

Supplementary information

Experimental VLP vaccine displaying a furin antigen elicits production of autoantibodies and is well tolerated in mice

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Supplementary data 1 (separate .xlsx file "Additional file 2")

A)

SpyTag-noro-VLP capsid protein 1 (557 amino acids, 60.9 kDa):

MKMASSDANPSDGSAAANLVPEVNNEVMALEPVVGAIAAPVAGQONVIDPWIRNNFVQAPGGFTVSPRNA
PGEILWSAPLGPDLNPLYSLHARMYNGYAGGFVQVILAGNAFTAGKVIFAAVPPNFPTEGLSPSQVTMFPHIVV
DVRQLEPVLIPDPVRNNFYHYNQSNPTIKLIAMLYTPLRANNAGDDVFTVSCRVLTRPSPDFDFIFLVPPTVESR
TKPFSVPVLTVEEMTNSRFPIPLEKLFTGPSSAFVVQPQNGRCTTDGVLGTTQLSPVNICFRGDVTHITGSRNYT
MNLASQNWNDYDPTTEEIPAPLGTPDFVVGKIQGVLTQTRTRTDGSTRGHKATVYTGSAFAPKLGRVQFETDTRD
FEANQNTKFTPVGVIQDGGTTHRNEPQQWVLPSSYGRNTHNVHLAPAVAPTFPGEQLLFFRSTMPGCSGYPN
MDLDCLLPQEWVQYFYQEAAPAQSDVALLRFVNPDTGRVLFECKLHKSGYVTVVAHTGOHDLVIPPNGYFRFDS
WVNQFYTLAPMGNGTGRRRAVTSGG**AHIVMVDAYKPTK**

B)

SpyCatcher-PEED (122 amino acids, 13.7 kDa):

M**HHHHHH**DYDIPTT**ENLYFQ**GSGDSATHIKFS**KR**DEDGKELAGATMELRDSSGKTISTWISDGQVKDFLYPGKY
TFVETAAPDGYEVATAITFTVNEOGQVT**VNGLE**PEEDGTRFHRQASK

SpyCatcher-DIIG (120 amino acids, 13.2 kDa):

M**HHHHHH**DYDIPTT**ENLYFQ**GSGDSATHIKFS**KR**DEDGKELAGATMELRDSSGKTISTWISDGQVKDFLYPGKY
TFVETAAPDGYEVATAITFTVNEOGQVT**VNGLE**DIIGASSDCSTC

SpyCatcher-SWG (122 amino acids, 13.5 kDa):

M**HHHHHH**DYDIPTT**ENLYFQ**GSGDSATHIKFS**KR**DEDGKELAGATMELRDSSGKTISTWISDGQVKDFLYPGKY
TFVETAAPDGYEVATAITFTVNEOGQVT**VNGLE**SWGPEDDGKTVDGP

SpyCatcher-P domain (241 amino acids, 27.0 kDa):

M**HHHHHH**DYDIPTT**ENLYFQ**GSGDSATHIKFS**KR**DEDGKELAGATMELRDSSGKTISTWISDGQVKDFLYPGKY
TFVETAAPDGYEVATAITFTVNEOGQVT**VNGLE**VAPQRCIVEILVEPKDIGKRLEVRKAVTACLGEPNHITRLEHV
QARLTLSYNRRGDLAIHLISPMGTRSTLLAARPHDYSADGFNDWAFMTTHSWDEDPAGEWVLEIENTSEANNY
GTLTKFTLVLYGTAPE

Supplementary figure 1. Primary amino acid sequences of the vaccine components used for immunizations. A) A noro-VLP capsid protein 1 (black) with a C-terminal SpyTag (light blue). Linker sequence between the viral protein and SpyTag is marked with grey. B) The SpyCatcher-conjugated (dark blue) target regions (black) with an N-terminal purification tag (his-tag, red) and TEV-endonuclease site (orange). Linker sequences are marked with grey. In A and B, the isopeptide bond forming amino acids of both SpyTag- and SpyCatcher-moiety are highlighted with bolded and underlined letters.

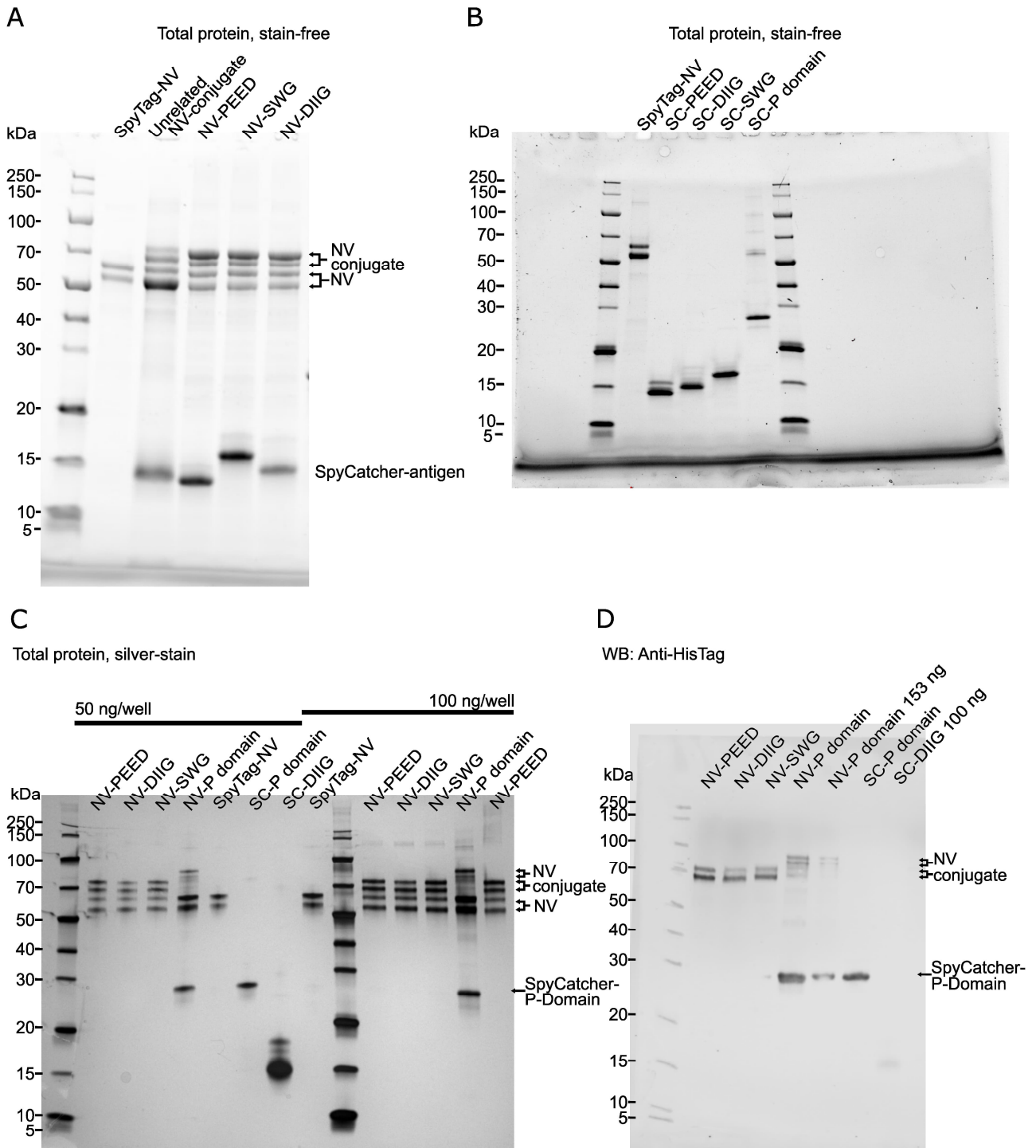
A)

PCSK1	VGGIRMLDGI-VTDAIEASSIG---FNP---GHVDIYSA	SWGPNDDGKTVE -GPGRLAQK	286
PCSK2	VAGIRMLDQPFMTDIEASSIS---HMP---QLIDIYSA	SWGPTDNGKTVD -GPRELTLQ	287
PCSK3/FURIN	IGGVRMLDGE-VTDAVEARSLG---LNP---NHIHIYSA	SWGPEDDGKTVD -GPARLAE	272
PCSK4	IGGVRMLDGT-ITDVIEAQSLS---LQP---QHIHIYSA	SWGPEDDGRTVD -GPGILTRE	277
PCSK5	IGGVRMLDGD-VTDMVEAKSVS---FNP---QHVHIYSA	SWGPDGDDGKTVD -GPAPLTRQ	290
PCSK6	IGGIRMLDGD-VTDVVEAKSLG---IRP---NYIDIYSA	SWGPDGDDGKTVD -GPGRLAKQ	324
PCSK7	IAGIRVLDGP-LTDSMEAVAFN---KHY---QINDIYSC	SWGPDGDDGKTVD -GPHQLGKA	306
	: . * : * : * * : * * : * * : * * * . * * * * * * : * : * * * : * * *		
PCSK8/MBTP1	LHIFRVFTNN----QVSYTSWFLDAFNAILKKIDVLNLSI-----	GGPDFM-DH	316
PCSK9	MRSRLRVLNCQ-GKGTVSGTLIG---LEF---IRKSQLVQPVGPLVLLPLAGGYSRVLNA		299

B)

Human	445	VAPQRKCIIDILTEPKDIGKRLEVRKTVTACLGEPNHITRLEHAQARLTLNRRGDLAI	503
		VAPQRKCI++IL EPKDIGKRLEVRK VTACLGEPNHITRLEH QARLTLNRRGDLAI	
Mouse	445	VAPQRKCIVEILVEPKDIGKRLEVRKAVTACLGEPNHITRLEHVQARLTLNRRGDLAI	503
Human	504	HLVSPMGTRSTLLAARPHDYSADGFNDWAFMTTHSWDEDPGSEWVLEIENTSEANNYGTL	563
		HL+SPMGTRSTLLAARPHDYSADGFNDWAFMTTHSWDEDP+GEWVLEIENTSEANNYGTL	
Mouse	504	HLISPMGTRSTLLAARPHDYSADGFNDWAFMTTHSWDEDPAGEWVLEIENTSEANNYGTL	563
Human	564	TKFTLVLYGTAPE 576	
		TKFTLVLYGTAPE	
Mouse	564	TKFTLVLYGTAPE 576	

Supplementary figure 2. A) A section of a multiple sequence alignment of the conventional PCSK family protein sequences from human. The SWG peptide antigen used in this study is highlighted in purple. The human sequences of unconventional PCSK proteins PCSK8/MBTP1 and PCSK9 are shown aligned below the conventional proteins. B) An alignment between the protein sequences of furin P domain of *Homo sapiens* and *Mus musculus*.

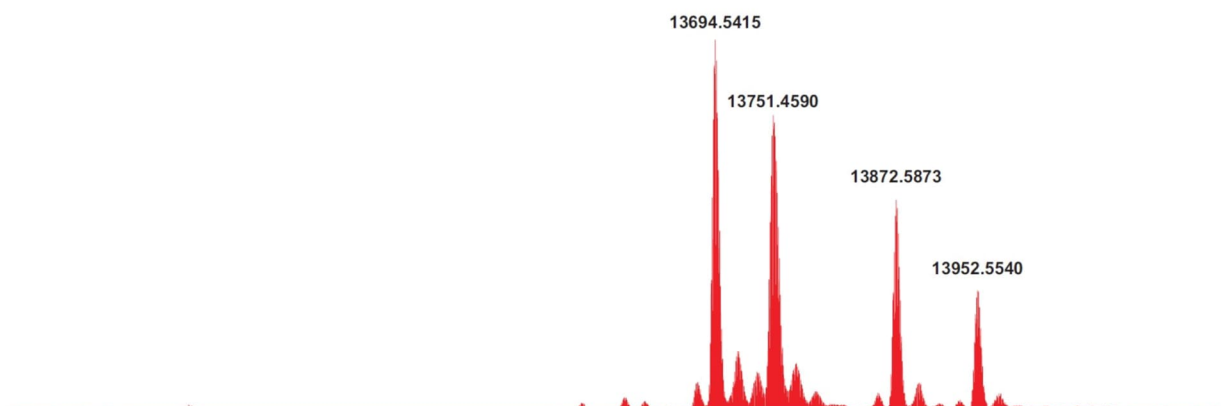


Supplementary **figure 3**. Characterization of noro-VLP (NV) based vaccine candidates against PCSK9 and furin. A) Stain-free SDS-PAGE gel showing the purified SpyCatcher PCSK antigens combined with SpyTag-NV before the removal of unreacted SpyCatcher-peptides. B) Uncropped gel from Figure 1C. C) Uncropped gel from Figure 1D. D) For Western blotting, we transferred the proteins onto nitrocellulose membrane using Trans-blot Turbo (Bio-Rad). The SpyCatcher-PCSK fusion proteins were identified based on the binding of an HisTag antibody (1:10 000, Thermo Fisher Scientific,

#ma1-21315) and IRDye 800CW goat anti-mouse IgG secondary antibody (1:20 000, LI-COR Biosciences, USA, #926–32210) with the Odyssey CLx instrument (LI-COR Biosciences, USA). All wells have between 500 and 2500 ng protein, unless otherwise mentioned. The molecular weight ladder is marked in the blot with a fluorescent pen.

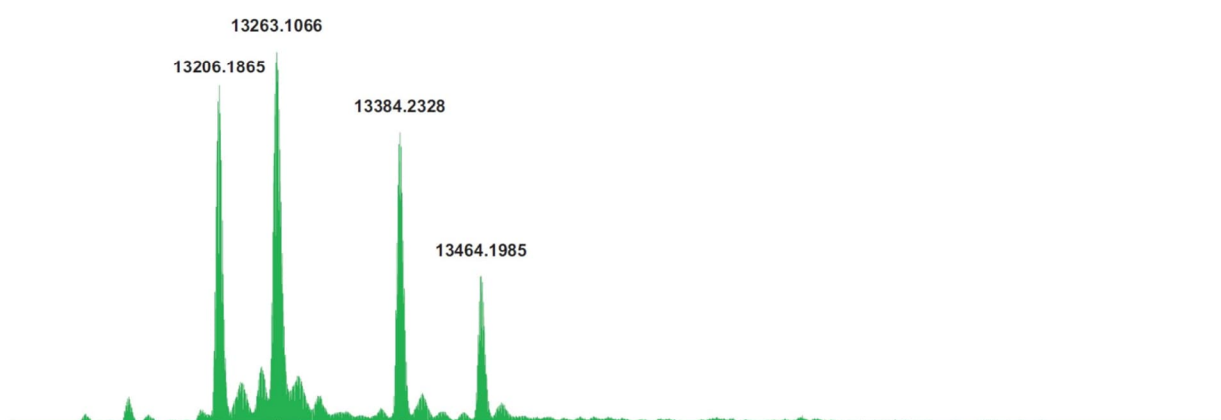
A SpyCatcher-PEED: Theoretical mass: 13695 Da

SpyCatcher-PEED.d: +MS, Deconvoluted



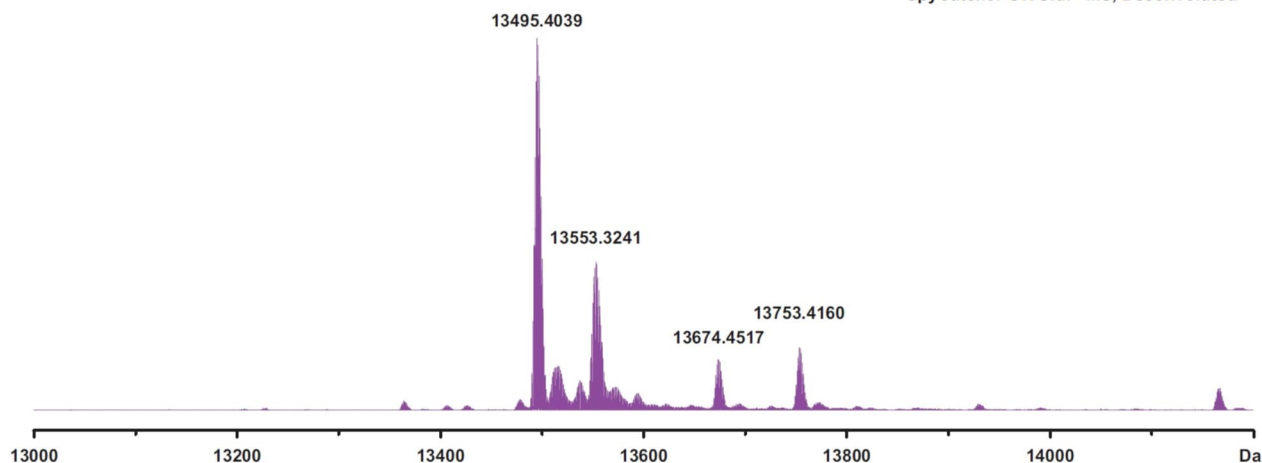
B SpyCatcher-DIIG: Theoretical mass: 13208 Da

SpyCatcher-DIIG.d: +MS, Deconvoluted

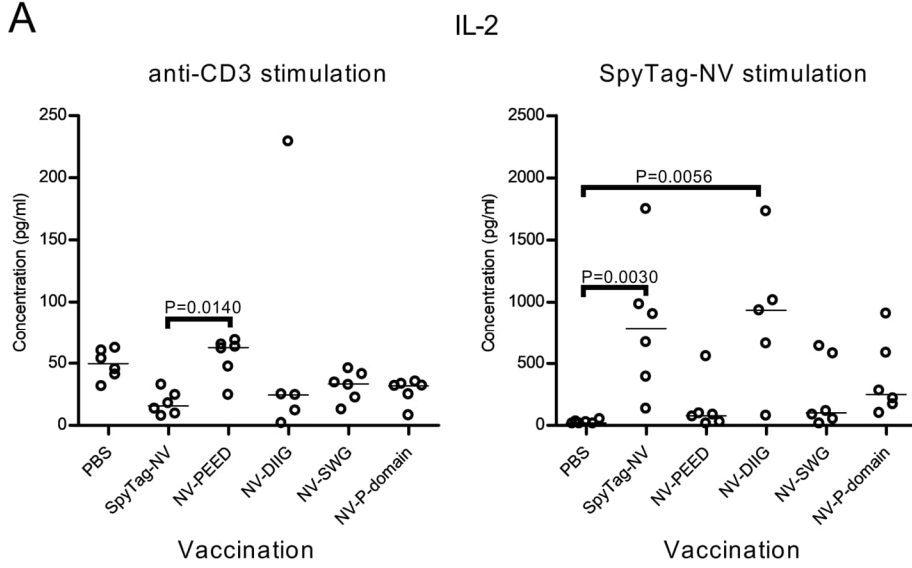
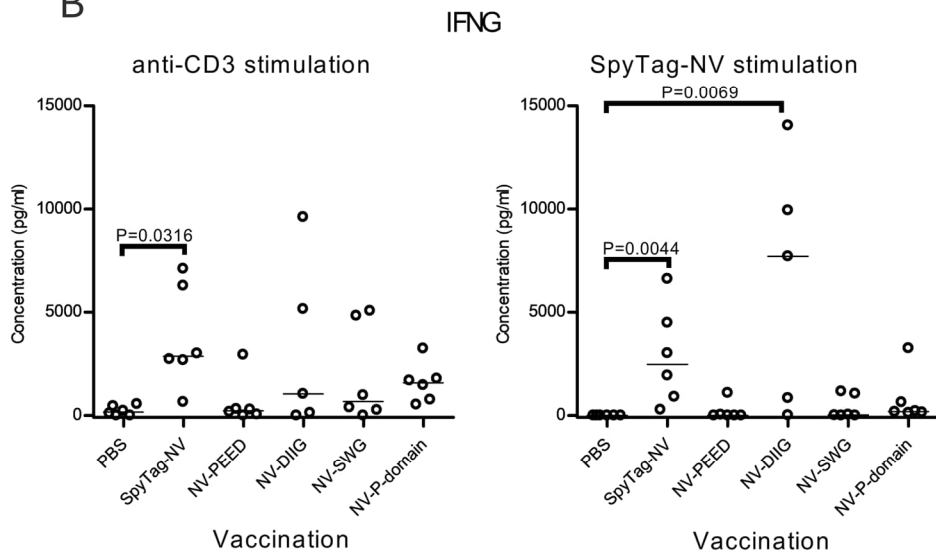
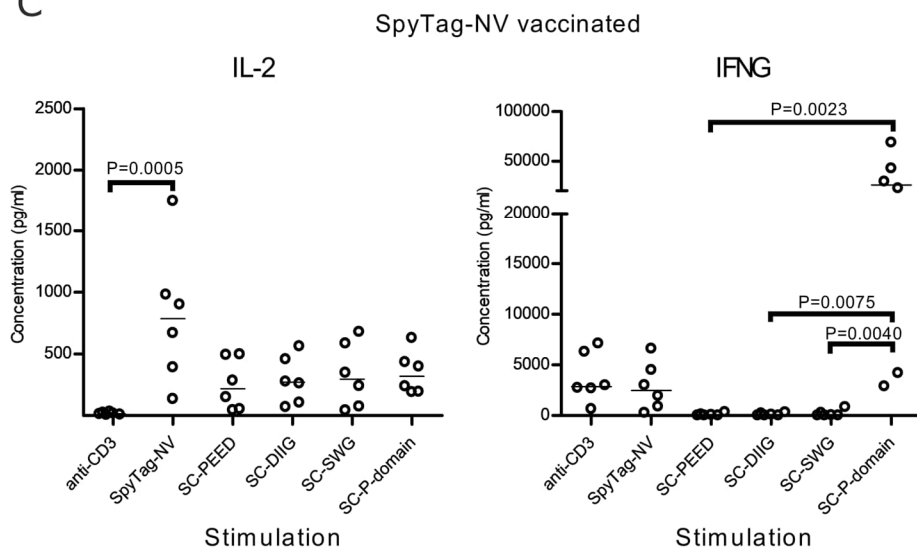


C SpyCatcher-SWG: Theoretical mass: 13497 Da

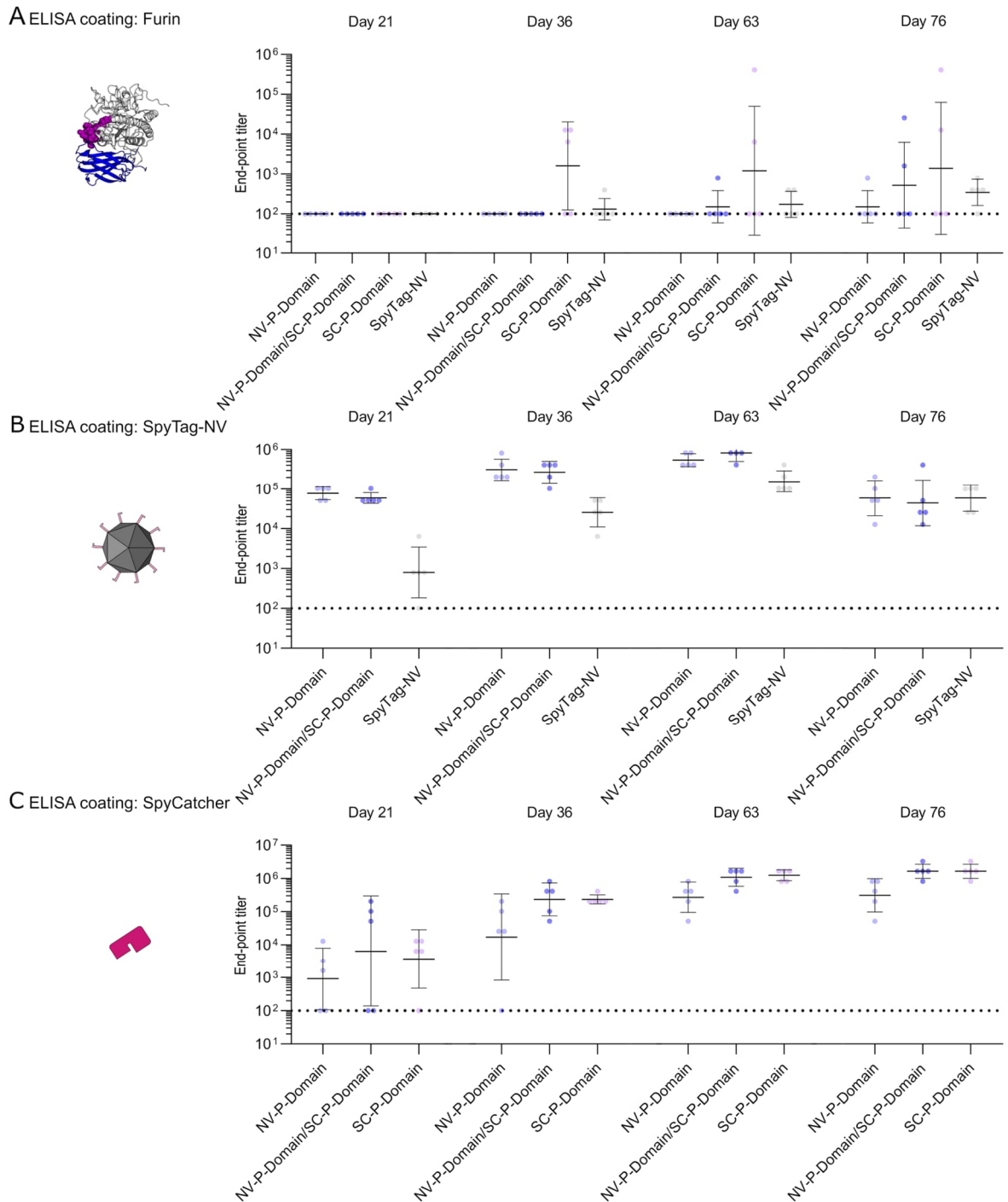
SpyCatcher-SWG.d: +MS, Deconvoluted



Supplementary figure 4. Deconvoluted masses of SpyCatcher-fused peptide antigens PEED (A) and DIIG (B) from PCSK9 and SWG (C) from furin were evaluated by matrix-assisted laser desorption/ionization mass spectrometry. The lowest measured masses correspond to the theoretical masses of the SpyCatcher-PEED, -DIIG and -SWG fusion proteins (13695, 13208 and 13497 Da, respectively). Note that the initiating methionine residues have been included in theoretical values.

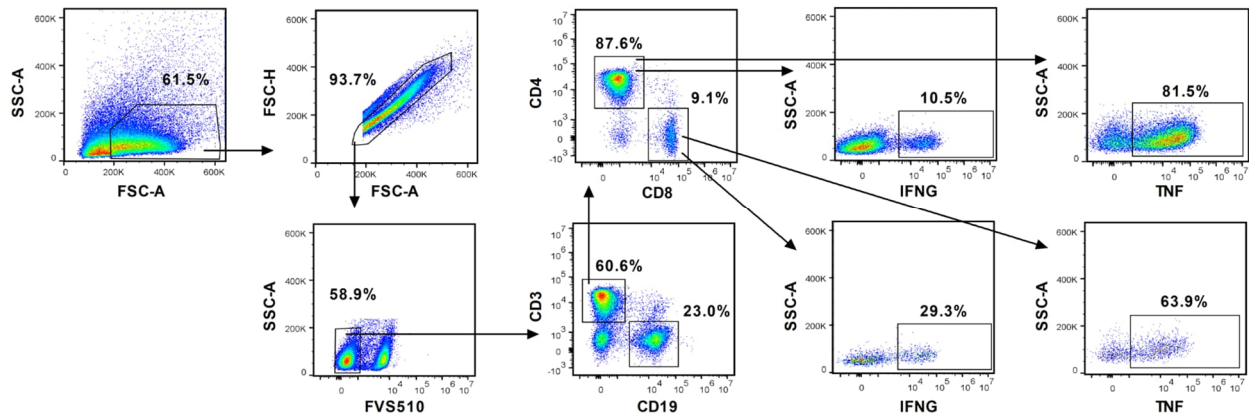
A**B****C**

Supplementary **figure 5**. T cell recall responses were evaluated from the culture media of *in vitro* stimulated splenocytes. Splenocytes were isolated from 13-week-old PBS, SpyTag-noro-VLP (NV) or antigen-conjugated noro-VLP immunized BALB/c mice (n=5-6, all females), and the cells were stimulated *in vitro* for 3 days at 37°C. The culture media was collected at 3 dps and the IL-2 and IFN- γ (IFNG) concentrations quantified using ELISA. Samples were stimulated using anti-CD3 antibody or SpyTag-NV for the evaluation of unspecific and SpyTag-NV mediated IL-2 (A) and IFN- γ (B) recall responses, respectively. In C), the antigen-specificity of the IL-2 and IFN- γ response is depicted for SpyTag-NV vaccinated mice. Scatter dot plot with median is shown for each group. Nonparametric Kruskal-Wallis one way ANOVA followed by Dunnet/Dunn's test was for the statistical evaluation of differences.

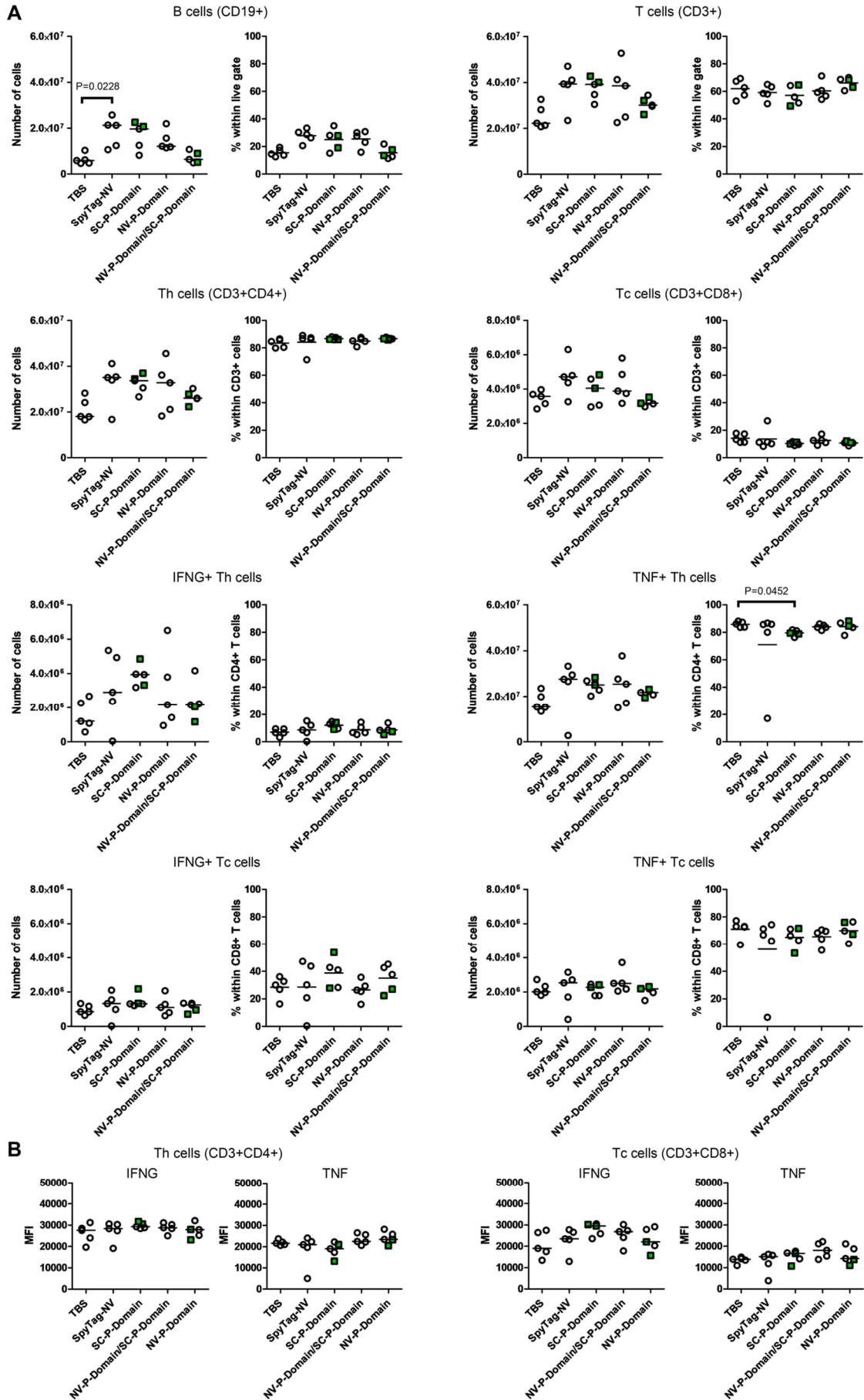


Supplementary **figure 6**. Adjuvanted vaccination with noro-VLP and SpyCatcher-fused furin P domain yields high anti-furin titers in individual mice. BALB/c mice (n=5, all females) were injected four times s.c. with alum adjuvanted P domain vaccine candidates, SpyTag-noro-VLP (NV), or TBS at days 0, 21, 42, and 63. Tail vein blood samples were collected at days 21, 36, 63, and on day 76 the animals were euthanized, and blood and spleen collected for analysis. Antibody titers were measured using ELISA. Plates were coated with the protein and the reciprocals of IgG antibody titers against recombinant furin (A), SpyTag-noro-VLP (B), and SpyCatcher (C) are shown. The end-point

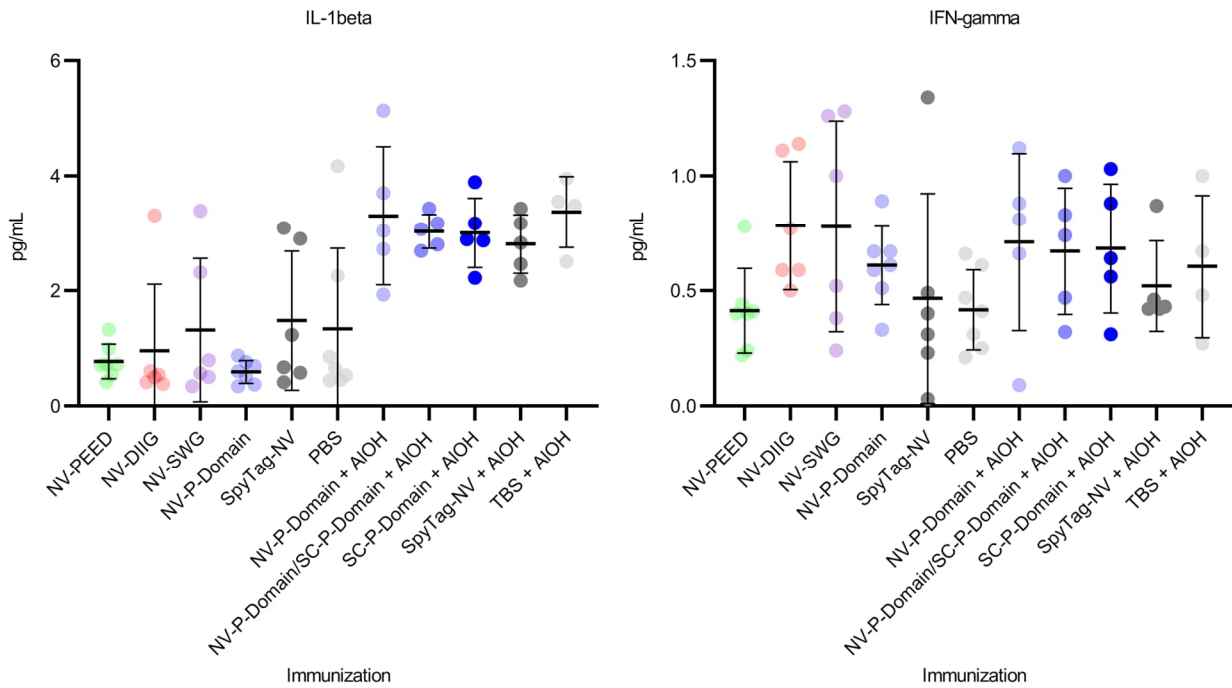
titer is the reciprocal of the lowest serum dilution that gives a significantly higher absorbance as compared to negative control (TBS) mouse group serum that is diluted 1:400. Mean titers \pm standard deviation are depicted. Each dot represents a serum sample from a single mouse. Undetectable antibody levels were denoted with the reciprocal titer 100 (vertical dashed line).



Supplementary figure 7. A representative gating strategy used for the flow cytometric analysis. Note that IFN- γ (IFNG) and TNF were analyzed separately from the samples. SSC = Side scatter, FSC = Forward scatter.



Supplementary figure 8. Complete dataset of lymphocyte populations and the IL-2 and IFN- γ producing T cells from the adjuvanted vaccination experiment. Splenocytes were isolated from 4-month-old BALB/c mice (n=5, all females) immunized with TBS, unconjugated SpyTag-noro-VLP (NV) or NV-furin-P domain, and the cells were analyzed using flow cytometry. A) The total cell numbers and the relative frequencies of the splenic lymphocyte populations were determined for each mouse. Target populations were identified based on their expression of CD19, CD3, CD4, CD8, IFN- γ (IFNG) and TNF. B) The mean fluorescence intensity (MFI) of IFN- γ ⁺ and TNF⁺ helper (Th) and cytotoxic (Tc) T cells. Nonparametric Kruskal-Wallis one way ANOVA followed by Dunnet/Dunn's test was used for the statistical evaluation of differences.



Supplementary figure 9. Mouse serum IL-1 β and IFN- γ concentrations were measured with the Meso Scale Discovery multiplex assay kit. The first six vaccination groups on the left-hand side of the graphs are from the first immunization experiment and the last five are the adjuvanted groups from the second immunization experiment. Only differences between groups were observed in IL-1 β levels, while IFN- γ serves as a representative example of the rest of the data. All data is available in Supplementary data 1. More specific information about the groups and doses is listed in Table 1.

Supplementary data 1 in **Additional file 2**. Mouse serum IFN- γ , IL-10, IL-12p70, IL-1 β , IL-2, IL-4, IL-5, IL-6, KC/gro, and TNF concentrations were measured with the Meso Scale Discovery multiplex assay kit. The sample origins, mouse groups and their abbreviations are shown in tab "Index". Mice encoded with "A" are from the first, unadjuvanted immunization experiment and the ones encoded "B" are from the second immunization experiment. The rest of the data sheets show the raw data from the measurement for each mouse, with the calculated concentration of the cytokine specified in the sheet title. Concentration calculation is based on the raw signal and a standard curve, according to the kit's instructions. 10 out of 64 serum samples were measured as duplicates, and for these, the standard deviation of concentration and standard deviation divided by mean concentration is shown in the tables. Concentrations below the fit curve range are calculated in the table, but calculated negative concentrations are shown as "NaN", instead. More specific information about the groups and doses is listed in Table 1.