Supporting Information

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Fig. 51. Anti-LILRB1 antibody and HLA-G homodimer induce similar proinflammatory cytokine gene profiles in decidual macrophages. (*A*) Decidual CD14⁺ lymphocytes were cultured for 5 h in the presence of anti-LILRB1 antibody (10 μ g/mL) and goat anti-mouse IgG secondary antibody (5 μ g/mL) (*Left*) or in a 1:1 ratio with untransfected 721.221 cells (None), 721.221 cells transfected with HLA-G monomer, or 721.221 cells transfected with HLA-G homodimer (*Right*). Total RNA was isolated and biotinylated cRNAs were used as probes. Hybridization was performed to a common cytokine Superarray membrane as described in *Materials and Methods*. The arrays were exposed to X-ray film, and 1 representative of 4 images is shown. (*B*) Expression was normalized to GAPDH. The average of 4 experiments is shown. Decidual CD14⁺ lymphocytes produce a proinflammatory cytokine profile upon (*Upper*) anti-LILRB1 stimulation (black bars) and isotype controls (gray bars), or (*Lower*) coculture with 721.221 (white bars), 721.221 cells transfected with HLA-G homodimer (OHS-021). For more information about gene names please visit http://www.sabiosciences.com/genetable.php?pcatn = ohs-021.

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RPS27A

AREG

SLURP1

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1	2	3	4	5	6	7	8
BMP3	BMP4	BMP5	BMP6	BMP7	BMP8B	CSF1	CSF2
9	10	11	12	13	14	15	16
CSF3	CYFIP2	LEFTY2	FAM3B	FAM3C	FGF10	FIGF	FLT3LG
17	18	19	20	21	22	23	24
LASS1	GDF10	GDF11	GDF15	GDF2	GDF3	GDF5	MSTN
25	26	27	28	29	30	31	32
GDF9	IFNA1	IFNA13	IFNA14	IFNA2	IFNA4	IFNA5	IFNA8
33	34	35	36	37	38	39	40
IFNB1	IFNG	IFNK	IFNW1	IK	IL10	IL11	IL12A
41	42	43	44	45	46	47	48
IL12B	IL13	TXLNA	IL15	IL16	IL17A	IL17B	IL17C
49	50	51	52	53	54	55	56
IL25	IL17F	IL18	IL19	IL1A	IL1B	IL1F10	IL1F5
57	58	59	60	61	62	63	64
IL1F6	IL1F7	IL1F8	IL1F9	IL2	IL20	IL21	IL22
65	66	67	68	69	70	71	72
IL24	IL26	IL3	IL4	IL5	IL6	IL7	IL8
73	74	75	76	77	78	79	80
IL9	INHA	INHBA	INHBB	LTA	LTB	MDK	NODAL
81	82	83	84	85	86	87	88
PDGFA	PDGFB	PTN	SPP1	TGFA	TGFB1	TGFB2	TGFB3
89	90	91	92	93	94	95	96
THPO	TNF	TNFRSF11B	TNFSF10	TNFSF11	TNFSF12	TNFSF13	TNFSF13B
97	98	99	100	101	102	103	104
TNFSF14	TNFSF15	TNFSF18	TNFSF4	CD40LG	FASLG	CD70	TNFSF8
105	106	107	108	109	110	111	112
TNFSF9	VEGFA	PUC18	VEGFB	Blank	AS1R2	AS1R1	AS1
113	114	115	116	117	118	119	120
GAPDH	B2M	HSP90AB1	HSP90AB1	ACTB	ACTB	BAS2C	BAS2C
121	122	123	124	125	126	127	128

ATP6AP1

BMP1

BMP10

BMP15

BMP2

Fig. S1. Continued.



Fig. 52. Anti-LILRB1 antibody and HLA-G homodimer induce similar proinflammatory cytokine profiles in CD14⁺ peripheral blood monocytes. (A) Peripheral CD14⁺ lymphocytes were cultured for 5 h in the presence of anti-LILRB1 antibody (10 µg/mL) and goat anti-mouse IgG secondary antibody (5 µg/mL) (*Left*) or in a 1:1 ratio with untransfected 721.221 cells (None), 721.221 cells transfected with HLA-G monomer, or 721.221 cells transfected with HLA-G homodimer (*Right*). Total RNA was isolated, and biotinylated cRNAs were used as probes. Hybridization was performed to a common cytokine Superarray membrane as described in *Materials and Methods*. The arrays were exposed to X-ray film, and 1 representative of 3 images is shown. (B) Expression was normalized to GAPPH. The average of 3 experiments is shown. Peripheral CD14⁺ lymphocytes produce a proinflammatory cytokine profile upon (*Upper*) anti-LILRB1 stimulation (black bars) and isotype controls (gray bars), or (*Lower*) coculture with 721.221 (white bars), 721.221 cells transfected with HLA-G homodimer (gray bars), or 721.221 cells transfected with HLA-G homodimer (black bars). *, *P* < 0.05; **, *P* < 0.01. Note the scale in *B Upper* is compressed 2-fold relative to that in Fig. S1B *Upper* because the mRNA levels for IL-8 in peripheral CD14⁺ cells.



Fig. S3. Anti-KIR2DL4 antibody and HLA-G homodimer up-regulate proinflammatory cytokine genes on decidual CD56⁺ lymphocytes. (A) Decidual CD56⁺ lymphocytes were cultured for 5 h in the presence of anti-KIR2DL4 antibody (10 μ g/mL) and goat anti-mouse IgG secondary antibody (5 μ g/mL) (*Left*) or in a 1:1 ratio with untransfected 721.221 cells (None), 721.221 cells transfected with HLA-G monomer, or 721.221 cells transfected with HLA-G homodimer (*Right*). Total RNA was isolated, and biotinylated cRNAs were used as probes. Hybridization was performed to a common cytokine Superarray membrane as described in *Materials and Methods*. The arrays were exposed to X-ray film and 1 representative of 4 images is shown. (*B*) Expression was normalized to GAPDH. The average of 4 experiments is shown. Decidual CD56⁺ lymphocytes produce a proinflammatory cytokine profile upon (*Upper*) anti-KIR2DL4 stimulation (black bars) and isotype controls (gray bars), or (*Lower*) coculture with 721.221 (white bars), 721.221 cells transfected with HLA-G monomer (gray bars), or 721.221 cells transfected with HLA-G homodimer (black bars).*, *P* < 0.05.

Table S1. Genes constitutively expressed in various cell types

PNAS PNAS

Gene Name	Peripheral	Decidual	Decidual
(Position)	CD14	CD14	CD56" NK cells
0052 (47)	wonocytes	Macrophages	
$\frac{\text{CSF3}(17)}{\text{OVEIDO}(40)}$		+	+
		+	+
BMP8B (14)		+/-	- /
LASS1 (25)			+/-
GDF3 (30)			+/-
FAM3C (21)		+	+
FGF10 (22)		+	
FLT3LG (24)		+/-	+
IL12A (48)		+/-	+
IL10 (46)			+
IFNB (41)			+/-
IFNG (42)			++
IL13 (50)	+/-		+/-
TXLNA (51)		++	+
IL15 (52)			+/-
IL17B (55)			+
IL17C (56)			+/-
IL18 (59)	+		
IL19 (60)			+/-
IL1A (61)		+/-	
IL1B (62)		+++	++
IL20 (70)	+		
IL26 (74)		+/-	++
IL6 (78)		+	+
IL7 (79)			+/-
IL8 (80)	+	++	
LTA (85)	++	++	+++
LTB (86)			+
PDGFB (90)	+	+/-	++
SPP1 (92)		+	
THPO (97)			+/-
TNF (98)	++	+++	++
TNFSF4 (108)			+/-
CD70 (111)	+	+	+
VEGFB (116)	++	++	+++

This chart was made by evaluating the intensity of the hybridization spots in the Superarray in Figs. S1-S3, which were obtained after incubation with untransfected 721.221 cells; this is a qualitative estimate. Gene position refers to the number shown in Fig. 1C. Differences between decidual CD14⁺ and CD56⁺ NK cells are highlighted. Scores are given as +++ for the highest intensity observed, whereas +/- indicates the lowest intensity observed.