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On the cover: Contrasting proteomics signatures of tumour lysis products (black) and inflammation markers (blue) observed in the terminal plasma of two contrasting B-cell cancer models. E $\mu$ -myc and E $\mu$ -TCL1 mice model the aggressive Burkitt's lymphoma and the slow-progressing chronic lymphocytic leukaemia, respectively. Integrating plasma proteomics with splenic B-cell tumour proteomics enabled the attribution and deconvolution of these signatures. E $\mu$ -TCL1 tumours demonstrated IL5-driven proliferation.

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