

Patient-derived anti- β 2GP1 antibodies recognize a peptide motif pattern and not a specific sequence of residues

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Table S1

Accession Number	Protein Name	Gene Name
<u>O60603</u>	Toll-like receptor 2	TLR2
<u>O75581</u>	Low-density lipoprotein receptor-related protein 6	LRP6
<u>O94813</u>	Slit homolog 2 protein	SLIT2
<u>P02749</u>	Beta-2-glycoprotein 1	APOH
<u>P04275</u>	von Willebrand factor	VWF
<u>P10643</u>	Complement component C7	C7
<u>P98155</u>	Very low-density lipoprotein receptor	VLDLR
<u>P98160</u>	Basement membrane-specific heparan sulfate proteoglycan core protein	HSPG2
<u>P98164</u>	Low-density lipoprotein receptor-related protein 2	LRP2
<u>Q07954</u>	Pro-low-density lipoprotein receptor-related protein 1	LRP1
<u>Q13705</u>	Activin receptor type-2B	ACVR2B
<u>Q14114</u>	Low-density lipoprotein receptor-related protein 8	LRP8
<u>Q9BXB4</u>	Oxysterol-binding protein-related protein 11	OSBPL11
<u>Q9BZF1</u>	Oxysterol-binding protein-related protein 8	OSBPL8
<u>Q9H244</u>	P2Y purinoceptor 12	P2RY12
<u>Q9NPF0</u>	CD320 antigen	CD320

Supplemental methods

Cell culture

Monocytes were isolated from blood buffy coats of healthy volunteers as previously described.^{1,2} Monocyte purity routinely consisted of >90% CD14⁺ cells, <1% CD3⁺ cells, and <1% CD19⁺ cells as assessed by flow cytometry. Cells were cultured in RPMI containing 10% Fetal Bovine Serum (FBS; Gibco BRL-Life Technologies). Each experiment was performed with at least three different preparations of monocytes.

HLA Typing

Patients were matched at the second field level typing (high-resolution typing, previously referred to as 4-digit typing) for the HLA-DRB1/B3, DQB1 and DPB1 loci by standard methods: PCR-SSO on microbeads arrays (Luminex Technology, LabType HD, OneLambda, Ingen, Chilly-Mazarin, France), PCR-SSP (Genovision, Milan Analytika AG, Rheinfelden, Switzerland) and SBT (Protrans, Endotell AG, Allschwil, Switzerland).

IgG purification

IgG fractions were isolated from patients plasma with Protein-G CL-4B Sepharose (GE Healthcare) as previously described.³ To assay for endotoxin and lipopeptide contamination of the IgG fractions, we depleted IgG from the aPLA and IgG-ctl (control) fractions by one step of affinity adsorption to Protein G-Sepharose and tested the remaining supernatant for its capacity for monocyte activation, as previously described⁴. In addition, endotoxin levels were measured by the Limulus Amebocyte Lysate Endochrome Assay (Charles River Laboratories), and were found to be below the detection limit (0.25 EU/mL) for all IgG fractions at the concentration used in the assays. Each experiment was performed with at least 2 different preparations of IgG.

Tetramer staining of class II-peptide

Staining was performed following the Benaroya research institute's protocol (<https://tetramer.benaroyaresearch.org/more-inforesources/protocols>). Briefly, we performed antigen specific amplification with domains I-II, *i.e* cells were cultured in the presence of antigens (10µg/ml) for 10 days (pulse), prior to tetramer staining as described below. 10 µg/mL of PE-labeled Class II tetramer were added to the cells for 3 hours at 37 °C in the dark. Subsequently, cells were stained with fluorochrome (AF647)-labeled anti-CD4 for 30 minutes on ice before flow cytometry analysis.

TNF production

The cells were treated with recombinant domains or peptides of β_2 GP1 at 10 µg/ml prior to incubation with aPLA or control IgG. Supernatants were collected for TNF quantification by ELISA (R&D System).

Western blot

Total cell lysates were prepared and subjected to Western blot analysis as described previously.³ The blots were probed with anti-cMyc (Zymed). Secondary antibodies conjugated to IR800CW (Rockland) were used. Antibody-bound proteins were detected and quantified by the Odyssey system (Li-Cor).

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