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Information about the contributions of each person named as having participated in the study

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3) Design & Methods. The following authors were responsible for specific investigations:

- Christina Bade-Doeding was responsible for eukaryotic and prokaryotic vector design, cloning, eukaryotic and prokaryotic protein expression, mass spectrometric analysis of the peptides, refolding, crystal seeding
- Alex Theodossis was responsible for crystallographic data collection, refinement of the structures, computational analysis
- Stephanie Gras was responsible for refolding, crystal seeding, crystallographic data collection
- Lars Kjer-Nielsen was responsible for prokaryotic vector design, cloning, prokaryotic protein expression
- Trevor Huyton was responsible for statistical analysis

4) Results. The following authors were responsible for specific portions of the results, including figures and tables:

- Christina Bade-Doeding was responsible for peptide analysis, Table 3: Polymorphic positions and corresponding peptide anchors within in the B*41 group, Supplementary Table 1: Ligands of the HLA-B*41 variants, Supplementary Table 2: List of promiscuous peptides, Supplementary Table 3: Differentially selected peptides by B*41 subtypes

- Alex Theodossis was responsible for crystallographic analysis, Table 1: Data collection and refinement statistics, Figure 1: The Crystal structures of B*41:03/AEMYGSVTEHPSPSPL and B*41:04/HEEAVSVDRVL, Figure 2: Peptides bound to B*41:03 and B*41:04, Figure 3: A conserved network of interactions in the B*41:03/16mer and B*41:04/11mer complexes, Supplementary Fig. 1: Stereographic representations of the Ag-binding cleft of B*41:03/AEMYGSVTEHPSPSPL and B*41:04/HEEAVSVDRVL, Supplementary Figure 2: Polymorphism alters the size and charge of the Ag-binding cleft in B*41:03 and B*41:04, Supplementary Table 4: Peptide contacts in the B*41:03/16-mer and B*41:04/11-mer structures, , Supplementary Table 5: Theoretically calculated ionisation states for selected titratable groups in the B*41:03 and B*41:04 structures
- Trevor Huyton was responsible for statistical analysis of the peptides, Table 2: Statistical analysis of the length of HLA-B*41-derived ligands

5) **Writing the manuscript.** The following authors were responsible for writing the manuscript:

- Christina Bade-Doeding was responsible for writing the manuscript
- Alex Theodossis was responsible for writing the manuscript
- Jamie Rossjohn was responsible for writing the manuscript
- Stephanie Gras was responsible for critically reviewing the manuscript
- Lars Kjer-Nielsen was responsible for critically reviewing the manuscript
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- Trevor Huyton was responsible for critically reviewing the manuscript

6) **Contributors Listed in Acknowledgments:**

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The atomic coordinates and structure factors (codes 3LN4 and 3LN5) have been deposited in the Protein Data Bank Japan (<http://www.pdbj.org/>).

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