

Rqn{ oqtr jku ou"kp" FPC" Tgrckt" I gpg"XRCCI"*Cti3; 6Vtr+"cpf"*Cti5; ; Inp+"cpf
vjgkt" Tqng"kp"vjg"uwuegrvkdknkv {"qh"Dcevgtkcn" Ogpkipkvku

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~}!^•c!i&c^á ~•^É ái•ciãã ~cã [}É æ)á !^! [á ~cã [} æ) ^ { ^ái ~ { É } : [çã^á c@^ [i* i}æ|æ ~c@ [i æ)á • [~i&^ æ!^ &!^áã^áÉ

and 30 s for codon 194 and of 2 min for codon 399 (temperatures given in Table 2), elongation for 1 min at 72°C and a final extension step of 10 min at 72°C. PCR products were run on a 1% agarose gels and visualized with SYBR green. The amplified fragments were digested with appropriate restriction endonucleases *PvuII* and *NciI* for

SNPs *XRCC1* Arg194Trp and Arg399Gln, respectively (Table 2) and the resulting cleavage products were separated by electrophoresis on 8% polyacrylamide gel for 1 h at 100 V and detected with silver staining according to the protocol described by Sanguinetti et al. [22].

Genetic Polymorphism	Exon	Primer sequence	Annealing Temp. (°C)	Restriction enzyme	Genotype	Fragments (bp)
YÜÖÖF CE!*FJIV:} ÇI•FÍJJİİGD	İ	Ø İ'ÖVVÖÖÖVÖVÖCEÖÖÖÖÖÖÖÖÖÖEH' Ü İ'ÖÖÖÖÖVÖVÖÖÖVÖVÖCEÖÖÖÖÖVÖVÖVÖVÖH' ÇFHİ Jàð	İİİİ°Ö	Úç [™]	CE!*BCE!* CE!*BV:} V:}BV:}	FHİ FHİÉİİÉİH İİÉİH
YÜÖÖF CE!*HJJÖ:} ÇI•GÍİİİD	F€	Ø İ'ÖÖÖÖVÖÖVÖVÖVÖVÖVÖVÖVÖVÖVÖVÖH' Ü İ'ÖÖÖÖVÖÖVÖVÖVÖVÖVÖVÖVÖVÖH' ÇİİF Jàð	İİ°Ö	p&ij	CE!*BCE!* CE!*BÖ:} Ö:}BÖ:}	I İFÉGİİÉFHG ÍJHÉİİFÉGİİÉFHG ÍJHÉGİİ

Table 2 SNPs in DNA repair genes, primers and conditions used for genotyping

Detection of FPG sensitive sites

The genomic DNA from patients was submitted to treatment with formamidopyrimidine DNA glycosylase (FPG) (New England Biolabs) as described by da Silva et al. [6]. This

Measurement of cytokines and chemokines in patients with BM

The main cytokines and chemokines involved in the BM inflammatory response were examined in the CSF samples from patients. BM patients carrying *XRCC1* Gln allele did not show statistical difference in the cytokines and chemokines levels (Figure 2). For SNP *XRCC1* Arg194Trp, no statistical difference was also observed in the concentration of cytokines and chemokines (data not shown).

Discussion

Bacterial meningitis has been known to be an important cause of mortality and morbidity. The disease evolution is mainly influenced by the host immune response determined by several genes that regulate the intensity of the inflammatory response to infection [8-11].



Figure 1A: FPG sensitive sites in genomic DNA in relation to *XRCC1*

difference

women with a family history of breast cancer from Rio de Janeiro. *Genet Mol Biol* 32:255-259

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