



Carriage of HLA-B*5701 and a Haplotypic Hsp70-Hom Variant is Associated with a Class I MHC-Restricted Hypersensitivity Response to Abacavir

Western Australian HIV Cohort Study

¹Annalisse Martin, ¹Coral Ann Almeida, ¹David Nolan, ¹Ian James, ^{2,3}Frank Christiansen, and ^{1,3}Simon Mallat

¹Centre for Clinical Immunology and Biomedical Statistics (CCIBS), Royal Perth Hospital and Murdoch University, Perth, Western Australia, ²Department of Pathology, University of Western Australia, Perth, Western Australia, ³Department of Clinical Immunology and Biochemical Genetics, Royal Perth Hospital, Western Australia.

Background - Susceptibility to a clinically significant drug hypersensitivity syndrome associated with abacavir use has a significant genetic component. We have shown that the presence of *HLA-B*5701* strongly predicts abacavir hypersensitivity (ABC HSR), particularly in combination with other allelic markers specific to the 57.1 ancestral haplotype (AH), and identified a potential susceptibility locus within a 300kb region between *MEGTL1* and *CA46* loci in the central MHC. Here we use fine recombination mapping to identify the susceptibility loci. **Methods** - 248 consecutive abacavir-exposed individuals were studied, representing full ascertainment of abacavir use in the Western Australian HIV Cohort study. 18 cases of definite ABC HSR (7.3%) and 230 tolerant controls were identified, utilising an updated clinical classification that included corroborative epicutaneous skin patch test. Patients were typed for genetic markers using standard molecular techniques. Intracellular measurement of TNF (three colour flow cytometry) and intracellular localisation of Hsp70 and HLA-B57 (confocal microscopy) were undertaken on abacavir exposed *ex vivo* polymorphic blood mononuclear cell (PBMC) cultures. **Results** - Recombinant mapping in patients with allelic markers of the 57.1 AH suggest a susceptibility locus within the Hsp70 gene cluster. *HLA-B*5701* was present in 94.4% of hypersensitive cases and 1.7% of controls (OR 960, *p*<0.00001). A haplotypic non-synonymous polymorphism of Hsp70-Hom (HspAIL, M493T) was found in combination with *HLA-B*5701* in 94.4% of hypersensitive cases and 0.4% of controls (OR 3893, *p*<0.00001). The Hsp70-Hom M493T allele was present in 22% of controls (OR 60, *p*<0.00001), suggesting that the combination of *HLA-B*5701* and Hsp70-Hom M493T conferred susceptibility. Individuals with ABC HSR exhibited a significantly higher proportion of monocytes expressing TNF in response to *ex vivo* abacavir stimulation, which was abrogated, on depletion of CD8⁺ T cells from whole blood. Increased intracellular expression of Hsp70 and HLA-B57 molecules in abacavir exposed *ex vivo* cultured PBMCs was observed in hypersensitive patients compared with controls. Hsp70 and HLA-B57 molecules co-localised within discrete vesicles.

Conclusions - These data indicate that the presence of *HLA-B*5701* and Hsp70-Hom M493T are predisposing factors in the development of ABC HSR, and implicates them in the generation of a Class I-restricted pathogenic immune response.

INTRODUCTION

Exposure to abacavir is accompanied by a rare and sometimes lethal hypersensitivity reaction that typically involve multiple organs and rapid and more severe recurrence on rechallenging^{1,2,3}. The presence of alleles carried on the 57.1 ancestral haplotype (AH), the *HLA-B*5701* allele in combination with *HLA-DR7*-*DO3*⁴ or independently⁵ and in association with TNF-238A^{6,7} has been shown to predispose patients to this reaction. The putative susceptibility region was mapped to a 300kb region between *CA46* and *MEGTL1* within the central MHC⁸. Further fine mapping using microsatellite, SNP and recombinant haplotype mapping techniques will help identify the most parsimonious region carrying the gene(s) contributing to abacavir hypersensitivity. Previous studies have indicated that the positive predictive value of *HLA-B*5701* testing (>70%) may be sufficient for use in clinical practice particularly in Caucasian populations⁹. Development of our idiosyncratic drug reactions involves MHC-restricted presentation of a drug or its reactive metabolite to the immune system^{10,9}. We therefore examined the association of MHC alleles including HLA-B57 and Hsp70 in *ex vivo* abacavir stimulated cultures of PBMCs.

INTRODUCTION

Exposure to abacavir is accompanied by a rare and sometimes lethal hypersensitivity reaction that typically involve multiple organs and rapid and more severe recurrence on rechallenging^{1,2,3}. The presence of alleles carried on the 57.1 ancestral haplotype (AH), the *HLA-B*5701* allele in combination with *HLA-DR7*-*DO3*⁴ or independently⁵ and in association with TNF-238A^{6,7} has been shown to predispose patients to this reaction. The putative susceptibility region was mapped to a 300kb region between *CA46* and *MEGTL1* within the central MHC⁸. Further fine mapping using microsatellite, SNP and recombinant haplotype mapping techniques will help identify the most parsimonious region carrying the gene(s) contributing to abacavir hypersensitivity. Previous studies have indicated that the positive predictive value of *HLA-B*5701* testing (>70%) may be sufficient for use in clinical practice particularly in Caucasian populations⁹. Development of our idiosyncratic drug reactions involves MHC-restricted presentation of a drug or its reactive metabolite to the immune system^{10,9}. We therefore examined the association of MHC alleles including HLA-B57 and Hsp70 in *ex vivo* abacavir stimulated cultures of PBMCs.

METHODS

Study Population - The first 270 participants of the Western Australian HIV Cohort Study prospectively observed 1/11 December 2001, who were included in our previous study¹¹ were included based on clinical criteria including the occurrence of clinical hypersensitivity response of multi-organ involvement that was present within 2 months of exposure. An appropriate control relationship between response resolution and cessation of abacavir use was also required to confirm diagnosis. Typically, in cases when the clinical diagnostic criteria were fulfilled but when an alternative explanation for these responses could not be ruled out, patients were included based on the presence of a single well-defined skin patch test response to abacavir. For each individual, any *ex vivo* abacavir stimulation were undertaken to identify the role of abacavir. **Questionnaire** - Every eighth individual exposed to abacavir since December 2001 was identified for a cohort study to the results of genetic analysis within the updated diagnostic scheme. However, individuals with HLA-B*5701 and HspAIL (M493T) were actively excluded from response to abacavir.

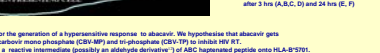
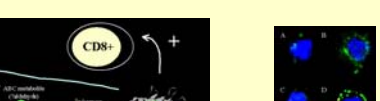
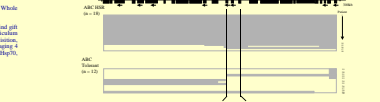
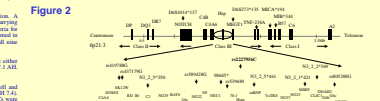
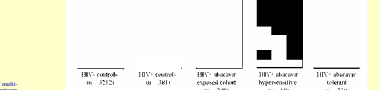
RESULTS

HLA-B*5701 - The frequency of HLA-B*5701 in the 270 participants was 10.4% (28/270). In the 270 individuals who were included in our previous study¹¹ were included based on clinical criteria including the occurrence of clinical hypersensitivity response of multi-organ involvement that was present within 2 months of exposure. An appropriate control relationship between response resolution and cessation of abacavir use was also required to confirm diagnosis. Typically, in cases when the clinical diagnostic criteria were fulfilled but when an alternative explanation for these responses could not be ruled out, patients were included based on the presence of a single well-defined skin patch test response to abacavir. For each individual, any *ex vivo* abacavir stimulation were undertaken to identify the role of abacavir. **Questionnaire** - Every eighth individual exposed to abacavir since December 2001 was identified for a cohort study to the results of genetic analysis within the updated diagnostic scheme. However, individuals with HLA-B*5701 and HspAIL (M493T) were actively excluded from response to abacavir.

DISCUSSION

Our study identifies a new susceptibility locus within the 57.1 AH region between *MEGTL1* and *CA46*. This region is highly polymorphic and contains several genes, including *Hsp70* and *HLA-B*5701*. We have identified a haplotypic variant of *Hsp70* (M493T) that is strongly associated with ABC HSR when carried on the 57.1 AH background. This finding suggests that the 57.1 AH region contains a susceptibility locus for ABC HSR, and that the M493T variant may be a marker for this locus. The M493T variant is a non-synonymous polymorphism of the Hsp70 gene, which is a stress-inducible protein that is involved in the heat shock response. The M493T variant has been shown to be associated with increased intracellular expression of Hsp70 in abacavir-exposed PBMCs. This finding is consistent with the hypothesis that ABC HSR is a hypersensitivity reaction to abacavir presented by HLA-B*5701. The M493T variant may also play a role in the regulation of Hsp70 expression. Further studies are needed to clarify the role of the M493T variant in ABC HSR.

CONTACT-Prof Simon Mallat, Executive Director Centre for Clinical Immunology & Biomedical Statistics Level 2, N Block, Royal Perth Hospital Wellington Street, WA, 6000, AUSTRALIA S.Mallat@murdoch.edu.au Ph:(61) 8 9224 2899, Fax:(61) 8 9224 2920 Website-http://www.maths.murdoch.edu.au/ccibs/



RESULTS

An updated restrictive definition of abacavir hypersensitivity individuals was used in this study. In a prospective study, 48 abacavir-recipients were HLA typed prior to exposure. All patients with HLA-B*5701, and DR7, DO3 were actively excluded. Two out of the 48 patients (4.2%) developed an ABC HSR reaction, both of whom carried the HLA-B*5701 allele, whilst the remaining 46 patients remained asymptomatic on abacavir therapy. Hence, the incidence of abacavir hypersensitivity among HLA-B*5701 negative individuals in this prospective cohort was 0% (95% confidence interval 0%-0.075%).

The distribution of MHC alleles was significantly different in the abacavir hypersensitive group compared to the abacavir tolerant group (Table 2, Fig 1). With regard to alleles that are specific to the 57.1 AH, HLA-B*5701 was present in 17/18 cases (94.4%) with abacavir hypersensitivity and 4/230 (1.7%) abacavir tolerant controls (OR=960, *p*<0.0001). CA46 was present in 14/18 cases (77.8%) and 7/230 (3.0%) controls (OR=111, *p*<0.0001), and the haplotypic combination of HLA-B*5701 and DO3 was present in 14/18 cases (77.8%) and 1/230 (0.4%) controls (OR=97, *p*<0.0001). Hence, HLA-B*5701 provided the strongest univariate association with susceptibility to abacavir hypersensitivity. Combining these markers, the previously described HLA-B*5701, CA46, HLA-DRB1*0701, DO3/haplotype (1) was present in 14/18 cases (77.8%) and none of the 230 controls (OR=1485, *p*<0.0001).

Recombinant haplotype analysis to map the putative susceptibility locus was carried out on the restricted patient sample recombinant for the 57.1AH in the MHC region (Fig 2). Polymorphic markers between CA46 and MEGT1 were identified by comparing the 57.1 AH with a subset of well-characterised EDV-variant control cell lines representative of common Caucasian AHs (data not shown). With reference to abacavir hypersensitive individuals, the boundary of the putative susceptibility region included the Hsp70 region of the 57.1AH. The translated sequences of Hsp70.1 and Hsp70.2 were similar in both abacavir hypersensitive individuals (cases 15-17) and controls with recombinant 57.1 haplotypes (cases 19-22), and were therefore excluded as candidate susceptibility loci. We observed that the Hsp70 Hom M493T allele, carried on the 57.1 AH, was detected in 94.4% of the hypersensitive group compared with 22.2% of tolerant controls (17/18 vs 51/230, *P*=0.000001, OR=593, *P*<0.000001). However, in combination with HLA-B*5701, the Hsp70 Hom M493T allele was strongly associated with abacavir hypersensitivity (17/18 vs 1/230, *P*=0.000001, OR=3893, *P*<0.000001) (Table 2).

The proportion of TNF-positive cells was higher in patients with abacavir hypersensitivity (*n*=8) than abacavir tolerant controls (*n*=2) with a median 13.4-fold increase (median 13.4, IQR=13.0) compared with a median 8.3-fold decrease in proportion of TNF positive cells with untyped or CD8-depleted cultures (Fig 4). Extracellular levels of TNF were higher in abacavir stimulated whole blood cultures of the ABC HSR individual compared with the tolerant control (Fig 5). In addition, TNF levels were augmented in the abacavir-stimulated blood of an abacavir hypersensitive patient when CD8+ T cells were depleted, compared with untyped or CD8-depleted cultures (Fig 4), suggesting the involvement of MHC class I molecules and CD8+ T cells in the development of this immune reaction.

A parsimonious model that suggests Hsp70 Hom M493T assisted cross-presentation of abacavir and/or its hapten by HLA-B*5701 to the immune system has been presented (Fig 5).

Co-localisation of HLA-B57 and Hsp70 was observed in discrete, punctate vesicles in peri-cell membrane and peri-nuclear region of a patient with ABC HSR but not in a tolerant control (Fig 6) studied using confocal microscopy CD14+ cells expressed (Fig 7) Hsp70 molecules. HLA-B57 and Hsp70 appear to co-localise within late endosomes (Fig 8) early endosomes (Fig 9) and the ER (Fig 10). A model describing the expression and subsequent compartmentalisation has been suggested (Fig 11).

CONCLUSIONS

- The putative HSR region includes a 14kb Hsp70.2 to Hsp70.1 gene interval.
- The association of HLA-B*5701 and Hsp70 Hom M493T is highly predictive of abacavir hypersensitivity.
- Co-localisation of HLA-B57 and Hsp70 occur in early, late endosomes and endoplasmic reticulum of CD14+ cells of ABC HSR individuals.

REFERENCES

Herberichsen et al. Clin Ther. 2001; 23:1600-1605. Walskley et al. AIDS. 1999; 13: 999-1000. Teague L et al. AIDS. 1999; 13: 1419-20. Mallat et al. Lancet. 2002; 359: 727-32. Herberichsen et al. Lancet. 2002; 359: 1121-22. Martin et al. 2004. HLA-Associated Immunobiology of the Human MHC. Proceedings of the 13th International Histocompatibility Workshop and Congress, in press. Nolan et al. J HIV Ther. 2003; 3: 44-41. Park et al. Toxicology. 2001; 150: 11-23. Zeman et al. J Clin Invest. 1998; 102: 1591-98. Whittips et al. AIDS. 2000; 14: 2223. Fourie et al. 1994 J Biol Chem. 269: 30470-70. Walsh et al. J Biol Chem. 2000; 142: 135. Martin et al. Proc Natl Acad Sci USA. 2000; in press. 14W et al. Mol Cell. 2001; 11: 3703-3715.

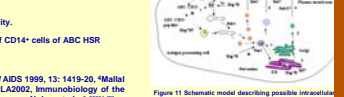
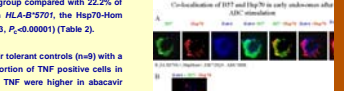
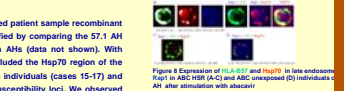
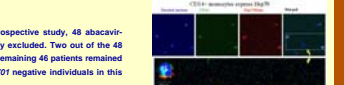


Figure 11 Schematic model describing possible intracellular Hsp70 and HLA-B57 expression and co-localisation. The model shows Hsp70 and HLA-B57 co-localising in discrete vesicles in the ER, endoplasmic reticulum, and late endosomes.