

# Inhibition of cleavage of human complement component C5 and the R885H C5 variant by two distinct high affinity anti-C5 nanobodies

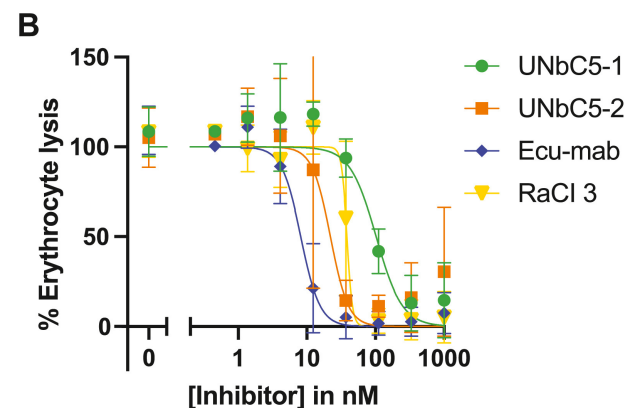
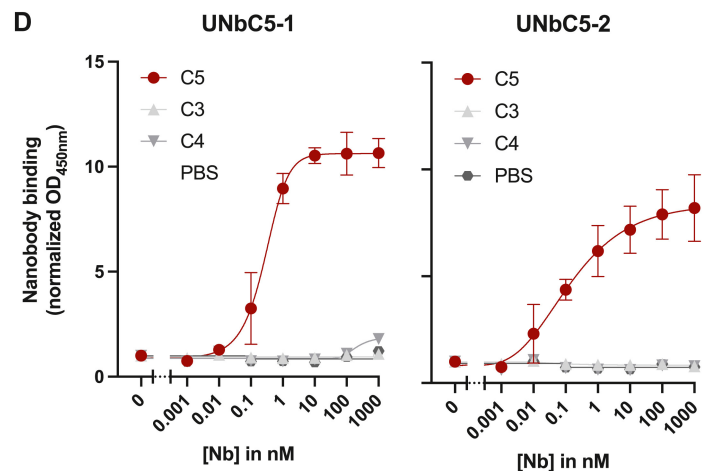
Eva M. Struijff<sup>1,†</sup>, Karla I. De la O Becerra<sup>2,†</sup>, Maartje Ruyken<sup>1</sup>, Carla J. C. de Haas<sup>1</sup>, Fleur van Oosterom<sup>1</sup>, Danique Y. Siere<sup>1</sup>, Joanne E. van Keulen<sup>1</sup>, Dani A. C. Heesterbeek<sup>1</sup>, Edward Dolk<sup>3</sup>, Raimond Heukers<sup>3</sup>, Bart W. Bardeol<sup>1</sup>, Piet Gros<sup>2</sup>, and Suzan H. M. Rooijackers<sup>1,\*</sup>

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Attention, la **Figure 1** originale contient un panel C mais nous vous demandons de commenter seulement les panels A, B et D représentés ici.

**A**

Clone	a.a.	Framework 1	CDR1	Framework 2	CDR2
UNbC5-1	1	EVQLVESGGGLVQAGGSLRRLSCAASGFTFD	DYAIG	WFRQAPGKERE	GVS CISTSDGSTYYADSVKG
UNbC5-2	1	EVQLVESGGGLVQPGGSLRRLSCAASGRFTFS	TNTMG	WFRQAPGQERE	FVA LISGNRILDYS
		Framework 3	CDR3	Framework 4	
UNbC5-1	67	RFTISSDNAKNTVYLQMNSLKPEDTAVVYCAA	DPYLP	IRGRGIESTDFGS	WGQGTQVTVSS
UNbC5-2	67	RFTISRDNKNTVYLQMNSLKPEDTGVYFCAA	EF----	RGRTL----	ASY WGQGTQVTVSS



**Figure 1. Identification of two C5-targeting nanobodies that interfere with complement.** *A*, protein sequence alignment of UNbC5-1 and UNbC5-2. Frameworks 1 to 4 and CDRs 1 to 3 are indicated. Conserved amino acids are indicated in *black* and unique amino acids are indicated in *red*. *B*, CP mediated hemolysis of antibody-coated sheep erythrocytes incubated with 2.5% normal human serum and a titration of our nanobodies UNbC5-1 and UNbC5-2 and known complement inhibitors RaCl3 and Ecu-mab. The  $A_{405}$  values of the supernatants were measured; the % erythrocyte lysis was calculated using the 0% lysis (buffer) and 100% lysis (milliQ water) control samples.

*D*, nanobody bind to complement protein C5 and not homologs C3 and C4. Microtiter plates were coated with complement proteins and incubated with increasing concentrations of UNbC5-1 and UNbC5-2. Nanobody binding was measured using polyclonal rabbit-anti-VHH QE19 antibodies and donkey-anti-rabbit-HRP antibodies, at an absorbance of 450 nm. Coating with PBS was taken along as a negative control. Data information: (*A*), sequences were aligned using T-coffee (42). *B–D*, data represent mean  $\pm$  SD of three individual experiments (*B* and *D*) curves were fitted and  $IC_{50}$  and  $EC_{50}$  values were obtained. CDR, complementarity determining regions; CP, classical pathway; HRP, horseradish peroxidase.