

TITLE	<b>Identification of overlapping B-cell and T-cell epitopes</b>
Authors	Dagmar Hildebrand, Elke Hoffner, Barbara Gonzales, <b><u>Thorsten Zacher</u></b> , Fiordiligie Casilag, Volker Stadler
Affiliations	<i>PEPperPRINT GmbH, Heidelberg, Germany</i>
Abstract text  (max. 2000 characters)	<p>Chimera peptide vaccines including assembled B-/T-cell epitopes are one approach to induce a humoral as well as a cellular immune response. Overlapping B-/T-cell epitopes with coincident core sequences could benefit a pursued minimal protective epitope subset. Epitope prediction tools can help to forecast immunogenicity. However, the prediction is limited and cannot replace laboratory screenings.</p> <p>Here we describe the use of peptide microarrays combined with functional T-cell activation assays to discover overlapping B-/T-cell epitopes. We applied this approach to unravel immunogenic epitopes in Epstein-Barr Virus (EBV) Nuclear Antigen-1 (EBNA-1) as a proof of principle. To identify B-/T-cell epitopes in EBNA-1, we analyzed sera and peripheral blood mononuclear cells from EBV-positive healthy donors. First, we mapped the humoral immune response and determined infection-elicited antibodies against EBNA-1, via screening sera with PEPperCHIP® EBV Peptide Microarrays, containing 5,549 linear peptides of immunogenic EBV antigens including 257 overlapping EBNA-1 peptides. After in-depth Analysing the IgG responses, we selected three overlapping peptides within EBNA-1 with a consensus motif that showed high antibody-binding. Subsequently, to discover potential overlapping T-cell epitopes within these, the three peptides were synthesized and utilized in IFN-<math>\gamma</math> ELISpot assays as single peptide or peptide minipools together with published immunogenic EBV-peptides. The high sensitivity of the ELISpot assay allowed discrimination between T-cell-activating / non-activating peptides and facilitated the identification of the recognized, immunogenic core sequence.</p> <p>In conclusion, our proof of principle study shows that the combination of peptide microarrays and ELISpot assays is suitable to identify overlapping B- / T-cell epitopes. As the approach is applicable to any other antigen, it could generally help to identify new peptide vaccine candidates.</p>