

Biology of trichomonosis

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Trichomonas vaginalis is emerging as a major pathogen of men and women and is associated with serious health consequences. Advances in diagnosis and treatment are presented. The complexity of trichomonad pathogenesis is illustrated in the interaction of this parasite with human cells, tissues and the immune system. It is now becoming evident that the interaction of trichomonads with the host is frequently modulated by environmental signals. The molecular biology of trichomonads is still in its infancy, but analysis of genes, genomic structure and transcriptional mechanisms suggest that trichomonads combine both prokaryotic and eukaryotic features. Evidence for the ancient divergence of trichomonads from other eukaryotic lineages is discussed. *Curr Opin Infect Dis* 13:37–45. © 2000 Lippincott Williams & Wilkins.

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Current Opinion in Infectious Diseases 2000, 13:37–45

Abbreviations

SLPI secretory leukocyte protease inhibitor
INR metazoan initiator element

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0951-7375

Introduction

Trichomonas vaginalis, despite its status as a stepchild of sexually transmitted infections, has numerically never been a minor sexually transmitted infection [1•]. Primarily considered to be a cause of vaginitis in women, trichomonads have been frequently dismissed as a nuisance parasite. Data, although slowly forthcoming, however, paint a very different picture. Recent publications indicate that this parasite's impact is not only limited to vaginitis but is also a major factor in promoting transmission of HIV [2•,3•], in causing low-weight and premature birth [4], and in predisposing women to atypical pelvic inflammatory disease [5], cervical cancer [6–8] and infertility [9]. This parasite may also contribute to long-lasting morbidity of women and children born to infected mothers [1•]. Finally, the recent finding of high rates of trichomonosis among men highlights the tremendous impact this parasite has on all sexually active persons, regardless of gender [10••]. Clearly, these new trends warrant a new interest and renewed efforts in understanding the biology of pathogenic trichomonads. The emerging themes of trichomonosis research over the past 2 years are thus the focus of this review. For a more comprehensive overview of the field, the reader is directed to the excellent recent reviews by Petrin *et al.* [11••] and Alderete [12••].

Clinical aspects of trichomonosis

Trichomonosis is now the most common non-viral sexually transmitted disease in the world [13••]. Trichomonal vaginitis has been reported in 50–75% of female sex workers, in up to 50% of female inmates, in 10–25% of women attending gynecological, sexually transmitted disease or family planning clinics and in 13% of pregnant women in the US [14–17]. Prevalence rates in sub-Saharan Africa have been reported to be as high as 46% in women of childbearing age [10••]. World wide, 107 million cases of trichomonosis are reported with the majority (92%) occurring in women [13••]. It should be noted that the incidence of trichomonosis in men is most likely grossly underestimated due the insensitivity of wet-mount preparations [18•,19]. Using a combination of wet-mount and polymerase chain reaction a recent study noted a prevalence of 20.8% and 12.2% in symptomatic and asymptomatic men, respectively [10••]. These surprising results highlight that trichomonosis can no longer be considered a cause of morbidity of women alone. It is estimated that annually 167 million people (84 million men and 83 million women) will become infected with *T. vaginalis*; of these 10–50% of women and 15–50% of men will be asymptomatic [13••].

Interestingly, the prevalence of trichomonosis in African American populations in the United States is significantly higher than that of any other ethnic or racial group [20]. No biological or behavioral reasons were known why ethnicity alone would alter the risk for acquisition of trichomonosis. A revolutionary study by Lauman and Yoosik-Youm may have solved this enigma [21••]. In their remarkable study, these authors found that the patterns of sexual networks can explain the increased risk for sexually transmitted infections among ethnic groups. In the case of African Americans, increased sexual contact between the core group and the periphery leads to the spread of infection into the population, whereas sexual segregation from other racial groups results in the maintenance of infection within this population [21••,22•].

Infection with *Trichomonas vaginalis* has major health consequences for women, including predisposition to HIV, association with cervical cancer and pelvic inflammatory disease, and complications of pregnancy [11••]. The vaginal epithelium is the primary site of infection. A hallmark and complication among patients with trichomonosis is the wide variations in symptomatology [15,23,24]. While some patients are characterized as asymptomatic, women with trichomonal vaginitis (70%) frequently experience vaginal discharge due to a leukocytic infiltration. The consistency of the discharge varies from patient to patient from thin and scanty to profuse and thick. The classical symptom of profuse frothy yellow discharge, however, occurs in only 10–30% of women. Similarly, the vagina and cervix of patients with trichomonosis may be erythematous and edematous, with general erosion of the cervical epithelium and punctate hemorrhages on the cervical wall, termed 'colpitis macularis' or 'strawberry cervix'. Although highly specific for trichomonosis, the strawberry cervix is nonetheless present in only a few (2–5%) women. Sharp abdominal pain has been documented among some patients with trichomonosis and may be indicative of infection in the upper urogenital region, possibly involving regional lymphadenopathy and salpingitis [25••].

In contrast to women, men become infected following recent contact with an infected partner but, for unknown reasons, may have a self-limiting infection [24]. Non-gonococcal non-chlamydial urethritis is the most common complaint of men infected with *T. vaginalis*. Incidence rates of trichomonal urethritis appear to be on the rise, especially in geographic areas where gonococcal infections are on a decline [1•]. The presence of trichomonads in the prostate gland in men with trichomonosis has been reported [26]. However, the role of infection in prostatitis and/or in infertility or numerous other reported sequelae is unclear.

Syndromic patient management algorithms of sexually transmitted disease syndromes have been used successfully for the diagnosis of many sexually transmitted infections but have performed poorly for *T. vaginalis* [27•]. Nevertheless diagnosis of trichomonosis is still often based on clinical presentation and especially the characteristics of the vaginal discharge [18•,28]. However, symptomatology and the presence of the classical signs of trichomonad infections are usually poor predictors of disease [18•]. A newly developed algorithm based on risk assessment, symptoms, speculum examination and microscopy with a sensitivity of 92.4% and a 100% positive predictive value, however, offers hope that syndromic management of *T. vaginalis* infections can be significantly improved [29].

Positive diagnosis of trichomonosis is generally established by wet-mount examination, but this method is only 30–80% sensitive compared to the gold standard of culture. Culturing is laborious, frequently inaccessible and often cost prohibitive for many clinical settings, and thus is not routinely used [5,30]. The InPouchTV culture system, equally sensitive as traditional culturing in many clinical settings, has been shown to be a cost-effective and sensitive alternative [31•]. Cost of diagnosis can be further reduced, by delaying inoculation into culture media, until wet-mount results become available. No significant difference in the sensitivity of the InPouchTV culture system between delayed and immediate inoculation of culture medium has been demonstrated [32•].

Other diagnostic techniques such as enzyme-linked immunosorbent assay [33], hybridization [34] and fluorescent antibody tests [19] have been used to detect this parasite and have reported sensitivities between 70% and 90%. The advent of polymerase chain reaction opened a new avenue for diagnosis. Several polymerase chain reaction diagnostic tests with sensitivities and specificities approaching 100% have been recently developed [18•,35]. Importantly, polymerase chain reaction diagnostics are easily adaptable to self-administered sampling, which in turn is useful due to the ease of specimen collection and patient compliance. In this regard, the tampon test has proved particularly useful. This test not only had high sensitivity and specificity, but was even useful on specimens where culture and wet mount failed to detect any trichomonads [36•,37•]. With further developments in polymerase chain reaction based diagnostics, speculum-free evaluation of vaginal infections should become a reality in the near future [38•].

Treatment options are still limited but exciting progress in the biochemical characterization of trichomonads hint at the development of new drugs, targeting novel and

unique trichomonad structures [39,40*,41*,42**,43**,44, 45,46*,47*,48,49,50*,51*]. Metronidazole (Flagyl) or derivatives thereof are the most commonly used drugs. Treatment consists of either a single 2 g dose of metronidazole or 400 mg of metronidazole twice daily for 5–7 days given orally [25**]. Vaginal application of metronidazole may provide symptomatic relief but will not cure the infection [52*]. In the past, metronidazole was contraindicated during pregnancy because of the concern about possible teratogenic effects [53–55]. The strong association of trichomonosis with premature delivery and low-birth weight, however, suggests the desirability of treatment. Indeed, the preponderance of evidence shows that the benefits of metronidazole treatment outweigh by far the risks of using this drug. First, teratogenic effects of metronidazole in animal species have not been demonstrated. Second, several studies have shown no or a very slight risk of metronidazole treatment during pregnancy [53–55]. Finally, a large study of 1706 pregnant women given metronidazole showed conclusively that oral metronidazole treatment during pregnancy is not significantly associated with congenital abnormalities [56**]. In spite of overwhelming medical evidence of the safety of metronidazole, treatment during the first trimester must be considered carefully because of possible legal liability.

Treatment failure is not uncommon and may be the result of non-compliance, reinfection or colonization with a metronidazole-resistant isolate of trichomonad. It appears that, at least since 1995, the majority of clinically resistant trichomonosis results from metronidazole resistant isolates [57]. Furthermore, the incidence of metronidazole-resistant trichomonosis may also be increasing [11**]. Treatment of such cases is often problematic. In many cases increased dosages of metronidazole may resolve the infection. Otherwise, a variety of treatment regimes have been utilized that may or may not work [58*,59]. A clear systematic and scientific examination of treatment alternatives does not exist.

It is clear that there is a need for a safe, systemic and non-imidazole based drug alternative for pregnant women, people with metronidazole-resistant trichomonosis, or for patients having allergic reactions to this group of drugs. Several promising leads toward this goal have recently been reported. Firstly, several new 5-nitroimidazole and 1-lactam-substituted nitroimidazole compounds have been shown to be highly effective against metronidazole-resistant trichomonads [60*]. Secondly, disulfiram and its first mammalian metabolite ditiocarb, both used in alcoholism treatment, have also been shown effective against metronidazole resistant trichomonads *in vitro* [61*]. Finally, the findings of alternative trichomonad keto-acid oxidoreductases [41*],

expression of chitin at the trichomonad cell surface [62*] and expression of a thymidine kinase [50*] may lead to the design of new antitrichomonal drugs. Particularly intriguing and promising is the ability of deoxyuridine analogs to inhibit *T. vaginalis* thymidine kinase, and consequently prevent proliferation of this parasite [50*].

Trichomonas vaginalis and HIV

The correlation between trichomonosis and increased risk of acquiring HIV (up to 5-fold) has been firmly established [63–68]. Even at such a modest increased risk, HIV community transmission can be increased significantly because of the high prevalence of trichomonosis within the general population, and in particular within risk groups [2*,69*]. Presently, the increased risk appears to be present for the receptive partner only. The mechanisms of acquisition enhancement are unknown, but suggestions have included erosion of the mucosa by the parasite and/or the inflammatory response, and recruitment of HIV target cells into the genital area [70*]. The complexity of these associations, however, is now becoming clear. Not only do trichomonads disrupt the epithelial barrier but also appear to recruit CD4 T-lymphocytes to the genital area [71*]. Increased secretion of cytokines (interleukins 1, 6, 8 and 10), known to increase HIV susceptibility, has now been demonstrated during trichomonosis [72,73]. Furthermore, the degradation of secretory leukocyte protease inhibitor, known to prevent HIV transmission through the oral mucosa, by trichomonad proteinase is yet another mechanism to enhance HIV transmission [42**]. Another important and unexpected factor in the risk for HIV acquisition is the demonstration of significantly increased viral loads in semen of men with trichomonal urethritis [10**]. Clearly, these new findings strongly suggest that aggressive screening and treatment for trichomonal infection of men and women in risk groups may significantly reduce community transmission of HIV.

Interactions between trichomonads and host cells

Receptor–ligand-type interactions are involved between trichomonads and host epithelial cells. The surface structures on vaginal epithelial cells recognized by *T. vaginalis* organisms are unknown. Four trichomonad surface proteins have been identified as mediating cytoadherence. The synthesis of the four adhesins was coordinately up-regulated by binding to epithelial cells and by iron [74–76,77*].

More recent work reveals that three of the four adhesins studied to date are each members of multi-gene families [78*]. Sequence analyses at both the nucleotide and amino acid levels revealed structural similarity of adhesins with known metabolic enzymes [79,80]. Analysis of the receptor-binding epitope for the adhesin AP33 identified

a 24-amino acid binding domain with the ability to inhibit parasite associations with host cells [43**].

The involvement of carbohydrate structures in cytoadherence has also been suggested, but remains problematic. A quantitative electron microscopy study showing the involvement of trichomonad surface-associated glycoconjugates in cytoadherence raises the possibility of the presence of lectin-like adhesins in *T. vaginalis* [81*]. A similar study, however, demonstrated that cell surface glycoconjugates are not involved in cytoadherence [82*]. The use of different target cells may be an explanation for these contradictory results. It should be noted that the protein adhesins are functional when tested with in-vivo relevant target cells.

Initial binding to host cells is followed by transformation of the parasite from its ellipsoid to amoeboid form [83]. The ability to undergo such a dramatic morphological transformation clearly requires a rearrangement of the cytoskeleton. Alpha-actinin, an actin binding protein that participates in rearrangements of actin fibers, has now been described in trichomonads. Its role in transformation is suggested by the finding that alpha-actinin, usually distributed throughout the cytoplasm, becomes localized to the periphery upon the change to the amoeboid form [84*].

Amoeboid transformation results in intimate contact of the parasite with the host cell. Extensive interdigitation and microchannel formation has been observed by electron microscopy [83]. Using a variety of electron microscopic techniques Furtado and Benchimol [85*] have extended these observations to include extensive plasma membrane fusion between *T. vaginalis* and vaginal epithelial cells. Although the roles of contact-dependent cytotoxins [86-89], proteinases [90] and phospholipase [91] have been well established in the lysis of host cells, membrane fusion may yet represent another mechanism leading to host cell damage.

It is reasonable to expect that cytoadherence to vaginal epithelial cells is not the only mechanism by which trichomonads adhere to host cell surfaces. For example, erosion of the vaginal epithelium as seen for colpitis macularis may allow access of parasites to the basement membrane and accompanying complex structures. Interestingly, the reports on the specific binding by *T. vaginalis* organisms to fibronectin [92] and laminin [92,93] may reflect associations with basement membrane sites. Importantly, the interaction of trichomonads with these extracellular matrix proteins appears to be mediated through the same binding domains as those utilized by the host. In the case of laminin, the amino acid sequence YIGSR is recognized by *T. vaginalis*, as it is by host cells [94*].

Finally, colonization of the vaginal epithelium requires trichomonads to breach the mucous covering. The interaction of trichomonads with mucus and subsequent penetration and parasitism of the underlying vaginal cells have been shown to be very complex. Mucin adhesins, proteinase activity and flagellar movement all appear to be necessary to traverse the mucous layer [95*].

Evasion of host immune responses

Cellular and humoral immune responses are evident in patients with trichomonosis, but are not protective [11**]. Although not found in all patients with trichomonosis, increased numbers of polymorphonuclear leukocytes can be detected readily in secretions [96]. Interestingly, while both leukocytes and macrophages in addition to antibody [97] and complement [98,99] can eliminate the parasites, it is clear that *T. vaginalis* has effectively neutralized the host immune surveillance system. Further, hydrogen peroxide-producing lactobacilli are considered protective normal vaginal flora [100*,101].

The numerous cysteine proteinases synthesized by *T. vaginalis* contribute significantly to immune evasion. The cysteine proteinases are cytotoxic [90] and hemolytic [88]. All subclasses of immunoglobulins are susceptible to the trichomonad cysteine proteinases [48,102,103]. Similarly, parasites are resistant to complement-mediated lysis by the action of at least one cysteine proteinase induced by high iron growth conditions. The proteinase degrades C3 deposited on *T. vaginalis* surfaces [98]. Secretory leukocyte protease inhibitor (SLPI) is yet another factor protecting mucosal surfaces. Again, trichomonad proteinases are able to degrade SLPI and render it non-functional. Interestingly, SLPI also has been shown to prevent HIV transmission, thus trichomonad proteinases may be partly responsible for the observed increase in risk of HIV acquisition in women with trichomonosis [42**]. Lastly, hydrogen peroxide readily neutralizes the cysteine proteinases, showing the protective effect of lactobacilli normal flora. However, displacement of the lactobacilli immediately following infection with *T. vaginalis*, through phagocytosis, may subvert this host protective effect [100*].

Leukocytic infiltration is frequently seen in women with trichomonosis. Although, leukocytes readily kill trichomonads *in vitro*, at parasite to leukocyte ratios of greater than 1:3000, they do not appear to be protective *in vivo*. However, at low parasite to leukocyte ratios, initial contact between parasite and leukocyte results in phagocytic pseudopod formation, internalization, and finally degradation of the immune cell in parasite phagocytic vacuoles [100*].

Phenotypic variation, another immune evasion mechanism for *T. vaginalis*, is defined on the basis of surface versus cytoplasmic expression of a repertoire of high M_r immunogens. A representative of these immunogens termed P270 has been extensively studied. From fluorescence experiments it has been observed that two types of isolates occur naturally during infections with *T. vaginalis*. Type I isolates were comprised of homogeneous populations of non-fluorescent trichomonads that synthesize and express P270 only in the cytoplasm. In contrast, type II isolates were heterogeneous with subpopulations of both fluorescent and non-fluorescent parasites and were able to switch between the fluorescent and non-fluorescent state [104–106].

The *p270* gene for one *T. vaginalis* isolate T068-II has now been sequenced [107*]. Consistent with earlier reports a significant portion of the *p270* gene has a 333-bp unit tandemly repeated 18 times that contains the epitope DREGRD detected by patient antibody. The non-repeat coding regions for the 5'- and 3'-ends are 69 nucleotides (23 amino acids) and 1183 nucleotides (395 amino acids), respectively. The start codon is immediately preceded by a 12-nucleotide sequence that has significant homology to the recently described Inr sequence of trichomonad promoters. A leader sequence and a transmembrane domain are also present. This gene appears to be highly conserved among trichomonads but the number of repeat units may vary [107*]. This single-copy gene was furthermore shown to be up-regulated by low iron conditions and highly phosphorylated under high iron conditions [108*].

Environmental adaptations

The vagina is one of the most complex sites of infection for a mucosal pathogen. This host environment is constantly changing under the influence of the menstrual cycle. The vagina is a nutrient-limiting site that cannot promote the 4–6-hour generation time seen during in-vitro growth of habituated parasites in a serum-based, trypticase and yeast extract complex medium [109]. It is thus not surprising that biological and biochemical properties of these in-vitro grown parasites may be different from those at the site of infection. Indeed, the availability of iron in growth media already has been shown to significantly affect the physiology of trichomonads [76,109,110]. Even a simple change in commonly accepted growth media appears to have a physiological consequence [111]. Further underscoring the tremendous adaptability of this parasite to challenging environmental pressures is the demonstration that trichomonads respond to changes in growth rate and carbon source with a change in mRNA levels and activity of glycolytic enzymes [112*]. In light of these findings, it may not be surprising that inconsistent results, depending on the growth condition under otherwise comparable experi-

mental set-ups, may be obtained. Even more important, these data also suggest that trichomonads possess signal transduction pathways that link changes in the environment with appropriate changes in transcriptional and post-transcriptional regulatory mechanisms. The exploration of such regulatory networks will be crucial in understanding this parasite–host interaction.

Molecular biology of trichomonads

Progress in our understanding of nucleic acid driven processes in *T. vaginalis* is excruciatingly slow. Despite this fact, several themes in this area of research seem to be emerging. First, all genes cloned from trichomonads are monocistronic. Second, genes are frequently present as multicopy genes, such as the trichomonad adhesins and the pyrophosphate-dependent phosphofructokinase. Third, homologues of genes have arisen either due to gene duplication, or are of polyphyletic origin. For example, the eukaryotic family of B DNA polymerases is suggested to have arisen by gene duplication; in contrast, the two forms of HSp70 may have arisen from two diverse origins. Finally, trichomonad genes appear to be devoid of introns. However, the discovery of a functional spliceosomal protein, PRP8, may indicate that at least some genes contain introns or that trichomonads may perform limited trans-splicing [113**].

New developments in this area have focused on transcription initiation and genomic structure. Despite considerable efforts, typical eukaryotic core promoter elements have not been found in trichomonads. An analysis of 33 genes has revealed that a consensus sequence resembling a metazoan initiator element (Inr) is located within 6–20 nucleotides upstream from the translational initiation codon. Transcriptional fusions and primer extension experiments established that the Inr invariably determines the start site of transcription. Interestingly, and in contrast to other metazoans, the trichomonad Inr is located very close to the translational start codon, resulting in unusually short 5'-untranslated regions for trichomonad mRNAs [114**].

The genomic structure of trichomonads has been examined by pulsed field gel electrophoresis. Six chromosomal elements that fall into three size classes have been identified. The size and number of chromosomes of 15 *T. vaginalis* isolates were found to be identical, suggesting that genetic heterogeneity among trichomonads does not involve major chromosomal rearrangements [115*].

Molecular phylogeny

The amitochondriate protist *Trichomonas vaginalis* has been suspected to be a member of the earliest branching eukaryotes. The finding of unusual trichomonad structures and biochemical pathways such as the unique post-

translational glutamylation of tubulins [116], the unique sulfur amino acid metabolism [46*], the spectroscopically atypical hydrogenosomal ferredoxin [45], the presence of novel iron superoxide dismutase [51*], and the centriole-like behavior of trichomonad flagella through the progression of the cell cycle [117*], all certainly confirm this assertion.

Phylogenies based on the sequence of the small rRNA subunit show that trichomonads emerge at the base of the eukaryotic tree [118,119,120*]. In contrast, phylogenies based on protein sequences often suggest a later divergence. Recently, phylogenetic analysis of DNA sequences of glyceraldehyde-3-phosphate dehydrogenase [121*], alpha-1 elongation factor [122*], DNA polymerase [123*], pyrophosphate-dependent phosphofructokinase [47*], heat-shock proteins [120*,124,125] and valyl-tRNA synthetase [126*] all support an early branching point for trichomonads. However, the presence of mitochondrial-like heat shock proteins (HSP60 and HSP70), in addition to the presence of nuclear-coded valyl-tRNA synthetase [126*], indicate that the branch point occurred some point after acquisition of the mitochondrial endosymbiont.

Unfortunately, most studies also show a lack of resolution in interphyla relationships among the early branching parabasalids, diplomonads and microsporidia. The reason for the inability to resolve clearly the evolutionary relationships among these organisms is unclear but may be due to recombination, lateral gene transfer, inversions, fusions, chimeric origins, gene duplication and convergent evolution. Data on alpha-1 elongation factor and HSP70, in addition, are also consistent with the notion of a simultaneous diversification of major eukaryotic lineages [120*,122*]. Importantly, taken together, these data strongly imply that the biochemistry of trichomonads is unique and sufficiently different from higher eukaryotes to provide ample targets for chemotherapeutic intervention.

Conclusion

Trichomonas vaginalis is a major pathogen of men and women, with tremendous health consequences for the infected individuals and their communities. Despite progress in our understanding of the evolutionary history, biochemistry, biology and pathogenesis of trichomonads that have led to the identification of many promising targets for drug development, treatment options are still very limited. A major reason for this is that preclinical and clinical studies are lagging behind these new and encouraging studies. A general lack of appreciation for the seriousness of this disease, once classified as a minor sexually transmitted infection, probably is yet another factor. This is particularly evident in the disturbing fact that the antiquated wet-

mount for diagnosis of trichomonosis is still widely used, despite the known and documented insensitivity of this test. It is obvious that an education program for medical professionals and the population as a whole is needed urgently.

Acknowledgements

This study was supported in part by Public Health Service grants G12-RR08124 and SO6-GM08012-26 (to M.W.L.) and AI-39803 and AI-43940 (to J.F.A.) from the National Institutes of Health.

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