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Symposium

## [S-3] S-3: Biomarkers and TDM of immunosuppressive drugs

Chairs: Uwe Christians, USA / Satohiro Masuda, Japan

2017年9月25日(月) 15:00 ~ 17:00 Main Hall (1F)

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### [S-3-4] Determinants of intra-graft tacrolimus concentrations in renal transplant recipients

Benedetta Sallustio<sup>1</sup>, Benjamin Noll<sup>2</sup>, Rong Hu<sup>3</sup>, Daniel Barratt<sup>4</sup>, Martin Basic<sup>5</sup>, Janet Coller<sup>6</sup>, Andrew Somogyi<sup>7</sup> (1.The Queen Elizabeth Hospital, 2.University of South Australia, 3.University of Adelaide, 4.University of Adelaide, 5.University of Adelaide, 6.University of Adelaide, 7.University of Adelaide)

キーワード : Tacrolimus, CYP3A5, P-glycoprotein, Kidney, tissue concentrations

#### Background

Whilst the calcineurin inhibitors (CNI), cyclosporine and tacrolimus, have significantly improved graft survival, their long-term use is limited by nephrotoxicity. Both are substrates for cytochrome P450 3A (CYP3A) and P-glycoprotein (P-gp). P-gp expression determines chronic tubulointerstitial damage in transplanted kidneys and intra-renal cyclosporine concentrations. Although tacrolimus has largely replaced cyclosporine, little is known regarding the factors that determine its intra-renal exposure, and hence potential nephrotoxicity.

#### Methods

This was a retrospective study in 134 transplant recipients from whom 239 matching blood and renal cortical biopsy samples had been collected between 2-2490 days post-transplantation. Trough blood ( $C_B$ ) and renal cortex ( $C_R$ ) tacrolimus concentrations were measured by LC-MS/MS. P-gp expression was assessed by immunohistochemistry in paraffin-embedded biopsy samples. Donor *CYP3A5* genotypes (\*1/\*1, \*1/\*3, \*3/\*3) were determined by TaqMan SNP Genotyping (ThermoFisher Scientific). Univariate and multivariate analyses were used to investigate the relationship between  $C_B$  and  $C_R$  and: tacrolimus dose, *CYP3A5* genotype, P-gp expression, acute CNI toxicity, generalized proximal tubular injury, rejection and delayed graft function.

#### Results

$C_B$  ranged from 2.6-52.3 mg/L and  $C_R$  from 33-828 pg/mg tissue (n=239). In univariate analyses, there was a weak but significant correlation between  $C_B$  and  $C_R$  (Spearman  $r = 0.44$ ,  $P < 0.0001$ ), and  $C_R/C_B$  was inversely correlated with time post-transplant (Spearman  $r = -0.16$ ,  $P = 0.03$ ). In the first month post-transplantation there was no effect of either graft P-gp expression ( $r = 0.01$ ) or donor *CYP3A5* expressor genotype ( $P = 0.81$ ) on  $C_R/C_B$ . However, median  $C_R/C_B$  was 1.7-fold higher in patients with acute CNI toxicity compared to those without ( $P = 0.004$ ). Using a linear multiple regression model (R studio), only rejection, time post-transplant, CNI toxicity and rejection:time post-transplant were significant predictors of  $C_R/C_B$  (multiple  $R^2 = 0.06$ , adjusted  $R^2 = 0.04$ ,  $P = 0.025$ ).

#### Conclusions

Blood tacrolimus concentrations only predict 20% of variability in renal tacrolimus concentrations. When adjusted for tacrolimus concentration in blood, graft tacrolimus exposure is not significantly affected by renal cellular clearance pathways (P-gp or CYP3A5). However, it is associated with CNI toxicity and rejection.