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FATORES DE VIRUI	LÊNCIA DE CEPAS FUMANTES E NÃ		ns ISOLADAS DE		

CURITIBA 2007

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FATORES DE VIRULÊNCIA DE CEPAS DE Candida albicans ISOLADAS DE FUMANTES E NÃO-FUMANTES

VIRULENCE FACTORS OF Candida albicans STRAINS ISOLATED FROM SMOKERS AND NON-SMOKERS

Dissertação apresentada ao Programa de Pós - Graduação em Odontologia da Pontifícia Universidade Católica do Paraná, como parte dos requisitos para obtenção do Título de Mestre em Odontologia, Área de Estomatologia

Orientador: Prof. Dr. Edvaldo Antonio Ribeiro Rosa

Co-orientadora: Dra. Maria Ângela Naval Machado

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1. LISTA DE ABREVIATURAS E SIGLAS

C. - Candida

et al. - E outros (abreviatura de "et alli")

g - Grama

spp. - Espécies

MgCl₂ - Cloreto de magnésio

mL - Mililitro

mm - Milímetro

NaCl - Cloreto de sódio

°C - Grau Celsius

pH - Potencial hidrogeniônico

Pz - Zona de precipitação

q.s.p - Quantidade suficiente para atingir determinado volume

μg/mL - Microgramas por mililitro

Sap - Secretoras de aspartil-protease

Z - Zimotipo

. RESUMO

O cigarro de tabaco é considerado fator predisponente para várias doenças, inclusive para a candidose bucal. Ele contem substâncias que podem ser degradadas por enzimas destes fungos, servindo como nutrientes. A produção de enzimas histolíticas pelas espécies de Candida é um fator importante no desenvolvimento de doença. As enzimas podem prover acesso aos nutrientes do organismo, facilitando penetração dos fungos nas células do hospedeiro. O objetivo deste estudo foi avaliar a carga fúngica bucal e a secreção de enzimas histolíticas das espécies de Candida isoladas de adultos saudáveis não-fumantes e fumantes . Cem adultos (42 não-fumantes e 58 fumantes) participaram do estudo. Os espécimes foram coletados do dorso da língua e da mucosa jugal e identificados utilizando-se o meio de cultura CHROMagar®Candida. A positividade do gênero Candida foi de 28,57% nos indivíduos não-fumantes e de 39,65% fumantes (P = 0,254). C. albicans foi a espécie mais prevalente nos dois grupos, sendo isolada em 78,57% nos não-fumantes e 72,41% nos fumantes. Outras espécies encontradas foram C. parapsilosis, C. glabrata e C. tropicalis. Dos indivíduos positivos foi selecionada aleatoriamente uma cepa de cada espécie, totalizando 43 cepas, que foram avaliadas quanto à produção das enzimas aspartil-protease (Sap), fosfolipase, condroitinase, esterase-lipase e hemolisina. Devido a pequena amostra de cepas não-albicans obtida, somente as cepas de Candida albicans foram analisadas estatisticamente. A atividade enzimática foi maior no grupo dos não-fumantes para todas as enzimas estudadas, com diferença estatística significativa na atividade da enzima fosfolipase (P = 0.013). Concluímos que o tabagismo não afetou significativamente a positividade de Candida spp. e nem a atividade enzimática das cepas de *C. albicans*.

Palavras chave: Candida albicans; fumantes; enzimas; fosfolipase.

3. ABSTRACT

The cigarette smoke contains substances that can be degraded by enzymes of Candida species, been used like nutrients and converted to carcinogen end products. The purpose of this study was evaluated the carriage of *Candida* species and analyse the extracellular proteolytic activities of *Candida* isolates from healthy adults non-smokers and smokers. One hundred adults (42 non-smokers and 58 smokers) were studied. The samples were collected from intra oral swabs and smears from dorsum of tongue, right and left jugal mucosa each. The identification of species of Candida was made using the culture medium CHROMagar®Candida. The carriage of the gender Candida was 28,57% in the non-smokers individuals and 39,65% in the smokers (P = 0,254). Candida albicans was the most prevalent one between the two groups, being isolated in 78.57% in non-smokers, and 72.41% in smokers. Other species found were C. parapsilosis, C. glabrata, and C. tropicalis. From the positive individuals was random one strain of C. albicans, comprised 32 strains that were evaluated regarding the proteolytic activity of the following enzymes: secreted aspartyl-protease (Sap), phospholipase, chondroitinase, esterase-lipase, and haemolysin. Regarding the hydrolytic enzymes, there was bigger enzymatic activity in the group of non-smokers, in all studied enzymes, with significant statistical difference between both groups in the activity of phospholipase, that was bigger in the strains of non-smokers (P = 0.013). The results obtained in this study have shown that the smoking do not alter significantly the carriage of Candida spp., neither the enzymatic activity of C. albicans strains, except the activity of phospholipase.

Keywords: Candida albicans, smokers, enzymes, phospholipase.

4. ARTIGO	EM PORT	ΓUGUÊS	

PÁGINA DE TÍTULO

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RESUMO

O cigarro de tabaco tem sido considerado como um fator predisponente para várias doenças, inclusive para a candidose bucal. Ele contem substâncias que podem ser degradadas por enzimas das espécies de candida, servindo como nutrientes. Por outro lado, a produção de enzimas histolíticas pelas espécies de Candida é um fator importante no desenvolvimento de doenças provocadas por fungos oportunistas. As enzimas podem prover acesso do fungo aos nutrientes do organismo, chegando mesmo a penetrar nas células do hospedeiro. O objetivo deste estudo foi avaliar a carga fúngica bucal e a secreção de enzimas histolíticas das espécies de Candida albicans isoladas de adultos saudáveis não-fumantes e fumantes. Cem adultos (42 não-fumantes e 58 fumantes) participaram do estudo. Os espécimes foram coletados do dorso da língua e da mucosa jugal (direita e esquerda). A identificação das espécies de Candida foi realizada utilizando o meio de cultura CHROMagar®Candida. A positividade do gênero Candida foi de 28,57% nos indivíduos não-fumantes e de 39,65% fumantes (P = 0,254). C. albicans foi a espécie mais prevalente nos dois grupos, sendo isolada em 78,57% nos não-fumantes e 72,41% nos fumantes. Outras espécies encontradas foram C. parapsilosis, C. glabrata e C. tropicalis. Dos indivíduos positivos foi selecionada aleatoriamente uma cepa de cada espécie, totalizando 43 cepas, que foram avaliadas quanto à produção das enzimas aspartil-protease (Sap), fosfolipase, condroitinase, esterase-lipase e hemolisina. A atividade enzimática foi maior no grupo dos nãofumantes para todas as enzimas estudadas, com diferença estatística significativa na atividade da enzima fosfolipase, sendo maior nas cepas de pacientes não-fumantes (P = 0.013). Concluímos que o tabagismo não afetou significativamente a positividade de Candida spp. e nem a atividade enzimática das cepas de C. albicans.

Palavras chave: Candida albicans; fumantes; enzimas; fosfolipase.

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Palavras chave: *Candida albicans;* fumantes; enzimas; fosfolipase. INTRODUCÃO

C. albicans é um fungo dimórfico, que está presente em aproximadamente 50% dos indivíduos saudáveis (37). Além da C. albicans, outras espécies como C. tropicalis, C. glabrata, C. krusei e C. parapsilosis são capazes de competir com a microbiota bucal e se tornarem patogênicas.

Segundo Almeida & Scully (1), os principais fatores que dificultam o crescimento desses fungos no homem são a elevada temperatura e a imunidade celular. A virulência da *C. albicans* está associada com o aumento da produção de tubo germinativo pelas leveduras em baixas temperaturas. Leveduras que crescem em temperatura ambiente são mais resistentes à ação dos leucócitos polimorfonucleares (2). O desenvolvimento

de micélio pelas espécies de *Candida* ssp. (dimorfismo) favorece as infecções fúngicas em decorrência da variabilidade antigênica da superfície e do formato que propicia maior aderência, dificultando a ação fagocitária pelo sistema imune (23).

O tabagismo pode estar associado à predisposição às candidoses bucais (15, 31, 45). A nicotina no tabaco pode causar mudanças estruturais e funcionais nos queratinócitos (25). O efeito do fumo nas células epiteliais, reduz a atividade anti-*Candida* das células e pode aumentar a susceptibilidade à candidose orofaringeana (42), além de causar um espessamento na camada queratinizada do epitélio, o que pode contribuir para a maior colonização pela levedura (4, 37, 44).

Pacientes fumantes relatam recidiva das infecções por *Candida* após o término da terapia antifúngica, por outro lado, há evidências clínicas de que algumas infecções desaparecem apenas com a cessação do vício de fumar (15).

A produção de enzimas histolíticas é um fator importante no desenvolvimento de doenças provocadas por fungos oportunistas, sobretudo em isolados clínicos de *Candida* spp. As enzimas podem prover acesso do fungo aos nutrientes do organismo, chegando mesmo a penetrar nas células do hospedeiro, via estruturas tipo haustório (14). Quando a hidrólise dos substratos ou ação das proteínas extracelulares afeta a função e a viabilidade do hospedeiro, as enzimas podem ser consideradas fatores de virulência que contribuem para o estabelecimento da infecção (7, 22, 32).

O objetivo deste estudo foi avaliar a carga fúngica bucal e os perfis de virulência das cepas de *Candida albicans* entre indivíduos não-fumantes e fumantes. Para isto foram realizados os seguintes testes: contagem de diferentes espécies de *Candida*, produção de aspartil-proteases secretoras (Sap), de fosfolipases, de condroitinase, de esterase/lipase e de hemolisinas.

MATERIAL E MÉTODOS

A participação dos indivíduos foi condicionada à assinatura de um Termo de Consentimento Livre e Esclarecido, em concordância com as diretrizes do Comitê de Ética em Pesquisa da Pontifícia Universidade Católica do Paraná (PUCPR).

Foram avaliados 100 indivíduos, 42 não-fumantes e 58 fumantes, e considerados fumantes aqueles com consumo igual ou superior a cinco cigarros por dia há pelo menos um ano, e não-fumantes aqueles que relataram nunca terem fumado (29). Indivíduos que fumavam menos de cinco cigarros por dia foram excluídos da pesquisa. Cada indivíduo foi avaliado por um único avaliador que realizou a anamnese e o exame físico intrabucal, inspecionando as mucosas, tecidos moles e dentes. Indivíduos com manifestações clínicas de candidose bucal; relato de doenças sistêmicas na anamnese (diabetes ou hipertensão); mulheres grávidas; indivíduos que tivessem utilizado antibióticos nos três meses que precederam as coletas ou fazendo uso de medicação sialorredutora, foram excluídos da amostra. A média de idade dos indivíduos não-fumantes foi de $19,95 \pm 2,13$ anos e dos fumantes de $22,83 \pm 5,75$ anos, sendo em sua maioria jovens. A média de cigarros consumidos por dia foi $12,79 \pm 5,73$ e do tempo de uso em anos foi de $6,19 \pm 5,10$.

Amostras de raspado superficial (*swabbing*) do dorso da língua e da mucosa jugal dos indivíduos foram recolhidas, semeadas em placas contendo CHROMagar®*Candida* (CHROMagar Microbiology, Biomerieux, Paris, França) e incubadas a 37°C, por 48 horas. Após a identificação fenotípica (24), as cepas foram transferidas para tubos de armazenamento contendo Ágar Sabouraud Dextrose (Difco Laboratories, Detroit, Mich., USA).

Um total de 43 cepas do gênero *Candida* (uma colônia/fenótipo por indivíduo) foram submetidas às provas de atividade enzimática. A cepa-padrão de *C. albicans* CBS562 proveniente do estoque do Laboratório de Estomatologia da PUCPR foi testada como controle. Os isolados foram crescidos em estufa a 37°C em 5mL de Caldo Sabouraud Dextrose (Difco Laboratories, Detroit, Mich., USA). Após 24 horas as células foram recolhidas por centrifugação e lavadas três vezes com água destilada estéril. Os *pellets* foram ressuspendidos em água estéril até o equivalente ao tubo 10 da escala de MacFarland e 5 μL inoculados com o auxílio de discos de papel filtro esterilizados, em cinco pontos eqüidistantes, nos meios de cultura para detecção semiquantitativa de aspartil-protease secretora (Sap), fosfolipases, condroitinase, esterase/lipase e hemolisinas. Todos os testes foram realizados em duplicata.

A secreção de Sap foi verificada seguindo o protocolo de Rüchel *et al.* (32), empregando a fração V da albumina bovina (Sigma Chem Co., St. Louis, Mo., USA) como substrato, Yeast Nitrogen Base (Difco Laboratories, Detroit, Mich., USA) e 2,5 mL de Protovit Plus[®] (Roche, São Paulo, Brasil) como co-fatores (22). A atividade enzimática foi evidenciada pela formação de um halo transparente ao redor da colônia dez dias após a inoculação. (10).

A atividade fosfolipásica foi pesquisada seguindo o método de Price *et al.*(28). Em 1.000 mL de água destilada estéril foram dissolvidos 10,0 g de peptona (Difco Laboratories, Detroit, Mich., USA), 30,0g de glucose, 57,3g de NaCl e 0,55g de CaCl₂ e 20,0g de Ágar (Oxoid, Fakola AG, Basel, Switzerland). O meio foi esterelizado em autoclave a 120°C por 15 minutos e resfriado a uma temperatura de 55°C. O *egg yolk* estéril sem telurito de potássio (Newprov Prod. Laborat., Inc., São José dos Pinhais, Paraná, Brasil) foi adicionado ao meio esterelizado. Após 4 dias da inoculação, as

placas foram observadas quanto à formação de uma zona de cor amarelada ao redor das colônias.

Para a pesquisa de condroitinase foi empregado o protocolo de Smith & Willett (41) modificado. O meio de cultura foi preparado usando-se neopeptona 10 g, glucose 40 g, Ágar noble 15 g (Sigma Chem Co., St. Louis, Mo., USA) e água destilada qsp 1.000 mL e autoclavado a 120°C por 15 minutos. Após resfriamento a 55°C, foi acrescentada uma solução de albumina bovina fração V a 10% e sulfato de condroitina a 4% (Sigma Chem Co., St. Louis, Mo., USA), dissolvidos em 10 mL de água destilada estéril e esterelizados em filtros de Millipore de 0,22 μm. O pH final foi ajustado entre 6,8 ± 0,2. O halo formado foi medido após 48 h de inoculação.

O teste de opacidade Tween[®] 80 (Sigma Chem Co., St. Louis, Mo., USA) foi utilizado para determinar a produção de esterase/lipase. O meio de cultura foi preparado com 15,0 g de SDA (Difco Laboratories, Detroit, Mich., USA) e 10 g CaCl₂, em 1.000 mL de água destilada (33). Após ser autoclavado, o meio foi resfriado a 50°C e 10 mL de Tween[®]80 (Sigma) autoclavado foi adicionado. As inoculações permaneceram em estufa a 37°C por sete dias consecutivos.

Para determinar a atividade hemolítica, as cepas foram semeadas em placas de ágar-sangue contendo 70 mL de sangue de ovelha fresco e desfibrinado (Newprov Prod. Laborat., Inc., São José dos Pinhais, Paraná, Brasil); 30 g de glucose (Sigma Chem Co., St. Louis, Mo., USA); qsp 1.000 mL de solução de 1% de Ágar-ágar (Oxoid, Fakola AG, Basel, Switzerland); pH final 5,6 (19). A formação do halo translúcido ao redor das colônias após 48 h de incubação a 37°C indicou atividade hemolítica positiva.

Após os períodos de observação foram feitas as leituras do tamanho das colônias e dos halos formados por um único pesquisador calibrado, utilizando transiluminação e um paquímetro digital (Mitutoyo, 6" digital caliper, w/abs, Japão) e calculadas as

atividades enzimáticas. Estas atividades foram determinadas de acordo com a técnica de Price *et al.* (28), dividindo-se o diâmetro da colônia (DC) pelo diâmetro do halo formado (DH). Este resultado é denominado Pz, que representa numericamente a atividade enzimática da cepa testada. Como menores valores Pz indicam maior atividade enzimática e vice-versa, optou-se por subtrair tais valores de um (1-Pz) (26). Os dados assim obtidos foram analisados quanto à sua normalidade de distribuição pelo teste de Kolmogorov-Smirnov. Para aspartil-protease secretora, condroitinase e hemolisinas houve distribuição normal dos valores e foi utilizado o teste *t* de Student. Nas enzimas onde não houve distribuição normal, fosfolipase e esterase/lipase, foi utilizado o teste não-paramétrico U de Mann-Whitney.

A atividade enzimática foi analisada estatisticamente somente nos isolados de *C. albicans*, devido à inexpressiva obtenção de outras espécies.

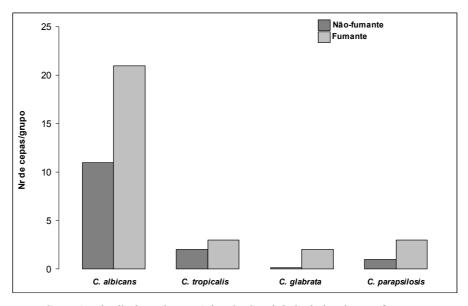
Além dos ensaios estatísticos, os valores 1-Pz também permitiram a classificação das cepas em zimotipos. Foram gerados dígitos a partir de um escore de atividade dado aos valores de Pz (7). Cepas não produtoras (Pz = 1,000) foram codificadas como 1, pois o diâmetro do halo foi considerado igual ao da colônia; cepas com atividade média (Pz ≤ 0,639) receberam código 2; cepas com atividade elevada (0,999 ≤ Pz ≥ 0,640) receberam código 3 (26). Os zimotipos foram gerados seguindo a ordenação dos dígitos oriundos da secreção das enzimas aspartil-protease secretora, fosfolipases, condroitinase, esterase/lipase e hemolisinas. A zimotipagem nos forneceu uma visão geral da capacidade produtora de cada cepa, associando as cinco enzimas pesquisadas.

RESULTADOS

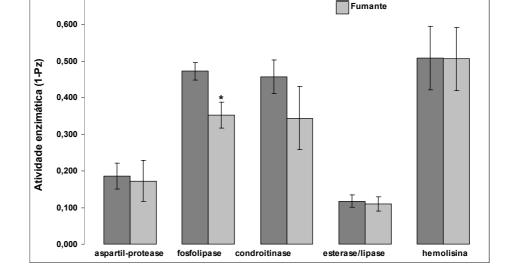
Isolamentos positivos de *Candida spp*. foram obtidos em 35 dos 100 indivíduos examinados, sendo 12/42 (28,57%) do grupo dos não-fumantes e 23/58 (39,65%) do grupo de fumantes, não havendo diferença estatística significante entre os grupos (P = 0,254).

Foram isoladas duas cepas de *C. tropicalis* 2/14 (14,28%) e uma de *C. parapsilosis* 1/14 (7,14%) dos indivíduos não-fumantes. Nos fumantes foram encontradas três cepas de *C. tropicalis* 3/29 (10,35%), duas de *C. glabrata* 2/29 (6,89%) e três de *C. parapsilosis* 3/29 (10,34%). O pequeno número isolado destas espécies não possibilitou análise estatística.

 $C.\ albicans$ foi prevalente em ambos os grupos, ocorrendo em 11/14 (78,57%) dos não-fumantes e 21/29 (72,41%) dos fumantes, não havendo diferença estatística significante entre eles (P=0,705) quando o teste t de Student foi aplicado. Não houve predomínio de colonização de nenhuma das espécies encontradas em ambos os grupos (Gráfico 1).



Graf. 1. Distribuição das espécies de *Candida* isoladas de não-fumantes (n=14) e fumantes.



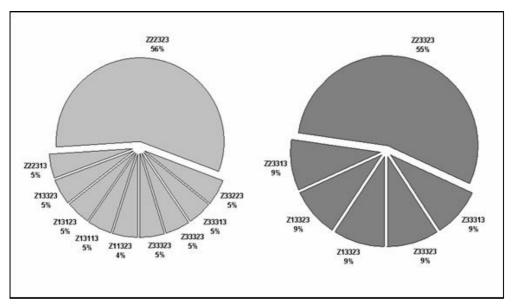
Graf. 2. Atividade das cepas de *C. albicans* isoladas de não-fumantes (n=11) e fumantes (n=21). * Teste U de Mann-Whitney (p

Das cepas de *C. albicans* isoladas de não-fumantes, 7/11 (63,63%) delas apresentaram atividade enzimática média para aspartil-protease, 2/11 (18,18%) apresentaram atividade elevada e em 2/11 (18,18%) não foi detectada atividade. Neste grupo, 11/11 (100%) das cepas apresentaram atividade elevada de fosfolipase, condroitinase e hemolisina. A atividade enzimática de esterase/lipase foi média em 9/11 (81,81%) cepas, e ausente em 2/11 (18,18%) dos isolados (Gráfico 2). A compilação dos escores de atividade enzimática das cepas de não-fumantes permitiu a obtenção de cinco zimotipos, sendo o Z32232 o mais prevalente.

Nas cepas isoladas de fumantes houve maior variação de atividade enzimática, sendo encontrados dez zimotipos. A aspartil-protease teve atividade elevada em 3/21 (14,28%) das cepas, assim como outras 3/21 não secretaram a enzima. Em 15/21 (71,42%) a atividade foi considerada média. Apenas uma 1/21 (4,76%) cepa não secretou fosfolipase, outras 2/21 (9,52%) apresentaram atividade média e 19/21 (90,47%) foram consideradas fortemente produtoras. Houve diferença estatística significante na produção desta enzima entre não-fumantes e fumantes (P = 0,013). Para condroitinase, 17/21 (80,95%) cepas apresentaram atividade elevada, 2/21 (9,52%) cepas apresentaram atividade média e as 2/21 cepas restantes não tiveram atividade detectável. Nenhuma cepa apresentou atividade elevada de esterase/lipase neste grupo.

A atividade foi média em 19/21 (90,47%) das cepas e ausente em 3/21 (14,28%) dos isolados. Na produção de hemolisina apenas uma cepa apresentou atividade média (4,76%), enquanto 20/21 (95,23%) apresentaram atividade elevada (Gráfico 2). O zimotipo predominante neste grupo foi o Z22323.

A cepa-padrão CBS562 apresentou o zimotipo coincidente com o mais encontrado nos fumantes, o Z23323, com produção de todas as cinco enzimas pesquisadas, tendo produção elevada em três delas (Gráfico 3).



Graf. 3. Distribuição dos zimotipos das cepas de *C. albicans* isoladas de não-fumantes (n=11) e fumantes (n=21).

DISCUSSÃO

Embora muitos artigos tenham sido publicados sobre a secreção de enzimas como fatores de virulência de *C. albicans* (6, 13, 17, 35, 36) até o nosso conhecimento, nenhum estudo anterior avaliou a atividade enzimática da espécie, ou do gênero, obtida de indivíduos saudáveis não-fumantes e fumantes.

No presente estudo, 28,57 % dos não-fumantes apresentou colonização por *Candida* spp. contra 39,65 % dos fumantes (p=0,254). Esses resultados são inferiores aos publicados por Rassol *et al.* (29), que isolaram cepas em 43% dos não-fumantes e 57% dos fumantes, embora estes autores também não tenham encontrado diferença estatística em os grupos. Em 1988, Odds (appud Scully) revisou 32 artigos sobre positividade de *Candida* spp. na boca de indivíduos saudáveis, e relatou que a média de positividade foi de 34,40%.

Em nosso estudo, *C. albicans* foi a espécie mais prevalente, em concordância com outros autores (22, 29, 37), sendo encontrada em 78,57% dos não-fumantes e 72,41% dos fumantes, quando esses se apresentavam colonizados. Embora tenha sido encontrado um percentual maior de isolamento dessa espécie no grupo dos não-fumantes, não há diferença estatística entre os grupos.

A positividade de produção de aspartil-proteases (Saps) pelas cepas encontradas no presente estudo está de acordo com os estudos de Bramono *et al.* (6), que encontraram atividade desta enzima nas cepas de *C. albicans, C. tropicalis, C. parapsilosis, C. glabrata, C. krusei e C. guilliermondii.* Mas, difere do encontrado por Dóstal *et al.* (10), que não encontraram atividade em cepas de *C. glabrata.*

Com relação às fosfolipases, não foi encontrada qualquer atividade nas cepas de C. glabrata e C. parapsilosis, o que contraria os achados de outros autores (11, 38), que afirmam que estas espécies são capazes de produzir tais enzimas. Tal fato pode ter ocorrido devido ao número reduzido de isolamentos dessa espécie nos indivíduos arrolados neste estudo.

Quando a secreção dessas fosfolipases foi avaliada em isolados de *C. albicans* provenientes de não-fumantes, detectamos uma significante elevação de atividade (*P* = 0,013) em relação aos isolados de fumantes. Por se tratar de uma enzima que parece estar diretamente relacionada com a virulência desta espécie, era esperado que no grupo dos fumantes esta secreção fosse maior. Tal fato pode ser explicado em função da seleção aleatória das colônias que deram origem aos isolados analisados. Como foi selecionada apenas uma colônia de *C. albicans* por indivíduo, podem ter sido excluídas do estudo outras cepas, com maiores atividades enzimáticas.

Ibrahim *et al.* (13) reportaram que cepas comensais com baixa produção de fosfolipase são capazes de colonizar a mucosa, sugerindo que a secreção destas enzimas não é um requisito primordial para a patogenicidade de *C. albicans*, opinião compartilhada por Mayser *et al.* (21). Como os indivíduos aqui analisados não apresentavam quaisquer sinais clínicos de candidose bucal, é razoável supor que essa maior secreção de fosfolipases por cepas de não-fumantes não deva ter maiores implicações na colonização.

A totalidade das cepas de *C. albicans* do grupo não-fumantes produziu condroitinase, em comparação com 90,47% produzidos por esta espécie no grupo dos fumantes. Das espécies "*não-albicans*" apenas *C. tropicalis* e uma cepa de *C. parapsilosis* produziram condroitinase. Ainda, as duas cepas de *C. glabrata* isoladas não produziram condroitinase. Este resultado está de acordo com Shimizu *et al.* (39) que observaram a produção da enzima em 97,8% das cepas de *C. albicans*, sendo esta produção seguida por *C. tropicalis*, *C. guilliermondii*, *C. parapsilosis* e *C. krusei*, e com

Chaffin *et al.* (9) que afirmam que hialuronidase e condroitinase podem ser produzidas por *C. albicans, C. tropicalis, C. guilliermondii, C. parapsilosis,* e *C. krusei*, mas em uma pequena proporção dos isolados.

No presente estudo foi avaliado também a secreção de esterase/lipase. Todas as espécies foram capazes de secretar esta enzima; porém, em diferentes proporções. Este achado está de acordo com as pesquisas de outros autores (6, 9, 33, 43). O papel exato da atividade lipolítica na patogenicidade das cepas ainda não foi esclarecido.

A atividade das hemolisinas está associada à capacidade que as cepas têm em destruir hemácias e utilizar o ferro como nutriente. Nossos resultados mostraram que todas as espécies encontradas produziram hemolisinas, em concordância com os estudos de Luo *et al.* (19), sendo fortemente produtoras na maioria das cepas.

Dentre as cepas de *C. albicans* examinadas, nenhuma apresentou elevados níveis de secreção para as cinco enzimas de forma concomitante, i.e. nenhuma apresentou valor 1-Pz maior que 0,640 para as cinco enzimas estudadas. Isto está em concordância com o estudo de Bramono *et al.* (6), que pesquisaram a secreção de aspartil-protease, lipase e α-glucosidase em cepas de *C. albicans*, e nenhuma exibiu altos níveis de atividade para as três enzimas simultaneamente.

Tem sido sugerido que o cigarro pode levar à alterações localizadas no epitélio que permitem a colonização por *Candida* (3, 25). Williams *et al.* (44) mostraram que uma maior queratinização favorece a aderência de *C. albicans*. Ainda, Barret-Bee *et al.* (5) propuseram que a fosfolipase teria um importante papel nessa aderência; proposição não compartilhada por Ghannoum *et al.* (12). Na mesma direção, Ray & Payne (30) mostraram que blastoporos de *C. albicans* podem degradar a queratina celular utilizando-se de proteases ácidas e Jayatilake *et al.* (14) propuseram que as fosfolipases

também podem ser participar desta destruição, o que, em última instância, favoreceria a colonização.

A despeito dos argumentos acima, talvez a menor secreção de enzimas por cepas obtidas junto a fumantes se deva a uma seleção clonal determinada pela qualidade da saliva. Como as Saps (8) e as fosfolipases (20, 34) são usualmente ativas somente em meios ácidos e já foi mostrado que fumantes apresentam saliva mais ácida e com menor capacidade tamponante que não-fumantes (17, 18, 27, 40), é razoável supor que no grupo dos fumantes não seja necessária uma alta taxa de secreção para que as cepas possam conduzir sua colonização.

De acordo com a literatura disponível, fumantes podem ter maior predisposição à colonização por *Candida spp.* devido à indução de um aumento na queratinização do epitélio (4, 25, 44); à diminuição da atividade dos leucócitos, do exudato gengival (42) e dos níveis de imunoglobulina A (42); sendo relatados os efeitos do fumo no indivíduo, mas não nos microorganismos.

Mais estudos são necessários para identificar o efeito, se é que existe, do cigarro sobre as espécies de *Candida* que fazem parte da microbiota bucal humana.

CONCLUSÃO

Concluímos com nosso estudo que a carga fúngica bucal não difere entre nãofumantes e fumantes. O perfil de virulência não diferiu entre não-fumantes e fumantes, exceto para a secreção das fosfolipases, pois houve uma menor secreção destas enzimas nas cepas de *Candida albicans* de indivíduos fumantes.

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5. ARTIGO EM INGLÊS

TITLE PAGE

VIRULENCE FACTORS OF *Candida albicans* STRAINS ISOLATED FROM SMOKERS AND NON-SMOKERS

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ABSTRACT

The cigarette smoke contains substances that can be degraded by enzymes of Candida species, been used like nutrients and converted to carcinogen end products. The purpose of this study was evaluated the carriage of *Candida* species and analyse the extracellular proteolytic activities of Candida isolates from healthy adults non-smokers and smokers. One hundred adults (42 non-smokers and 58 smokers) were studied. The samples were collected from intra oral swabs and smears from dorsum of tongue, right and left jugal mucosa each. The identification of species of Candida was made using the culture medium CHROMagar®Candida. The carriage of the gender Candida was 28,57% in non-smokers individuals and 39,65% in smokers (P = 0,254). Candida albicans was the most prevalent one between the two groups, being isolated in 78.57% in the non-smokers, and 72.41% in the smokers. Other species found were C. parapsilosis, C. glabrata, and C. tropicalis. From the positive individuals was random one strain of C. albicans, comprised 32 strains that were evaluated regarding the proteolytic activity of the following enzymes: secreted aspartyl-protease (Sap), phospholipase, chondroitinase, esterase-lipase, and haemolysin. Regarding the hydrolytic enzymes, there was bigger enzymatic activity in the group of non-smokers, in all studied enzymes, with significant statistical difference between both groups in the activity of phospholipase, that was bigger in the strains of non-smokers (P = 0.013). The results obtained in this study have shown that the smoking do not alter significantly the carriage of Candida spp., neither the enzymatic activity of C. albicans strains, except the activity of phospholipase.

Keywords: Candida albicans, smokers, enzymes, phospholipase.

INTRODUCTION

Candida albicans is present in the inoffensive form in the oral cavity of 50% of healthy people (37). Besides *C. albicans*, other species such *C. tropicalis*, *C. glabrata*, *C. krusei*, and *C. parapsilosis* are able to compete with the oral microbiota and become pathogenic. These forms of candidosis are mycotic infections relatively common and well known, however, not completely understood (22).

According to Almeida & Scully, (1) the main obstacles for these fungi to grown in humans are the high temperature and the cellular immunity. The virulence of *C. albicans* is associated to the increase of the production of the germinative tube by yeasts under low temperatures. Yeasts that grow under room temperature are more resistant to the actions of polymorph nuclear leukocytes (2). The development of mycelium by strains of *Candida* (dimorphism) also favours the infection as result of the antigenic variability of the surface and of the mycelial format that propitiates more adherence hindering the phagocytary action by the immune system (23).

The smoking is also associated to the predisposition to the oral candidosis (15, 31, 42, 45). The nicotine in the tobacco may cause structural and functional changes in the keratinocytes (25). The effects of tobacco on the epithelial cells reduces the anti-Candida activity of cells and may increase the susceptibility to oropharyngeal candidosis (42), and can cause thickening of the keratinised layer of the epithelium, that may contribute for a more pronounced colonization by yeast (37, 44).

Smoking subjects report recurrence of infections caused by *Candida* after the end of antifungal therapy; on the other hand, there are clinic evidences that some infections disappeared after ceasing the smoking habit (15).

The production of histolytic enzymes is an important factor in the development of diseases caused by opportunistic fungi, especially in clinic isolates of *Candida* spp. The enzymes may promote the access of the fungus to the nutrients of the organism, even penetrating the cells of the host through haustorium-like structures (14). When the hydrolysis of the substrata or the action of the extracellular proteins affects the function and viability of the host, the enzymes may be considered factors of virulence that contribute for the establishment of infection (7, 22, 32).

This study was carried aiming at assessment of the carriage of oral *Candida* species and the profiles of virulence of the strains of *Candida* among the smoking and non-smoking populations. Thus, there were performed the following tests: counting of the different species of *Candida*, productions of secreted aspartyl-proteases (Saps), phospholipases, chondroitinase, esterase/lipase, and of hemolysins.

MATERIAL AND METHODS

The participation of individuals was conditioned to the agreement of term of free and aware consentient with accordance to policies of the Research Ethics Committee of the *Pontifical Catholic University of Paraná (PUCPR)*.

There were examined 100 individuals, 42 non-smoking and 58 smoking and there was considered chronic smoker that individual whose daily consumption has been equal or superior to five cigarettes for, at least, one year, and considered non-smoker that one who has never smoked before (29). Individuals whose daily consumption was lower to five cigarettes per day, were excluded of the study. Each individual was evaluated by only one professional who performed the anamnesis and a physical exam, survey of the

health conditions of the mucosa and other soft tissues as well as the teeth. Individuals with clinical manifestations of oral candidosis; relates of systemic diseases like diabetes, hypertension; pregnancy; or individuals used of antibiotics in the last three months before the collects, or using of sialoreductor drugs were excluding conditions of the sample. The medium age of the non-smokers individuals was 19.95 ± 2.13 years and of the smokers was 22.83 ± 5.75 years.

Sample of superficial scrapings (*swabbing*) of the back of the tong and jugal mucosa of subjects were taken, inoculated in dishes containing CROMAgar[©]Candida (CHROMagar Microbiology, Biomerieux, Paris, France) and incubated at 37°C, for 48 hours. After the phenotypic identification (24), the strains were transferred to storing tubes containing Sabouraud Dextrose Agar (Difco Laboratories, Detroit, Mich., USA).

A total of 43 strains from the genus *Candida* (a colony/phenotype from each individual) which were submitted to checks of enzymatic activity. There were also tested the type-strain of *C. albicans* CBS562 from the archive of *Laboratory of Stomatology* of *PUCPR*. The isolates were grown at 37°C in 5mL of Sabouraud Dextrose Broth (Difco Laboratories, Detroit, Mich., USA). After 24 hours, the cells collect through centrifugation and washed three times with sterile distilled water. The pellets were resuspended in sterile water until the equivalent to the tube #10 of the scale of MacFarland and inoculated, in equidistant points, in the culture media for the semi-quantitative detection of secreted aspartyl-protease (Sap), phospholipases, chondroitinase, esterase/lipase, and haemolysins. All the tests were made in duplicate.

The secretion of Sap was verified following the protocol of Ruchel *et al.* (32), employing the fraction V of bovine albumin (Sigma Chem Co., St. Louis, Mo., USA) as substratum, Yeast Nitrogen Base (Difco Laboratories, Detroit, Mich., USA) and 2.5ml of Protovit[®]Plus (Roche Laboratories, São Paulo, Brazil) as cofactors (22). The

enzymatic activity was evidenced by the formation of a translucent around the colony, 10 days after the inoculation. (10).

The phospholipasic activity was researched according to the method of Price *et al.* (28). In 1.000 mL of sterile distilled water, there were dissolved 10.0g of peptone (Difco Laboratories, Detroit, Mich., USA) 30.0g of glucose, 57.3g of NaCl, 0.55g CaCl₂, and 20.0g of agar (Oxoid, Fakola AG, Basel, Switzerland). The medium was sterilized in autoclave at 120°C for 15 minutes and cooled to a temperature of 55°C. The sterile egg yolk without potassium tellurite (Newprov Prod. Laborat., Inc., São José dos Pinhais, Paraná, Brazil) was added to the sterilized medium. After 4 days of inoculation, the dishes were observed regarding the formation of an area of yellow colour around the colonies.

Concerning the research of chondroitinase, there was employed the modified protocol of Smith & Willett (41). The culture medium was prepared using neopeptone 10g, glucose 40g, agar noble 15g (Sigma Chem Co., St. Louis, Mo., USA) and distilled water till 1.000mL and autoclaved at 120°C for 15 minutes. After cooling to 55°C, there was added 10mL of sterile solution of 10% fraction V bovine albumin plus 4% chondroitin sulphate (Sigma). The final pH was adjusted among 6.8 ± 0.2 . The inoculations were kept in stove for 48 hours.

The Tween[®]80 (Sigma) opacity test was used for determining the production of esterase/lipase. The medium of culture was prepared with 15.0g of SDA (Difco) and 10g CaCl₂, in 1.000 mL of distilled water (33). After autoclaved, the medium was cooled to 50°C and 10mL of Tween[®]80 autoclaved (Sigma) were added. The inoculations were kept in stove for seven consecutive days.

For determining the haemolytic activity, the strains were inoculated in dishes of Blood-Agar with 70 mL of fresh and defibrinated blood of sheep (Newprov); 30g of

glucose (Sigma); till 1.000 mL of 1% Agar-agar solution (Oxoid); final pH 5,6 (19). The formation of a translucent halo around the colonies after 48 hours of incubation has pointed positive haemolytic activity.

After the daily observation, there were performed the reading of the size of the colonies and size of halos formed for each enzyme by unique observer using transillumination and a digital caliper (Mitutoyo, Japan) and calculated the enzymatic activities. These enzymatic activities (Pz) were determined according to the technique of Price $et\ al.$ (28), through the ratio among the diameter of the colony (CD) and the diameter of the formed halo (HD). This result is denominated Pz, and represents the enzymatic activity of each strain tested. As little values of Pz points a more expressive enzymatic activity and vice-versa, there was opted for subtracting such values from 1 (1-Pz) (26). The data therefore obtained were analysed with regard to their distribution by the test of Kolmogorov-Smirnov. For secreted aspartyl-protease, chondroitinase, and haemolysins, there was a normal distribution of the values, and there was used the Student's t test. In the enzymes, which did not present normal distribution (phospholipase and esterase/lipase), there was used the non-parametric Mann-Whitney U test.

The enzymatic activity was analysed only in isolates of *C. albicans* due to the inexpressive obtaining of other species.

Besides the statistic assays, the values of 1-Pz also allowed the classification of strains in zymotypes. Each digit was generated from a score of activity (7). Non-producer strains (Pz = 1.000) were codified as 1, besides the diameter of the formed halo was considered the same of the diameter of the colony; strains with intermediary activity (Pz \leq 0,639) received code 2; strains with high enzymatic activity (0,999 \leq Pz \geq 0.640) received code 3 (26). The zymotypes were generated following the ordination of

the digits resultant of the secreted aspartyl-protease, phospholipases, chondroitinase, esterase/lipase, and haemolysins. The zimotyping gave us an overall of the enzymatic activity that each strain associated the activity of the five enzymes studied.

RESULTS

Positive isolations of *Candida* were obtained in 43 out of the 100 collected sampled, being 12/42 (28,57%) from the check group and 29/58 (39,65%) from the group of smokers, don't having statistic difference among the groups (P = 0,254).

C. albicans was the most prevalent in both groups, occurring in 11/14 (78.57%) of non-smokers and 21/29 (72.42%) of smokers, not presenting statistic difference among the groups (P = 0.705) when the Student's t test was performed. Regarding the non-albicans species, there was no colonial predominance of any species in either group, as observed in graphic 1.

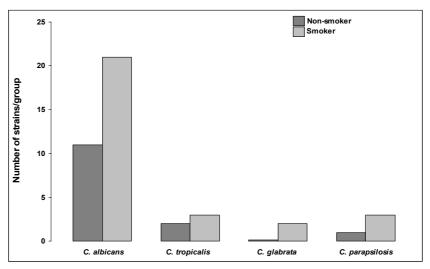


FIG. 1. Distribution of yeast species isolated from non-smokers (n = 14) and smokers (n = 29). No disparities were found when the Test for Differences Between Two Proportions was employed.

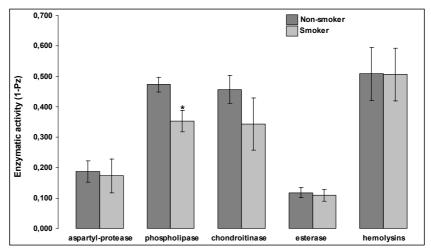


FIG. 2. Overall view of extracellular enzymatic activity of C. albicans strains isolated from non-smokers (n = 11) and smokers (n = 21), * Mann-Whitney U test; P = 0.0131.

From the strains of *C. albicans* isolated from non-smokers, 7/11 (63,64%) of them have presented mean enzymatic activity for secreted aspartyl-protease, 2/11 (18.18%) have presented high activity, and in 2/11 (18.18%), there was not detected activity. In this group, 11/11 (100%) of the strains have presented high activity of phospholipase, chondroitinase, and haemolysin. The enzymatic activity of esterase/lipase was medium in 9/11 (81.81%) strains and absent in 2/11 (18.18%) of the isolates. (Graphic 2).The compilation of scores of enzymatic activity allowed the obtaining of six zymotypes, being the Z23323 the most prevalent one.

In the strains isolated from smokers, there was a larger variation of the enzymatic activity, being found ten zymotypes (Figure 3). Secreted aspartyl-protease presented high activity in 3/21 (14,28%) of the strains, and the other 3/21 have not secreted the enzyme. In 15/21 (71,41%), the activity was considered medium. Only 1/21 (4.76%) strain has not secreted phospholipase, other 2/21 (9.52%) have presented intermediary activity, and 19/21 (90,47%) were considered strong producers. The enzymatic activity of this enzyme showed difference between non-smokers and smokers (P = 0,013). For chondroitinase, 17/21 (80.95%) strains have presented high activity, 2/11 (9.52%) strains have presented intermediary activity, and 2/11 strains have not presented a

detectable activity. No strain has presented high activity of esterase/lipase in this group. The activity was intermediary in 19/21 (85.71%) of strains and absent in 3/21 (14.29%) of the isolates. In the production of haemolysin, only one strain has presented intermediary activity (4.76%) while 20/21 (95.23%) have presented high activity. The predominant zymotype was the Z22323.

The standard-strain CBS562 has presented a coinciding zymotype to the one found in the smokers, the Z23323, with production of all the researched enzymes, having elevated production in three of them (Graphic 3).

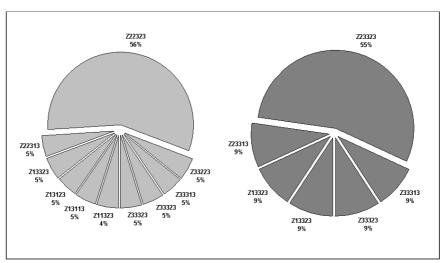


Fig 3. Distribution of zymotypes of C. albicans strains isolated from non-smokers (n = 11) and smokers (n = 21).

DISCUSSION

Despites the considerable source of information regarding the factors that determine the virulence of *C. albicans* (6, 13, 17, 35, 36), in our knowledge, no previous study has evaluated the enzymatic activity of the species, not even of the genus that was obtained from smoking and non-smoking individuals.

In the present study, 28,57% of smokers presented themselves positive for the colonization by *Candida* spp. against 39,65% of non-smokers (P > 0.05). These results

are less inferior to the ones published by Rassol *et al.* (29) that isolated yeasts in 43% of non-smokers and 57% of smokers. Nevertheless, not even these authors obtained statistic differences among groups, what would take us to an inference that the age might be a factor of predisposition to the more determinant colonization than the smoking habit.

C. albicans was the most prevalent species, being found in 78.54% of non-smokers and 72.41% of smokers, when colonized. Although there was found a higher percentage of isolation of this species, there is no statistical difference among the groups.

The production of secreted aspartyl-proteases (Saps) by the strains found in the present study corroborates with the studies of Bramono *et al.* (6), which found the activity of this enzyme in the strains of *C. albicans, C. tropicalis, C. parapsilosis, C. glabrata, C. krusei*, and *C. guilliermondii*. But, differently from the findings of Dóstal *et al.* (10), there was not found activity in strains of *C. glabrata*.

Regarding the phospholipases, there was not found any activity in the strains of *C. glabrata* and *C. parapsilosis*, what contradict the findings of other authors (11, 38), who affirm that these species are able to produce such enzymes. Such fact might have happened due to the reduced number of isolations of this species in the individuals who participated in this study.

When the secretion of these phospholipases is evaluated in isolates of C. albicans from non-smokers, it is noticeable a significant activity elevation (P = 0.013) in relation to the isolates of non-smokers. As it seems to be an enzyme directly related to the virulence of this species, it was expected higher secretion in the group of smokers. Such fact may be partially explained in function to random selection of the colonies that originated the analysed isolates. As there was selected only a colony of C. albicans per

individual, other strains with higher enzymatic activity might have been excluded from the study.

Ibrahim *et al.* (13) reported that commensal strains with low production of phospholipase are able to colonize mucosas, suggesting that the secretion of these enzymes is a prerequisite for the colonization of those type of epithelium, opinion have shared with Mayser *et al.* (21). As the individuals here analysed have not presented any clinic sign of oral candidosis, it is reasonable to suppose that this higher secretion of phospholipases by the strains of non-smokers might not have bigger implications in the simple event of colonization.

The totality of strains of *C. albicans* of the non-smoking group produced chondroitinase, in comparison to 90% produced by this species in the smoking group. Among the non-albicans only *C. tropicalis* and one strain of *C. parapsilosis* produced chondroitinase. More, the two strains of *C. glabrata* have not produced chondroitinase. This result corroborates with Shimizu *et al.* (39) who observed the production of the enzyme in 97.8% of the strains of *C. albicans*, being this production followed by *C. tropicalis*, *C. guilliermondii*, *C. parapsilosis*, and *C. krusei*; it also corroborates with Chaffin *et al.* (9) who affirmed that hyaluronidase and chondroitinase may be produced by *C. albicans*, *C. tropicalis*, *C. guilliermondii*, *C. parapsilosis*, and *C. krusei*, but in a small proportion of the isolates.

In the present study, there was evaluated also the secretion of esterase/lipase. All species were able to secrete this enzyme; however, in different proportions. This result corroborates with reported by another authors (6, 9, 33, 43) that this enzyme is present in a series of species of *Candida*; however, its correlation with the pathogenic behaviour of the strains was not established yet.

The activity of haemolysins is associated to the ability that these strains have for destroying red blood cells and use the iron as nutrient. Our results have shown that all species found have produced haemolysins, in concordance with the sstudy of Luo *et al.* (19), being strongly produced in the majority of strains.

Among the strains of examined C. albicans, none of them has presented high levels of secretion for the five enzymes, concomitantly, i.e. no one has presented a value 1-Pz higher than 0.640 for the five enzymes studied. This agrees with the study of Bramono $et\ al$. (6) who researched the secretion of secreted aspartyl-protease, lipase, and α -glucosidase in strains of C. albicans, and none of them has shown high levels of activity for those three enzymes, simultaneously.

The literature suggest that tobacco's cigarettes can cause changes in the oral epithelium that allowed the *Candida* colonization (3, 25). Willians *et al.* (44) showed that a higher keratinisation favours the adherence of *Candida albicans*. Moreover, Barret-Bee *et al.* (5) proposed that the phospholipase could have an important role in this adherence; a proposition not shared with Ghannoun *et al.* (12). In the same direction, Ray & Payne (30) showed that blastopores of *C. albicans* can degrade cellular keratin using acidic proteases and Jayatilake *et al.* (14) proposed that phospholipases can also be important in this destruction, what, without further appeal, would favour the colonization.

Concerning the arguments above, maybe, the lowest secretion of the enzymes per strains obtained from smokers is due to a clonal selection determined by the quality of the saliva. As the Saps (8) and the phospholipases (20, 34) are usually active only under acidic conditions and there was already shown that smokers present a more acidic saliva and smaller buffering ability than the non-smokers (17, 18, 27, 40), it is

reasonable to suppose that in this second group, it is not necessary a high rate of secretion for the strains to manage their colonization.

According with the available literature, smokers may show a higher oral Candidal colonization due to induction of increased epithelial keratinisation (4, 25, 44); to a reduction: in function of polymorphonuclear leukocytes, in gingival exudates and in salivary immunoglobulin A levels (42).

Researches of *Candida* species, especially *C. albicans*, his virulence and the effects of chemical substances about them, can facilitate the diagnosis, change the needs of treatment e help in the prevention of oral candidosis (6).

CONCLUSION

The results obtained in this study showed that in the two populations here analysed, the oral fungal load does not differ between smoker and non-smoker. Regarding the hydrolytic enzymes, there was a higher tendency of enzymatic secretion in the strains of *Candida albicans* isolated from non-smokers.

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6. ANEXOS		

ANEXO I - Material e Métodos

O projeto de pesquisa foi submetido e aprovado pelo Comitê de Ética em Pesquisa da PUCPR, sob registro no CEP nº 658/06 (Anexo V).

Amostra

Neste estudo foram avaliados 100 indivíduos, 42 não-fumantes e 58 fumantes. Foram considerados fumantes aqueles com consumo igual ou superior a cinco cigarros por dia há pelo menos um ano, e não-fumantes aqueles que relataram nunca terem fumado (29). Indivíduos que fumavam menos de cinco cigarros por dia foram excluídos da pesquisa. Cada indivíduo foi avaliado por um especialista em Estomatologia que realizou a anamnese e o exame físico intrabucal, inspecionando as mucosas, tecidos moles e dentes. Indivíduos com manifestações clínicas de candidose bucal; relato de doenças sistêmicas na anamnese, como diabetes ou hipertensão; mulheres grávidas; indivíduos que tivessem utilizado antibióticos nos três meses que precederam as coletas ou fazendo uso de medicação sialorredutora, foram excluídos da amostra. A média de idade dos indivíduos não-fumantes foi de 19,95 ± 2,13 anos e dos fumantes de 22,83 ± 5,75 anos. A média de cigarros consumidos por dia foi $12,79 \pm 5,73$ e do tempo de uso em anos foi de 6.19 ± 5.10 . Amostras de raspado superficial (swabbing) do dorso da língua e da mucosa jugal dos indivíduos foram recolhidas, semeadas em placas contendo CHROMagar®Candida (CHROMagar Microbiology, Biomerieux, Paris, França) e incubadas a 37°C, por 48 horas. Após a identificação fenotípica (24), as cepas foram transferidas para tubos de armazenamento contendo Ágar Sabouraud Dextrose (DAS, Difco Laboratories, Detroit, Mich., USA).

Ensaios enzimáticos

Um total de 43 cepas do gênero *Candida* (1 colônia/fenótipo por indivíduo) foram submetidas às provas de atividade enzimática. A cepa-padrão de *C. albicans* CBS562 proveniente do estoque do Laboratório de Estomatologia da PUCPR foi testada como controle. Os isolados foram crescidos em estufa a 37°C em 5mL de Caldo Sabouraud Dextrose (Difco Laboratories, Detroit, Mich., USA). Após 24 horas as células foram recolhidas por centrifugação e lavadas três vezes com água destilada estéril. Os *pellets* foram ressuspendidos em água estéril até o equivalente ao tubo 10 da escala de MacFarland e 5 μL inoculados com o auxílio de discos de papel filtro esterilizados, em cinco pontos eqüidistantes, nos meios de cultura para detecção semi-quantitativa de aspartil-protease secretora (Sap), fosfolipases, condroitinase, esterase/lipase e hemolisinas. Todos os testes foram realizados em duplicata.

A secreção de Sap foi verificada seguindo o protocolo de Ruchel *et al.* (32). Uma suspensão contendo Ágar -15 g (Oxoid, Fakola AG, Basel, Suíça) e Yeast Nitrogen Base - 11,7 g (YNB, Difco Laboratories, Detroit, Mich., USA) e água destilada – 900 mL foi esterilizada por autoclave a 120°C durante 15 minutos . Após resfriamento a 55°C, esta suspensão foi acrescida de albumina bovina fração V - 2,0 g (Sigma Chem Co., St. Louis, Mo., USA), e 2,5 mL de Protovit Plus[®] (Roche, São Paulo, Brasil) dissolvidos em 100 mL de água destilada estéril e esterilizados em filtros de Millipore de 0,22 μm. O pH foi ajustado (entre 4,0 e 4,5), e a solução imediatamente vertida em placas de *petri* estéreis de 90 mm de diâmetro. As placas permaneceram em estufa a 37°C por 10 dias (10).

A atividade fosfolipásica foi pesquisada seguindo o método de Price *et al.*(28). Em 1.000 mL de água destilada estéril foram dissolvidos 10,0g de peptona (Difco Laboratories, Detroit, Mich., USA), 30,0g de glucose, 57,3g de NaCl e 0,55g CaCl₂ e 20,0g de Ágar (Oxoid). O meio foi esterelizado em autoclave a 120°C por 15 minutos e

resfriado a uma temperatura de 55°C. O *egg yolk* estéril sem telurito de potássio a 50% (Newprov Prod. Laborat., Inc., São José dos Pinhais, Paraná, Brasil) foi adicionado ao meio esterelizado. Após 4 dias de inoculação, as placas foram observadas quanto à formação de uma zona de cor amarelada ao redor das colônias,

Para a pesquisa de condroitinase foi empregado o protocolo de Smith & Willett (41) modificado. O meio de cultura foi preparado usando-se neopeptona 10g, glucose 40g, Ágar noble 15g (Sigma Chem Co., St. Louis, Mo., USA) e água destilada qsp 1.000mL e autoclavado a 120°C por 15 minutos. Após resfriamento a 55°C, foi acrescentada uma solução de albumina bovina fração V - 1,0g (Sigma) e sulfato de condroitina - 400mg dissolvidos em 10mL de água destilada estéril e esterilizados em filtros de *Millipore* de 0,22 μm. O pH final foi ajustado entre 6,8 ± 0,2. O halo formado foi medido após 48h de inoculação.

O teste de opacidade Tween[®] 80 (Sigma Chem Co., St. Louis, Mo., USA) foi utilizado para determinar a produção de esterase/lipase. O meio de cultura foi preparado com 15,0 g de SDA (Difco Laboratories, Detroit, Mich., USA) e 10 g CaCl₂, em 1.000 mL de água destilada (33). Após ser autoclavado, o meio foi resfriado a 50°C e 10 mL de Tween[®]80 (Sigma) autoclavado foi adicionado. As inoculações permaneceram em estufa a 37°C por sete dias consecutivos.

Para determinar a atividade hemolítica, as cepas foram semeadas em placas de Ágar-sangue usando o protocolo de Luo *et al.* Modificado (19), utilizando 70mL de sangue de ovelha fresco e desfibrinado (Newprov Prod. Laborat.); 30g de glucose; qsp 1.000mL de solução de Ágar-ágar 1% (Oxoid); pH final 5,6. A formação de um halo translúcido ao redor das colônias após 48h de incubação indicou atividade hemolítica positiva.

Obtenção dos resultados

Após os períodos de observação foram feitas as leituras do tamanho das colônias e dos halos formados por um único pesquisador calibrado, utilizando transiluminação e um paquímetro digital (Mitutoyo, 6" digital caliper, w/abs, Japão) e calculadas as atividades enzimáticas. Estas atividades foram determinadas de acordo com a técnica de Price *et al.* (28), dividindo-se o diâmetro da colônia (DC) pelo diâmetro do halo formado (DH). Este resultado é denominado Pz, que representa numericamente a atividade enzimática da cepa testada. Como menores valores Pz indicam maior atividade enzimática e vice-versa, optou-se por subtrair tais valores de um (1-Pz) (26). Os dados assim obtidos foram analisados quanto à sua normalidade de distribuição pelo teste de Kolmogorov-Smirnov. Para aspartil-protease secretora, condroitinase e hemolisinas houve distribuição normal dos valores e foi utilizado o teste *t* de Student. Nas enzimas onde não houve distribuição normal, fosfolipase e esterase/lipase, foi utilizado o teste não-paramétrico U de Mann-Whitney (Gráfico 2).

A atividade enzimática foi analisada estatisticamente somente nos isolados de *C. albicans*, devido à inexpressiva obtenção de outras espécies.

Além dos ensaios estatísticos, os valores 1-Pz também permitiram a classificação das cepas em zimotipos. Foram gerados dígitos a partir de um escore de atividade dado aos valores de Pz (7). Cepas não produtoras (Pz =1,000) foram codificadas como 1, pois o diâmetro do halo foi considerado igual ao da colônia; cepas com atividade média (Pz≤0,639) receberam código 2; cepas com atividade elevada (0,999≤Pz ≥0,640) receberam código 3. Os zimotipos foram gerados seguindo a ordenação dos dígitos oriundos da secreção das enzimas aspartil-protease secretora, fosfolipases, condroitinase, esterase/lipase e hemolisinas. A zimotipagem nos forneceu uma visão geral da capacidade produtora de cada cepa, associando as cinco enzimas pesquisadas.

Questionário

Questionário	Data: / / .	
Voluntário nº:	-	
Nome:		
	Período:	
Data de nasc.:	Idade:anos e	meses
Há quanto tempo foi sua última	refeição/ingestão de líquidos?	
Fumante?	Quantos cigarros por dia?	
Qual marca?	Fumante há	quanto
tempo?		
Diabético?	Compensado?	
_		
Hipertenso?	Depressivo?	
Toma algum medicamento?	Qual (is)?	
	Há quanto tempo?	
Boca seca?Quanto	os litros de líquido ingere por dia?	
Transpiração		
noturna?		_
Usuário de prótese dentária?		
Usuário de aparelho ortodôntico	o?	

ANEXO II – Análise estatística

Indivíduos não-fumantes

Cepa	espécie	aspartil	fosfolipase	condroitinase	esterase	hemolisina
NF1	C. albicans	0,660	0,520	0,435	0,848	0,472
NF2	C. albicans	1,000	0,400	0,503	0,931	0,234
NF3	C. albicans	1,000	0,468	0,579	1,000	0,572
NF4	C. albicans	1,000	0,466	0,548	0,921	0,401
NF5	C. albicans	0,941	0,501	0,478	0,943	0,703
NF7	C. albicans	0,624	0,445	0,578	0,969	0,633
NF9	C. albicans	0,725	0,500	0,633	1,000	0,333
NF11	C. albicans	0,744	0,472	0,624	1,000	0,311
NF12	C. albicans	0,588	0,500	0,368	1,000	0,575
NF13	C. albicans	0,687	0,538	0,472	0,827	0,532
NF14	C. albicans	0,803	0,444	0,420	1,000	0,363
NF1d	C. albicans	0,788	0,456	0,510	1,000	0,548
NF2d	C. albicans	0,919	0,425	0,476	0,896	0,235
NF3d	C. albicans	1,000	0,534	0,518	0,944	0,419
NF4d	C. albicans	1,000	0,516	0,610	0,920	0,441
NF5d	C. albicans	0,957	0,454	0,429	1,000	0,460
NF7d	C. albicans	0,609	0,451	0,568	0,939	0,639
NF9d	C. albicans	0,766	0,529	0,625	0,715	0,357
NF11d	C. albicans	0,722	0,450	0,591	0,752	0,274
NF12d	C. albicans	0,576	0,474	0,378	1,000	0,657
NF13d	C. albicans	0,663	0,572	0,679	0,790	0,600
NF14d	C. albicans	0,751	0,571	0,385	1,000	0,238

Indivíduos fumantes

GF1 C. albicans 0,799 0,464 0,338 0,794 0,356 GF2 C. albicans 0,575 0,425 0,432 1,000 0,325 GF3 C. albicans 0,637 0,580 0,449 0,794 0,347 GF6 C. albicans 0,891 0,462 0,538 0,828 0,439 GF6 C. albicans 0,741 0,500 0,471 1,000 0,397 GF7 C. albicans 0,430 0,457 0,649 0,885 0,508 GF9 C. albicans 0,627 0,476 0,429 0,748 0,239 GF10 C. albicans 1,000 0,791 0,531 0,633 0,360 GF12 C. albicans 1,000 0,791 0,531 0,633 0,360 GF15 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 0,833 0,475 0,476 0,855 0,513	Cepa	espécie	aspartil	fosfolipase	condroitinase	esterase	hemolisina
GF3 C. albicans 0,637 0,580 0,449 0,794 0,347 GF5 C. albicans 0,891 0,462 0,538 0,828 0,439 GF6 C. albicans 0,861 0,462 0,538 0,828 0,439 GF7 C. albicans 0,741 0,500 0,471 1,000 0,397 GF8 C. albicans 0,430 0,457 0,649 0,855 0,508 GF9 C. albicans 0,627 0,476 0,429 0,748 0,239 GF10 C. albicans 1,000 0,791 0,531 0,633 0,360 GF14 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 0,814 0,478 0,666 0,938 0,412	GF1	C. albicans	0,799	0,464	0,338	0,794	0,356
GF5 C. albiceans 0,891 0,462 0,538 0,828 0,439 GF6 C. albiceans 0,865 0,814 0,440 1,000 0,519 GF7 C. albicans 0,740 0,500 0,471 1,000 0,397 GF8 C. albicans 0,632 0,476 0,429 0,748 0,239 GF10 C. albicans 0,632 0,615 0,654 0,967 0,344 GF12 C. albicans 0,632 0,615 0,654 0,967 0,344 GF14 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 1,000 0,516 0,407 0,796 0,307 GF16 C. albicans 1,000 0,516 0,407 0,796 0,307 GF17 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 1,000 0,473 0,626 0,835 0,510	GF2	C. albicans	0,575	0,425	0,432	1,000	0,325
GF6 C. albicans 0,865 0,814 0,440 1,000 0,519 GF7 C. albicans 0,741 0,500 0,471 1,000 0,397 GF8 C. albicans 0,430 0,457 0,649 0,855 0,508 GF9 C. albicans 0,627 0,476 0,429 0,748 0,239 GF10 C. albicans 1,000 0,791 0,531 0,633 0,360 GF12 C. albicans 0,632 0,615 0,654 0,967 0,344 GF14 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 0,888 0,475 0,476 0,885 0,513 GF16 C. albicans 0,881 0,477 0,476 0,835 0,513 GF16 C. albicans 1,000 0,473 0,626 0,833 0,412 GF17 C. albicans 1,000 0,491 1,000 1,000 0,494	GF3	C. albicans	0,637	0,580	0,449	0,794	0,347
GF7 C. albicans 0,741 0,500 0,471 1,000 0,397 GF8 C. albicans 0,430 0,457 0,649 0,855 0,508 GF9 C. albicans 0,627 0,476 0,429 0,748 0,239 GF10 C. albicans 1,000 0,791 0,531 0,633 0,360 GF12 C. albicans 1,000 0,516 0,407 0,796 0,307 GF14 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 0,838 0,475 0,476 0,855 0,513 GF16 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 0,802 0,492 0,395 1,000 0,494 GF19 C. albicans 0,802 0,492 0,393 0,500 GF20	GF5	C. albicans	0,891	0,462	0,538	0,828	0,439
GF8 C. albicans 0,430 0,457 0,649 0,855 0,508 GF9 C. albicans 0,627 0,476 0,429 0,748 0,239 GF10 C. albicans 1,000 0,791 0,531 0,633 0,360 GF12 C. albicans 0,632 0,615 0,654 0,967 0,344 GF14 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 0,838 0,475 0,476 0,855 0,513 GF16 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 1,000 0,473 0,626 0,835 0,506 GF18 C. albicans 1,000 0,473 0,626 0,835 0,506 GF19 C. albicans 0,802 0,492 0,395 1,000 0,494 GF19 C. albicans 0,883 0,531 0,391 0,763 0,251	GF6	C. albicans	0,865	0,814	0,440	1,000	0,519
GF9 C. albicans 0.627 0.476 0.429 0.748 0.239 GF10 C. albicans 1,000 0,791 0,531 0,633 0,360 GF12 C. albicans 0,632 0,615 0,654 0,967 0,344 GF14 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 0,838 0,475 0,476 0,855 0,513 GF16 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 1,000 0,473 0,626 0,835 0,506 GF18 C. albicans 1,000 0,471 1,000 1,000 0,494 GF19 C. albicans 0,802 0,492 0,395 1,000 0,494 GF19 C. albicans 0,802 0,492 0,395 1,000 0,500 GF20 C. albicans 0,883 0,531 0,391 0,763 0,2251	GF7	C. albicans	0,741	0,500	0,471	1,000	0,397
GF10 C. albicans 1,000 0,791 0,531 0,633 0,360 GF12 C. albicans 0,632 0,615 0,654 0,967 0,344 GF14 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 0,838 0,475 0,476 0,855 0,513 GF16 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 1,000 0,473 0,626 0,835 0,506 GF18 C. albicans 0,802 0,492 0,395 1,000 0,494 GF19 C. albicans 1,000 0,491 1,000 1,000 0,500 GF20 C. albicans 0,883 0,531 0,391 0,763 0,251 GF21 C. albicans 0,794 0,477 0,610 0,954 0,672 GF22 C. albicans 1,000 0,506 1,000 0,932 0,00	GF8	C. albicans	0,430	0,457	0,649	0,855	0,508
GF12 C. albicans 0,632 0,615 0,654 0,967 0,344 GF14 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 0,838 0,475 0,476 0,885 0,513 GF16 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 1,000 0,473 0,626 0,835 0,506 GF18 C. albicans 1,000 0,473 0,626 0,835 0,506 GF19 C. albicans 1,000 0,491 1,000 1,000 0,494 GF19 C. albicans 1,000 0,491 1,000 1,000 0,500 GF20 C. albicans 0,794 0,477 0,610 0,954 0,672 GF21 C. albicans 1,000 0,586 1,000 0,934 0,326 GF22 C. albicans 0,851 0,500 0,792 1,000 0,487	GF9	C. albicans	0,627	0,476	0,429	0,748	0,239
GF14 C. albicans 1,000 0,516 0,407 0,796 0,307 GF15 C. albicans 0,838 0,475 0,476 0,855 0,513 GF16 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 1,000 0,473 0,626 0,835 0,506 GF18 C. albicans 0,802 0,492 0,395 1,000 0,494 GF19 C. albicans 1,000 0,491 1,000 1,000 0,500 GF20 C. albicans 0,883 0,531 0,391 0,763 0,251 GF21 C. albicans 0,794 0,477 0,610 0,954 0,672 GF22 C. albicans 1,000 0,586 1,000 0,934 0,326 GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF24 C. albicans 0,872 0,609 0,357 0,891 0,464	GF10	C. albicans	1,000	0,791	0,531	0,633	0,360
GF15 C. albicans 0,838 0,475 0,476 0,855 0,513 GF16 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 1,000 0,473 0,626 0,835 0,506 GF18 C. albicans 0,802 0,492 0,395 1,000 0,494 GF19 C. albicans 1,000 0,491 1,000 1,000 0,500 GF20 C. albicans 0,883 0,531 0,391 0,763 0,251 GF21 C. albicans 0,794 0,477 0,610 0,954 0,672 GF22 C. albicans 1,000 0,586 1,000 0,934 0,326 GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF24 C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,795 0,659 0,413 0,788 0,469	GF12	C. albicans	0,632	0,615	0,654	0,967	0,344
GF16 C. albicans 0,814 0,478 0,666 0,938 0,412 GF17 C. albicans 1,000 0,473 0,626 0,835 0,506 GF18 C. albicans 0,802 0,492 0,395 1,000 0,494 GF19 C. albicans 1,000 0,491 1,000 1,000 0,500 GF20 C. albicans 0,883 0,531 0,391 0,763 0,251 GF21 C. albicans 0,794 0,477 0,610 0,954 0,672 GF22 C. albicans 1,000 0,586 1,000 0,934 0,326 GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF24 C. albicans 0,851 0,500 0,792 1,000 0,484 GF1d C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,688 0,588 0,633 1,000 0,296	GF14	C. albicans	1,000	0,516	0,407	0,796	0,307
GF17 C. albicans 1,000 0,473 0,626 0,835 0,506 GF18 C. albicans 0,802 0,492 0,395 1,000 0,494 GF19 C. albicans 1,000 0,491 1,000 1,000 0,500 GF20 C. albicans 0,883 0,531 0,391 0,763 0,251 GF21 C. albicans 0,794 0,477 0,610 0,954 0,672 GF22 C. albicans 1,000 0,586 1,000 0,934 0,326 GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF27 C. albicans 1,000 1,000 0,355 0,941 0,484 GF1d C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,755 0,659 0,413 0,788 0,469 GF3d C. albicans 0,797 0,562 0,695 0,873 0,444	GF15	C. albicans	0,838	0,475	0,476	0,855	0,513
GF18 C. albicans 0.802 0.492 0.395 1,000 0,494 GF19 C. albicans 1,000 0,491 1,000 1,000 0,500 GF20 C. albicans 0,883 0,531 0,391 0,763 0,251 GF21 C. albicans 0,794 0,477 0,610 0,954 0,672 GF22 C. albicans 1,000 0,586 1,000 0,934 0,326 GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF27 C. albicans 1,000 1,000 0,353 0,941 0,484 GF1d C. albicans 0,688 0,588 0,633 1,000 0,992 GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF3d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,675 0,828 0,600 1,000 0,321	GF16	C. albicans	0,814	0,478	0,666	0,938	0,412
GF19 C. albicans 1,000 0,491 1,000 1,000 0,500 GF20 C. albicans 0,883 0,531 0,391 0,763 0,251 GF21 C. albicans 0,794 0,477 0,610 0,954 0,672 GF22 C. albicans 1,000 0,586 1,000 0,934 0,326 GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF27 C. albicans 1,000 1,000 0,353 0,941 0,484 GF1d C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,688 0,588 0,633 1,000 0,296 GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505	GF17	C. albicans	1,000	0,473	0,626	0,835	0,506
GF20 C. albicans 0,883 0,531 0,391 0,763 0,251 GF21 C. albicans 0,794 0,477 0,610 0,954 0,672 GF22 C. albicans 1,000 0,586 1,000 0,934 0,326 GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF27 C. albicans 1,000 1,000 0,353 0,941 0,484 GF1d C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,688 0,588 0,633 1,000 0,296 GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,437 0,470 0,660 0,937 0,514	GF18	C. albicans	0,802	0,492	0,395	1,000	0,494
GF21 C. albicans 0,794 0,477 0,610 0,954 0,672 GF22 C. albicans 1,000 0,586 1,000 0,934 0,326 GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF27 C. albicans 1,000 1,000 0,353 0,941 0,484 GF1d C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,688 0,588 0,633 1,000 0,296 GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514	GF19	C. albicans	1,000	0,491	1,000	1,000	0,500
GF22 C. albicans 1,000 0,586 1,000 0,934 0,326 GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF27 C. albicans 1,000 1,000 0,353 0,941 0,484 GF1d C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,688 0,588 0,633 1,000 0,296 GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321	GF20	C. albicans	0,883	0,531	0,391	0,763	0,251
GF23 C. albicans 0,851 0,500 0,792 1,000 0,487 GF27 C. albicans 1,000 1,000 0,353 0,941 0,484 GF1d C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,688 0,588 0,633 1,000 0,296 GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,639 0,533 0,679 0,790 0,321	GF21	C. albicans	0,794	0,477	0,610	0,954	0,672
GF27 C. albicans 1,000 1,000 0,353 0,941 0,484 GF1d C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,688 0,588 0,633 1,000 0,296 GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,825 0,693 0,372 1,000 0,444 GF12d C. albicans 1,000 0,576 0,385 0,886 0,296 <tr< td=""><td>GF22</td><td>C. albicans</td><td>1,000</td><td>0,586</td><td>1,000</td><td>0,934</td><td>0,326</td></tr<>	GF22	C. albicans	1,000	0,586	1,000	0,934	0,326
GF1d C. albicans 0,872 0,609 0,357 0,891 0,464 GF2d C. albicans 0,688 0,588 0,633 1,000 0,296 GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,825 0,693 0,372 1,000 0,444 GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 <t< td=""><td>GF23</td><td>C. albicans</td><td>0,851</td><td>0,500</td><td>0,792</td><td>1,000</td><td>0,487</td></t<>	GF23	C. albicans	0,851	0,500	0,792	1,000	0,487
GF2d C. albicans 0,688 0,588 0,633 1,000 0,296 GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,825 0,693 0,372 1,000 0,444 GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,916 0,477 0,588 0,835 0,886 0,2	GF27	C. albicans	1,000	1,000	0,353	0,941	0,484
GF3d C. albicans 0,755 0,659 0,413 0,788 0,469 GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,825 0,693 0,372 1,000 0,444 GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,817 0,501 0,652 0,952 0,488	GF1d	C. albicans	0,872	0,609	0,357	0,891	0,464
GF5d C. albicans 0,797 0,562 0,695 0,873 0,444 GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,599 0,450 0,500 1,000 0,321 GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488	GF2d	C. albicans	0,688	0,588	0,633	1,000	0,296
GF6d C. albicans 0,675 0,828 0,600 1,000 0,505 GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,825 0,693 0,372 1,000 0,444 GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584	GF3d	C. albicans	0,755	0,659	0,413	0,788	0,469
GF7d C. albicans 0,789 0,500 0,692 0,747 0,327 GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,825 0,693 0,372 1,000 0,444 GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586	GF5d	C. albicans	0,797	0,562	0,695	0,873	0,444
GF8d C. albicans 0,437 0,470 0,606 0,937 0,514 GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,825 0,693 0,372 1,000 0,444 GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,672 0,490 0,641 0,944 0,612	GF6d	C. albicans	0,675	0,828	0,600	1,000	0,505
GF9d C. albicans 0,599 0,450 0,500 1,000 0,321 GF10d C. albicans 0,825 0,693 0,372 1,000 0,444 GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612	GF7d	C. albicans	0,789	0,500	0,692	0,747	0,327
GF10d C. albicans 0,825 0,693 0,372 1,000 0,444 GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 <td>GF8d</td> <td>C. albicans</td> <td>0,437</td> <td>0,470</td> <td>0,606</td> <td>0,937</td> <td>0,514</td>	GF8d	C. albicans	0,437	0,470	0,606	0,937	0,514
GF12d C. albicans 0,639 0,533 0,679 0,790 0,321 GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542 <td>GF9d</td> <td>C. albicans</td> <td>0,599</td> <td>0,450</td> <td>0,500</td> <td>1,000</td> <td>0,321</td>	GF9d	C. albicans	0,599	0,450	0,500	1,000	0,321
GF14d C. albicans 1,000 0,576 0,385 0,886 0,296 GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542	GF10d	C. albicans	0,825	0,693	0,372	1,000	0,444
GF15d C. albicans 0,788 0,450 0,556 1,000 0,497 GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542	GF12d	C. albicans	0,639	0,533	0,679	0,790	0,321
GF16d C. albicans 0,916 0,477 0,588 0,835 0,384 GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542	GF14d	C. albicans	1,000	0,576	0,385	0,886	0,296
GF17d C. albicans 0,817 0,501 0,652 0,952 0,488 GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542	GF15d	C. albicans	0,788	0,450	0,556	1,000	0,497
GF18d C. albicans 0,822 0,538 0,449 0,904 0,584 GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542	GF16d	C. albicans	0,916	0,477	0,588	0,835	0,384
GF19d C. albicans 1,000 0,476 1,000 1,000 0,586 GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542	GF17d	C. albicans	0,817	0,501	0,652	0,952	0,488
GF20d C. albicans 0,906 0,500 0,567 0,900 0,326 GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542	GF18d	C. albicans	0,822	0,538	0,449	0,904	0,584
GF21d C. albicans 0,672 0,490 0,641 0,944 0,612 GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542	GF19d	C. albicans	1,000	0,476	1,000	1,000	0,586
GF22d C. albicans 1,000 0,675 1,000 0,685 0,393 GF23d C. albicans 0,902 0,471 0,750 0,858 0,542		C. albicans	0,906	0,500	0,567	0,900	0,326
GF23d C. albicans 0,902 0,471 0,750 0,858 0,542	GF21d	C. albicans	0,672	0,490	0,641	0,944	0,612
		C. albicans	1,000	0,675	1,000		0,393
GF27d C. albicans 1,000 1,000 0,349 0,911 0,548		C. albicans	0,902	0,471	0,750	0,858	0,542
	GF27d	C. albicans	1,000	1,000	0,349	0,911	0,548

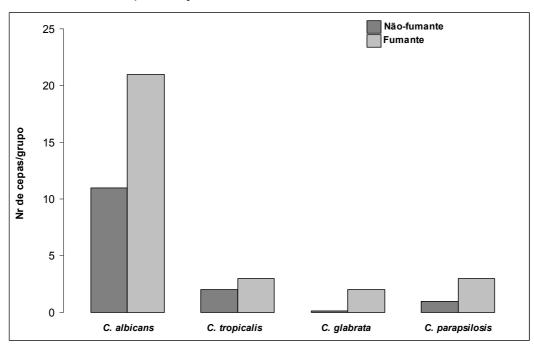
ANEXO II - Análise Estatística

Com a finalidade de se eliminar a hipótese nula de que as diferenças observadas entre os grupos são devidas ao acaso foi aplicado o teste de normalidade de Kolmogorov-Smirnov. Nas enzimas aspartil-protease, condroitinase e hemolisina, houve distribuição normal dos valores, portanto foi utilizado o teste "t" de Student. Nas enzimas onde não houve distribuição normal, fosfolipase e esterase/lipase, foi utilizado o teste não paramétrico U de Mann-Whitney. De acordo com este teste, para a atividade enzimática de fosfolipase, p=0,01, sendo estatisticamente significante, pois p<0,05.

Testes de Normalidade		Kolmogorov-Smirne	ov		
	Grupos	Statistic	df	Sig.	
condroitinase	Não Fumantes	0,114295346	22	0,2000	
	Fumantes	0,113708348	42	0,1992	
hemolisina	Não Fumantes	0,107259444	22	0,2000	
	Fumantes	0,118425577	42	0,1514	
fosfolipase	Não Fumantes	0,147174136	22	0,2000	
	Fumantes	0,218649254	42	0,0000	
esterase	Não Fumantes	0,205855777	22	0,0161	
	Fumantes	0,145593267	42	0,0255	
aspartil	Não Fumantes	0,151696582	22	0,2000	
	Fumantes	0,130404093	42	0,0701	
Grupos	Variáveis	Count	Mean	Median	Std Deviation
Não Fumantes	condroitinase	22	0,518	0,514	0,092
	hemolisina	22	0,454	0,450	0,149
	fosfolipase	22	0,486	0,473	0,046
	esterase	22	0,927	0,944	0,088
	aspartil	22	0,796	0,758	0,153
Fumantes	condroitinase	42	0,575	0,561	0,185
	hemolisina	42	0,432	0,444	0,104
	fosfolipase	42	0,563	0,500	0,140
	esterase	42	0,894	0,907	0,098
	aspartil	42	0,811	0,820	0,155
Teste U de Mann-Whitr	ney				
Ranks					
	Grupos	N	Mean Rank	Sum of Ranks	
fosfolipase	Não Fumantes	22	24,52272727	539,5	
	Fumantes	42	36,67857143	1540,5	
	Total	64			
esterase	Não Fumantes	22	36,59090909	805	
	Fumantes	42	30,35714286	1275	
	Total	64			

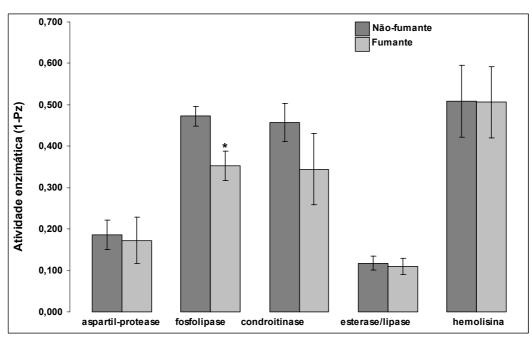
ANEXO III – GRÁFICOS

GRÁFICO 1. Distribuição das espécies isoladas



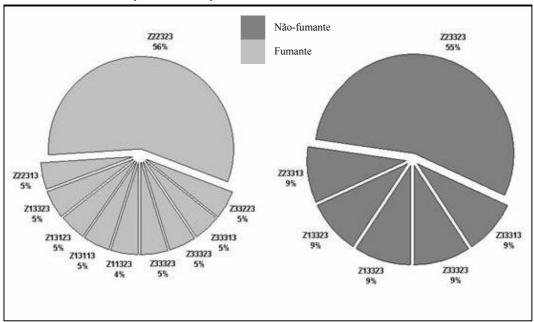
Graf. 1. Distribuição das espécies de *Candida* isoladas de não-fumantes (n=14) e fumantes.

GRÁFICO 2. Atividade enzimática das cepas de C. albicans



Graf. 2. Atividade das cepas de C. albicans isoladas de não-fumantes (n=11) e fumantes (n=21). Teste U de Mann-Whitney (p = 0.0131).

GRÁFICO 3. Distribuição dos zimotipos de *C. albicans*



Graf. 3. Distribuição dos zimotipos das cepas de *C. albicans* isoladas de não-fumantes (n=11) e fumantes (n=21).

ANEXO IV - Imagens

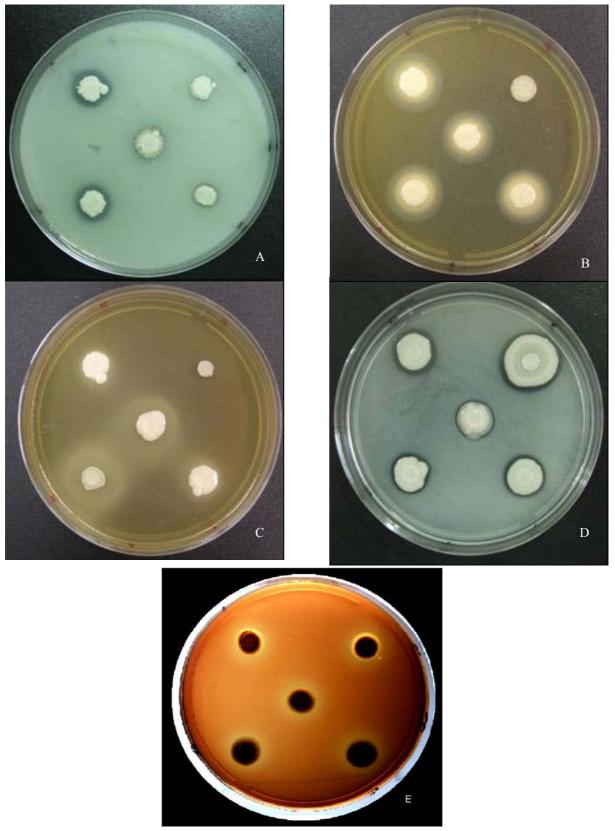


FIGURA 1. Halos de digestão característicos de cada enzima. A- aspartil-protease; B- fosfolipase; C-condroitinase; D- esterase/lipase e E- hemolisina.

ANEXO V – Termo de aprovação do Comitê de Ética em Pesquisa da PUCPR



Curitiba, 19 **de dezembro** de 2006. Of. **658**/06/CEP-PUCPR 2ªvia solicitada

Ref. "Diferenças nos perfis de virulência de cepas de cândida ssp. Isoladas de populações de tabagistas e não tabagistas"

Prezado (a) Pesquisador (es),

Venho por meio deste informar a Vossa Senhoria que o Comitê de Ética em Pesquisa da PUCPR, no dia 03 de agosto de 2005 aprovou o Projeto Intitulado "Diferenças nos perfis de virulência de cepas de cândida ssp. Isoladas de populações de tabagistas e não tabagistas", pertencente ao Grupo III, sob o registro no CEP n° 658, e será encaminhado a CONEP para o devido cadastro. Lembro ao senhor (a) pesquisador (a) que é obrigatório encaminhar relatório anual parcial e relatório final a este CEP.

Atenciosamente,

Prof^a M. Sc Ana Cristina Miguez Ribeiro Coordenadora do Comitê de Ética em Pesquisa - PUCPR

Ilma Sra Ana Claudia Santos de Azevedo Izidoro

Rua Imaculada Conosição, 1155. Prado Velho. CEP 80215.901 Curtiba. Paraná. Brasil. Caixa. Postal. 17315. CEP 80242.980. Telefax. (41) 3271.2292. www.pucpr.br

ANEXO VI – Termo de Consentimento Livre e Esclarecido

TÍTULO DO PROJETO: DIFERENÇAS NOS PERFIS DE VIRULÊNCIA DE CEPAS DE

CANDIDA spp. ISOLADAS DE POPULAÇÕES DE TABAGISTAS E NÃO-TABAGISTAS

Investigador: Cirurgiã Dentista Ana Claudia Santos de Azevedo Izidoro

Endereço: Pontificia Universidade Católica do Paraná - Centro de Ciências Biológicas e da

Saúde - Curso de Odontologia

Rua Imaculada Conceição 1155, CEP 80215-901, Curitiba - Paraná

Introdução

O pesquisador que conduz este projeto determinou que você atende aos requisitos iniciais para a participação no estudo. As informações a seguir descrevem o estudo no qual você terá o papel de participante. O pesquisador ou membro da equipe explicará os procedimentos e responderá a qualquer pergunta que você possa ter sobre este termo de consentimento informado e/ou sobre o estudo. Leia cuidadosamente este documento e não hesite em fazer perguntas sobre o objetivo do estudo ou sobre as informações fornecidas abaixo que você não compreenda. Nenhuma garantia pode ser feita quanto aos

resultados do estudo.

Finalidade do estudo

A Candida albicans é um microorganismo que pode estar presente na mucosa bucal, na saliva ou ainda dentro de bolsas periodontais ao redor dos dentes de indivíduos com periodontite. Entretanto não se sabe ao certo a virulência desse microorganismo em pacientes fumantes e não-fumantes. O presente estudo tem por objetivo avaliar a produção de enzimas produzidas por este microorganismo e se microorganismos que estão presentes em pacientes fumantes têm a mesma virulência dos encontrados em

pacientes não fumantes.

Descrição do estudo e procedimentos

Será preenchida uma ficha padronizada contendo dados pessoais, informações sobre o estado de saúde geral, tabagismo, tempo transcorrido após o último tratamento odontológico e uso de medicações.

Riscos e benefícios

De acordo com a metodologia exposta anteriormente acreditamos que o exame empregado neste estudo não seja capaz de produzir qualquer dano ou risco aos pacientes examinados. A coleta de saliva e de material na mucosa jugal não causa nenhum desconforto.

Receberei esclarecimentos sobre o que é candidose e como tratá-la e orientações sobre higiene bucal.

Perguntas relacionadas a este estudo

Você tem o direito de fazer perguntas com relação a este estudo a qualquer momento. Caso tenha novas perguntas, você poderá entrar em contato com Ana Claudia Santos de Azevedo Izidoro, no telefone 041-3274-5066.

Participação voluntária e direito de recusa/afastamento

Sua participação neste estudo é voluntária e não remunerada. Você poderá se recusar a participar ou poderá descontinuar sua participação a qualquer momento durante o estudo, sem penalidades ou perda de benefícios.

Confidencialidade		
Os registros de sua par	ticipação neste es	tudo serão mantidos em confidência, de acordo com as leis
municipais, estaduais e fede	erais. O pesquis	ador e o Comitê de Ética em Pesquisa da Pontifícia
Universidade Católica do Para	aná (CEP-PUCPF	R) poderão inspecionar e ter acesso aos dados confidenciais
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objetivos específicos desta pe	esquisa, de forma	a clara e detalhada. Recebi informações específicas sobre
cada procedimento no qual e	starei envolvido,	dos desconfortos, tanto quanto dos beneficios esperados.
Todas as minhas dúvidas fora	m respondidas co	m clareza e sei que poderei solicitar novos esclarecimentos
a qualquer momento. Além	disso, sei que	novas informações obtidas durante o estudo, me serão
fornecidas e que terei liberda	ade de retirar me	eu consentimento de participação na pesquisa, se assim o
desejar.		
Fui informado que caso exista	am danos à minh	a saúde, causados diretamente pela pesquisa, terei direito a
tratamento médico e indeniz	zação conforme	estabelece a lei. Também sei que, caso existam gastos
adicionais, estes serão abso	rvidos pelo orça	mento da pesquisa. Autorizo a liberação dos registros
médicos/odontológicos ao pe	squisador e ao (Comitê de Ética em Pesquisa da Pontificia Universidade
Católica do Paraná (CEP-PUC	CPR). Declaro ain	da que recebí cópia do presente Termo de Consentimento
Assinatura da paciente	Nome	(Local, Data e Hora)
Assinatura do pesquisador	Nome	(Local, Data e Hora)
Este formulário foi lido	para	(nome do paciente) em
/ pelo		(nome do pesquisador) enquanto eu estava
presente.		
Assinatura da testemunha	-	Nome

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APPLIED AND ENVIRONMENTAL MICROBIOLOGY

2006 INSTRUCTIONS TO AUTHORS*

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Applied and Environmental Microbiology (AEM) pub lishes descriptions of all aspects of applied microbial research, basic research on microbial ecology, and re-search of a genetic and molecular nature that focuses on microbial topics of practical value. Research must address salient microbiological principles, fundamental microbial processes, or basic questions in applied or environmental microbiology. Topics that are considered include microbiology in relation to foods, agriculture, industry, biotechnol-ogy, public health, plants, and invertebrates and basic biological properties of bacteria, fungi, algae, protozoa, and other simple eukaryotic organisms as related to micro-bial ecology. Manuscripts should report new and significant findings that advance the understanding of microbiology and upon which other scientists may build.

The microbial ecology section covers a wide range of topics on the ecology of microorganisms, including culture-independent molecular assessments that provide new insights on (i) the structure-function relationships of microorganisms, (ii) the impact of in situ conditions on community structure, and (iii) the effect of changes in microbial community composition on ecosystem func-tion. Archival phylogenetic snapshots that do not provide such insights are not acceptable for publication in AEM.

The plant microbiology section covers manuscripts dealing with all aspects of plant-microorganism interactions, including symbiotic and rhizosphere bacteria and

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Manuscripts submitted to the mycology section should be clearly of a microbiological nature and may deal with basic biology, biochemistry, genetics, or physiology of fungi, molds, yeasts, or algae. Papers dealing purely with taxonomy or phylogeny, with fungal or algal structure, or with metabolism/alteration of metabolites/toxins by animal, plant, or insect cells, tissues, or organisms are not suitable. Documentation of the distribution/occurrence of toxins or metabolites in natural samples (foods, cere-

als, grains, soils, etc.) is suitable if the work includes studies involving the isolation, occurrence, or enumera-tion of the responsible microbes in these samples. The chemical or biochemical elucidation of metabolite or toxin structures is suitable if the work includes aspects of the enzymology or biosynthesis of these compounds. Invertebrate microbiology manuscripts should address

interactions between invertebrates and microorganisms, ranging from commensalism and mutualism to parasit-ism and pathogenicity. Manuscripts describing work dealing with the metabolites or toxins from animal, plant, or insect cells or the physiology of such cells are not suitable for AEM unless it affects a microbial community or individual microorganisms.

ASM publishes a number of different journals cover-ing various aspects of the field of microbiology. Each journal has a prescribed scope which must be considered in determining the most appropriate journal for each manuscript. The following guidelines may be of assistance.

(i) AEM will consider manuscripts describing properties of enzymes and proteins that are produced by either wildtype or genetically engineered microorganisms and that are significant or have potential significance in industrial or environmental settings. Studies dealing with basic biological phenomena of enzymes or proteins or in which enzymes have been used in investigations of basic biological functions are more appropriate for the Journal of Bacteriology.

(ii) AEM will consider papers which describe the use of antimicrobial agents as tools for elucidating aspects of applied and environmental microbiology. Other papers dealing with antimicrobial agents, including manuscripts dealing with the biosynthesis and metabolism of such agents, are more appropriate for Antimicrobial Agents and Chemotherapy.

(iii) Papers on the biology of bacteriophages and other viruses are more appropriate for the Journal of Virology or the Journal of Bacteriology. AEM does, however, con-sider manuscripts dealing with viruses in relation to environmental, public health, or industrial microbiology.

 (iv) Manuscripts dealing with the immune system or with topics of basic medical interest or oral microbiology are more appropriate for Infection and Immunity. Reports of clinical investigations and environmental biol-ogy applied to hospitals should be submitted to the *Jour-*nal of Clinical Microbiology.

(v) AEM and Eukaryotic Cell (EC) accept manuscripts on population dynamics and the ecology of eukaryotic microbes. Studies of microbial communities and of microbial populations with identified economic or ecological significance, e.g., plant pathogens or symbionts, are usually more appropriate for AEM. Studies of single species of eukaryotes, especially "model" organisms or those without identified economic or ecological importance, are usually more appropriate for EC.

(vi) Manuscripts dealing with the purification and char-

^{*} Shading indicates material that has been added or significantly

acterization of enzymes or cloning of genes that have already been extensively described for other organisms will be considered for publication only if they offer experimentally supported new insights into the biological role, properties, or applications of these enzymes. Descriptions of genes or enzymes that differ only in minor ways from the prototypes are not suitable for AEM.

To best serve its readership, the journal must accept only those papers that are most significant to the field of applied and environmental microbiology. Thus, the editors will reject manuscripts that, while scientifically sound, represent only incremental extensions of other studies, are mainly confirmatory, or do not pursue a question in sufficient depth.

Questions about these guidelines may be directed to the editor in chief of the journal being considered.

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Note that a manuscript rejected by one ASM journal on scientific grounds or on the basis of its general suitability for publication is considered rejected by all other ASM journals.

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Manuscripts submitted to the journal must represent reports of original research, and the original data must be available for review by the editor if necessary.

All authors of a manuscript must have agreed to its submission and are responsible for its content (initial submission and any subsequent versions), including appropriate citations and acknowledgments, and must also have agreed that the corresponding author has the authority to act on their behalf in all matters pertaining to publication of the manuscript. The corresponding author is responsible for obtaining such agreements and for informing the coauthors of the manuscript's status throughout the submission, review, and publication process. For Authors' Corrections and Retractions, signed letters of agreement from all of the authors must be submitted (see p. 12).

By submission of a manuscript to the journal, the authors guarantee that they have the authority to publish the work and that the manuscript, or one with substantially the same content, was not published previously, is not being considered or published elsewhere, and was not rejected on scientific grounds by another ASM journal.

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the paper as supporting material.

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See p. 14 for nucleic acid sequence formatting instructions.

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The URLs of the databases mentioned above are as follows: Gene Expression Omnibus (GEO), http://www .ncbi.nlm.nih.gov/geo; ArrayExpress, http://www.ebi.ac.uk /arrayexpress; and Center for Information Biology Gene Expression Database (CIBEX), http://cibex.nig.ac.jp.

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- Elder, B. L., and S. E. Sharp. 2003. Cumitech 39, Competency assessment in the clinical laboratory. Coordinating ed., S. E. Sharp. ASM Press, Washington. D.C.
- Fitzgerald, G., and D. Shaw. In A. E. Waters (ed.), Clinical microbiology, in press. EFH Publishing Co., Boston, Mass.* {Chapter title is optional.}
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- Green, P. N., D. Hood, and C. S. Dow. 1984. Taxonomic status of some methylotrophic bacteria, p. 251–254. In R. L. Crawford and R. S. Hanson (ed.), Microbial growth on C₁ compounds. Proceedings of the 4th International Symposium. American Society for Microbiology, Washington, D.C.
- Odell, J. C. April 1970. Process for batch culturing. U.S. patent 484,363,770. {Include the name of the patented item/process if possible.}
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Maximum height: 91/16 inches (23.0 cm)

Contrast. Illustrations must contain sufficient contrast to withstand the inevitable loss of contrast and detail inherent in the printing process. See also the section on color illustrations below.

Labeling and assembly. All final lettering, labeling, tooling, etc., MUST be incorporated into the figures. It cannot be added at a later date. If a figure number is included, it must appear well outside the boundaries of the image itself. (Numbering may need to be changed at the copyediting stage.) Each figure must be uploaded as a separate file, and any multipanel figures must be assembled into one file; i.e., rather than uploading a separate file for each panel in a figure, assemble all panels in one piece and supply them as one file.

Fonts. To avoid font problems, set all type in one of the following fonts: Helvetica, Times Roman, European PI, Mathematical PI, or Symbol. All fonts other than these five must be converted to paths (or outlines) in the application with which they were created. For font use in PowerPoint images, refer to the Cadmus digital art website, http://cjs.cadmus.com/da.

Compression. Images created with Macintosh applications may be compressed with Stuffit. Images created with Windows applications may be compressed with WINZIP or PKZIP.

Color illustrations. Because the process of placing ink on paper by using printing presses is different from that used to produce a photo print or a laser print and the color rendition on images viewed on a monitor depends to some extent on monitor resolution, some differences in color and contrast between the image you submit and the image printed in the journal or published online will be evident. (Figures showing red or green fluorescence and those with a significant range of colors may be difficult or impossible to reproduce exactly.) Color illustrations must be saved as either TIFF or EPS files, according to the application used (see charts above). The mode of the TIFF or EPS file must be CMYK, not RGB. Graphics in the RGB color space are intended for display on a monitor only and will not separate correctly for print-

The cost of printing in color must be borne by the author. The current color costs may be accessed from the submission form in Rapid Review and, for accepted manuscripts, will be included in the acceptance letter sent out by ASM. Adherence to the following guidelines, in ad-dition to the general ones above, will help to minimize costs and to ensure color reproduction that is as accurate as possible.

Include only the significant portions of illustrations so that the number of printed pages containing color fig-ures is minimized. The individual panels of a single figure must be assembled in a single file, including any necessary labels. Optimal color reproduction will be obtained if the composites comprise panels containing similar colors of similar lightness or darkness. If necessary, make unlike panels into separate figures/files; this will increase the cost, but the color rendition will be more accurate since the two panels will be "scanned" separately.

Drawing

Submit graphs, charts, complicated chemical or mathematical formulas, diagrams, and other drawings as finished products not requiring additional artwork or typesetting. No part of the graph or drawing may be handwritten. All elements, including letters, numbers, and symbols, must be easily readable, and both axes of a graph must be labeled. Keep in mind that the journal is published both in print and online and that the same electronic files submitted by the authors are used to produce both.

When creating line art, please use the following guidelines:

- All art MUST be submitted at its intended publication size. For acceptable dimensions, see the Size section on p. 12.
- Avoid using screens (i.e., shading) in line art. It can be difficult and time-consuming to reproduce these images without moiré patterns. Various pattern backgrounds are preferable to screens as long as the patterns are not imported from another application. If you must use images containing screens,
 - Generate the image at line screens of 85 lines per inch or lower.
 - When applying multiple shades of gray, differentiate the gray levels by at least 20%.
 - Never use levels of gray below 20% or above 70% as they will fade out or become totally black upon scanning and reduction.
- Use thick, solid lines that are no finer than 1 point in thickness.
- No type should be smaller than 6 points at the final publication size.
- Avoid layering type directly over shaded or textured areas.
- Avoid the use of reversed type (white lettering on a black background).
- Avoid heavy letters, which tend to close up, and unusual symbols, which the printer may not be able to reproduce in the legend.
- If colors are used, avoid using similar shades of the same color and avoid very light colors.

In figure ordinate and abscissa scales (as well as table column headings), avoid the ambiguous use of numbers with exponents. Usually, it is preferable to use the Système International d'Unités (SI) symbols (μ for 10⁻⁶, m for 10⁻³, k for 10³, M for 10⁶, etc.). A complete listing of SI symbols can be found in the International Union of Pure and Applied Chemistry (IUPAC) "Manual of Symbols and Terminology for Physicochemical Quantities and Units" (Pure Appl. Chem. 21:3–44, 1970). Thus, a representation of 20,000 cpm on a figure ordinate is to be made by the number 20 accompanied by the label kcpm.

When powers of 10 must be used, the journal requires that the exponent power be associated with the number shown. In representing 20,000 cells per ml, the numeral on the ordinate would be "2" and the label would be "10⁴ cells per ml" (not "cells per ml × 10^{-4} "). Likewise, an enzyme activity of 0.06 U/ml would be shown as 6 accompanied by the label 10^{-2} U/ml. The preferred designation would be 60 mU/ml (milliunits per milliliter).

Presentation of Nucleic Acid Sequences

Nucleic acid sequences of limited length which are the primary subject of a study may be presented freestyle in the most effective format. Longer nucleic acid sequences must be presented as figures in the following format to conserve space. Print the sequence in lines of approximately 100 to 120 nucleotides in a nonproportional (monospace) font that is easily legible when published with a line length of 6 inches (ca. 15.2 cm). If possible, lines of nucleic acid sequence should be further subdivided into blocks of 10 or 20 nucleotides by spaces within the sequence or by marks above it. Uppercase and lowercase letters may be used to designate the exon-intron structure, transcribed regions, etc., if the lowercase letters remain legible at a 6-inch (ca. 15.2-cm) line length. Number the sequence line by line: place numerals, representing the first base of each line, to the left of the lines. Minimize spacing between lines of sequence, leaving room only for annotation of the sequence. Annotation may include boldface, underlining, brackets, boxes, etc. Encoded amino acid sequences may be presented, if necessary, immediately above or below the first nucleotide of each codon, by using the single-letter amino acid symbols. Comparisons of multiple nucleic acid sequences should conform as nearly as possible to the same

Figure Legends

Legends should provide enough information so that the figure is understandable without frequent reference to the text. However, detailed experimental methods must be described in the Materials and Methods section, not in a figure legend. A method that is unique to one of several experiments may be reported in a legend only if the discussion is very brief (one or two sentences). Define all symbols used in the figure and define all abbreviations that are not used in the text.

Tables

Tables that contain artwork, chemical structures, or shading must be submitted as illustrations in an acceptable format at the modification stage. The preferred format for regular tables is MS Word; however, WordPerfect and Acrobat PDF are also acceptable. Note that a straight Excel file is not currently an acceptable format. Excel files must be either embedded in a Word or WordPerfect document or converted to PDF before being uploaded. If your modified manuscript contains PDF tables, select "for reviewing purposes only" at the beginning of the file upload process.

Tables should be formatted as follows. Arrange the data so that columns of like material read down, not across. The headings should be sufficiently clear so that the meaning of the data is understandable without reference to the text. See the Abbreviations section (p. 18) of these Instructions for those that should be used in tables. Explanatory footnotes are acceptable, but more extensive table "legends" are not. Footnotes should not include detailed descriptions of the experiment. Tables must include enough information to warrant table format; those with fewer than six pieces of data will be incorporated into the text by the copy editor. Table 1 is an example of a well-constructed table.

TABLE 1. Distribution of protein and ATPase in fractions of dialyzed membranes^a

	erane Fraction	ATPase	
Membrane		U/mg of protein	Total U
Control	Depleted membrane	0.036	2.3
	Concentrated supernatant	0.134	4.82
E1 treated	Depleted membrane	0.034	1.98
	Concentrated supernatant	0.11	4.6

[&]quot;Specific activities of ATPase of nondepleted membranes from control and

Cover Photographs and Drawings

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NOMENCLATURE

Chemical and Biochemical Nomenclature

The recognized authority for the names of chemical compounds is Chemical Abstracts (CAS, Columbus, Ohio) and its indexes. The Merck Index, 13th ed. (Merck & Co., Inc., Whitehouse Station, N.J., 2001), is also an excellent source. For biochemical terminology, including abbreviations and symbols, consult Biochemical Nomenclature and Related Documents (1978; reprinted for The Biochemical Society, London, England) and the instructions to authors of the Journal of Biological Chemistry and the Archives of Biochemistry and Biophysics (first issues of each year).

Do not express molecular weight in daltons; molecular weight is a unitless ratio. Molecular mass is expressed in

For enzymes, use the recommended (trivial) name assigned by the Nomenclature Committee of the International Union of Biochemistry (IUB) as described in Enzyme Nomenclature (Academic Press, Inc., New York, N.Y., 1992) and at http://www.chem.qmul.ac.uk/iubmb /enzyme/. If a nonrecommended name is used, place the proper (trivial) name in parentheses at first use in the abstract and text. Use the EC number when one has been assigned, and express enzyme activity either in katals (preferred) or in the older system of micromoles per minute.

Nomenclature of Microorganisms

Binary names, consisting of a generic name and a specific epithet (e.g., Escherichia coll), must be used for all microorganisms. Names of categories at or above the genus level may be used alone, but specific and subspecific epithets may not. A specific epithet must be preceded by a generic name, written out in full the first time it is used in a paper. Thereafter, the generic name should be abbreviated to the initial capital letter (e.g., E. coli), provided there can be no confusion with other genera used in the paper. Names of all taxa (kingdoms, phyla, classes, orders, families, genera, species, and subspecies) are printed in italics and should be underlined (or italicized) in the manuscript; strain designations and numbers are not. Vernacular (common) names should be in lowercase roman type (e.g., streptococcus, brucella). For Salmonella, genus, species, and subspecies names should be rendered in standard form: Salmonella enterica at first use, S. enterica thereafter; Salmonella enterica subsp. arizonae at first use, S. enterica subsp. arizonae thereafter. Names of serovars should be in roman type with the first letter capitalized: Salmonella enterica serovar Typhimurium. After the first use, the serovar may also be given without a species name: Salmonella serovar Typhimurium. For other information regarding serovar designations, see Identification and Serotyping of Salmonella and an Update of the Kaufmann-White Scheme (A.C. McWhorter-Murlin and F. W. Hickman-Brenner, Centers for Disease Control and Prevention, Atlanta, Ga., 1994) and Antigenic Formulas of the Salmonella Serovars (M. Y. Popoff and L. Le Minor, WHO Collaborating Centre for Reference and Research on Salmonella, Institut Pasteur, Paris, France, 1997). For a summary of the current standards for Salmonella nomenclature and the Kaufmann-White criteria, see the articles by Brenner et al. (J. Clin. Microbiol. 38:2465-2467, 2000) and McQuiston et al. (J. Clin. Microbiol. 42:1923–1932, 2004).

The spelling of bacterial names should follow the Approved Lists of Bacterial Names (Amended) & Index of the Bacterial and Yeast Nomenclatural Changes (V. B. D. Skerman et al. ed., ASM Press, Washington, D.C., 1989) and the validation lists and notification lists published in the International Journal of Systematic and Evolutionary Microbiology (formerly the International Journal of Systematic Bacteriology) since January 1989. In addition, two sites on the World Wide Web list current approved bacterial names: Bacterial Nomenclature Up-to-Date (http://www.dsmz.de/microorganisms/main.php?contentleft_id=14) and List of Prokaryotic Names with Standing in Nomenclature (http://www.bacterio.cict.fr). If there is reason to use a name that does not have standing in nomenclature, the name should be enclosed in quotation marks in the title and at its first use in the abstract and the text and an appropriate statement concerning the nomenclatural status of the name should be made in the text. "Candidatus" species should always be set in quotation marks.

For guidelines regarding new names and descriptions of new genera and species, see the articles by Tindall (Int. J. Syst. Bacteriol. 49:1309–1312, 1999) and Stackebrandt et al. (Int. J. Syst. Evol. Microbiol. 52:1043–1047, 2002). To validate new names and/or combinations, authors must submit three copies of their published article to the International Journal of Systematic and Evolutionary Microbiology.

It is recommended that a strain be deposited in at least two recognized culture collections in different countries when that strain is necessary for the description of a new taxon (Int. J. Syst. Evol. Microbiol. 50:2239–2244, 2000).

Since the classification of fungi is not complete, it is the responsibility of the author to determine the accepted binomial for a given organism. Sources for these names include *The Yeasts: a Taxonomic Study*, 4th ed. (C. P. Kurtzman and J. W. Fell, ed., Elsevier Science Publishers B.V., Amsterdam, The Netherlands, 1998), and *Ainsworth and Bisby's Dictionary of the Fungi*, 9th ed. (P. M. Kirk, P. F. Cannon, J. C. David, and J. A. Stalpers, ed., CABI Publishing, Wallingford, Oxfordshire, United Kingdom, 2001).

Names used for viruses should be those approved by the International Committee on Taxonomy of Viruses (ICTV) and published in Virus Taxonomy: Classification and Nomenclature of Viruses, Seventh Report of the International Committee on Taxonomy of Viruses (M. H. V. van Regenmortel et al., ed., Academic Press, San Diego, Calif., 2000). In addition, the recommendations of the ICTV regarding the use of species names should generally be followed: when the entire species is discussed as a taxonomic entity, the species name, like other taxa, is italic and has the first letter and any proper nouns capitalized (e.g., Tobacco mosaic virus, Murray Valley encephalitis virus). When the behavior or manipulation of individual viruses is discussed, the vernacular (e.g., tobacco mosaic virus, Murray Valley encephalitis virus) should be used. If desired, synonyms may be added parenthetically when the name is first mentioned. Approved generic (or group) and family names may also be used.

Microorganisms, viruses, and plasmids should be given designations consisting of letters and serial numbers. It is generally advisable to include a worker's initials or a descriptive symbol of locale, laboratory, etc., in the designation. Each new strain, mutant, isolate, or derivative should be given a new (serial) designation. This designation is a serial of the serial of the

nation should be distinct from those of the genotype and phenotype, and genotypic and phenotypic symbols should not be included. Plasmids are named with a lowercase "p" followed by the designation in uppercase letters and numbers. To avoid the use of the same designation as that of a widely used strain or plasmid, check the designation against a publication database such as Medline.

Genetic Nomenclature

To facilitate accurate communication, it is important that standard genetic nomenclature be used whenever possible and that deviations or proposals for new naming systems be endorsed by an appropriate authoritative body. Review and/or publication of submitted manuscripts that contain new or nonstandard nomenclature may be delayed by the editor or the Journals Department so that they may be reviewed by the Genetics and Genomics Committee of the ASM Publications Board.

Before submission of manuscripts, authors may direct questions on genetic nomenclature to the committee's chairman: Maria Costanzo (e-mail: maria@genome stanford.edu). Such a consultation should be mentioned in the manuscript submission letter.

Bacteria. The genetic properties of bacteria are described in terms of phenotypes and genotypes. The phenotype describes the observable properties of an organism. The genotype refers to the genetic constitution of an organism, usually in reference to some standard wild type. The guidelines that follow are based on the recommendations of Demerce et al. (Genetics 54:61-76, 1966).

(i) Phenotypic designations must be used when mutant loci have not been identified or mapped. They can also be used to identify the protein product of a gene, e.g., the OmpA protein. Phenotypic designations generally consist of three-letter symbols; these are not italicized, and the first letter of the symbol is capitalized. It is preferable to use Roman or Arabic numerals (instead of letters) to identify a series of related phenotypes. Thus, a series of nucleic acid polymerase mutants might be designated Pol1, Pol2, Pol3, etc. Wild-type characteristics can be designated with a superscript plus (Pol⁺), and, when necessary for clarity, negative superscripts (Pol-) can be used to designate mutant characteristics. Lowercase superscript letters may be used to further delineate phenotypes (e.g., Str for streptomycin resistance). Phenotypic designations should be defined.

(ii) Genotypic designations are also indicated by three-letter locus symbols. In contrast to phenotypic designations, these are lowercase italic (e.g., ara his rps). If several loci govern related functions, these are distinguished by italicized capital letters following the locus symbol (e.g., araA araB araC). Promoter, terminator, and operator sites should be indicated as described by Bachmann and Low (Microbiol. Rev. 44:1-56, 1980), e.g., lacZp, lacAt, and lacZo.

(iii) Wild-type alleles are indicated with a superscript plus (ara⁺ his⁺). A superscript minus is not used to indicate a mutant locus; thus, one refers to an ara mutant rather than an ara strain.

(iv) Mutation sites are designated by placing serial isolation numbers (allele numbers) after the locus symbol (e.g., araA1 araA2). If it is not known in which of several related loci the mutation has occurred, a hyphen is used instead of the capital letter (e.g., ara-23). It is essential in papers reporting the isolation of new mutants that allele numbers be given to the mutations. For Escherichia coli, there is a registry of such numbers: E. coli Genetic Stock Center, Department of Biology, Yale University, New Haven, CT 06511-5188. For the genus Salmonella, the registry is Salmonella Genetic Stock Center, Department of Biology, University of Calgary, Calgary, Alberta T2N 1N4, Canada. For the genus Bacillus, the registry is Bacillus Genetic Stock Center, Ohio State University, Columbus, OH 43210.

(v) The use of superscripts with genotypes (other than + to indicate wild-type alleles) should be avoided. Designations indicating amber mutations (Am), temperaturesensitive mutations (Ts), constitutive mutations (Con), cold-sensitive mutations (Cs), production of a hybrid protein (Hyb), and other important phenotypic properties should follow the allele number [e.g., araA230(Am) hisD21(Ts)]. All other such designations of phenotype must be defined at the first occurrence. If superscripts must be used, they must be approved by the editor and defined at the first occurrence in the text.

Subscripts may be used in two situations. Subscripts may be used to distinguish between genes (having the same name) from different organisms or strains, e.g., his E. coli or his K-12 for the his genes of E. coli or strain K-12 in another species or strain, respectively. An abbreviation may also be used if it is explained. Similarly, a subscript is also used to distinguish between genetic elements that have the same name. For example, the promoters of the gin operon can be designated ginAp1 and ghtAp2. This form departs slightly from that recommended by Bachmann and Low (e.g., desCIp).

(vi) Deletions are indicated by the symbol Δ placed before the deleted gene or region, e.g., Δπp.4432, Δ(aroPaceE)419, or Δhis(dhuA hisJ hisQ)1256. Similarly, other symbols can be used (with appropriate definition). Thus, a fusion of the ara and lac operons can be shown as Φ(ara-lac)95. Likewise, Φ(araB'-lacZ'+)96 indicates that the fusion results in a truncated araB gene fused to an intact lacZ gene, and $\Phi(malE-lacZ)97(Hyb)$ shows that a hybrid protein is synthesized. An inversion is shown as IN(mD-mE)1. An insertion of an E. coli his gene into plasmid pSC101 at zero kilobases (0 kb) is shown as pSC101 Ω(0kb::K-12hisB)4. An alternative designation of an insertion can be used in simple cases, e.g., galT236 :: Tn5. The number 236 refers to the locus of the insertion, and if the strain carries an additional gal mutation, it is listed separately. Additional examples, which utilize a slightly different format, can be found in the papers by Campbell et al. and Novick et al. cited below. It is important in reporting the construction of strains in which a mobile element was inserted and subsequently deleted that this fact be noted in the strain table. This can be done by listing the genotype of the strain used as an intermediate in a table footnote or by making a direct or parenthetical remark in the genotype, e.g., (F-), \(\Delta Mu cts \), or mal::ΔMu cts::lac. In setting parenthetical remarks within the genotype or dividing the genotype into constituent elements, parentheses and brackets are used without special meaning; brackets are used outside parentheses. To indicate the presence of an episome, parentheses (or brackets) are used (λ, F⁺). Reference to an integrated episome is indicated as described above for inserted elements, and an exogenote is shown as, for example, W3110/F'8(gal⁺).

For information about the symbols in current use, consult Berlyn (Microbiol, Mol. Biol. Rev. 62:814-984. 1998) for E. coli K-12. Sanderson and Roth (Microbiol, Rev. 52:485-532, 1988) for Salmonella serovar Typhimurium, Holloway et al. (Microbiol. Rev. 43:73-102, 1979) for the genus Pseudomonas, Piggot and Hoch (Microbiol. Rev. 49:158-179, 1985) for Bacillus subtilis. Perkins et al. (Microbiol. Rev. 46:426-570, 1982) for Neurospora crassa, and Mortimer and Schild (Microbiol. Rev. 49:181-213, 1985) for Saccharomyces cerevisiae. For yeasts, Chlamydomonas spp., and several fungal species, symbols such as those given in the Handbook of Microbiology (A. I. Laskin and H. A. Lechevalier, ed., CRC Press, Inc., Cleveland, Ohio, 1974) should be used.

Conventions for naming genes. It is recommended that (entirely) new genes be given names that are mnemonics of their function, avoiding names that are already assigned and earlier or alternative gene names, irrespective of the bacterium for which such assignments have been made. Similarly, it is recommended that, whenever possible, homologous genes present in different organisms receive the same name. When homology is not apparent or the function of a new gene has not been established, a provisional name may be given by one of the following methods. (i) The gene may be named on the basis of its map location in the style yaaA, analogous to the style used for recording transposon insertions (zef) as discussed below. A list of such names in use for E. coli has been published by Rudd (Microbiol. Mol. Biol. Rev. 62: 985-1019, 1998). (ii) A provisional name may be given in the style described by Demerec et al. (e.g., usg, gene upstream of folc). Such names should be unique, and names such as orf or genX should not be used. For reference, the E. coli Genetic Stock Center's database includes an updated listing of E. coli gene names and gene products. It is accessible on the Internet (http://cgsc.biology.yale.edu /cgsc.html). The Center's relational database can also be searched via Telnet; for access, send a request to berlyn @cgsc.biology.yale.edu. A list can also be found in the work of Riley (Microbiol. Rev. 57:862-952, 1993). For the genes of other bacteria, consult the references given above.

"Mutant" versus "mutation." Keep in mind the distinction between a mutation (an alteration of the primary sequence of the genetic material) and a mutant (a strain carrying one or more mutations). One may speak about the mapping of a mutation, but one cannot map a mutant. Likewise, a mutant has no genetic locus, only a phenotype. "Homology" versus "similarity." For use of terms that describe relationships between genes, consult the articles by Theissen (Nature 415:741, 2002) and Fitch (Trends Genet. 16:227-231, 2000). "Homology" implies a relationship between genes that share a common evolutionary origin; partial homology is not recognized. When sequence comparisons are discussed, it is more appropriate to use the term "percent sequence similarity" or "percent sequence identity," as appropriate.

Strain designations. Do not use a genotype as a name (e.g., "subsequent use of leuC6 for transduction"). If a strain designation has not been chosen, select an appropriate word combination (e.g., "another strain containing the leuC6 mutation").

"Natural" versus "artificial" transformation. Natural transformation is a process whereby the recipient cell has the inherent capacity to take up and integrate exogenous DNA into its genome. As such, natural transformation is part of the biology of the recipient cell line and should not be confused with processes through which integration of DNA is forced upon recipient cells.

Viruses. The genetic nomenclature for viruses differs from that for bacteria. In most instances, viruses have no phenotype, since they have no metabolism outside host cells. Therefore, distinctions between phenotype and genotype cannot be made. Superscripts are used to indicate hybrid genomes. Genetic symbols may be one, two, or three letters. For example, a mutant strain of λ might be designated A Aam11 int2 red114 c1857; this strain carries mutations in genes cI, bit, and red and an amber-suppressible (am) mutation in gene A. A strain designated λ at 434 would represent a hybrid of phage λ which carries the immunity region (irran) of phage 21 and the attachment (att) region of phage 434. Host DNA insertions into viruses should be delineated by square brackets, and the genetic symbols and designations for such inserted DNA should conform to those used for the host genome. Genetic symbols for phage λ can be found in reports by Szybalski and Szybalski (Gene 7:217-270, 1979) and Echols and Munialdo (Microbiol, Rev. 42:577-591, 1978).

Eukaryotes. For information about the genetic nomenclature of eukaryotes, see the Instructions to Authors for Eukaryotic Cell and Molecular and Cellular Biology.

Transposable elements, plasmids, and restriction enzymes. Nomenclature of transposable elements (insertion sequences, transposons, phage Mu, etc.) should follow the recommendations of Campbell et al. (Gene 5: 197–206, 1979), with the modifications given in section viabove. The Internet site where insertion sequences of eubacteria and archaea are described and new sequences can be recorded is http://www-is.biotoul.fr/s.html.

The system of designating transposon insertions at sites where there are no known loci, e.g., zef-123::Tn5, has been described by Chumley et al. (Genetics 91:639–655, 1979).

The nomenclature recommendations of Novick et al. (Bacteriol. Rev. 40:168–189, 1976) for plasmids and plasmid-specified activities, of Low (Bacteriol. Rev. 36:587–607, 1972) for F' factors, and of Roberts et al. (Nucleic Acids Res. 31:1805–1812, 2003) for restriction enzymes, DNA methyltransferases, homing endonudeases, and their genes should be used when possible. The nomenclature for recombinant DNA molecules constructed in vitro follows the nomenclature for insertions in general. DNA inserted into recombinant DNA molecules should be described by using the gene symbols and conventions for the organism from which the DNA was obtained.

Tetracycline resistance determinants. The nomenclature for tetracycline resistance determinants is based on the proposal of Levy et al. (Antimicrob. Agents Chemother. 43:1523–1524, 1999). The style for such determinants is, e.g., Tet B; the space helps distinguish the determinant designation from that for phenotypes and proteins (TetB). The above-referenced article shows the correct format for genes, proteins, and determinants in this family.

ABBREVIATIONS AND CONVENTIONS

Verb Tense

ASM strongly recommends that for clarity you use the past tense to narrate particular events in the past, including the procedures, observations, and data of the study that you are reporting. Use the present tense for your own general conclusions, the conclusions of previous researchers, and generally accepted facts. Thus, most of the abstract, Materials and Methods, and Results will be in the past tense, and most of the introduction and some of the Discussion will be in the present tense.

Be aware that it may be necessary to vary the tense in a single sentence. For example, it is correct to say "White (30) demonstrated that XYZ cells grow at pH 6.8," "Figure 2 shows that ABC cells failed to grow at room temperature," and "Air was removed from the chamber and the mice died, which proves that mice require air." In reporting statistics and calculations, it is correct to say "The values for the ABC cells are statistically significant, indicating that the drug inhibited...."

For an in-depth discussion of tense in scientific writing, see p. 207-209 in How To Write and Publish a Scientific Paper, 5th ed.

Abbreviations

General. Abbreviations should be used as an aid to the reader rather than as a convenience to the author, and therefore their use should be limited. Abbreviations other than those recommended by the IUPAC-IUB (Biochemical Nomenclature and Related Documents, 1978) should be used only when a case can be made for necessity, such as in tables and figures.

It is often possible to use pronouns or to paraphrase a long word after its first use (e.g., "the drug" or "the substrate"). Standard chemical symbols and trivial names or their symbols (folate, Ala, Leu, etc.) may also be used. It is strongly recommended that all abbreviations except those listed below be introduced in the first paragraph in Materials and Methods. Alternatively, define each abbreviation and introduce it in parentheses the first time it is used; e.g., "cultures were grown in Eagle minimal essential medium (MEM)." Generally, elimi-

nate abbreviations that are not used at least three times in the text (including tables and figure legends).

Not requiring introduction. In addition to abbreviations for Système International d'Unités (SI) units of measurement, other common units (e.g., bp, kb, and Da), and chemical symbols for the elements, the following should be used without definition in the title, abstract, text, figure legends, and tables: DNA (deoxyribonucleic acid); cDNA (complementary DNA); RNA (ribonucleic acid); cRNA (complementary RNA); RNase (ribonuclease); DNase (deoxyribonuclease); rRNA (ribosomal RNA); mRNA (messenger RNA); tRNA (transfer RNA); AMP, ADP, ATP, dAMP, ddATP, GTP, etc. (for the respective 5' phosphates of adenosine and other nucleosides) (add 2'-, 3'-, or 5'- when needed for contrast); ATPase, dGTPase, etc. (adenosine triphosphatase, deoxyguanosine triphosphatase, etc.); NAD (nicotinamide adenine dinucleotide); NAD+ (nicotinamide adenine dinucleotide, oxidized); NADH (nicotinamide adenine dinucleotide, reduced); NADP (nicotinamide adenine dinucleotide phosphate); NADPH (nicotinamide adenine dinucleotide phosphate, reduced); NADP+ (nicotinamide adenine dinucleotide phosphate, oxidized); poly(A), poly(dT), etc. (polyadenylic acid, polydeoxythymidylic acid, etc.); oligo(dT), etc. (oli-godeoxythymidylic acid, etc.); UV (ultraviolet); PFU (plaque-forming units); CFU (colony-forming units); MIC (minimal inhibitory concentration); Tris [tris(hydroxymethyl)aminomethane]; DEAE (diethylaminoethyl); EDTA (ethylenediaminetetraacetic acid); EGTA [ethylene glycol-bis(β-aminoethyl ether)-N,N,N',N'-tetraacetic acid]; HEPES (N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid); PCR (polymerase chain reaction); and AIDS (acquired immunodeficiency syndrome). Abbreviations for cell lines (e.g., HeLa) also need not be defined.

The following abbreviations should be used without definition in tables:

amt (amount) approx (approximately) avg (average) concn (concentration) diam (diameter) expt (experiment) exptl (experimental) ht (height) mo (month) mol wt (molecular weight) no. (number) prepn (preparation) SD (standard deviation)

SE (standard error) SEM (standard error of the mean) sp act (specific activity) sp gr (specific gravity) temp (temperature) tr (trace) vol (volume) vs (versus) wk (week) wt (weight) yr (year)

Reporting Numerical Data

Standard metric units are used for reporting length, weight, and volume. For these units and for molarity, use the prefixes m, μ, n, and p for 10⁻³, 10⁻⁶, 10⁻¹ 10⁻¹², respectively. Likewise, use the prefix k for , respectively. Likewise, use the prefix k for 103. Avoid compound prefixes such as mµ or µµ. Parts per million (ppm) may be used when that is the common measure for the science in that field. Units of temperature are presented as follows: 37°C or 324 K.

When fractions are used to express such units as enzymatic activities, it is preferable to use whole units, such as g or min, in the denominator instead of fractional or multiple units, such as µg or 10 min. For example, "pmol/min" is preferable to "nmol/10 min," and "µmol/ g" is preferable to "nmol/µg." It is also preferable that an unambiguous form, such as exponential notation, be used; for example, "µmol g⁻¹ min⁻¹" is preferable to "µmol/g/min." Always report numerical data in the ap-plicable SI units.

Representation of data as accurate to more than two significant figures must be justified by presentation of appropriate statistical analyses.

For a review of some common errors associated with statistical analyses and reports, plus guidelines on how to avoid them, see the article by Olsen (Infect. Immun. 71:6689-6692, 2003).

For a review of basic statistical considerations for virology experiments, see the article by Richardson and Overbaugh (J. Virol. 79:669-676, 2005).

Statistics

If biological variation within a treatment (coefficient of variation, the standard deviation divided by the mean) is small (less than 10%) and the difference among treatment means is large (greater than 3 standard deviations), it is not necessary to report statistics. If the data do not meet these criteria, however, the authors must include an appropriate statistical analysis (e.g., Student's t test, analysis of variance, Tukey's test, etc.). Statistics should represent the variation among biological units (e.g., replicate incubations) and not just the variation due to method of analysis.

Phylogenetic trees based on nucleotide or amino acid sequence alignments must be supported by appropriate statistical analyses of tree stability (e.g., bootstrap analysis), and nonsupported branches (e.g., bootstrap coefficients below 50%) should be collapsed. A copy of the alignment should be available for examination by the editor or the reviewers upon request.

For a review of some common errors associated with statistical analyses and reports, plus guidelines on how to avoid them, see the article by Olsen (Infect. Immun. 71:6689-6692, 2003).

For a review of basic statistical considerations for virology experiments, see the article by Richardson and Overbaugh (J. Virol. 79:669-676, 2005).

Equations

In mathematical equations, indicate the order of operations clearly by enclosing operations in parentheses, brackets, and braces, in that order: $(a+b) \times c$ or $a+(b \times c)$, $100 \times \{[(a/b) \times c] + d\}$ or $100 \times \{a/[(b \times c) + d]\}$. Italicize (by underlining) variables and constants (but not numerals), and use roman type for designations: E₀, E_h, M_t , K_{re} , K_x , a+2b=1.2 mM, Ca^{2+} $V_{max}=\exp(1.5x+y)$, BOD = $2.7x^2$.

Isotopically Labeled Compounds

For simple molecules, isotopic labeling is indicated in the chemical formula (e.g., ¹³CO₂, ³H₂, and H³⁵SO₄). Brackets are not used when the isotopic symbol is attached to the name of a compound that in its natural state does not contain the element (e.g., ³²S-ATP) or to a word that is not a specific chemical name (e.g., ¹³¹I-

labeled protein, 14C-amino acids, and 3H-ligands).

For specific chemicals, the symbol for the isotope introduced is placed in brackets directly preceding the part of the name that describes the labeled entity. Note that configuration symbols and modifiers precede the isotopic symbol. The following examples illustrate correct usage.

[14C]urea L-[methyl.14C] methionine [2,3-14] serine [a-14C] hysine [y-12P] ATP UDP-[U-²⁴C]glucose E coli [²²P]DNA fructose 1,6-[1-²²P]bisphosphate

AEM follows the same conventions for isotopic labeling as the *Journal of Biological Chemistry*, and moredetailed information can be found in the instructions to authors of that journal (first issue of each year).