SUPPLEMENTARY MATERIAL

TITLE: Biochemical and biophysical characterization of purified native CD20 alone and in complex with rituximab and obinutuzumab

AUTHORS:

Morgane AGEZ¹, Elodie DESUZINGES MANDON¹, Thomas IWEMA¹, Reto GIANOTTI², Florian LIMANI², Sylvia HERTER², Ekkehard MÖSSNER², Eric A KUSZNIR³, Sylwia HUBER³, Matthias LAUER³, Philippe RINGLER⁴, Claudia FERRARA², Christian KLEIN² & Anass JAWHARI¹

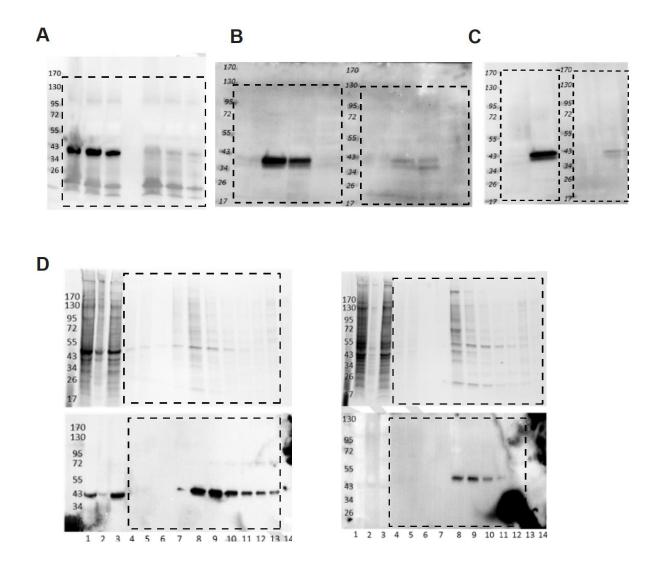


Figure S1: Original gels or Western-blot used for Figure 1A-D. A rectangle represents what is shown in the main figures.

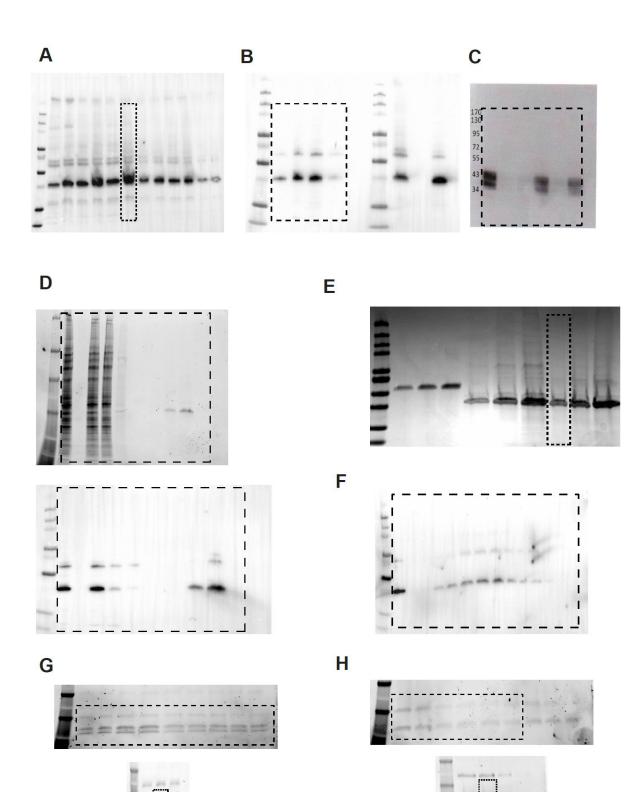


Figure S2: Original gels or Western-blot used for the main Figures. A-E correspond to Figure 2A-E. F corresponds to Figure 3A. G-H correspond to Figure 5A-B. A rectangle represents what is shown in the main figures.

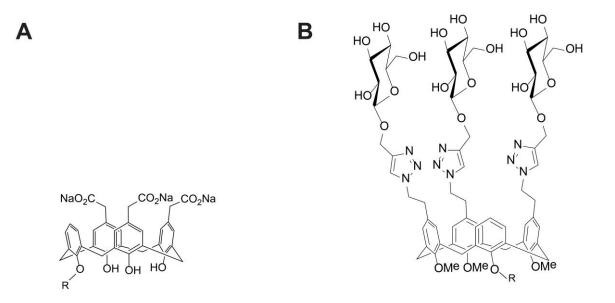


Figure S3: Structure of two calixarene based detergents used for solubilization and purification of CD20. **A**- 5,11,17-tris[(carboxy)methyl]-25-monooctyloxy-26,27,28trihydroxycalix[4]arene. Also called CALX-R2 in this manuscript. **B**- 5,11,17-tris(2-[4-(β -Dglucopyranosyl)-1H-1,2,3-triazol-1-yl]ethyl),-25- monoheptoxy-26,27,28trimethoxycalix[4]arene; respectively, also called CALX-173-GK. R corresponds to an aliphatic tail of seven carbons.

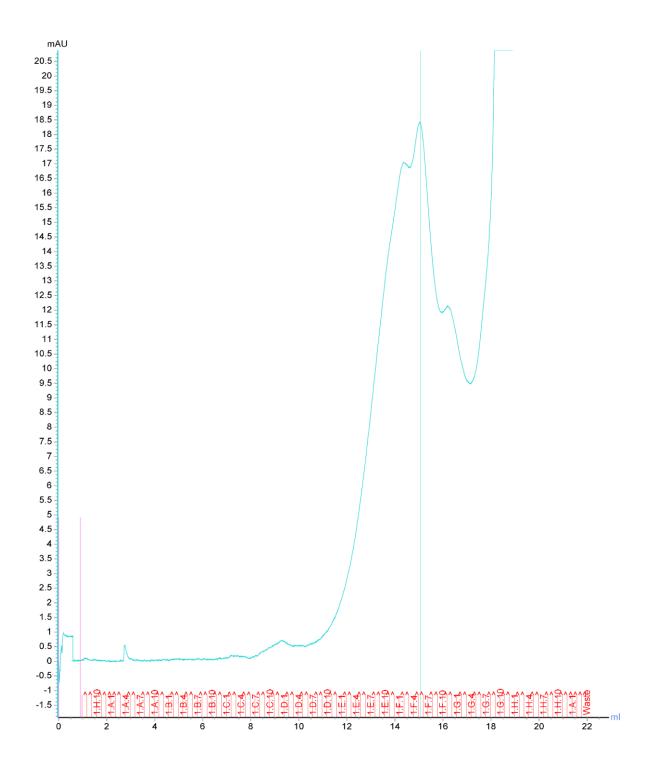


Figure S4: Full length size exclusion chromatogram of purified CD20.

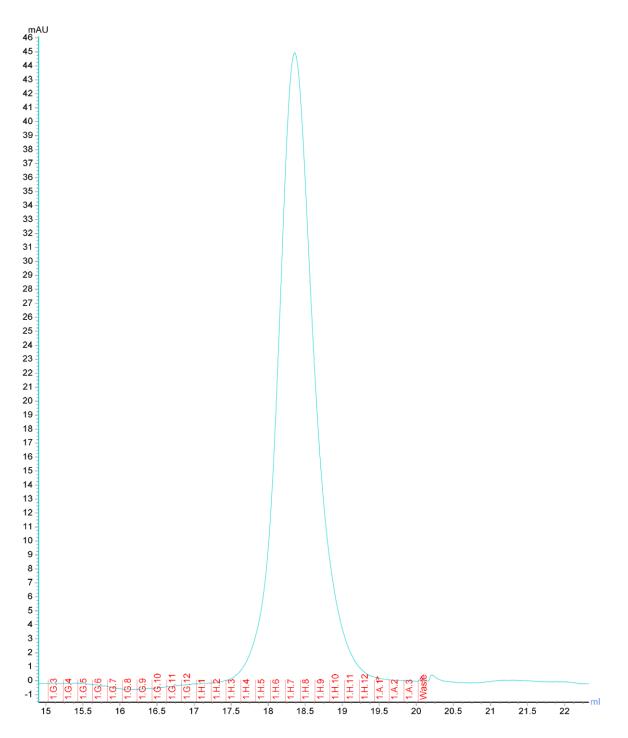


Figure S5: Full length size exclusion chromatogram of the rituximab.

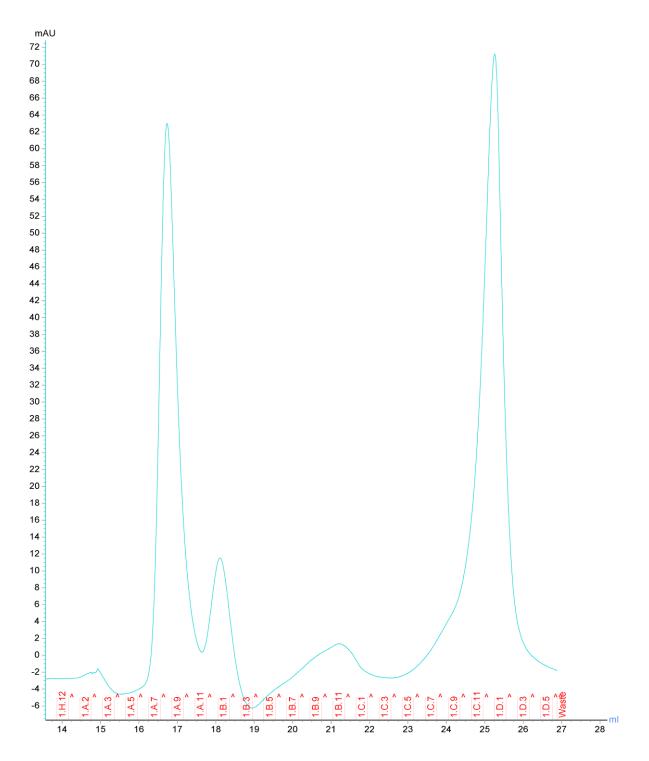


Figure S6: Full length size exclusion chromatogram of the obinutuzumab.

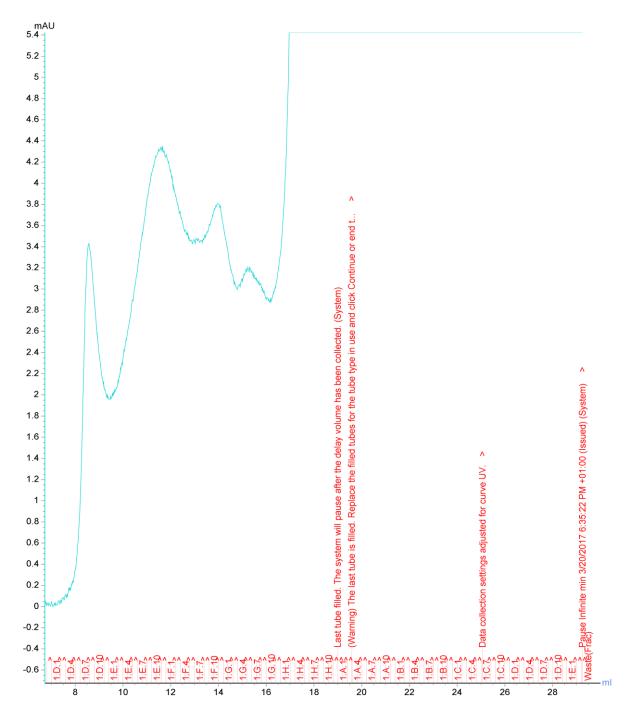


Figure S7: Full length size exclusion chromatogram of CD20/rituximab complex.

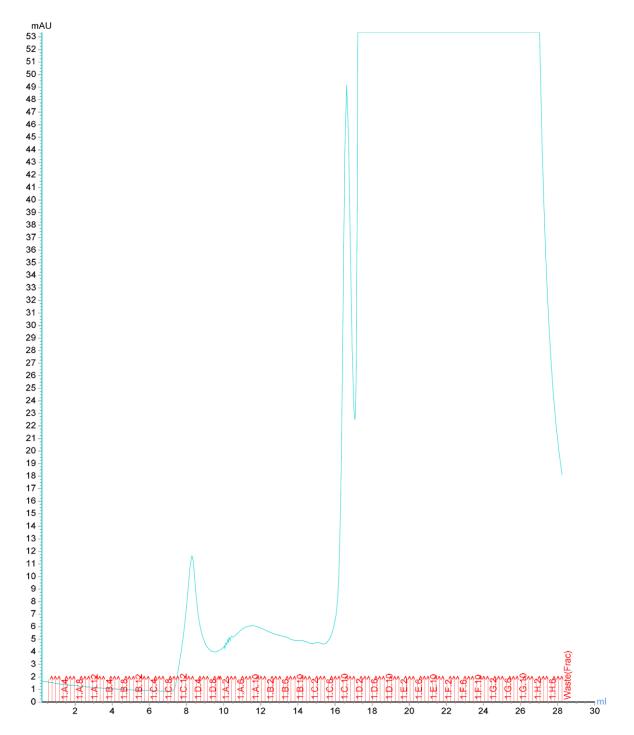


Figure S8: Full length size exclusion chromatogram of CD20/ obinutuzumab complex.

Sample	conc. [mg/ml]	Detergent in buffer	Detection system	Peak	sw [S]	f/f0	Area [%]	vbar [ml.g ⁻¹]	MW _{expec.} [kDa]	MW _{calc.} [kDa]	Interpretation
CALX-173- GK alone	5,000	CALX-173-GK	Interference	1	2,9	1,18	100	0,764	no	38	detergent micelle
CD20	0,500	CALX-173-GK	Absorbance	1	3,5	1,35	30	0,741	34	~ 54	probably monomeric CD20
				2	4,6		70			> 80	higher oligomers
				3	5,8						
				4	7,7						
				5	10,0						
CD20	0,500	CALX-173-GK	Interference	1	3,5	1,35	27	0,741	34	~ 61	probably monomeric CD20
				2	4,4		73			> 84	higher oligomers
				3	5,8						
				4	7,4						
				5	8,6						
				6	9,9						
DDM alone	20,000	DDM	Interference	1	2,9	1,00	100	0,836	50	53	detergent micelle
CD20	0,044	CALX-173-GK	Absorbance	1	3,6	1,35	46	0,741	34	~ 57	probably monomeric CD20
				2	5,8		17			>100	higher
				range	6-12		37				oligomers
CD20	0,044	DDM	Absorbance	1	3,1	1,42	72	0,741	34	~ 48	probably monomeric CD20
				2	5,0		28			99	probably dime

Table S9: Summary of AUC calculations of the molecular mass of CD20 species.

Expected molecular mass (MW_{expec}) was calculated from protein amino acid sequence. The partial specific volume (vbar) of the protein alone was taken into account as it is impossible to determine exact detergent/protein ratio due to simultaneous absorbance of the protein and CALX-173-GK detergent. Therefore, the calculated molecular masses should be interpreted as "estimates".