Mechanisms underlying the lack of endogenous processing and CLIP-mediated binding of the invariant chain by HLA-DP^{84Gly}

Mark Anczurowski^{1,2}, Yuki Yamashita¹, Munehide Nakatsugawa¹, Toshiki Ochi¹, Yuki Kagoya¹, Tingxi Guo^{1,2}, Chung-Hsi Wang^{1,2}, Muhammed A. Rahman¹, Kayoko Saso¹, Marcus O. Butler^{1,2,3}, Naoto Hirano^{*1,2}

¹Tumor Immunotherapy Program, Campbell Family Institute for Breast Cancer Research, Campbell Family Cancer Research Institute, Princess Margaret Cancer Centre, University Health Network, Toronto, Ontario, Canada M5G 2M9 ²Department of Immunology, University of Toronto, Toronto, Ontario, Canada M5S 1A8; ³Department of Medicine, University of Toronto, Toronto, Ontario, Canada M5S 1A8

*Correspondence should be addressed to N.H.

Contact Information

Correspondence and requests for materials should be addressed to N.H. (naoto.hirano@uhnresearch.ca).

Naoto Hirano, MD, PhD

Princess Margaret Cancer Centre

610 University Avenue, Toronto, ON M5G 2M9, Canada

Phone: (416) 946-2190

Fax: (416) 946-6529

E-mail: naoto.hirano@uhnresearch.ca

Supplementary Figure 1



Supplementary Figure 1: CLIP derived from endogenous Ii is presented by HLA-DP4^{84DEAV87}, but not DP4, yet can associate with both DP4 and DP4^{84DEAV87}-expressing HEK293 cells when exogenously pulsed to the cell surface as a synthetic peptide. (a) Surface class II and CLIP expression, along with intracellular Ii expression, were analyzed on HEK293 transfectants by flow cytometry following staining with specific mAbs. HEK293 cells were retrovirally transduced with *DPA1*01:03* (DPA1) and either wild-type or mutated *DPB1*0401* (DPB4) with substitution mutations of the 84-87 region of the DP β chain made as indicated. (b) Parental HEK293 or HEK293 cells stably expressing DP4 or DP4^{84DEAV87} were pulsed with 100 μ M of CLIP, TT₉₄₇₋₉₆₇, or an equivalent volume of DMSO for 18 hrs. Surface CLIP expression was then analyzed by flow cytometry on the indicated HEK293 transfected cells following staining with CLIP-specific mAbs.

Supplementary Figure 2



Supplementary Figure 2: Expression data for cells used in T cell activation (Fig. 4) (a) and flow cytometry experiments (Fig. 5d) (b). Surface class II, CLIP and Δ NGFR expression, along with intracellular Ii and HLA-DM expression as indicated, were analyzed on the indicated K562 transfectants by flow cytometry following staining with specific mAbs. K562 cells were retrovirally transduced with *DPA1*01:03* (DPA1) and either wild type or mutated *DPB1*0401* (DPB4) with substitution mutations of the 84-87 region of the DP β chain made as indicated to generate DP4 or DP4^{84DEAV87} respectively.







Supplementary Figure 3: Transfection efficiencies for cross-linked cells lysed in Figure 3b and 3c (**a**), as well as for stimulator cells used in ELISPOT analysis in Figure 4 (**b**) and Figure 6 (**c**). Transduction efficiencies are measured by GFP expression in cells by flow cytometry.

Supplementary Figure 4



Supplementary Figure 4: Uncropped western blotting images from Figure 5c. Outlined regions represent portion of each image displayed in the main figures.