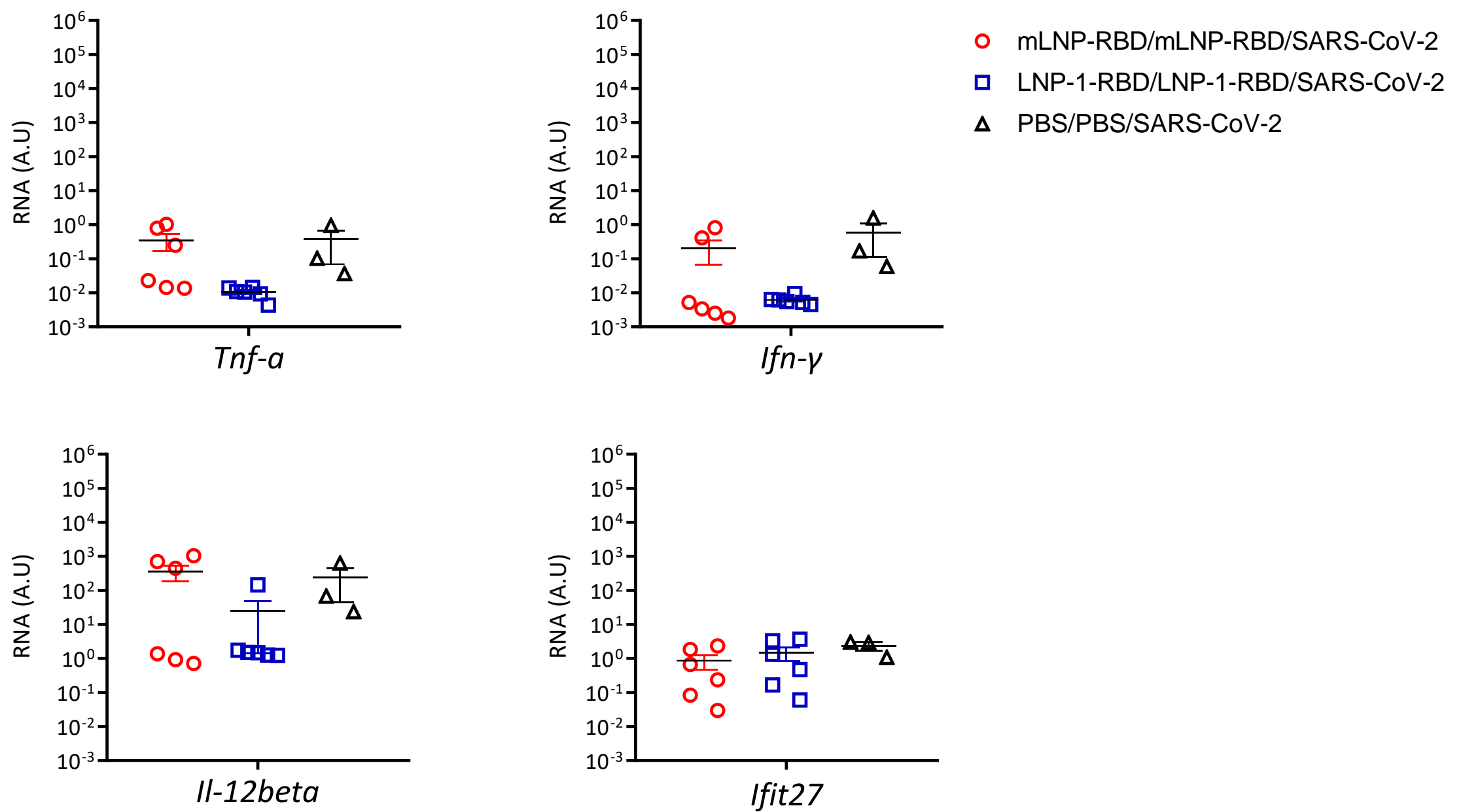


Supplementary Figure 1. Time-course expression of SARS-CoV-2 RBD protein after transfection of 293T cells with different naked mRNAs. (A) Scheme of the structure of the three different mRNAs expressing the full receptor binding domain (RBD: aas 330-532) of the SARS-CoV-2 spike protein (RBD-mRNA and RBD-mRNA*) or its highly immunogenic motif (RBDepi-mRNA). **(B)** Detection of RBD expression in transfected 293T cells at 3, 6 and 24 h post-transfection (h.p.t.) by flow cytometry using a rabbit polyclonal anti-SARS-CoV-2 spike/RBD antibody (5 μ g/mL). Geometric Mean Fluorescence Intensity (gMFI) values on the “live cells” gate were used to calculate the RBD score by applying the formula: No. RBD⁺ cells \times gMFI/No. live cells. Cells transfected with naked LUC-mRNA were used as negative control. Mean with standard error of the mean (SEM) is represented. A multiple *t* test was performed. ** $p < 0.005$; *** $p < 0.001$. **(C)** Kinetics of RBD expression in cellular pellets and supernatants from transfected 293T cells by western-blotting using a rabbit polyclonal anti-SARS-CoV-2 spike/RBD antibody (upper panels). Ponceau staining (lower panels) was used as loading control. All blots derive from the same experiment and were processed in parallel.



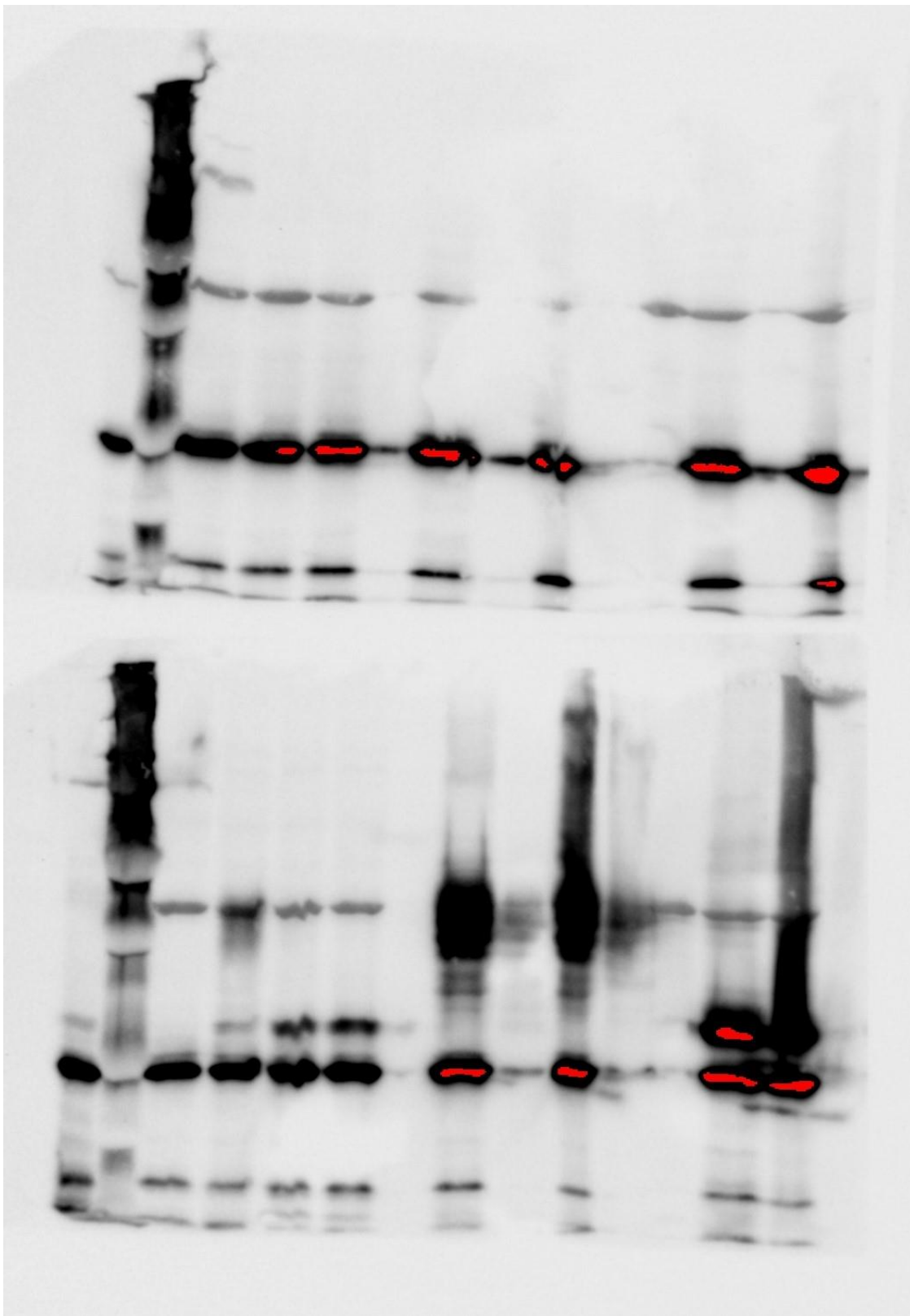
Supplementary Figure 2. mLNP-RBD and LNP-1-RBD formulations do not change significantly *Tnf-α*, *Ifn-γ*, *Il-12beta* and *Ifit27* gene expression profiles in lungs from vaccinated and challenged transgenic K18-hACE2 mice. Proinflammatory cytokines and chemokines were detected by RT-qPCR in lungs from individual mice at 14 days (groups 1 and 2) or 7 days (group 3) after SARS-CoV-2 challenge. Mean RNA levels (in A.U.) with standard error of the mean (SEM) from duplicates of each lung sample is represented; relative values are referred to uninfected mice (group 4). An ordinary one-way ANOVA of transformed data followed by Tukey's multiple comparison test was performed. ns > 0.05.

Supplementary Table 1

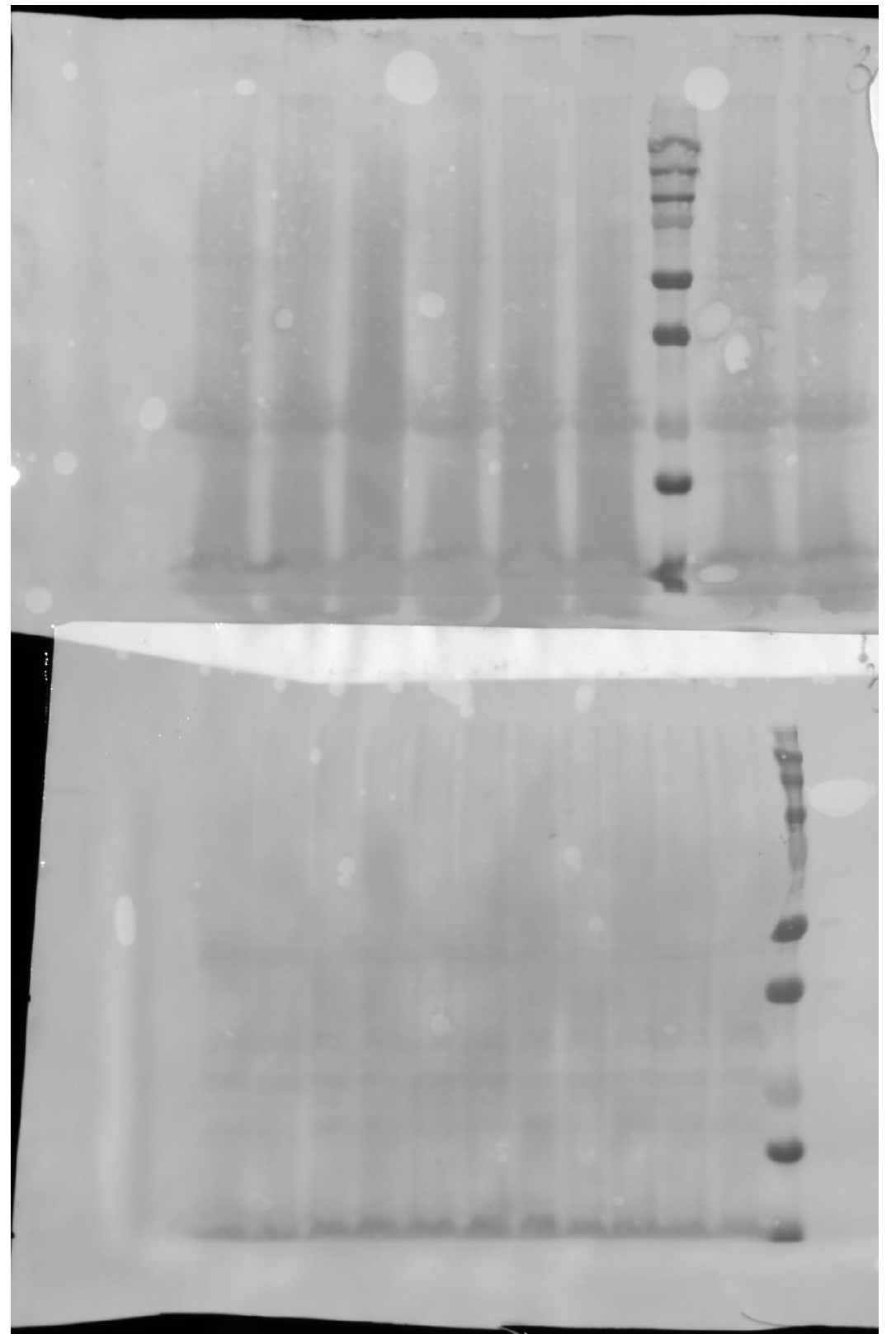
Target	Taqman probes
<i>Il-24</i>	Mm00474102_m1
<i>Ccl2</i>	Mm00441242_m1
<i>Ip-10 (Cxcl10)</i>	Mm00445235_m1
<i>Ifn-beta1</i>	Mm00439552_g1
<i>Cxcl5</i>	Mm00436451_g1
<i>Fcgr4</i>	Mm00519988_m1
<i>Ccl12</i>	Mm01617100_m1
<i>Timp-1</i>	Mm01341361_m1
<i>Il-10</i>	Mm00439616_m1
<i>Il-6</i>	Mm00446190_m1
<i>Ifn-γ</i>	Mm01168134_m1
<i>Tnf-α</i>	Mm00443258_m1
<i>Il-12beta</i>	Mm01288989_m1
<i>Ifit27</i>	Mm00508912_m1
	Primers/probe sets
<i>28s</i>	PROBE[JOE] 5' TAGTAGCTGGTTCCTCCGAAGTTTCCT 3' Rev 5' CGAGAGCAGCTATCCT 3' Fwd 5' GGCGAAAGACTAATCGAACCAT 3'
	Kit
<i>RdRp</i>	NZYTEch SARS-CoV-2 One-Step RT-PCR Kit II,
<i>N</i>	RdRp and N genes, IVD (MD04871)

Supplementary Table 1. Taqman probes used for the detection of proinflammatory cytokines and chemokines. Primers/probe sets used for the detection of 28S (housekeeping gene). Detection kit used for the quantification of the total SARS-CoV-2 genomic (*RdRp*) and subgenomic (*N*) RNA copy numbers.

Uncropped blots of Figure 1b

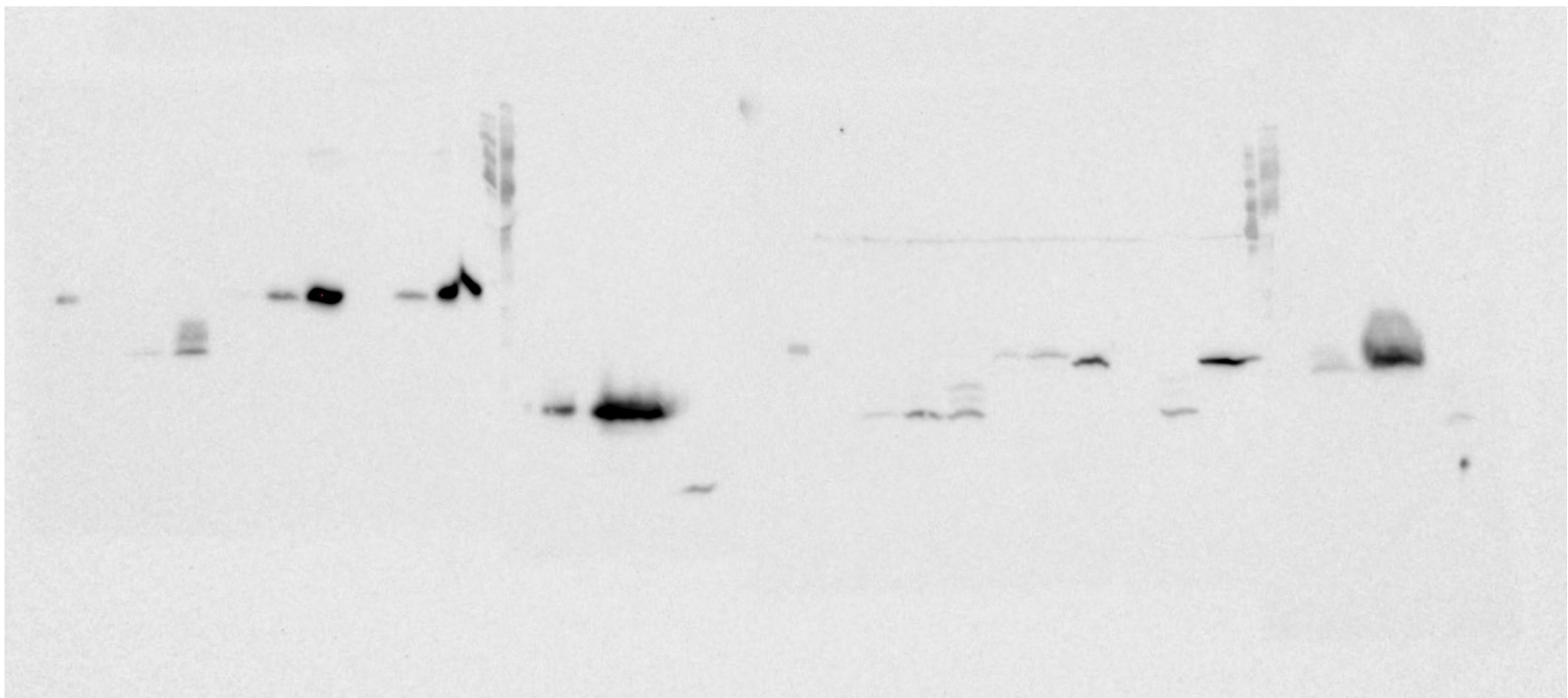


RBD staining

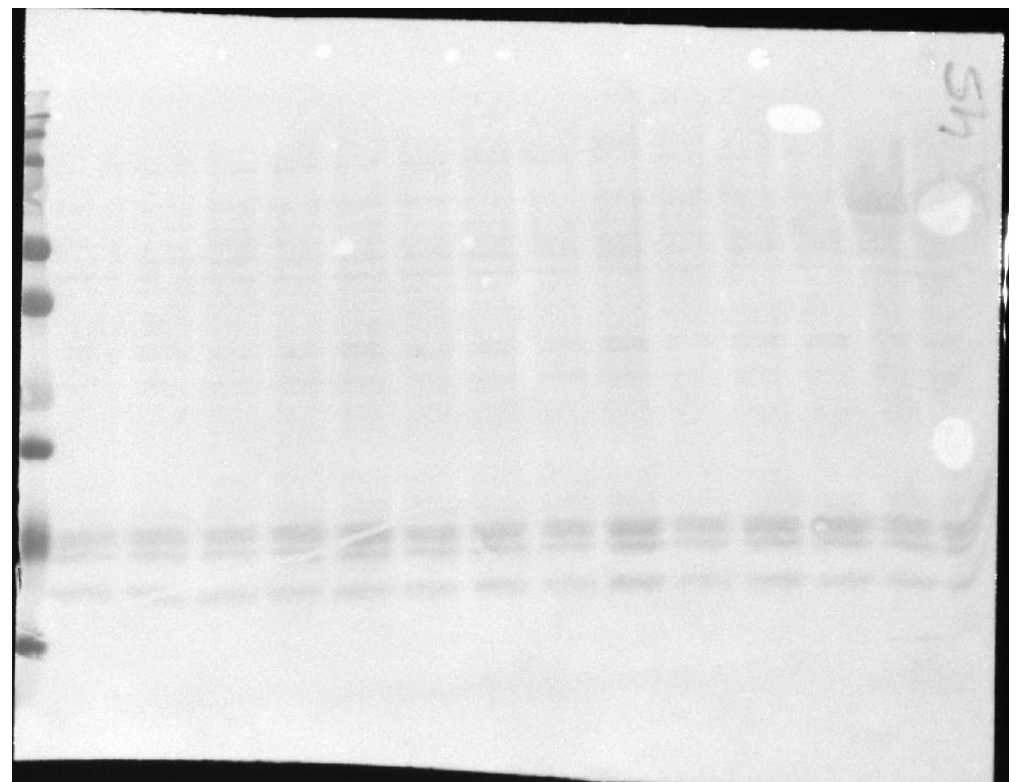
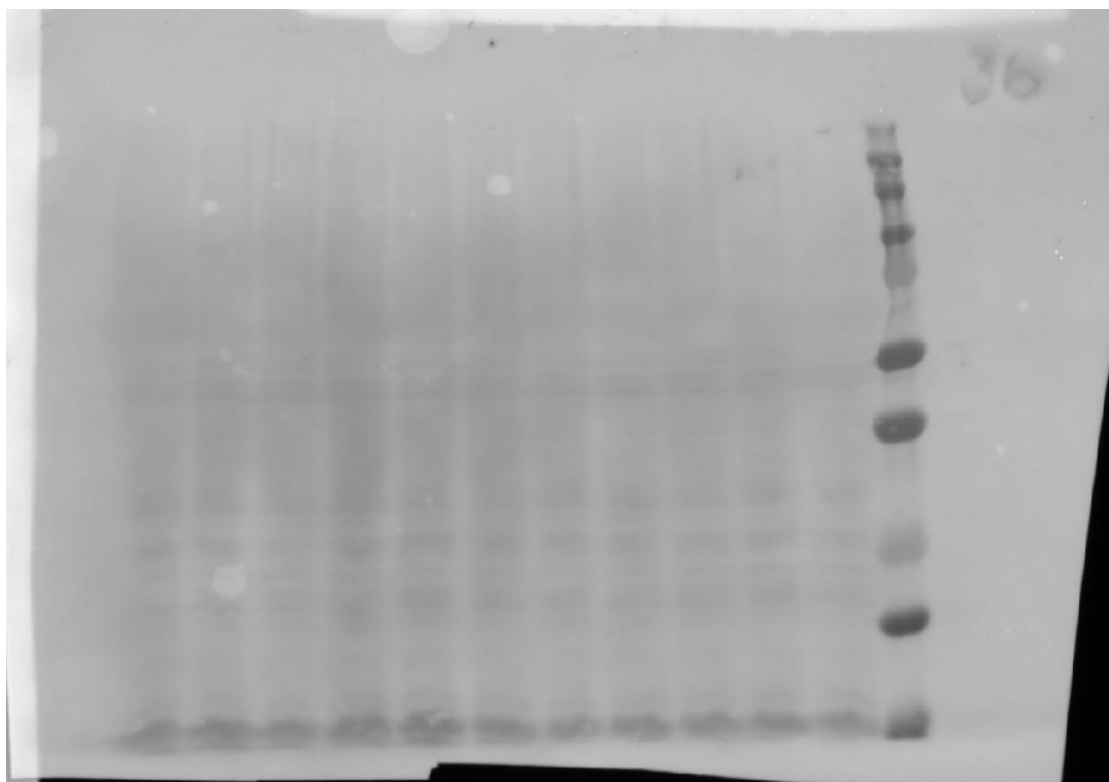


Ponceau staining

Uncropped blots of Supplementary Figure 1c



RBD staining



Ponceau staining