

# SigN SEMINAR

*Hosted by Dr Chong Shu Zhen*

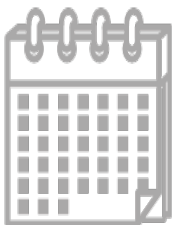


## Maximilien Evrard

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### Resolving circulating and tissue-resident memory T cells using high-dimensional protein screens

Memory T cells are essential gatekeepers ensuring host protection against microbial and cancerous threats. Paradigmatically, memory CD8<sup>+</sup> T cells can be broadly divided into circulating ( $T_{\text{CIRCUM}}$ ) and tissue-resident memory T ( $T_{\text{RM}}$ ) cell populations based on their trafficking properties. Despite well-defined migratory and transcriptional differences between these T cell subsets, the phenotypic and functional delineation of  $T_{\text{CIRCUM}}$  and  $T_{\text{RM}}$  cells, particularly across tissues, remains elusive. Here, we utilised a comprehensive screening platform of cell surface molecules combined with a machine learning prediction pipeline (InfinityFlow) to profile the expression of >250 protein markers in  $T_{\text{CIRCUM}}$  and  $T_{\text{RM}}$  cells across multiple anatomical sites. High-dimensional analysis provided detailed phenotypic definition and revealed unappreciated heterogeneity within and between  $T_{\text{CIRCUM}}$  and  $T_{\text{RM}}$  cell lineages in distinct tissue microenvironments. Importantly, we devised strategies that allowed for the selective ablation of distinct T cell subsets and identified stable markers allowing disentanglement of  $T_{\text{CIRCUM}}$  and  $T_{\text{RM}}$  cells and the characterisation of their effector profiles during inflammation. Together, these data and analytical framework provide an in-depth resource to enable the identification, phenotypic interrogation, and functional classification of memory T cells at steady-state and in disease contexts.



**3<sup>rd</sup> October 2022 (Monday)**  
**10 AM – 11 AM (Singapore Time)**  
**SigN Seminar Room, Immunos, Level 4**

**8A Biomedical Grove, Immunos, #04-06, Singapore 138648**

*Seminar is  
open for all  
to attend.*

*Registration  
is not  
required.*

