Li et al. "Chemically Modified Human Serum Albumin Potently Blocks Entry of Ebola Pseudoviruses and Virus-like Particles."

Supplementary Materials:

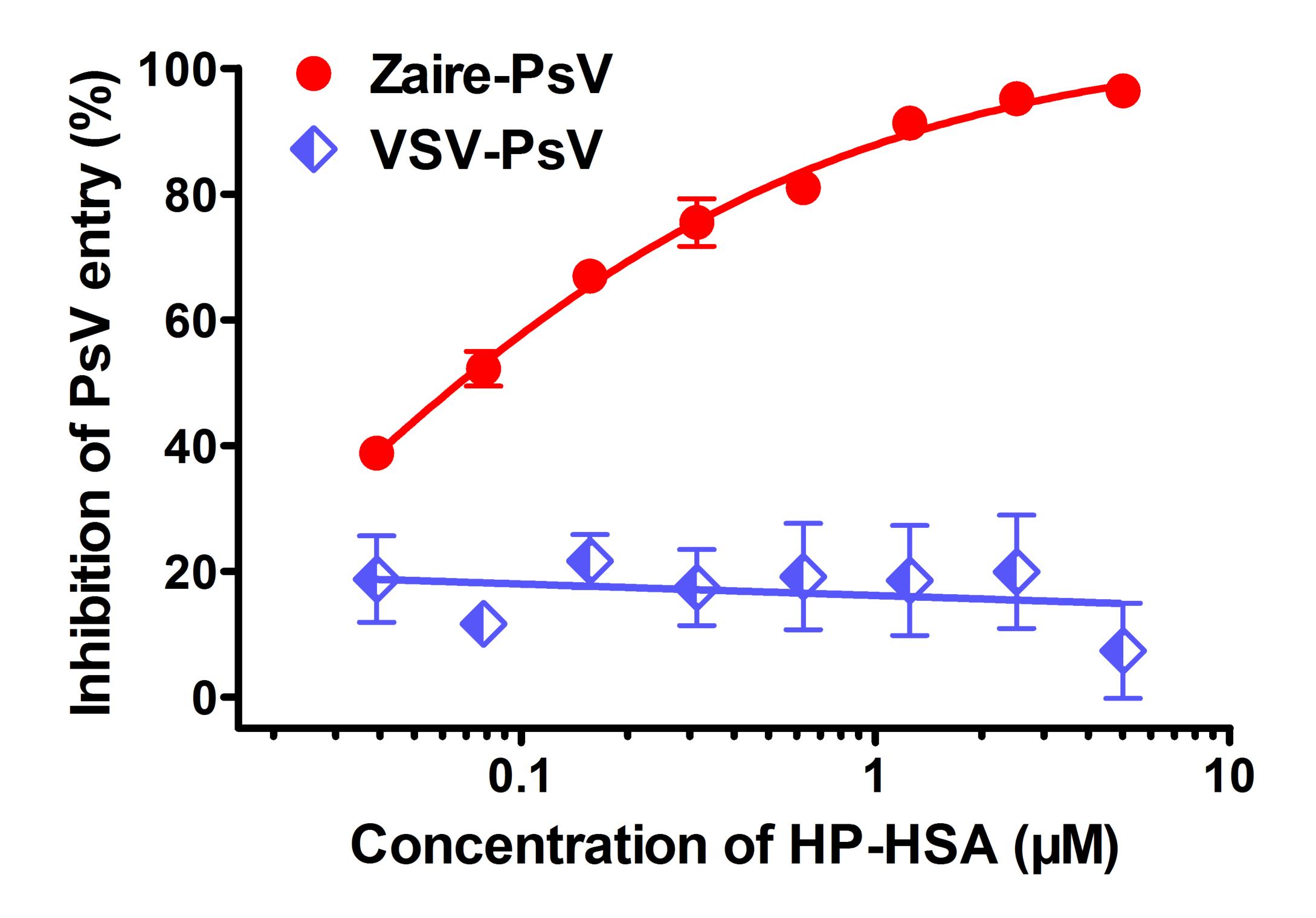


FIG S1 Comparison of entry inhibitory activity of HP-HSA on Zaire pseudovirus and VSV pseudovirus. HuH-7 cells were served as target cell. The experiments were performed in triplicate, and the error bars indicate standard deviations.

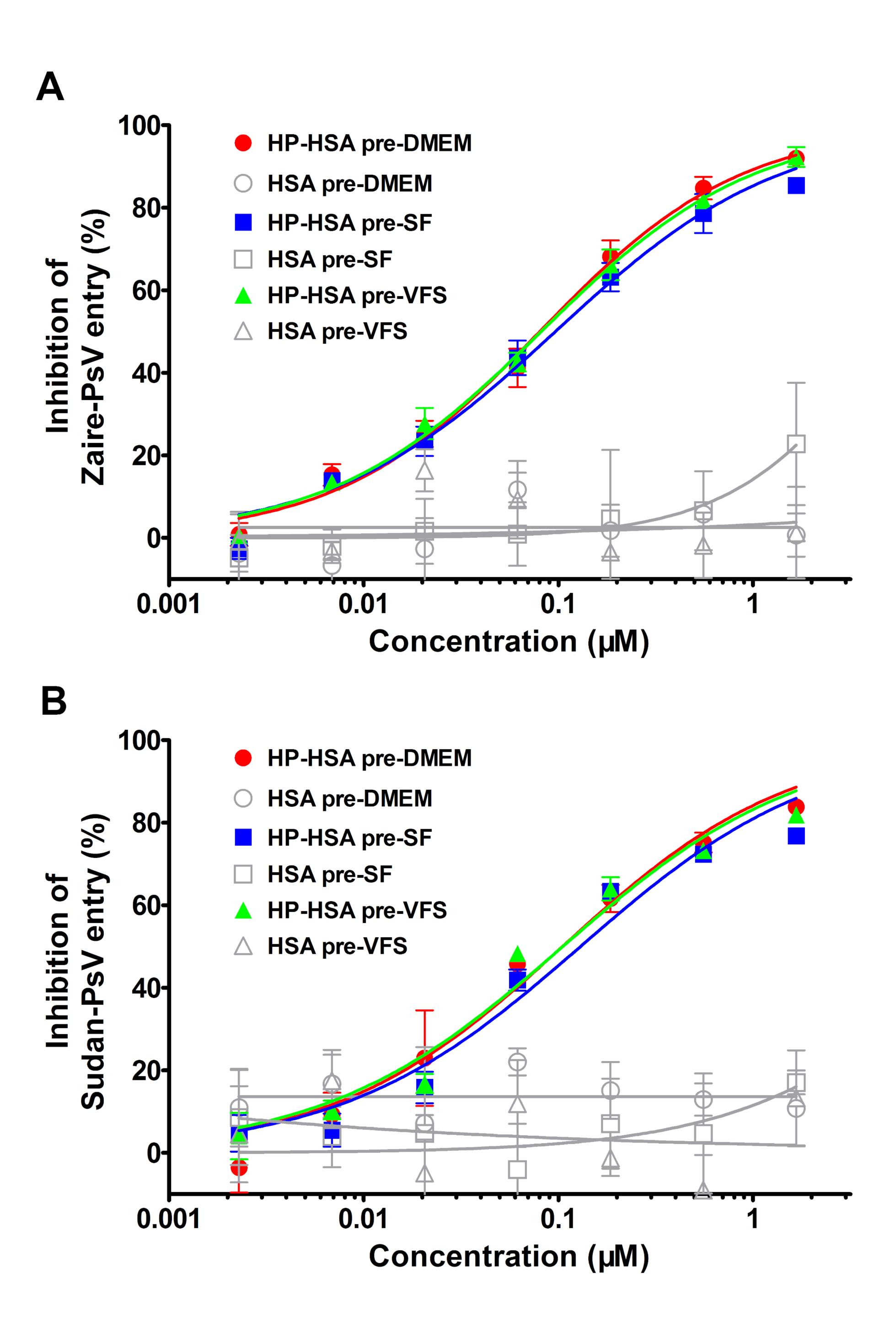


FIG S2 The entry inhibitory activity of HP-HSA in human seminal fluid and VFS against EBOV-PsVs. HP-HSA and HSA were preincubated with DMEM, human seminal fluid (SF) or vaginal fluid simulant (VFS), respectively, before gradient dilution ("pre-" means "preincubated with"). The inhibitory activities of the premixed samples was evaluated with Zaire-PsV (A) and Sudan-PsV (B). All the experiments were performed in triplicate, and the error bars indicate standard deviations.

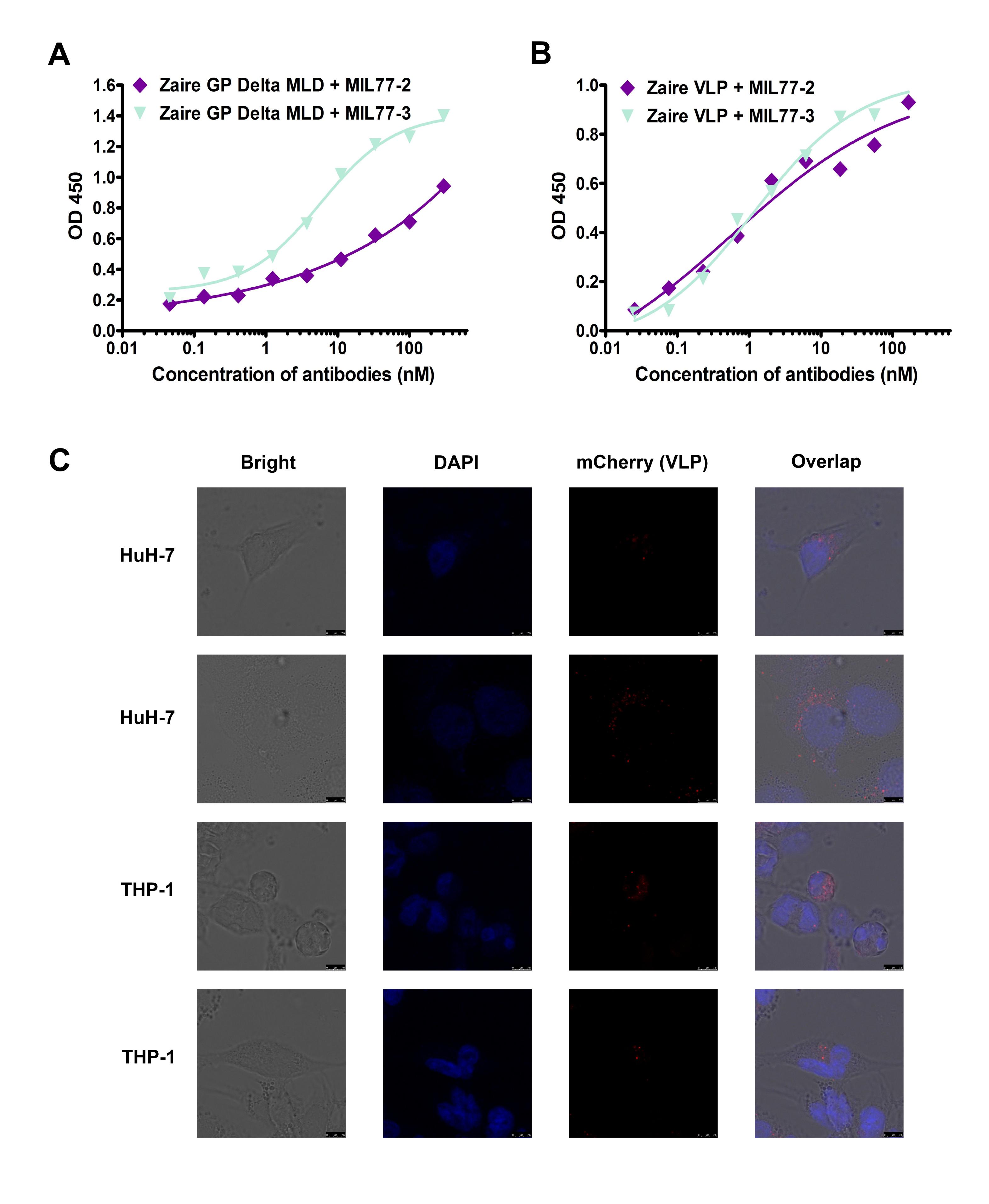


FIG S3 The purified soluble Zaire GP (with a deletion of 314 - 462 aa and 637 - 676 aa) (A) and concentrated VLPs (B) were recognized by antibodies MIL77-2 and MIL77-3. The images of cells infected with mCherry-labeled Zaire-VLPs were captured by laser scanning confocal microscope and displayed as single channel images or merged images (C).

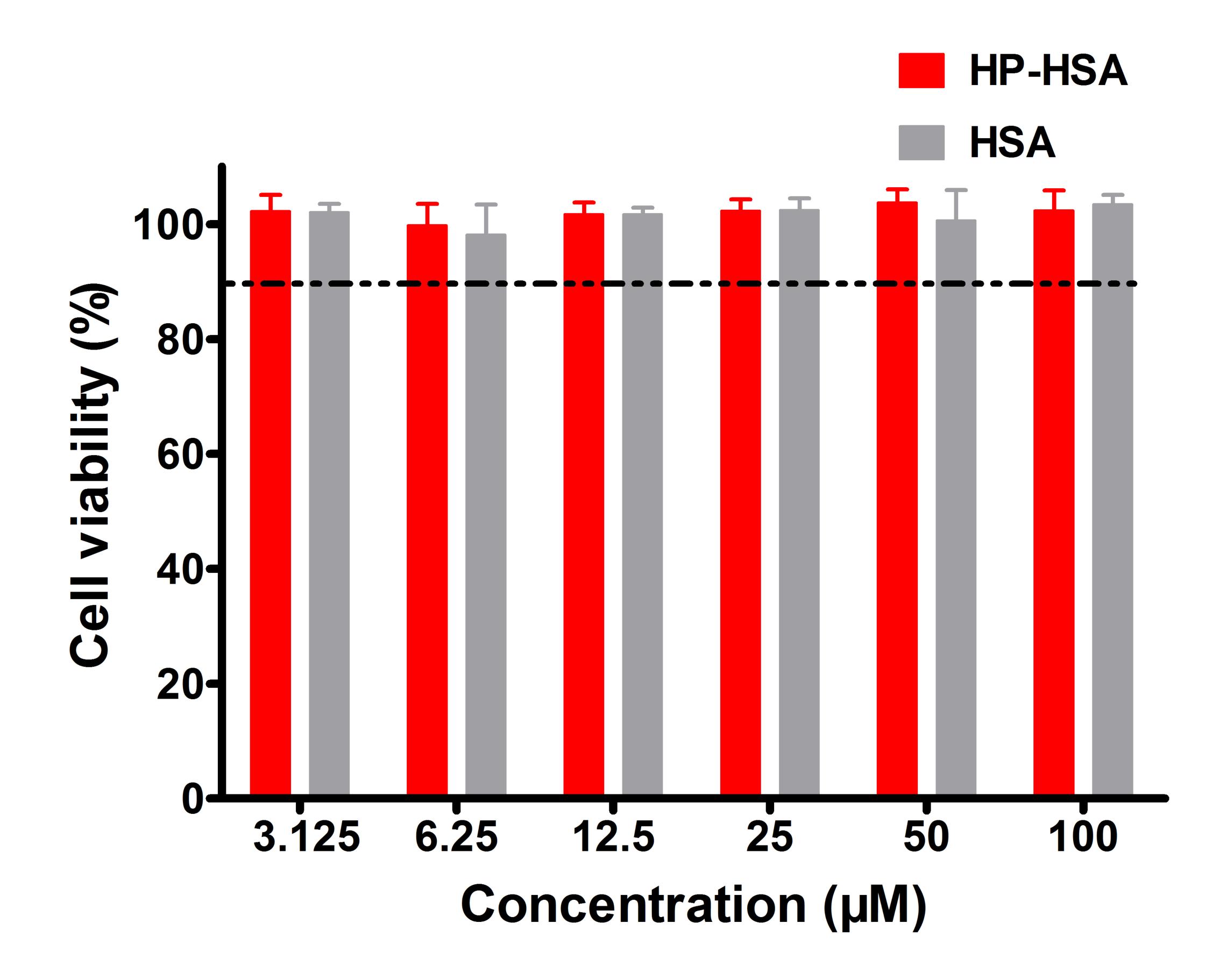


FIG S4 The cytotoxicity of HP-HSA to differentiated human THP-1 cells was measured by CCK-8 cytotoxicity assay. The unmodified HSA was served as control. All the experiments were performed in triplicate, and the error bars indicate standard deviations.

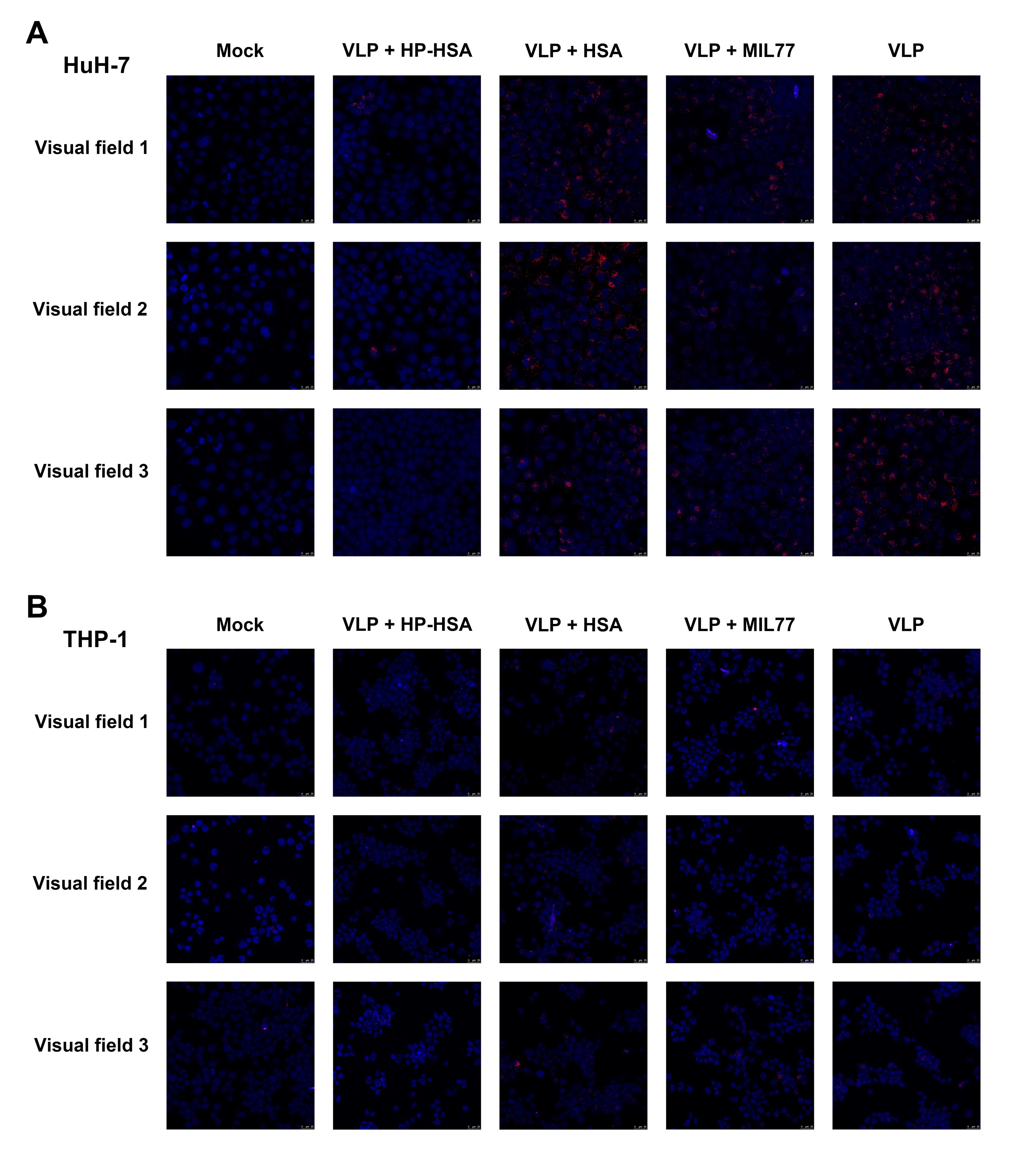


FIG S5 The raw images (merged by DAPI channel and mCherry channel) applied for the analysis of Figure 6A. The visual fields of different samples were arranged in (A) for HuH-7 cell and (B) for differentiated THP-1 cell.

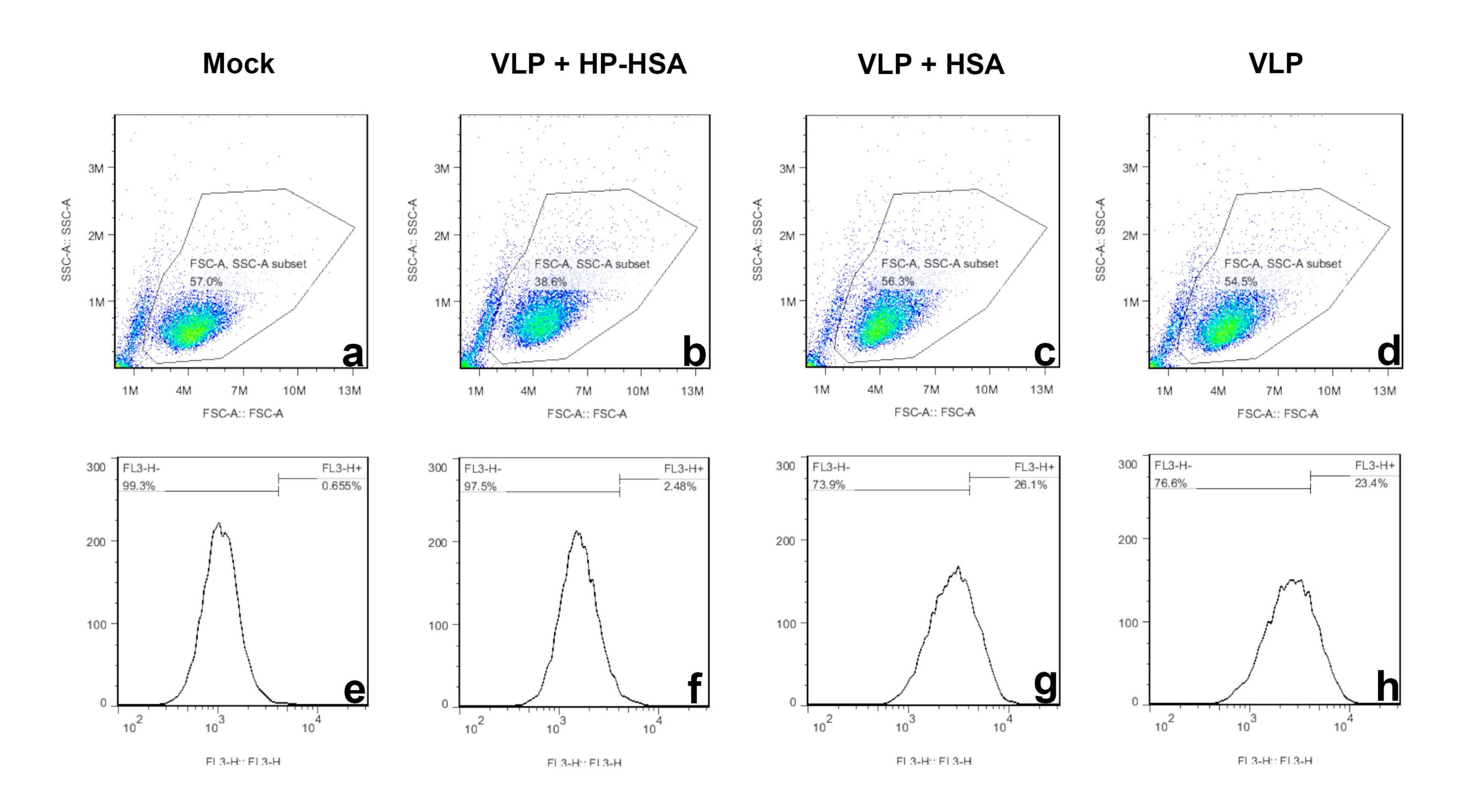


FIG S6 The raw data of flow cytometry for the Figure 6B. The intact cells from different samples gated in (a-d) were further used to count the number of mCherry positive cells (e-h).

DATASET.S1 The raw data of fluorescent images applied for Fig. 6A-a (HuH-7 cells) output by Image J.

Sample	Visual Field	Area	IntDen	MinThr	MaxThr	CellNum	Area/CellNum	IntDen/CellNum
Mock	1	0	N/A	0	200	151	0.00	N/A
	2	0	N/A	0	200	97	0.00	N/A
	3	0	N/A	0	200	113	0.00	N/A
VLP + HP-HSA	1	1687	256189	0	200	147	11.48	1742.78
	2	2441	354513	0	200	156	15.65	2272.52
	3	291	43434	0	200	231	1.26	188.03
VLP + HSA	1	19247	2920827	0	200	179	107.53	16317.47
	2	24550	3404812	0	200	125	196.40	27238.50
	3	12236	1882591	0	200	143	85.57	13164.97
VLP + MIL77	1	13352	2079565	0	200	169	79.01	12305.12
	2	4669	758151	0	200	167	27.96	4539.83
	3	9848	1502177	0	200	187	52.66	8033.03
VLP	1	17053	2624626	0	200	219	77.87	11984.59
	2	19930	2983948	0	200	256	77.85	11656.05
	3	25642	3834110	0	200	148	173.26	25906.15
Area: the fluorescend	ce positve area ir	n each image						
IntDen: Intergated de	ensity							
MinThr: minimum th	reshold							
MaxThr: maximum t								
CellNum: the cell nu	mbers of each in	nage						

	2	367	62821	0	200	100	3.67	628.21
	3	679	97894	0	200	249	2.73	393.15
VLP + HP-HSA	1	637	96462	0	200	255	2.50	378.28
	2	318	49723	0	200	193	1.65	257.63
	3	329	54737	0	200	150	2.19	364.91
VLP + HSA	1	1330	192409	0	200	158	8.42	1217.78
	2	2083	335501	0	200	233	8.94	1439.92
	3	1840	261216	0	200	176	10.45	1484.18
VLP + MIL77	1	2111	304144	0	200	189	11.17	1609.23
	2	1271	173336	0	200	170	7.48	1019.62
	3	1277	196571	0	200	152	8.40	1293.23
VLP	1	1279	193118	0	200	184	6.95	1049.55
	2	1190	181284	0	200	132	9.02	1373.36
	3	1122	176951	0	200	152	7.38	1164.15
Area: the fluorescence	positve area i	n each image						
IntDen: Intergated den	sity							
MinThr: minimum thre	eshold							
MaxThr: maximum thr	reshold							
CellNum: the cell num	bers of each in	mage						

75

35

473

463

642

290

262

308

VLP + HSA

VLP

IntDen: Intergated density

MinThr: minimum threshold

MaxThr: maximum threshold

Area: the fluorescence positve area in each image

CellNum: the cell numbers of each image

11889

65585

59640

88770

41041

33548

44502

5406

	3	1277	196571	0	200	152	8.40	1293.23
VLP	1	1279	193118	0	200	184	6.95	1049.55
	2	1190	181284	0	200	132	9.02	1373.36
	3	1122	176951	0	200	152	7.38	1164.15
Area: the fluorescen	ce positve area ir	each image						
IntDen: Intergated de	ensity							
MinThr: minimum th	nreshold							
MaxThr: maximum	threshold							
CellNum: the cell nu	ımbers of each in	nage						
DATASET.S3 The				,	-	<u> </u>		
Sample	Visual Field	Area	IntDen	MinThr	MaxThr	CellNum	Area/CellNum	
Mock		7 11 Ou			WIGATIII	Ocinitanii	/ ti da/ ddiii taiii	IntDen/CellNum
IVIOCK	1	8	1543	0	200	116	0.07	13.30
IVIOCK	1 2			0				
IVIOCK	1 2 3	8	1543	0 0 0	200	116	0.07	13.30

200

200

200

200

200

200

200

200

0

105

113

103

110

97

132

99

91

113.23

47.84

636.75

542.18

915.15

310.92

338.87

489.03

0.71

0.31

4.59

4.21

6.62

2.20

2.65

3.38

Mock	1	73	12132	0	200	114	0.64	106.42
	2	367	62821	0	200	100	3.67	628.21
	3	679	97894	0	200	249	2.73	393.15
VLP + HP-HSA	1	637	96462	0	200	255	2.50	378.28
	2	318	49723	0	200	193	1.65	257.63
	3	329	54737	0	200	150	2.19	364.91
VLP + HSA	1	1330	192409	0	200	158	8.42	1217.78
	2	2083	335501	0	200	233	8.94	1439.92
	3	1840	261216	0	200	176	10.45	1484.18
VLP + MIL77	1	2111	304144	0	200	189	11.17	1609.23
	2	1271	173336	0	200	170	7.48	1019.62
	3	1277	196571	0	200	152	8.40	1293.23
VIP	1	1270	103118	0	200	18/	6.05	10/0 55

IVIIII I I I I I I I I I I I I I I I I	116311010	,						
MaxThr: maximum t	threshold							
CellNum: the cell nu	ımbers of each in	nage						
DATASET.S2 The	raw data of flu	orescent images	applied for F	ig. 6A-b (differentiat	ed THP-1 cells)	output by Image J.	
		_				-		
Sample	Visual Field	Area	IntDen	MinThr	MaxThr	CellNum	Area/CellNum	IntDen/CellNum
Mock	1	73	12132	0	200	114	0.64	106.42
	2	367	62821	0	200	100	3.67	628.21
	3	679	97894	0	200	249	2.73	393.15
VLP + HP-HSA	1	637	96462	0	200	255	2.50	378.28
	2	318	49723	0	200	193	1.65	257.63
	3	329	54737	0	200	150	2.19	364.91
VLP + HSA	1	1330	192409	0	200	158	8.42	1217.78
	2	2083	335501	\cap	200	233	2 Q <i>1</i>	1/30 02

		Z44 I	334313	U	200	130	13.03	2212.32
	3	291	43434	0	200	231	1.26	188.03
VLP + HSA	1	19247	2920827	0	200	179	107.53	16317.47
	2	24550	3404812	0	200	125	196.40	27238.50
	3	12236	1882591	0	200	143	85.57	13164.97
VLP + MIL77	1	13352	2079565	0	200	169	79.01	12305.12
	2	4669	758151	0	200	167	27.96	4539.83
	3	9848	1502177	0	200	187	52.66	8033.03
VLP	1	17053	2624626	0	200	219	77.87	11984.59
	2	19930	2983948	0	200	256	77.85	11656.05
	3	25642	3834110	0	200	148	173.26	25906.15
Area: the fluorescence	e positve area	in each image						
IntDen: Intergated der	nsity							
MinThr: minimum thr	eshold							
MaxThr: maximum th	reshold							
CellNum: the cell num	nbers of each	image						
					. 4 . 4			
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	3	0	N/A	0	200	113	0.00	N/A
VLP + HP-HSA	1	1687	256189	0	200	147	11.48	1742.78
	2	2441	354513	0	200	156	15.65	2272.52
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VLP + MIL77	1	13352	2079565	0	200	169	79.01	12305.12
	2	4669	758151	0	200	167	27.96	4539.83
	3	9848	1502177	0	200	187	52.66	8033.03
VLP	1	17053	2624626	0	200	219	77.87	11984.59
	2	19930	2983948	0	200	256	77.85	11656.05
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Area: the fluorescence	e positve area	in each image						
IntDen: Intergated der	nsity							
MinThr: minimum thre	eshold							
MaxThr: maximum th	reshold							
CellNum: the cell num	nbers of each	image						

	3	0	N/A	0	200	113	0.00	N/A
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	2	2441	354513	0	200	156	15.65	2272.52
	3	291	43434	0	200	231	1.26	188.03
VLP + HSA	1	19247	2920827	0	200	179	107.53	16317.47
	2	24550	3404812	0	200	125	196.40	27238.50
	3	12236	1882591	0	200	143	85.57	13164.97
VLP + MIL77	1	13352	2079565	0	200	169	79.01	12305.12
	2	4669	758151	0	200	167	27.96	4539.83
	3	9848	1502177	0	200	187	52.66	8033.03
VLP	1	17053	2624626	0	200	219	77.87	11984.59
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MaxThr: maximum th	reshold							
CellNum: the cell num	bers of each	image						