease, but the species of *Ureaplasma* spp., the sexual history of the patient, and the titer of bacteria present all appear to be key risk factors for the development of disease. A large prospective case-controlled study considering the species and load of *Ureaplasma*, the presence of other microorganisms, the number of PMNs as a marker of inflammation, and the number of sexual partners will be crucial to confirm or refute the role of *Ureaplasma* spp. in the development of NGU in men. If a clear link is identified, then current qualitative diagnostic methods may not be appropriate for determining a causal role for *Ureaplasma* spp. in cases of NGU. In the meantime, in light of the evidence presented in this review, we recommend that among cases of symptomatic NGU in which classic etiological agents have been ruled out, a risk-based approach taking into account patients with a younger age, a low number of partners, and high titers of *U. urealyticum* should be considered for treatment.

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