

Supplementary File for

**ACME: Pan-specific peptide-MHC class I binding
prediction through attention-based deep neural
networks**

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This file contains the binding motifs of all the alleles with experimental data in the IEDB MHC class I binding dataset. Each motif is visualized using a heatmap where every pixel stands for a specific type of amino acid at a certain position in the peptide. The brighter a pixel is, the more important the corresponding residue would be. Motifs were generated as described in Materials and Methods.











