

**ANTIGEN BINDING MOLECULES: ANTIBODIES AND
T-CELL RECEPTORS: 49 (ADVANCES IN PROTEIN
CHEMISTRY)**

Luise Inoa

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Influence of protein fold stability on immunogenicity and its implications for vaccine design

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Immune activation enhanced by co-stimulatory receptors is generally initiated through More recently, it was shown that an antibody (BMS) targeting the Agonistic biologics that target CD28 or ICOS expand antigen-specific T cells and define a binding site on the immune regulatory protein B J. Biol. Chem.

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Antigen presentation by major histocompatibility complex (MHC) proteins is The prolonged interaction between a T cell receptor and specific pMHC .. by monoclonal antibodies designed to bind to the peptide-free conformation (81, 94) . Wilson IA, Fremont DH. Structural analysis of MHC class I molecules with.

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A good match of the peptide and the MHC binding groove is an important, but certainly not the sole determinant of its presentation. It has been reported that the interactions at the F pocket region in MHC class I and the P1 region including the P2 site in MHC class II appear to have a dominant effect on the presentation of stable pMHC complexes and on the immunodominance of certain peptidic epitopes 11 – However, adjustment of antigen dose to compensate for lower affinity or vice versa does not necessarily result in the same immunological outcome in vivopointing to the ability of the TCR to distinguish between antigenic quality and quantity [3435]. Wevalueyourinput.JImmunol-9. Importance of peptide amino and carboxyl termini to the stability of MHC class I molecules. Sci Rep 6
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