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CD1d lipid-antigen recognition by the $\gamma\delta$ TCR

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The T lymphocytes repertoire is divided into two major lineages, $\alpha\beta$ and $\gamma\delta$ T cells, which are defined by their T cell receptor (TCR) gene-segment usage. To date, the key discoveries on human CD1d restricted T cells have focussed on the type I Natural Killer T cells (NKT) subset that express an invariant TCR α chain (V α 24J α 18) which pairs with a β chain (V β 11). The structural basis for the recognition of CD1d-lipid antigen by type I NKT cells is also now well established [1, 2]. However, there are other subsets of NKT cells that exhibit reactivity towards lipid-antigen presenting molecules (CD1d) but that do no express the typical V α 24J α 18 TCR. We identify human NKT cell subsets that express V δ 1+ $\gamma\delta$ TCRs that recognize CD1d presenting the lipid-antigen α -galactosylceramide (α -Galcer). Here, we describe the first crystal structure of a CD1d/ $\gamma\delta$ TCR ternary complex [3] and provide structural insights into the binding mode of a $\gamma\delta$ TCR with CD1d- α Galcer. The $\gamma\delta$ TCR binds orthogonally over the A' pocket of CD1d, that is in clear contrast with the typical type I parallel docking mode in which the $\alpha\beta$ TCR is perched over the F' pocket of CD1d. The germ line-encoded CDR1 δ loop dominates the contacts with the CD1d molecule while the CDR3 γ loop represents the main structural determinant for the antigen specificity. These findings highlight the emergence of diverse populations of NKT TCRs that exhibit different binding mode and lipid antigen specificity.

[1] NA. Borg, KS. Wun, L. Kjer-Nielsen et al. Nature, 2007, 448, 1137-1145., [2] KS. Wun, NA. Borg, L. Kjer-Nielsen et al. J. Exp. Med., 2008, 205, 939-949., [3] AP. Uldrich, J. Le Nours, DG. Pellicci et al. Nature Immunology, 2013, 14, 44-49.

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