

CONCLUSION GENERALE

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Le but de ce travail était d'évaluer la place de la PCR *Aspergillus* dans le diagnostic de l'aspergillose invasive.

La première étude réalisée sur les lavages broncho-alvéolaires a établi clairement la nécessité d'inclure la PCR *Aspergillus* dans le diagnostic de l'aspergillose invasive. Pour que ce test soit utile, il faut cependant que la réponse soit rendue dans les 24 à 48 heures après la réception du prélèvement. Or, à l'heure actuelle, seules les nouvelles technologies permettant de détecter les produits d'amplification en temps réel offrent cette possibilité. En effet, ces techniques plus rapides permettent d'augmenter la fréquence des analyses et par conséquent la diversité des recherches par PCR.

Dans le cadre de pathologies opportunistes où la rapidité du diagnostic étiologique est souvent cruciale, l'association de la recherche de cibles différentes est un atout appréciable. On peut ainsi envisager de rechercher par PCR, en association, *Aspergillus*, Cytomégalovirus, Toxoplasme, *Pneumocystis carinii*, le virus *herpes simplex*... sur le même prélèvement. Le développement de puces à ADN applicables au secteur de la microbiologie (ce qui est loin d'être encore le cas) rendra la batterie infectieuse investiguée encore plus large.

Cependant, si la recherche au niveau des lavages broncho-alvéolaires est importante dans le cadre de l'apparition de troubles respiratoires chez un patient immunodéprimé, la détection de l'agent infectieux, avant l'apparition de signes cliniques notables permettra de traiter les pathologies infectieuses de façon plus efficace. Dans le cadre des aspergilloses invasives, la détection de l'antigène galactomannane est pour le moment le seul paramètre qui semble toutefois utile selon les données de la littérature. La PCR *Aspergillus* ne fait pas l'unanimité des publications. L'étude réalisée ici sur modèle animal a permis de montrer que la PCR *Aspergillus* était outil complémentaire, à réaliser en parallèle avec l'antigénémie galactomannane, dans la mesure où les deux tests ne sont pas toujours positifs en même temps.

En effet, peu d'études portent sur le suivi de patients immunodéprimés par PCR *Aspergillus* avec détection simultanée de l'antigène galactomannane. Peu de centres, en effet, réalisent la recherche d'ADN *Aspergillus* par PCR, arguant le risque important de contamination des réactifs utilisés en PCR et donc de faux positifs. La commercialisation de kits PCR complets permettra sans doute de

remédier à ce problème. Cependant, pour la plupart des microbiologistes, la priorité de la détection des champignons in vasifs vient après celle de la recherche des bactéries et des virus.

Les perspectives immédiates de ce travail sont donc la développement d'une PCR en temps réel et sa comparaison aux deux techniques précédemment développées. Son évaluation sur tous les liquides biologiques sera nécessaire à la fois en terme de diagnostic ponctuel que dans le cadre de suivi de patients à risque en parallèle avec d'autres recherches de pathogènes opportunistes.

D'autre part, une meilleure connaissance du génome d'*Aspergillus* qui fait l'objet d'un séquençage dans le cadre d'un projet européen, permettra sans doute d'améliorer la définition des cibles à utiliser lors du choix des amorces et donc de rendre les techniques PCR encore plus performantes.

Ainsi, le diagnostic et le traitement de la pathologie aspergillaire sont en pleine évolution. D'une part l'arsenal thérapeutique s'ouvre à de nouvelles molécules moins toxiques que le « gold standard » l'amphotéricine B et néanmoins tout aussi efficaces sinon davantage. D'autre part les outils moléculaires qui doivent désormais faire partie de la routine des laboratoire hospitaliers ouvre la perspective d'un diagnostic plus rapide ce qui augmente les chances de succès d'une chimiothérapie plus précoce et mieux ciblée.

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6. BIBLIOGRAPHIE

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ANNEXES

**ANNEXE 1 : Valeurs des CMI des 4 antifongiques vis à vis des 106
souches d'*Aspergillus fumigatus* .**

Tableau A. Valeurs des CMI des 106 souches d'*A. fumigatus* vis à vis des 4 antifongiques testés.

N° souche	Amphotéricine B		Itraconazole		Voriconazole		Posaconazole	
	0,03125 à 16 µg/ml		0,03125 à 16 µg/ml		0,03125 à 16 µg/ml		0,004 à 16 µg/ml	
	Lecture visuelle	630 nm	Lecture visuelle	630 nm	Lecture visuelle	630 nm	Lecture visuelle	630 nm
	48 H	48 H	48 H	48 H	48 H	48 H	48 H	48 H
	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)
1	1	0,5	0,5	0,5	0,25	0,25	0,01563	0,01563
2	1	1	0,25	0,5	0,25	0,25	0,01563	0,01563
3	0,25	0,5	0,25	0,5	0,25	0,25	0,03125	0,03125
4	0,5	0,5	0,25	0,25	0,25	0,25	0,01563	0,01563
5	1	1	0,25	0,25	0,25	0,25	0,01563	0,01563
6	1	0,5	0,25	0,5	0,25	0,5	0,03125	0,03125
7	1	1	0,25	0,25	0,25	0,25	0,01563	0,01563
8	1	0,5	0,5	0,5	0,5	0,5	0,01563	0,01563
9	1	1	0,5	0,5	0,25	0,25	0,03125	0,03125
10	0,5	0,5	0,25	0,25	0,25	0,25	0,03125	0,03125
11	0,25	0,25	0,25	0,25	0,0625	0,0625	0,01563	0,01563
12	0,5	0,5	0,25	0,25	0,25	0,5	0,01563	0,01563
13	1	1	0,25	0,25	0,25	0,25	0,01563	0,03125
14	1	1	0,25	0,25	0,25	0,25	0,03125	0,0625
15	1	1	0,5	0,5	0,25	0,25	0,03125	0,01325
16	2	2	0,5	0,5	0,25	0,25	0,01563	0,01563
17	1	1	0,5	0,5	0,25	0,25	0,03125	0,03125
18	1	1	0,25	0,25	0,25	0,25	0,01563	0,01563
19	1	1	0,5	0,5	0,25	0,25	0,01563	0,01563
20	1	1	0,5	0,5	0,25	0,25	0,01563	0,03125
21	1	1	0,5	0,5	0,125	0,25	0,03125	0,0625
22	1	1	1	1	0,125	0,125	0,03125	0,03125
23	2	2	0,25	0,25	0,25	0,25	0,03125	0,03125
24	1	1	0,5	0,5	0,5	0,25	0,01563	0,01563
25	1	1	0,25	0,25	0,25	0,25	0,03125	0,03125
26	0,5	0,5	0,25	0,25	1	1	0,03125	0,0625
27	1	1	0,25	0,25	0,25	0,25	0,03125	0,03125
28	0,5	0,5	0,5	0,25	0,25	0,25	0,01563	0,01563
29	0,5	0,5	0,5	0,5	0,25	0,25	0,01563	0,03125
30	0,5	0,5	0,5	0,5	0,25	0,25	0,03125	0,03125
31	1	1	0,5	0,5	0,25	0,25	0,01563	0,01563
32	0,5	0,5	0,5	0,5	0,25	0,5	0,01563	0,01563
33	1	1	0,5	0,5	0,25	0,25	0,03125	0,03125
34	0,5	0,5	0,5	0,5	0,25	0,5	0,03125	0,03125

N° de souche en vert: environnement; N° en rouge: invasion; N° en noir: colonisation.

Tableau A. Valeurs des CMI des 106 souches d'*A. fumigatus* vis à vis des 4 antifongiques testés.

	Amphotéricine B		Itraconazole		Voriconazole		Posaconazole	
	0,03125 à 16 µg/ml		0,03125 à 16 µg/ml		0,03125 à 16 µg/ml		0,004 à 16 µg/ml	
	Lecture visuelle	630 nm	Lecture visuelle	630 nm	Lecture visuelle	630 nm	Lecture visuelle	630 nm
N° souche	48 H CMI (µg/ml)	48 H CMI (µg/ml)	48 H CMI (µg/ml)	48 H CMI (µg/ml)	48 H CMI (µg/ml)	48 H CMI (µg/ml)	48 H CMI (µg/ml)	48 H CMI (µg/ml)
35	2	2	0,5	0,5	0,25	0,25	0,01563	0,01563
36	0,5	0,5	0,5	0,5	0,25	0,25	0,03125	0,03125
37	0,5	0,5	0,5	0,5	0,25	0,25	0,01563	0,01563
38	1	2	0,5	0,5	0,5	0,5	0,01563	0,01563
39	0,5	0,5	0,5	0,25	0,5	0,5	0,03125	0,03125
40	1	1	0,5	0,5	1	1	0,03125	0,03125
41	0,25	0,25	0,5	0,25	0,25	0,25	0,01563	0,01563
42	1	1	1	1	0,25	0,25	0,03125	0,03125
43	1	1	0,125	0,25	0,5	1	0,03125	0,03125
44	0,25	0,5	0,25	0,25	0,25	0,25	0,01563	0,01563
45	0,5	0,5	0,25	0,125	0,25	0,25	0,01563	0,01563
46	0,5	0,5	0,25	0,25	0,5	0,5	0,01563	0,01563
47	0,25	0,25	0,25	0,25	0,5	0,5	0,03125	0,03125
48	0,5	0,5	0,25	0,25	0,25	0,25	0,01563	0,01563
49	0,5	0,5	0,5	0,5	0,25	0,25	0,03125	0,03125
50	0,5		0,25	0,125	0,25	0,125	0,0625	0,0625
51	0,5	0,5	0,25	0,25	0,25	0,25	0,01563	0,01563
52	0,5	0,5	0,5	0,5	0,25	0,25	0,03125	0,03125
53	1	1	1	1	0,25	0,25	0,01563	0,03125
54	0,5	0,5	0,0625	0,0625	0,25	0,25	0,01563	0,01563
55	0,25	0,25	0,25	0,25	0,125	0,0625	0,03125	0,03125
56	1	1	0,25	0,25	0,25	0,25	0,125	0,125
57	0,5	0,5	0,25	0,25	0,5	0,5	0,03125	0,03125
58	1	1	0,5	1	0,25	0,25	0,01563	0,01563
59	0,5	0,5	0,5	0,25	0,25	0,25	0,01563	0,01563
60	1	0,5	1	1	0,5	0,5	0,03125	0,0625
61	0,5	0,5	0,25	0,5	0,25	0,25	0,03125	0,03125
62	1	0,5	0,25	0,25	0,5	0,5	0,03125	0,03125
63	1	1	0,5	0,5	0,125	0,125	0,01563	0,01563
64	1	0,5	0,5	0,5	0,25	0,25	0,0312	0,0312
65	1	0,5	0,25	0,25	0,125	0,125	0,0156	0,0156
66	1	1	0,5	0,5	0,5	0,5	0,0156	0,0156
67	1	1	0,5	1	0,5	0,5	0,01563	0,01563

N° de souche en vert: environnement; N° en rouge: invasion; N° en noir: colonisation.

Tableau A. Valeurs des CMI des 106 souches d'*A. fumigatus* vis à vis des 4 antifongiques testés.

N° souche	Amphotéricine B		Itraconazole		Voriconazole		Posaconazole	
	0,03125 à 16 µg/ml		0,03125 à 16 µg/ml		0,03125 à 16 µg/ml		0,004 à 16 µg/ml	
	Lecture visuelle	630 nm	Lecture visuelle	630 nm	Lecture visuelle	630 nm	Lecture visuelle	630 nm
	48 H	48 H	48 H	48 H	48 H	48 H	48 H	48 H
	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)
68	1	1	1	1	0,25	0,25	0,0078	0,0078
69	0,5	0,5	0,25	0,25	0,25	0,25	0,01563	0,01563
70	1	1	1	1	0,25	0,25	0,01563	0,01563
71	0,25	0,25	0,25	0,25	0,125	0,125	0,0078	0,0078
72	0,5	0,5	0,25	0,25	0,25	0,25	0,0078	0,01563
73	0,5	0,5	0,5	0,5	0,5	0,5	0,01563	0,0156
74	1	2	0,25	0,5	0,25	0,25	0,01563	0,01563
75	1	0,5	0,5	0,5	0,5	0,5	0,03125	0,03125
76	0,5	0,5	0,5	0,5	0,5	0,5	0,01563	0,03125
77	1	1	0,5	0,5	0,25	0,5	0,03125	0,03125
78	1	1	0,5	0,5	0,25	0,25	0,01563	0,01563
79	1	1	1	0,5	0,125	0,125	0,01563	0,01563
80	1	1	0,125	0,125	0,25	0,25	0,01563	0,01563
81	1	1	0,5	0,5	0,5	0,5	0,01563	0,01563
82	1	1	0,125	0,25	0,25	0,25	0,03125	0,01563
83	0,5	0,5	1	1	0,5	0,5	0,03125	0,03125
84	1	1	1	1	0,5	0,5	0,03125	0,03125
85	0,5	0,5	0,25	0,25	0,125	0,25	0,03125	0,01563
86	0,5	0,5	0,5	0,5	0,5	0,5	0,03125	0,03125
87	0,5	0,5	0,125	0,125	0,125	0,125	0,0078	0,0078
88	0,5	0,5	0,25	0,5	0,25	0,25	0,0078	0,0078
89	1	1	0,25	0,5	0,125	0,25	0,0078	0,0078
90	0,5	0,5	0,5	0,5	0,25	0,25	0,0078	0,0078
91	0,5	0,5	0,25	0,25	0,25	0,25	0,0078	0,0078
92	1	0,5	0,25	0,5	0,25	0,25	0,0078	0,0078
93	0,25	0,25	1	1	0,5	0,5	0,01563	0,01563
94	0,5	0,5	0,5	0,5	0,25	0,25	0,01563	0,01563
95	0,5	0,5	0,5	0,5	0,25	0,25	0,01563	0,01563
96	0,25	0,125	0,5	0,5	0,5	0,5	0,125	0,125
97	0,5	0,5	0,5	0,5	0,25	0,25	0,03125	0,03125
98	0,5	0,5	0,5	0,5	0,125	0,25	0,0078	0,0078
99	0,5	0,5	0,5	0,5	0,25	0,25	0,01563	0,01563
100	0,25	0,25	0,25	0,25	0,125	0,25	0,03125	0,03125
101	0,5	0,5	0,5	0,5	0,5	0,5	0,01563	0,01563

N° de souche en vert: environnement; N° en rouge: invasion; N° en noir: colonisation.

Tableau A. Valeurs des CMI des 106 souches d'*A. fumigatus* vis à vis des 4 antifongiques testés.

N° souche	Amphotéricine B		Itraconazole		Voriconazole		Posaconazole	
	0,03125 à 16 µg/ml		0,03125 à 16 µg/ml		0,03125 à 16 µg/ml		0,004 à 16 µg/ml	
	Lecture visuelle	630 nm	Lecture visuelle	630 nm	Lecture visuelle	630 nm	Lecture visuelle	630 nm
	48 H	48 H	48 H	48 H	48 H	48 H	48 H	48 H
	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)	CMI (µg/ml)
102	0,5	0,5	0,5	0,5	0,25	0,25	0,03125	0,01563
103	0,5	0,5	0,125	0,125	0,125	0,125	0,01563	0,01563
104	0,5	0,5	0,25	0,25	0,25	0,125	0,01563	0,01563
105	0,5	0,5	0,5	0,5	0,5	0,5	0,03125	0,03125
106	0,5	0,5	0,25	0,25	0,25	0,25	0,0078	0,0078
moyenne des 106 souches	0,666693	0,642561	0,374985	0,387448	0,270408	0,283073	0,020028	0,020797

N° de souche en vert: environnement; N° en rouge: invasion; N° en noir: colonisation.

ANNEXE 2 :Lettre d'approbation du COMITE D'ETHIQUE.



**Université de Liège
Centre Animalier Universitaire
Commission d'éthique**

Liège, le 21 juin 2001

**Dr. HAYETTE
Microbiologie Médicale
Tour de Pathologie - B23**

Cher (ère) Collègue,

Votre dossier intitulé " Diagnostic d'aspergillose pulmonaire invasive chez le lapin: antigénémie versus Polymerase chain reaction (PCR)" a été examiné par la Commission d'Ethique en date du 6 avril 2001 et a reçu le numéro 51.

La Commission a émis un avis favorable .

Nous vous prions de croire, Cher (ère) Collègue, en l'expression de nos sentiments les meilleurs.

**Professeur B. NICKS.
Président de la Commission d'Ethique.**

ANNEXE 3 : Article publié dans une revue internationale

(Journal of Clinical Microbiology)

Detection of *Aspergillus* Species DNA by PCR in Bronchoalveolar Lavage Fluid

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GENEVIÈVE CHRISTIAENS,¹ PIERRETTE MELIN,¹ AND PATRICK DE MOL¹

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The usefulness of a nested PCR assay for detection of *Aspergillus* sp. DNA was evaluated in 177 bronchoalveolar lavage (BAL) fluid specimens. This test was accurate both to diagnose culture-negative BAL fluid specimens from patients with invasive pulmonary aspergillosis and to confirm culture-positive samples. However, it did not differentiate between infection and colonization.

Diagnosis in the early stages of invasive pulmonary aspergillosis (IPA) is very difficult, as clinical and radiological signs are nonspecific and the sensitivity of fungal cultures is low (40 to 60%), even when combined with direct microscopic examination (2, 5). Circulating-antigen detection may contribute to the diagnosis. However, up to 8% false positives are reported (11). Other antigens are under investigation as diagnostic tools (6), but attention has now turned to molecular methods. The role of PCR assay of bronchoalveolar lavage (BAL) fluid for diagnosing IPA does not clearly emerge in the literature (1, 3, 7, 9, 10, 14). Therefore, we have evaluated a nested-PCR-based amplification of *Aspergillus fumigatus* DNA that targets the genes encoding alkaline proteases of the fungus to determine the role of PCR in diagnosing IPA from BAL fluid under routine conditions.

Clinical and reference strains were tested to assess the specificity of the method: *A. fumigatus* (three strains), *Aspergillus clavus* (three strains), *Aspergillus niger* (two strains), *Aspergillus nidulans* (two strains), *Aspergillus terreus*, *Aspergillus glaucus*, *Fusarium oxysporum*, *Fusarium solani* (ATCC 10154), *Paecilomyces* spp., *Penicillium* spp., *Pseudallescheria boydii* (two strains), *Trichoderma harzianum* (two strains), *Rhizopus rhizodiformis*, *Mucor* spp., *Candida albicans* (ATCC 10231), and *Candida glabrata* (ATCC 90030). The fungal strains were cultivated on Sabouraud dextrose agar and incubated at 37 or 38°C for up to 5 days, depending on the species. The clinical isolates were identified by macroscopic, microscopic, and culture characteristics (12).

All patients undergoing bronchoscopy at the University Hospital of Liège (Liège, Belgium) during a 12-month period (1997 to 1998) were included in the study. There were 74 immunosuppressed and 103 nonimmunosuppressed patients. Patients were referred to as immunosuppressed if they were under long-term corticotherapy for chronic obstructive pulmonary disease (COPD) ($n = 18$) or other diseases ($n = 13$) or if they had hematological malignancy ($n = 16$), organ transplantation ($n = 5$), AIDS ($n = 3$), or cancer ($n = 19$). Nonimmu-

nosuppressed patients had bronchoscopy for investigation of severe pneumonia. Medical, radiological, histopathological, and microbiological records and autopsy findings were reviewed to assess IPA. Three groups were defined (A, proven or probable aspergillosis [$n = 10$]; B, colonization [$n = 5$]; and C, no evidence of aspergillosis [$n = 162$]) according to the following criteria: proven, histology with hyphal tissue invasion and *Aspergillus*-positive culture of one or more respiratory specimens or of a lung biopsy; probable, *Aspergillus*-positive culture from two or more respiratory samples and positive clinical or radiological findings (pulmonary infiltrates, nodular opacity, cavitation, or persistent fever under broad-spectrum antimicrobial chemotherapy) or positive histology without *Aspergillus*-positive culture; colonization, *Aspergillus*-positive cultures from one or more respiratory samples without clinical or radiological evidence (see above) of respiratory tract infection due to this pathogen; and no evidence of aspergillosis, *Aspergillus*-negative cultures and no clinical or radiological findings (see above) of aspergillosis.

After reception in the laboratory, each BAL fluid specimen was homogenized and separated into two parts: one part (1 to 5 ml) was stored at -20°C until it was analyzed by PCR, and the second part was included in the routine procedure and tested for the presence of bacteria, yeasts, fungi, parasites, and viruses. A PCR assay was performed a minimum of 1 week after sampling, and the clinicians were unaware of the results. The technique used for DNA extraction from fungal culture was adapted from that of Tang et al. (13), except that DNA was resuspended in 30 μl of water containing RNase A at a concentration of 50 $\mu\text{g}/\text{ml}$. The same technique was used for DNA extraction from yeasts except for the culture, which was made on Sabouraud dextrose agar and incubated for 48 h at 37°C. DNA extraction from BAL fluid was adapted from techniques already published by Tang and colleagues (14) except for two details: (i) 200 μg of proteinase K (Sigma, St. Louis, Mo.) was added to the 500 μl of BAL fluid and buffer and (ii) DNA was dried in a dry-heating block and resuspended in 30 μl of distilled H_2O containing RNase A (50 $\mu\text{g}/\text{ml}$). Positive-displacement pipettes were used throughout, and DNA extraction buffer was extracted in parallel in order to preclude contamination.

We developed a nested PCR using as external primers alp 11

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TABLE 1. Clinical data and aspergillosis detection for group A (proven or probable aspergillosis) and group B (colonization) patients

Case no.	Underlying disease ^a	IPA group (disease ^b)	BAL fungal culture	Autopsy performed	BAL PCR results ^c	Outcome ^d
Group A (n = 10)						
1	Hepatic transplantation	Proven (DIS)	<i>A. fumigatus</i>	Yes	+	Died
2	COPD (corticosteroid dependent)	Proven (CNA)	<i>A. fumigatus</i>	Yes	+	Died (AmB)
3	Chronic lymphoid leukemia + COPD	Probable	<i>A. fumigatus</i>	No	+	Died
4	RA (corticosteroid dependent)	Proven	Negative	Yes	+	Died
5	COPD (corticosteroid dependent)	Proven (DIS)	<i>A. fumigatus</i>	Yes	+	Died
6	Gastric carcinoma	Proven	<i>A. fumigatus</i>	Yes	+	Died
7	COPD (corticosteroid dependent)	Proven (CNA)	<i>A. fumigatus</i>	Yes	+	Died
8	Chronic myeloid leukemia blast-crisis	Probable	<i>A. fumigatus</i>	No	+	Alive (Vor)
9	Chronic lymphoid leukemia blast-crisis	Proven	Negative	Yes	+	Died
10	COPD (corticosteroid dependent)	Probable	<i>A. fumigatus</i>	No	+	Died
Group B (n = 5)						
1	Mitral valvulopathy	None	<i>A. fumigatus</i>	No	+	Alive (AmB/It)
2	COPD (corticosteroid dependent)	None	<i>A. fumigatus</i>	No	+	Died
3	COPD (corticosteroid dependent)	None	<i>A. fumigatus</i>	Yes	+	Died
4	COPD (no corticoid)	None	<i>A. fumigatus</i>	No	+	Alive
5	Myelodysplasia blast-crisis	None	<i>A. flavus</i>	Yes	+	Died

^a RA, rheumatoid arthritis.

^b DIS, disseminated aspergillosis; CNA, chronic necrotizing aspergillosis.

^c +, positive.

^d Vor, treatment with voriconazole; AmB/It, treatment with amphotericin B and itraconazole; AmB, treatment with amphotericin B.

5'-AGCACCGACTACATCTAC-3') and alp 12 (5'-GAGATGGTGTGGTGGC-3'). These primers were derived from the sequence of cloned fragments of genes encoding the alkaline proteases (Alp) of *A. fumigatus* and *A. flavus* (14). As internal primers, we chose Alp13 (5'-CTGGCATAACAACGCCGCTG-3') and Alp14 (5'-TTGTTGATCGCAACC-3'), expected to amplify a fragment of 527 bp. The primers were synthesized by Eurogentec, Liège, Belgium. The PCR mixtures were identical for both steps except for MgCl₂. They were carried out in a 50-μl volume containing 10 mM Tris-Cl at pH 8.3, 50 mM KCl, and 1.5 mM MgCl₂ (2.25 mM for the second step), with 20 pmol of both primers, 2.5 mM deoxynucleoside triphosphate (buffer and deoxynucleoside triphosphate were provided by Takara, Otsu, Japan), and 1.25 U of *Taq* polymerase (Takara). A 5-μl volume of DNA was added to the mixture. Positive and negative controls were amplified in parallel to assess the validity of the procedure. Thermal cycling conditions (GeneAmp PCR system 2400; Perkin-Elmer Cetus, Norwalk, Conn.) were identical for both PCRs: 5 min at 94°C; 30 cycles of 30 s at 94°C, 45 s at 63°C, and 2 min at 72°C; and a final extension step of 10 min at 72°C. For the nested PCR, 5 μl of the first amplified product was added to a new reaction mixture and amplified under the same conditions. The final amplified products were analyzed on 1.5% agarose gels stained with ethidium bromide and visualized by UV transillumination. Each sample was investigated for the presence of inhibitors by amplification of the β-globin gene (8).

No band of the expected size (527 bp) was detected with *A. niger*, *A. nidulans*, *A. glaucus*, *A. terreus*, or *A. flavus* or with the other fungal species. However, a band of 690 nm, corresponding to the sequence amplified by the first PCR, was observed for *A. flavus*. The sensitivity of the nested PCR was 25 fg of genomic DNA for *A. fumigatus* by ethidium bromide staining and 10 pg for *A. flavus*. Some BAL fluids (n = 5) were excluded because of lack of amplification of the β-globin gene. The PCR results are reported in Table 1. The sensitivity, specificity, and

positive and negative predictive values of this PCR test for diagnosing IPA were 100, 96, 62, and 100%, respectively. All BAL fluids from patients with IPA were PCR positive. Those from the five cases of *Aspergillus* colonization were PCR and culture positive. In all, six PCR-positive cases were not associated with IPA among the 177 BAL fluid specimens, representing 3.4% false-positive results. Only one false-positive PCR result was induced in group C. Among the three patients with COPD and proven IPA (group A), two patients (cases 2 and 7) presented with chronic necrotizing aspergillosis and one (case 5) presented with disseminated aspergillosis. In two cases (4 and 9), the diagnosis of proven IPA was made at autopsy by histology and *A. fumigatus*-positive culture of a lung biopsy specimen.

In the present study, the PCR always confirmed the culture results, except for two culture-negative BAL fluid specimens, for which diagnosis of aspergillosis was missed and made at autopsy. In both cases, the PCR could have contributed to diagnosing IPA before death. Only one PCR-positive BAL result was observed for a patient with no evidence of aspergillosis: a burn patient who was intubated for 1 month. This false-positive result could be explained by contamination of the BAL fluid during sampling or the PCR process, or it could have been due to the colonization of the respiratory tract by *Aspergillus* spores during intubation. Some authors have reported up to 35% PCR-positive results for patients with no risk or low risk for aspergillosis and who did not develop IPA (1, 3, 9, 10, 14, 15). However, other studies performed in neutropenic patients (4) or in nonimmunosuppressed patients (7) reported no false-positive PCR results for BAL fluids. Among our population, only 3.4% positive PCR results not associated with IPA were observed. Half of the patients (3 out of 6) had COPD. These patients may have had a relatively high level of tracheobronchial colonization, which differs from patients with hematological malignancies, who may have had minimum involvement before developing severe invasive fungal infection.

our study, the major risk factor associated with IPA was corticotherapy associated with COPD. However, the major risk factor for aspergillosis is known to be a prolonged neutropenia, accounting for its high frequency in patients with acute leukemia (2). In fact, in our institution, patients with hematological malignancies benefit from a good follow-up and are often treated early in cases where there is a suspicion of infectious disease.

In conclusion, nested PCR of BAL fluid is an accurate test to diagnose culture-negative patients with IPA, but it does not differentiate between infection and colonization. It is an appropriate method to exclude *Aspergillus* sp. infection in patients at risk of IPA and should be included in routine laboratory practice for this immunocompromised population.

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ANNEXE 4 : Posters présentés dans un congrès international

**ICAAC (Intersciences conference on antimicrobial agents and
chemotherapy)**

ICAAC 1999

ICAAC 2001

Introduction

Aspergillus fumigatus is an opportunistic nosocomial pathogen which causes severe to fatal infection in neutropenic patients. The risk of invasive aspergillosis (IA) increases with the duration of neutropenia and can reach 70% after 5 weeks of neutropenia (1).

The diagnostic of IA is one of the most difficult and the gold standard is still based on culture and/or histology results which sensitivity is very low. Recently, molecular biology methods were developed to diagnose aspergillosis in urine, serum, blood and BAL.

We used a nested PCR-based amplification of *A. fumigatus* DNA in BAL fluids that targets the genes-encoding alkaline proteases of the fungus to evaluate the usefulness of PCR to diagnose IA.

Abstract

The usefulness of a nested PCR for detection of *Aspergillus fumigatus* DNA was evaluated in bronchoalveolar lavage (BAL) fluid during a period of two years (1996-1998). The aim of the study was to assess the role of PCR in diagnosing invasive pulmonary aspergillosis (IPA). **Methods:** a nested PCR-based amplification of fragments of genes-encoding alkaline proteases from *Aspergillus fumigatus* was used to test 167 BAL samples. All samples were checked for the absence of amplification inhibitors. Medical, radiological, microbiological records and autopsy findings were reviewed for assessing invasive aspergillosis. All successive patients investigated by BAL were included in the study. They were distributed in three groups: A: proven or probable aspergillosis (n=11); B: colonization (n=9); C: no evidence of IPA (n=154). PCR results were compared to culture detection as gold standard and to clinical data. **Results:** BAL fluids from 10 patients of group A were PCR positive. One case was falsely negative. Among group B, one case was PCR positive, and the second one PCR negative but had negative BAL cultures (only culture positive sputum). No false positive was detected among group C. Comparing to culture, sensitivity was 81%, specificity, 100%, positive predictive value, 100%, and negative predictive value, 99%. **Conclusions:** *Aspergillus fumigatus* PCR in BAL fluid was an accurate test to diagnose culture negative patients with IPA and to confirm culture positive samples; however it doesn't make difference between infection and colonization. 2. It is an appropriate test to exclude *Aspergillus* infection in patients at risk of invasive illness.

Introduction

Aspergillus fumigatus is an opportunistic nosocomial pathogen which causes severe to fatal infection in neutropenic patients. The risk of invasive aspergillosis (IA) increases with the duration of neutropenia and can reach 70% after 5 weeks of neutropenia (1).

The diagnostic of IA is one of the most difficult and the gold standard is still based on culture and/or histology results which sensitivity is very low. Recently, molecular biology methods were developed to diagnose aspergillosis in urine, serum, blood and BAL.

We used a nested PCR-based amplification of *A. fumigatus* DNA in BAL fluids that targets the genes-encoding alkaline proteases of the fungus to evaluate the usefulness of PCR to diagnose IA.

Abstract

The usefulness of a nested PCR for detection of *Aspergillus fumigatus* DNA was evaluated in bronchoalveolar lavage (BAL) fluid during a period of two years (1996-1998). The aim of the study was to assess the role of PCR in diagnosing invasive pulmonary aspergillosis (IPA). **Methods:** a nested PCR-based amplification of fragments of genes-encoding alkaline proteases from *Aspergillus fumigatus* was used to test 167 BAL samples. All samples were checked for the absence of amplification inhibitors. Medical, radiological, microbiological records and autopsy findings were reviewed for assessing invasive aspergillosis. All successive patients investigated by BAL were included in the study. They were distributed in three groups: A: proven or probable aspergillosis (n=11); B: colonization (n=9); C: no evidence of IPA (n=154). PCR results were compared to culture detection as gold standard and to clinical data. **Results:** BAL fluids from 10 patients of group A were PCR positive. One case was falsely negative. Among group B, one case was PCR positive, and the second one PCR negative but had negative BAL cultures (only culture positive sputum). No false positive was detected among group C. Comparing to culture, sensitivity was 81%, specificity, 100%, positive predictive value, 100%, and negative predictive value, 99%. **Conclusions:** *Aspergillus fumigatus* PCR in BAL fluid was an accurate test to diagnose culture negative patients with IPA and to confirm culture positive samples; however it doesn't make difference between infection and colonization. 2. It is an appropriate test to exclude *Aspergillus* infection in patients at risk of invasive illness.

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Methods

- Clinical data: all patients undergoing bronchoscopy at the university hospital of Liege between 1996 and 1998 were included in the study and distributed in three groups:
 - A: proven or probable aspergillosis (n=11)
 - B: colonization (n=2)
 - C: no evidence of aspergillosis (n=154)
- BAL fluid specimens: all lavages collected by the microbiological laboratory were included in the study and stored at -20°C before processing. Microbiological examination was directly performed on every sample and results recorded.
- DNA extraction: adapted from Tang et coll. (2).
 - BAL fluid (250µl) + extraction buffer (250µl) containing proteinase K (100 µg); 65°C for 60 min
 - extraction once with phenol:chloroform:isomylalcol (25:24:1) and then with chloroform:isomylalcol (24:1).
 - Precipitation of DNA by ethanol
- Nested PCR: The target DNA corresponds to genes-encoding alkaline proteases from *A. fumigatus* (2).
 - External primers: alp11 and alp12 (2).
 - Internal primers: alp13 (5'-CTGGATACAGCCGGCT-3') alp14 (5'-TTGTTGATCGCAACC-3')
- Product length after amplification: 527 base pairs.
- Mix, (50µl volume): 10mM TRIS-Cl at pH 8.3, 50 mM KCl and 1.5 mM MgCl2 with 100 pmol of both primers and 1.25 U Taq polymerase (Takara Taq, Japan).
- Thermal cycling conditions: 30 cycles at 94°C for 30 sec., 63°C for 45sec., and 72°C for 2 min.
- Products of PCR : analyzed on 1.5% agarose gels, stained with ethidium bromide and visualized by UV transillumination.

Results

case	UNDERLYING DISEASE	IPA	CULTURE	HISTOLOGY	PCR	SURVIVAL
GROUP A: IPI or probable IPI (n=11)						
1	Hepatic transplantation	yes	yes	yes	yes	no
2	COPD corticoids dependent	yes	yes	yes	yes	no
3	Hematologic malignancy	yes	yes	no histology	yes	no
4	Cardiac transplantation	yes	yes	yes	yes	no
5	Atrialitis	yes	yes	yes	yes	no
6	COPD corticoids dependent	yes	yes	yes	yes	no
7	Gastric carcinoma	yes	yes	yes	yes	no
8	COPD corticoids dependent	Probable	yes	yes	yes	no
9	Bone marrow transplantation	yes	yes	no histology	yes	no
10	Hematologic malignancy	yes	yes	yes	yes	no
11	Hematologic malignancy	yes	yes	no histology	yes	no
GROUP B: colonization (n=2)						
case	UNDERLYING DISEASE	IPA	CULTURE	HISTOLOGY	PCR	SURVIVAL
1	Mitral valveulopathy	no	yes	no histology	yes	yes
2	COPD (no corticoids)	no	yes	no histology	no	yes
GROUP C: no evidence of aspergillosis (n=154)						
UNDERLYING DISEASE						
164	Respiratory disease	0	0	0	0	0

Sensitivity and specificity of PCR

IPA	PCR-	PCR+	Sensitivity:
10	1	10	91%
NO IPA			
154	155	1	100%
166	156	1	Predictive values
187	187	0	negative
			positive

Conclusions

- 1. *A. fumigatus* PCR in BAL fluid was an accurate test to diagnose culture negative patients with IPA and to confirm culture positive samples. However it doesn't make difference between infection and colonization.
- 2. This PCR is an appropriate method to exclude *Aspergillus* infection in patients at risk of invasive illness.

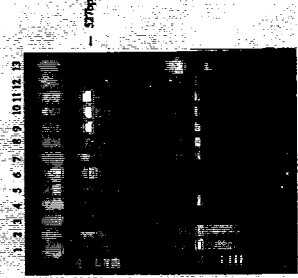


Figure 1. Amplification products of ethidium bromide stained agarose gel. Above: lanes 1-3, 7, 8: negative BAL; Lanes 4, 5: patient 7 from group A; Lanes 6, 9: patient 8 from group A; Lanes 12, 13: negative sample (water); Below: beta-globin (reference sample). Lanes 14-16: patient 9 from group A.



Comparative *in vitro* activity of amphotericin B, itraconazole, voriconazole and posaconazole against *Aspergillus fumigatus*.

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Abstract

Background. New azoles have been successfully used as treatment of invasive aspergillosis. The purpose of this study was to compare the *in vitro* activity of posaconazole (Posa) with that of amphotericin B (AmB), itraconazole (itra) and voriconazole (Vor) against *A. fumigatus* isolates according to NCCLS method (M38-P), and to compare visual and spectrophotometric readings for MIC determination. **Methods.** A total of 108 *A. fumigatus* isolates were selected as follows: 81 clinical isolates from colonized patients, 18 from patients with invasive aspergillosis and 9 environmental isolates. Their *in vitro* susceptibility was evaluated by the NCCLS microdilution method (M38-P) in RPMI 1640 medium. Determination of results was made by visual and spectrophotometric readings (630 nm) after 48 hours incubation at 35°C. Three *A. fumigatus* reference strains (HEM 5734, 6149 and 13935) were included as control.

Results. 1. Geometric mean MICs/MIC₅₀ (µg/ml) obtained by visual reading were respectively 0.691 (AmB), 0.370.5 (itra), 0.270.5 (Vor) and 0.020.03 (Posa). 2. MIC values were comparable by spectrophotometric and by visual readings for all antifungal agents tested (p > 0.05) and did not depend on the isolate's origin (p > 0.05). 3. Posaconazole had the lowest MICs (p < 0.001). 4. The itraconazole-resistant reference strain did not give cross resistance with voriconazole and posaconazole. **Conclusion:** Among azoles, posaconazole had a better *in vitro* activity against *A. fumigatus* than did voriconazole or itraconazole. Spectrophotometric reading could replace the less standardized visual reading for NCCLS microdilution method and MIC values obtained were comparable among all *A. fumigatus* isolates.

Introduction

New azoles have been recently introduced in the treatment of invasive aspergillosis. Clinical studies are in progress to compare the efficiency and toxicity of these molecules to the gold standard, amphotericin B.

The purpose of our study was first to determine the MICs of two new azoles, voriconazole (Vor) and posaconazole (posa), against *Aspergillus fumigatus* isolates from diverse origins and to compare the results to amphotericin B (AmB) and itraconazole (itra) MIC values.

Secondly, the aim of the study was to compare the visual reading with a spectrophotometric one in order to enhance the standardisation of the method. E-test was also performed for posaconazole and compared with NCCLS method in a parallel direction.

Materials and methods

Strains: 106 isolates of *A. fumigatus* have been selected as follows: 81 clinical isolates from colonized patients, 18 from patients with invasive aspergillosis and 7 environmental isolates. Three *A. fumigatus* (HEM 5734, 6149 and 13935) reference strains have been included as controls.

Culture: the strains have been cultivated on Sabouraud dextrose agar following the recommendations of the NCCLS M38-P microdilution methodology (1). **NCCLS method:** RPMI 1640 medium (Gibco BRL, Life Technologies, USA) buffered with MOPS (Sigma, St Louis, USA) has been used. All antifungals were dissolved in DMSO before diluting them in the RPMI medium. The concentrations tested were as follows: 0.031 to 16 µg/ml for AmB, itra and Vor; 0.004 to 16 µg/ml for Posa. The drugs were kindly provided as pure powder by their manufacturers: AmB (Bristol-Myers Squibb, USA), itra (Janssen Pharmaceutica, Beerse, Belgium), Vor (Pfizer, New-York, USA), and Posa (Schering Plough, Kenilworth, USA). Determining MICs: reading has been performed according to the NCCLS recommendations (visual reading) with a score 0 for AmB and score 2 for the azoles. A spectrophotometric reading at 630 nm was also performed and compared to the visual reading with the same scoring system.

E-test posaconazole: E-test (AB-biodisk, Solna, Sweden) for posaconazole was performed on RPMI 1640 buffered agar (1.5%). Reading was performed after 24h incubation at 35°C.

Results

MICs geometric means (µg/ml) of *A. fumigatus* isolates (n=106).

Amphotericin B		Itraconazole		Voriconazole		Posaconazole	
Visual	630 nm	Visual	630 nm	Visual	630 nm	Vfs.	630h
0.67	0.64	0.37	0.39	0.27	0.28	0.02	0.02
						0.02	0.01

1. There is no significative difference between visual and spectrophotometric readings (p > 0.05).
2. Among azoles, posaconazole has lower MICs than voriconazole and itraconazole (p < 0.001).
3. E-test and NCCLS results for posaconazole are in agreement (p > 0.05).

MICs geometric means (µg/ml) by visual reading

Origin of isolates	Amphotericin B	Itraconazole	Voriconazole	Posaconazole
Environmental (n=7)	0.74	0.37	0.25	0.02
Colonising (n=81)	0.61	0.43	0.43	0.02
Invasive (n=18)	0.67	0.36	0.27	0.02

3. There is no significative difference according to origin of isolates (p > 0.05).

Discussion and conclusions

The standardisation of the NCCLS M38-P method can be improved by use of spectrophotometric reading. This method avoids the variability due to the reader giving a better reproducibility. However, automatic MICs calculation is needed to speed up the procedure.

Antifungal susceptibility is not a marqueur of pathogenicity for *A. fumigatus* since there is no significative difference between isolates MICs from different origins. Since authors have reported *A. fumigatus* itraconazole-resistant isolates (2), it seems useful to test the susceptibility before treatment particularly for azoles.

Among azoles posaconazole has the lowest MICs values. It is a very promising drug and the ongoing clinical trials will determine if the drug can successfully cure invasive aspergillosis without serious side effects.

Nevertheless the NCCLS method could better be replaced by an easier method for routine laboratory. According to our study and to others authors (3), E-test method could be used for *A. fumigatus* as this methodology gives good correlations with NCCLS method.

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ANNEXE 5 : Formules de tampons

Bleu de bromothymol

Bleu de bromothymol 250mg
Ficoll..... 15ml
H2O QSP100ml

Dissoudre le bleu dans l'eau puis ajouter la glycérol. Mélanger une nuit par agitateur magnétique.
Conserver à 4°C. A utiliser tel quel en ajoutant 2µl de bleu à 10 µl de produit amplifié.

Bromure d'éthidium (solution stock).

Bromure d'éthidium 1 g
H2O qsp..... 100ml.

Mélanger jusqu'à dissolution complète. Conserver à température ambiante et à l'abri de la lumière.

EDTA 0.5mM

Disodium éthylène diaminetetracétate-2H2O186.12 g
H2Oppiqsp 1l

Ajuster à pH 8 avec NaOH (soit environ 20g) et ajuster le volume à 1 litre.

N.B. Utiliser un agitateur magnétique pour dissoudre la poudre. Le sel d'EDTA ne devient soluble que lorsque le pH est ajusté à 8.

Marqueurs de poids moléculaire (Eurogentec, Belgium).

Marqueur IV (ΦX174 /Hae III) 75µg

Marqueur IV 150µl
Bleu de Bromothymol 120µl Concentration finale du marqueur 0.1µg/µl
H2O ppi 480ml

Marqueur V (Φ X174 /Hinc II digest) 75 μ g

Marqueur V	150 μ l	
Bleu de Bromothymol	120 μ l	<u>Concentration finale du marqueur 0.1μg/μl</u>
H2O ppi	480 μ l	

Tampon d'extraction selon Tang et coll. pour extraction d'*Aspergillus sp.* dans les LBA.

200 mM Tris-HCl pH8	soit 2.42g
0.5M NaCl	soit 2.92g
0.01M EDTA	soit 0.37g
1% SDS	soit 1 ml de solution à 10%
H2O ppi	QSP100ml

Préparer en tube Falcon de 50ml. Stériliser sur filtre 0.2 μ .

Tampon d'extraction selon Yamakami et coll. pour extraction d'*Aspergillus sp.* dans le sang.

100mM KCl.....	soit 0,745 g
20mM Tris-HCl (pH 8.3)	soit 24.22 mg
5mM MgCl2	soit 0.5 ml d'une solution à 1 M (Sigma Aldrich).
0.9% Tween 20	soit 900 μ l
0.2 mg/ml de gélatine	soit 20mg
H2O ppi	qsp100ml

Filtrer sur filtre 0.2 μ .

Tampon de lyticase

50mM Tris pH 7.5	soit 0.302 mg
1 mM EDTA.....	soit 0.015 mg
20% β -mercaptoéthanol	97 μ l
H2O ppi	qsp 50 ml

Filtrer sur filter 0.2 μ .

TBE 5x concentré (Tampon tris-Acide borique-EDTA)

Tris base 54 g
Acide borique 27.4 g
EDTA à 0.5mM pH 8 20ml
H2O ppi qsp 1l
Conserver à température ambiante.

ANNEXE 6 : Protocoles d'extraction de l'ADN

1. Technique modifiée de Yamakami et coll. ()

La composition du tampon de lyse est donnée dans les annexes « tampons ».

Cette technique consiste à mélanger 100 µl d'échantillon avec 100 µl de tampon. La protéinase K est ajoutée extemporanément à la concentration de 200 µg (à la place de 60µg/ml recommandée par les auteurs) soit 20 µl d'une solution à 10 µg/µl. Le mélange est incubé 60 minutes à 65°C puis 30 minutes à 95°C et centrifugé à 12.000 g pendant 10 minutes à +4°C.

Le surnageant est utilisé pour réalisation de la PCR.

2. Technique d'extraction sur colonne QIAamp DNA mini kit, tissue protocole (QIAamp DNA Mini Kit, QIAgen, Allemagne).

1. Ajouter 180 µl de solution ATL au culot de leucocytes et 20 µl de protéinase K (Qiagen). Mélanger par vortex et incuber 3 heures à 55°C.

2. Ajouter 200µl de solution AL, mélanger par vortex et incuber 10 min. à 70°C. Centrifuger quelques secondes pour concentrer tout le liquide dans le fond du tube.

3. Ajouter 200 µm d'éthanol, bien mélanger par vortex. Centrifuger quelques secondes pour concentrer tout le liquide dans le fond du tube.

4. Placer une colonne dans un tube de 2 ml (Qiagen), puis déposer tout le liquide sur la colonne. Fermer le bouchon. Centrifuger 1 min à 8000 rpm.

5. Sortir la colonne du tube et la placer dans un nouveau tube de 2 ml. Déposer 500 µl de solution AW1 (Qiagen). Centrifuger 1 min à 8000 rpm.

6. Sortir la colonne du tube et la placer dans un nouveau tube de 2 ml. Déposer 500 µl de solution AW2 (Qiagen). Centrifuger 1 min à 14.000 rpm.

7. Sortir la colonne du tube recentrifuger 1 min à 14.000 pour éliminer toute trace de AW2.

8. Sortir la colonne du tube. Ajouter 50 µl de solution AE et laisser 5min à température ambiante. Centrifuger 1 min à 14.000 rpm et garder l'éluat. Celui-ci peut être congelé à -20°C ou traité directement en PCR.

3. Technique Amplicor® Whole Blood (Roche).

Le kit Amplicor® Whole Blood est composé de deux réactifs :

BLD WS permet de lyser les globules rouges et d'isoler les leucocytes à partir du sang. Il s'agit d'une solution de phosphate de sodium contenant moins de 0.4% de détergent et de 0.05 % d'azide de sodium.

BLD EXT permet la lyse des leucocytes. Il s'agit d'un tampon Tris-HCl contenant 0.01% de protéinase K, 7.5 mg de MgCl₂, 1% de détergent et 0.05% d'azide sodium.

Technique de lyse des globules rouges :

Le sang total prélevé sur EDTA a été soumis à une lyse des globules rouges par le kit de préparation des échantillons Amplicor® Sang Total (Roche diagnostics, Brandburg, USA), permettant de lyser les globules rouges et blancs en deux étapes séparées et grâce à deux réactifs différents.

1. On mélange 1 volume (500 µl) de sang total à 2 volumes (1 ml) de solution de lyse BLD WS. On mélange 10 à 15 fois par retournement, puis on laisse à température ambiante pendant 5 minutes pour faciliter la lyse. Ensuite, les tubes sont centrifugés pendant 3 minutes à vitesse maximale (15.000 trs/min). Le culot de globules blancs se distingue par sa couleur plus claire. Le surnageant est prélevé et éliminé.

2. Deux volumes de solution de lavage sont ajoutés au culot. On mélange sur Vortex pour remettre en suspension et on mélange de nouveau par retournement 10 à 15 fois. Puis on centrifuge comme précédemment 3 minutes à vitesse maximale (15.000 tr/min).

3. L'étape 2 est répétée encore une fois de façon à éliminer au maximum l'hémoglobine présente dans le prélèvement.

Le culot peut être directement traité pour extraction de l'ADN ou être congelé à -80°C .

Technique de lyse du culot leucocytaire :

Le culot cellulaire est décongelé à température ambiante quelques minutes s'il n'a pu être traité directement.

La méthodologie consiste à ajouter 200 μl de réactif BLD EXT au culot décongelé et de placer le tube fermé au bain marie à 60°C pendant 45 minutes. Le tube est ensuite maintenu à 100°C pendant 30 minutes pour éliminer complètement la protéinase K.

L'échantillon est ensuite centrifugé quelques secondes, pour concentrer l'échantillon au fond du tube. On peut le garder à $+4^{\circ}\text{C}$ jusqu'à la réalisation de la PCR.

4. Technique d'extraction par TriPure.

(TriPure isolation reagent[®], Boehringer Mannheim, Allemagne).

1. Ajouter 1 ml de TriPure à 100 μl de solution à extraire. Incuber 5 min à température ambiante.

2. Ajouter 0.2 ml de chloroforme (0.2 ml par ml de TriPure). Mélanger vigoureusement par vortex. Incuber 15 min à température ambiante.

Centrifuger 15 min à 12.000 g à 4°C .

On obtient trois phases :

- phase supérieure aqueuse (incolore) contenant le RNA.
- interface contenant le DNA de couleur blanche.
- phase inférieure organique (rouge) contenant les protéines.

3. Eliminer la phase supérieure et ajouter 0.3 ml d'éthanol pur pour séparer DNA et protéines (soit 0.3 ml d'éthanol pur par ml de TriPure). Mélanger par inversion et incuber 3 minutes à T°C ambiante.
4. Centrifuger 5 min à 2000g à 4°C. Eliminer le surnageant (protéines).
5. Laver le culot avec une solution de citrate de sodium 0.1M contenant 10 % d'éthanol (1ml de solution par ml de TriPure). Incuber 30 min à température ambiante en mélangeant de temps en temps.
6. Centrifuger 5 min à 2000g à 4°C. Eliminer le surnageant.
7. Répéter l'étape de lavage deux fois.
8. Laver le culot dans une solution d'éthanol à 75% (1.5 à 2 ml par ml de TriPure). Incuber 20 min à T°C ambiante en mélangeant de temps en temps.
Centrifuger 5 min à 2000g à 4°C et éliminer le surnageant.
9. Sécher le culot (dans un bloc chauffant). Dissoudre le culot dans de l'eau. Conserver à 4°C ou congeler à -20°C jusqu'à réalisation de la PCR.

ANNEXE 7 : Milieux de culture

RPMI 1640 (NCCLS M38-P)

Dissoudre 10.4 g de RPMI 1640 (avec glutamine, et sans bicarbonate)(Difco) dans 900 ml d'eau distillée.

Ajouter 2 g de NaHCO₃ (Sigma) et 34.53g de tampon MOPS (Sigma). Ajuster le pH à 7 à l'aide de NaOH 1N. Compléter avec de l'eau pour atteindre un volume final d'un litre.

Filtrer la solution (filtres Nalgène, Nalge Nunc corporation, New York, USA).

Conserver à +4 °C au maximum un mois.

SABOURAUD DEXTROSE AGAR (Oxoid)

Sabouraud dextrose agar 65g

Eau distillée QSP..... 1litre

Peser la poudre et compléter à 1 litre avec de l'eau distillée. Placer sur un barreau magnétique dans le récipient et placer sur un agitateur magnétique. Agiter jusqu'à formation d'une suspension homogène. Chauffer sur un bec bunsen en agitant fréquemment puis distribuer dans les tubes selon les quantités désirées. Autoclaver 30 minutes à 120°C, puis sortir les tubes encore chauds les incliner et les laisser refroidir. Boucher complètement une fois refroidis et conserver à température ambiante.

ANNEXE 8 : Répertoire des tableaux

N°	TITRE	Page ou verso (V)
Tableaux CHAPITRE I		
1	Position systématique des champignons d'intérêt médical	V8
2	Incidence de l'aspergillose invasive d'après Denning (42)	V20
3	Incidence des infections chez les transplantés d'organe solide d'après Patel et Paya (121).	V23
4	Molécules d' <i>Aspergillus fumigatus</i> ayant un rôle potentiel dans la virulence .	V27
5	Profils caractéristiques de l'aspergillose invasive selon les groupes de patients d'après Denning (42)	V35
6	Efficacité des différents types de prélèvements pour le diagnostic d'une aspergillose pulmonaire invasive d'après Kappe et Rimek (74)	V47
7	Efficacité des techniques de détection dans le diagnostic d'une aspergillose pulmonaire invasive (d'après Kappe et Rimek (74)).	V47
8	Analyse réalisées en biologie moléculaire dans le domaine de la microbiologie par les 18 centres de diagnostic moléculaire (CDM)	V58
9	Résultats comparatifs des techniques PCR appliquées au diagnostic des aspergilloses pulmonaires invasives dans les lavages broncho-alvéolaires	V59
10	Résultats comparatifs des techniques PCR appliquées au diagnostic des aspergilloses pulmonaires invasives sur sang périphérique	V60
11	Traitement de 1 ^{ère} et de 2 ^e ligne de l'aspergillose invasive d'après Denning (42).	67
Tableaux CHAPITRE II		
1	Amorces Alp 11, 12, 13, 14	V72
2	Souches fongiques	V72
3	Préparation des mix pour la PCR nichée	79
4	Paramètres de la PCR nichée	80
5	Amorces β -globine	V79
6	Préparation du mix pour la PCR β -globine	81
7	Paramètres de la PCR β -globine	81
8	Résultats concernant les lavages broncho-alvéolaires des groupes A	V84
9	Résultats concernant les lavages broncho-alvéolaires des groupes B	V84

10	Résultats concernant les lavages broncho-alvéolaires des groupes C	V85
11	Comparaison des résultats de la culture pour recherche d' <i>Aspergillus</i> sp. versus PCR dans les groupes A, B et C.	V 85
12	Résultats des cultures des lavages bronchoalvéolaires du groupe C	V85
13	Sensibilité, spécificité, valeurs prédictives positive et négative.	86

Tableaux CHAPITRE III

1	Caractéristiques des amorces Asp5, Asp8, Asp 1 et Asp7.	V91
2	Protocole d'immunodépression des lapins	93
3	Préparation des mix pour la PCR nichée	102
4	Paramètres de la PCR nichée	102
5	Comparaison des différents protocoles d'extraction testés.	105
6	Apparition et degré de la neutropénie chez les lapins	107
7	Moyenne de la concentration en hémoglobine et du nombre de plaquettes des lapins d'expérience et des lapins témoins.	108
8	Résultats histologiques des 5 organes prélevés exprimés en %.	109
9	Résultats de l'analyse histologique et de la culture fongique des organes des 14 lapins	V109
10	Résultats obtenus en PCR versus antigénémie	111
11	Résultats PCR et antigène galactomannane exprimés en %	112
12	Précocité de la positivité de la PCR et du galactomannane	115
13	Listes des publications comparant galactomannane et PCR <i>Aspergillus</i> .	V118

V : verso de la page

N°	Tableaux CHAPITRE IV	Page ou verso (V)
1	Moyenne géométrique des CMI des isolats d' <i>A. fumigatus</i> : comparaison entre les lectures visuelle et spectrophotométrique	126
2	CMI ₉₀ des isolats d' <i>A. fumigatus</i> comparaison entre les lectures visuelle et spectrophotométrique	127
3	Moyenne géométrique des CMI des souches d' <i>A. fumigatus</i> par lecture visuelle	127
4	Moyenne des CMI obtenues par lecture visuelle pour les souches contrôles tout au long de l'étude	128
5	Résistance de 3 souches d' <i>A. fumigatus</i> vis à vis de l'amphotéricine B	128

ANNEXE 9 : Répertoire des Figures

N°	TITRE DES FIGURES	Page ou verso (V)
Figures CHAPITRE I		
1	Représentation schématique du conidiophore et des têtes aspergillaires	V12
2	Représentation schématique de la formation de phialospores.	V12
3	Représentation schématique de la reproduction des ascomycètes	V13
4	Ultrastructure d'une cellule fongique	V14
5	Structure moléculaire de la paroi fongique d' <i>A. fumigatus</i>	V14
6	Représentation temporelle de la survenue des infections chez les greffés de moelle allogénique	V22
7	Rôle de l'immunité naturelle et sa modulation par les agents immunosuppresseurs	V30
8	Etapes nécessaires au développement d'un test diagnostique (galactomannane)	V51
9	Différentes étapes de la PCR	V57
10	Algorithme décisionnel	V61
11	Structure moléculaire de l'amphotéricine B	V63
12	Formulations lipidiques d'amphotéricine B	V63
13	Représentation schématique de la structure moléculaire de l'Ambisome	V63
14	Représentation schématique de la structure moléculaire de l'Abelcet	V63
15	Représentation schématique de la structure moléculaire de l'Amphocil	V63
16	Structure moléculaire de l'itraconazole	V65
17	Structure moléculaire du voriconazole	V65
18	Structure moléculaire du posaconazole	V65
19	Structure moléculaire des échinocandines naturelles	V65
Figures CHAPITRE II		
1	Séquence de Reichard et coll. (130)	V70'

Figures CHAPITRE III

1	Séquence de Melchers et coll. (100)	V90'
2	Représentation graphique de l'évolution de l'index galactomannane	110
3	Représentation graphique des résultats PCR versus galactomannane	113
4	Représentation graphique des résultats cumulés PCR versus galactomannane	114

ANNEXE 10 : Répertoire des Photos

N°	TITRE DES PHOTOS	Page ou verso (V)
Photos CHAPITRE I		
1	<i>Aspergillus fumigatus</i> sure gélose au malt	V10
2	<i>Aspergillus fumigatus</i> sur milieux de Sabouraud et Czapeck	V10
3	A. flavus sur milieux de Sabouraud et Czapeck	V10
4	A. nidulans sur milieux de Sabouraud et Czapeck	V10
5	A. niger sur milieux de Sabouraud et gélose au malt	V10'
6	A. terreus sur milieux de Sabouraud et Czapeck	V10'
7	A. versicolor sur milieu de Sabouraud et gélose au malt	V10'
8	Coupe histologique de filaments mycéliens d' <i>Aspergillus</i> sp. colorés à l'HES	V50
9	Coupe histologique de parenchyme pulmonaire d'aspergillose invasive colorée au Grocott	V50
Photos CHAPITRE II		
1	Electrophorèse des produits d'amplification en fonction de la concentration en MgCl ₂ (1 ^{ème} étape de la PCR nichée)	V82
2	Electrophorèse des produits d'amplification en fonction de la concentration en MgCl ₂ (2 ^{ème} étape de la PCR nichée)	V82
3	Spécificité de la PCR	V83

Photos CHAPITRE III		
1	Cubicule de lapin	V92
2	Médicaments injectés	V92
3	Partie du cou rasée pour injection intra-trachéale	V92
4	Injection intra-trachéale	V92
5	Distribution de gaz isoflurane	V94
6	Cage d'induction	V94
7	Table de travail avec maintien du lapin sous anesthésie gazeuse	V94
8	Prélèvement de sang au niveau de l'oreille du lapin.	V94
9	Sensibilité de la PCR sur sang total	V103
10	Prélèvement de poumon : aspect macroscopique pathologique	V108
11	Prélèvement de foie: aspect macroscopique pathologique	V108

V= verso de la page