#### FEATURE ARTICLE



# Future perspective: high-throughput construction of new ultrasensitive cytokine and virion liquid chips for high-throughput screening (HTS) of anti-inflammatory drugs or clinical diagnosis and treatment of inflammatory diseases

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#### Abstract

Pathogen-host cell interactions play an important role in many human infectious and inflammatory diseases. Several pathogens, including Escherichia coli (E. coli), Mycobacterium tuberculosis (M. tb), and even the recent 2019 novel coronavirus (2019-nCoV), can cause serious breathing and brain disorders, tissue injury and inflammation, leading to high rates of mortality and resulting in great loss to human physical and mental health as well as the global economy. These infectious diseases exploit the microbial and host factors to induce serious inflammatory and immunological symptoms. Thus the development of anti-inflammatory drugs targeting bacterial/viral infection is an urgent need. In previous studies, YojI-IFNAR2, YojI-IL10RA, YojI-NRP1,YojI-SIGLEC7, and YojI-MC4R membrane-protein interactions were found to mediate E. coli invasion of the blood-brain barrier (BBB), which activated the downstream anti-inflammatory proteins NACHT, LRR and PYD domains-containing protein 2(NLRP2), using a proteomic chip conjugated with cell immunofluorescence labeling. However, the studies of pathogen (bacteria/virus)-host cell interactions mediated by membrane protein interactions did not extend their principles to broad biomedical applications such as 2019-nCoV infectious disease therapy. The first part of this feature article presents in-depth analysis of the cross-talk of cellular anti-inflammatory transduction signaling among interferon membrane protein receptor II (IFNAR2), interleukin-10 receptor subunit alpha (IL-10RA), NLRP2 and [Ca<sup>2+</sup>]-dependent phospholipase A2 (PLA2G5), based on experimental results and important published studies, which lays a theoretical foundation for the high-throughput construction of the cytokine and virion solution chip. The paper then moves on to the construction of the novel GPCR recombinant herpes virion chip and virion nano-oscillators for profiling membrane protein functions, which drove the idea of constructing the new recombinant virion and cytokine liquid chips for HTS of leading drugs. Due to the different structural properties of GPCR, IFNAR2, ACE2 and Spike of 2019-nCoV, their ligands will either bind the extracellular domain of IFNAR2/ACE2/Spike or the specific loops of the GPCR on the envelope of the recombinant herpes virions to induce dynamic charge distribution changes that lead to the variable electron transition for detection. Taken together, the combined overview of two of the most innovative and exciting developments in the immunoinflammatory field provides new insight into high-throughput construction of ultrasensitive cytokine and virion liquid chips for HTS of anti-inflammatory drugs or clinical diagnosis and treatment of inflammatory diseases including infectious diseases, acute or chronic inflammation (acute gouty arthritis or rheumatoid arthritis), cardiovascular disease, atheromatosis, diabetes, obesity, tissue injury and tumors. It has significant value in the prevention and treatment of these serious and painful diseases.

**Keywords** Bacterial/viral infection  $\cdot$  Signaling cross-talk  $\cdot$  Anti-inflammatory  $\cdot$  High-throughput construction  $\cdot$  Cytokine/virion liquid

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Abbreviation	S
2019-nCoV	2019 Novel coronavirus
(BBB)	Blood-brain barrier
BSA	Bovine serum albumin
CNF1	Cytotoxic necrotizing factor 1
DR-18	Decoy-resistant IL-18
E. coli	Escherichia coli
FACS	Fluorescence-activated cell sorting
G2A	G protein-coupled receptor 132
GlcNAc	N-acetylglucosamine
GPCR	G protein-coupled receptor
GRASP55	Golgi reassembly stacking protein 55
HBMEC	Human brain microvascular endothelial cells
HPLC-MS	High-performance liquid
	chromatography-mass spectrometry
HSV-1	Herpes simplex virus-1
HTS	High-throughput screening
IFNAR2	Interferon alpha/beta receptor subunit 2
IL	Interleukin
IL-10RA	Interleukin-10 receptor subunit alpha
IL-18 BP	IL-18 binding protein
ISGs	Interferon-stimulated genes
LPS	Lipopolysaccharide
M. tb	Mycobacterium tuberculosis
MS	Mass spectrometry
m/z	Mass/charge
Ni	Nickel
NK	Natural killer
NLRP2	NACHT, LRR and PYD
	domains-containing protein 2
OmpA	Outer-membrane protein A
ORF	Open reading frame
OS	Oscillatory shear force
PEG	Polyethylene glycol
PLA2G5	[Ca <sup>2+</sup> ]-dependent phospholipase A2
PS	Pulsating shear force
QCM	Quartz microbalance
Rab8a	Ras-related protein Rab-8a
ROS	Reactive oxygen species
Th1	T-helper 1
VCAM-1	Vascular cell adhesion molecule

#### Introduction

Infectious diseases such as bacterial meningitis induced by *E. coli* are accompanied by serious inflammation and are an important cause of mortality and morbidity in neonates and children. In infants, the rate of mortality from these serious diseases is as high as 10-30% [1, 2], and mortality rates are even higher in deadly viral diseases such as that seen with the COVID-19 pandemic.

E. coli invasion of the blood-brain barrier, which exploits microbial and host factors, is the essential step for bacterial meningitis to occur. Although such diseases are usually accompanied by a serious inflammatory response, which has received much attention, the detailed molecular mechanism related to the inflammatory and immunological symptoms has not yet been revealed. In a previous study, we found that YojI-IFNAR2, YojI-IL10RA, YojI-NRP1, YojI-SIGLEC7 and YojI-MC4R membrane-protein interactions mediated the E. coli-human brain microvascular endothelial cell (HBMEC) interaction, through the use of human and E. coli proteomic chips [HuProtarray (version II)] conjugated cell labeling, fluorescence-activated cell sorting (FACS) flow cytometry separation technology, and confocal immunofluorescence techniques at Johns Hopkins University medical school (Fig.1, Table 1) [1-5] It was further reported that invasion of HBMEC by E. coli causing meningitis also requires outermembrane protein A (OmpA), invasion of IbeA, IbeB, IbeC and cytotoxic necrotizing factor 1 (CNF1) and rearrangement of the actin cytoskeleton. It has also been shown that surfaceexposed loops of OmpA contribute to E. coli binding to HBMECs and that OmpA interacts with HBMECs through N-acetylglucosamine (GlcNAc) residues [6]. However, both the IFNAR2 and 1L10RA binding the YojI of E. coli activated the downstream NLRP2 protein (NLR protein family), which induced the anti-inflammatory response of HBMEC. It is speculated that it is involved in the type 2 interferon signaling pathway and suppression of autophagy to inhibit the secretion of IL- $1\beta$  and IL-18 cytokines [3–5, 7–13]. The newly discovered NLRP2 protein induced by E. coli infection and the NLR protein family are important molecular mechanisms for in vivo inflammation and cytokine secretion [1-7].

However, the identification of IFNAR2 and IL10RA as a YojI-interacting protein receptor is a very interesting discovery, and the type II interferon signaling pathway (IFN- $\alpha/\beta$ ) which binds IFNAR2 is the most important antiviral cellular defense mechanism (Table 1) [1]. Both IFNAR2 and IL10RA may inhibit autophagy to increase the cytokine secretion of IL-1 $\beta$  and IL-18, because autophagy can target pro-IL-1 $\beta$  for degradation, and inhibition of autophagy prevents degradation of pro-IL-1 $\beta$  and pro-IL-18, leaving more cytokines freely available in the cytosol for processing and secretion. In addition, inhibition of autophagy leads to activation of the NLRP3 inflammasome and caspase-1-dependent processing of IL-1 $\beta$  and IL-18 cytokines [3–5, 7, 8, 15, 16].

Interleukins (IL) are a group of cytokines secreted by many cell types and can act on a variety of cells by activating and regulating immune cells and mediating the activation, proliferation and differentiation of T and B lymphocytes. They can eliminate infectious pathogens through the induction of secondary mediators and the recruitment of additional immune cells to the site of infection, leading to pyroptosis of the infected cells. Interleukins can also induce the assembly of cytokines, enzymes



**Fig. 1** The applications of the proteomic chip and virion display array in the pathogen (bacteria/virus)–host cell interaction field: **a** High-throughput chip assay for investigating *E. coli* interaction with the blood–brain barrier using microbial and human proteomic microarrays.

that regulate local and systemic inflammation responses. IL-1 $\beta$  and IL-18 can induce IFN- $\gamma$  in T cells and natural killer (NK) cells, and assemble pro-inflammatory cytokines and cell adhesion molecules to amplify the inflammatory response, which is referred to as cytokine storm. It activates the expression of a large number of interferon-stimulated genes (ISGs) to inhibit viral replication, and performs the tumor-killing function by immune cells and cellular autophagy. It has recently been found that the IL-18

**b** Construction of the virion display array for profiling functional membrane proteins and HTS of molecular drugs. The recombinant virions can be immunized mice for further antibody and cytokine production

cytokine is a potential immunotherapy target for tumors, which can be directionally engineered by synthetic biology and yeast assembly to become IL-18 mutant (decoy-resistant IL-18, DR-18). It does not bind the natural pseudoreceptor (IL-18BP) that is secreted by many tumor types, which can serve as a tumor marker, but maintains the signaling transduction function to bind the receptor (IL-18Ra/R $\beta$ ) to activate NK and CD8 T cells, and replication of immune cells to clear away tumor cells [7, 15].

 Table 1
 List of the highlighted hits of human membrane proteins identified by YojI on the HuProt chip (\*denotes membrane proteins which bind to both YojI and IbeC)

Gene symbol	Protein size (a.a.)	Function
*IFNAR2	515	Interferon alpha/beta receptor 2; associated with IFNAR1 to form the type I interferon receptor. Receptor for interferons alpha and beta. Involved in IFN-mediated association of STAT1, STAT2 and STAT3 activation [1, 2, 5, 6, 14].
*IL10RA	578	Cell surface receptor for the cytokine IL10 that participates in IL10-mediated anti-inflammatory functions, limiting excessive tissue disruption caused by inflammation. Upon binding to IL10, induces a conformational change in IL10RB, allowing IL10RB to bind IL10 as well. In turn, the heterotetrameric assembly complex, composed of two subunits of IL10RA and IL10RB, activates the kinases JAK1 and TYK2 that are constitutively associated with IL10RA and IL10RB, respectively. These kinases then phosphorylate specific tyrosine residues in the intracellular domain in IL10RA, leading to the recruitment and subsequent phosphorylation of STAT3. Once phosphorylated, STAT3 homodimerizes, translocates to the nucleus and activates the expression of anti-inflammatory genes. In addition, IL10 RA-mediated activation of STAT3 inhibits starvation-induced autophagy [1, 2].



**Fig. 2** Anti-inflammatory mechanism triggered by YojI-IFNAR2 and YojI-IL10RA membrane protein interactions. **a** YojI-IFNAR2 and YojI-IL10RA membrane protein interaction induces the anti-inflammatory responses of HBMECs, because the *NLRP2* gene inhibits inflammasome-dependent autophagy. NLR oligomerization activates

NF-kB activity for the transcription of IL-1 $\beta$ , and mediates caspase-1 cleavage of pro-inflammatory factor IL-1 $\beta$  to the mature IL-1 $\beta$ . **b** IFN- $\alpha/\beta$  binding to IFNAR triggers inflammasome activation and inflammatory responses

As such, IL-1 $\beta$  and IL-18 are important proinflammatory cytokines and inflammatory markers, which can be engineered as new targets of immunotherapy for cancer (Fig. 2). Cytokines play a direct role in the rapid activation of the immune system to respond to emergency signals, and were the first immunotherapy drug approved by the US Food and Drug Administration (FDA) [15].

To date, thirty-eight members are known. IL is the most common and powerful cytokine factor family and includes IL-1 $\beta$ , IL-1a, IL-18, IL-18BP, IL-33, IL-36Ra, IL-36 $\beta$ , IL-36 $\gamma$ , IL-37 and IL-38. IL-1 $\beta$  and IL-18 are the main secreted forms of IL-1 and play an important role in infectious (bacteria/virus), inflammatory and autoimmune diseases (Table 2) [6–8, 14–19]. If these cytokines are subjected to ionized fragments, due to their different mass/charge (m/z) ratios, they will run to form distinct semicircle trajectories in an applied magnetic field, which can be differentiated by high-performance liquid chromatography-mass spectrometry (HPLC-MS) (Table 3).

In the downstream events, YojI binding of the IL10RA receptor activates *NLRP2* gene expression, as identified by the microarray assay. According to the reported literature, the NLRP2 protein belongs to the NOD-like receptor (NLR) signaling family, along with NLRP1, NLRP3 and NLRP7, but its function is speculated to be contrary to NLRP1,3,7 in a few cell types [8, 16]. The NLR proteins, which lead to NF-kB activation of both transcription of IL-1 $\beta$  and cleavage of pro-1 $\beta$  by caspase-1, can be spontaneously oligomerized to form NLR1,3,7-ASC-caspase-1 inflammasomes, which promote the secretion of IL-1 $\beta$ . We

have conducted a pre-experiment that has already knocked down the *NLRP2* gene in HBMEC to promote a large quantity of IL- $1\beta$  secretion (data not shown here). Thus, IFNAR2 and IL10RA inhibit autophagy to increase cytokine secretion, possibly through an NLRP2 protein-mediated inflammasome inhibition mechanism (Fig. 2a) [6–8, 14–19].

In our recent experiment, results confirmed that knockdown of the *NLRP2* gene by ShRNA lentivirus combined with lowconcentration puromycin (8  $\mu$ g/mL) treatment induced HBMEC secretion of a large quantity of IL-1 $\beta$ . The NLRP2 protein, in contrast to NLRP3, can inhibit the cellular inflammatory response. Therefore, studies on the ligand–membrane protein interactions associated with an intracellular antiinflammatory mechanism are very meaningful for the development of recombinant virion and cytokine chips for clinical inflammatory disease detection and therapy (Fig. 2b) [20, 22–26].

#### Membrane protein virion chips and virion nano-oscillators have been constructed for profiling membrane protein functions and HTS of small-molecule drugs

In previous studies, CD4 and G protein-coupled receptor (GPCR) virion chips and virion nano-oscillators for profiling membrane protein functions were constructed by the author of the present study and Hu et al. in the USA (Fig. 3) [1, 2]. Both single (i.e., CD4) and multi-pass

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Table 2

IL-8

Gene name	Protein name	Sequence	Function
$ILI\beta$	Interleukin-1 beta (mature form: 117-269, 153AA)	APVRSLNCTLRDSQQKSLVMSGPYELKALHLQGQDM EQQVVFSMSFVQGEESNDKIPVALGLKEKNLYLSCV LKDDKPTLQLESVDPKNYPKKKMEKRFVFNKIEIN NKLEFESAQFPNWYISTSQAENMPVFLGGTKGGQ DIT DFTMQFVSS	Potent proinflammatory cytokine. Initially discovered as a major endogenous pyrogen, induces prostaglandin synthesis, neutrophil influx and activation, T-cell activation and cytokine production, B cell activation and antibody production, and fibroblast prolif- eration and collagen production. Promotes Th17 differentiation of T cells. Synergizes with IL 12/interleukin-12 to induce IFNY synthesis from T-helper 1 (Th1) cells [3–6].
IL 18	Interleukin-18 (mature form: 37-193aa, 117AA)	YFGK LESKLSVIRNLNDQVLFI DQ GNRPLFEDMT DSDCRDNAPR TIFIISMYKD SQPRGMAVTI SVKCEKISTL SCENKIISFK EMNPPDNIKD TKSDIIFFQR SVPGHDNKMQ FESSSYEGYF LACEKERDLF KLILKKEDEL GDRSIMFTVQNED	A proinflammatory cytokine primarily involved in polarized T-helper 1 (Th1) and natural killer (NK) cell immune responses to perform the tumor-killing effect. Upon binding to IL18R1 and IL18RAP, forms a signaling termary complex which activates NF-kappa-B, triggering synthesis of inflammatory mediators. It can be applied for immunotherapy for cancer. Synergizes with IL12/interleukin-12 to induce IFNY synthesis from T-helper 1 (Th1) cells and natural killer (NK) cells [18].
IL6	Interleukin-6 (mature form: 30-212aa, 183AA)	VPPGEDSKDVA APHRQPLTSS ERIDKQIRYI LDGISALRKE TCNKSNMCES SKEALAENNL NLPKMAEKDG CFQSGFNEET CL VKIITGLL EFEVYLEYLQ NRFESSEEQA RAVQMSTKVL IQFLQKKAKN LDAITTPDPT TNASLLTKLQ AQNQWLQDMT THLILRSFKEFLQSSLRALRQM	Cytokine with a wide variety of biological functions. It is a potent inducer of the acute phase response. Plays an essential role in the final differentiation of B-cells into Ig-secreting cells Involved in lymphocyte and monocyte differentiation. Acts on B and T cells, hepatocytes, hematopoietic progenitor cells and cells of the CNS. Required for the generation of T(H)17 cells. Also acts as a myokine. It is discharged into the bloodstream after muscle contraction and acts to increase the breakdown of fats and to improve insulin resistance. It induces myeloma and plasmacytoma growth and induces nerve cell differentiation [6, 19].
IL 8	Interleukin-8 (99AA)	MTSKLAVALL AAFLISAALC EGAVLPRSAK ELRCQCIKTY SKPFHPKFIKELRVIESGPH CANTEIIVKL SDGRELCLDP KENWVQRVVE KFLKRAENS	IL-8 is a chemotactic factor that attracts neutrophils, basophils and T cells, but not monocytes. It is also involved in neutrophil activation. It is released from several cell types in response to an inflammatory stimulus. IL-8(6-77) has a 5–10-fold higher activity on neutrophil activation, IL-8(5-77) has increased activity on neutrophil activation and IL-8(7-77) has a higher affinity to receptors CXCR1 and CXCR2 as compared to IL-8(1-77) [6, 19].

Analytical approaches	Working principle	Applied field	Advantages	Disadvantages
HPLC-MS	Liquid chromatography-mass spectrometry, which uses liquid chromatography as separa- tion system and mass spectrometry as detection system. After the sample is separated from the mobile phase in the mass spectrum, the ion fragments are separated according to the mass number by the mass analyzer of the mass/charge (m/z) spectrum, which is obtained by the detector	Drug metabolism, cytokines, antibody, enzyme, polypeptide, polysaccharide, polymer analysis and molecular biology	Can analyze multiple cytokines and drugs at the same time. Analyzes biological macromolecular compounds with very large molecular weight; the determination range of protein analysis molecular weight is more than 10 <sup>5</sup> [1, 2, 20, 21].	
FACS	After being dyed with specific fluorescent dye, the cells to be tested are put into the sample tube and enter the flow chamber filled with sheath fluid under gas pressure. Under the restriction of sheath fluid, cells are arranged in a single row and ejected from the nozzle of the flow chamber to form a cell column. The latter intersects the incident laser beam vertically, and the cells in the liquid column are excited by the laser to produce fluorescence. A series of optical systems (lenses, apertures, filters, detectors, etc.) in the instrument collect signals of fluorescence, light scattering, light absorption or cell electrical impedance. The computer system collects, stores, displays and analyzes various signals, and generates statistical	Cell surface protein analysis, immune cytochemical typing, exogenous and endogenous gene expression, membrane potential and cell permeability	Can analyze multiple components of cytokines and drugs at the same time, with strong fluorescence signal [1, 2, 20].	
ELISA, Char MI	analysis of various indicators. Sandwich principle of molecular hybridization	High-precision detection of single drug, proteins and cytokines [1, 2]		It cannot provide molecular structure, and it is difficult to detect and analyze multiple components of drugs at the same time; cannot effectively exclude a false positive, so it has certain limitations.
8%12% SDS-P- AGE	Polyacrylamide gel as a support medium for a common electrophoresis technology. It has a molecular sieve effect, which is used to separate cytokines and oligonucleotides.		In this method, the mobility of cytokines is determined mainly by their molecular weight, not by charge and molecular shape [1].	It cannot determine the specific cytokine.



**Fig. 3** Application of the single (IFNAR2) and multi-transmembrane protein (G protein-coupled receptor, GPCR) recombinant herpes virions (IFNAR2-HSV-1 and GPCR-HSV-1) to construct virion nano-oscillators

with the perfusion system of a microfluidic device for HTS of molecular drugs. Pulsating shear force (PS) and oscillatory shear force (OS) can be loaded to gain variable binding kinetics to characterize the drug activity

(i.e., GPCRs) membrane proteins were exquisitely displayed on the envelopes of herpes simplex virus-1 (HSV-1) with correct orientation and conformation [1, 2,20]. Single recombinant virion (~185 nm diameter) reacted with polyethylene glycol (PEG) linkers (63 nm in length) to form the virion-PEG complex to be tethered to a goldcoated chip via PEG linkers. With a similar constructive method, a total of 332 (98.5%) GPCR open reading frames (ORFs) of a total of 337 non-odorant GPCRs were subcloned into the UL27 locus (gB) of the herpes virus genome and used to fabricate the VirD-GPCR array for high-content study of their pharmacological activity and preferred drug targets (Fig. 3) [2, 21]. Among the 1400 FDA-approved drugs, 475 (34%) drugs targeting 108 non-odorant GPCRs have been identified. Although there are about 70% known to have the ligand, 30% of the GPCRs are still without an identified ligand [20, 21, 26-31]. Thus, the structure of un-characterized GPCRs and the abovementioned membrane proteins including IFNAR2 and IL10RA or Spike/ACE2 of 2019-nCoV can be further incorporated into the herpes virus envelope or highly expressed extracellular transmembrane domain to induce synthetic antibody to block bacterial/viral infection [1, 2, 9, 27, 29–31]. These recombinant virions are versatile players in the biomedical field. Various methods using a combined proteomics and genomics approach are rapidly being developed; however, utilization of the recombinant membrane protein herpes virions for inflammatory disease diagnosis and treatment has not yet been reported. In order to construct a new recombinant virion solution chip to establish a multi-task platform for clinical biomedical applications, we proposed the recombinant virion solution chip integrated with the properties of an electrochemical sensor for use in clinical diagnosis of inflammatory diseases as well as HTS of anti-inflammatory drugs [20, 23–26]. Because the molecular drug-receptor interactions on these recombinant virions will trigger charge distribution and electron transition to produce electrochemical signaling, IFNAR2 and IL10RA membrane proteins can be applied with detection of electrical signals for HTS of anti-inflammatory molecular drugs (Fig. 3) [1, 6, 14, 20, 21]. Different shear forces such as pulsating (PS) and oscillatory shear force (OS) can also be loaded to gain variable binding kinetics to characterize the drug activity (Fig. 3) [6, 20].

#### The*NLRP2* gene can be knocked down by ShRNA lentivirus for high-throughput construction of a cytokine solution chip

It is reported that the NLRP2 protein can suppress inflammation [3–5, 16, 32]; thus knockdown of the *NLRP2* gene in HBMECs may activate the inflammasome and transcription factor NF-kB and autophagosome to promote the expression and secretion of the IL-1 $\beta$ , IL-18, IL-1 $\alpha$ , IL-6, TGF $\beta$ , IL-8 and IL-33 cytokines.

There are two steps in IL-1 $\beta$  secretion. First, inflammatory signals, such as lipopolysaccharide (LPS), stimulate synthesis and promote accumulation of the cytosol of pro-IL-1 $\beta$ . Additional signals, such as ATP, ion, membrane perturbations or reactive oxygen species (ROS), are also required for inflammasome activation and assembly, leading to caspase-1 activation and processing of pro-IL-1 $\beta$  and eventual secretion of the active cytokines.

Findings showed that autophagy has a positive contribution to the biogenesis and secretion of the pro-inflammatory cytokine IL-1 $\beta$  via an export pathway that depends on Atg5 and inflammasomes, and at least one of the two mammalian Golgi reassembly stacking protein 55(GRASP55) and Ras-related protein Rab-8a (Rab8a) is required [9–13, 33]. Additionally, PLA2G5 is required for phospholipid hydrolysis and release of IL-1 $\beta$  and IL-18. Knockdown of the *NLRP2* gene induces autophagy to promote inflammasome-dependent IL-1 $\beta$  secretion [9–12, 33, 34]. IL-1 $\beta$  lacks any known signal sequence, and the pathways of its secretion in HBMECs are speculated to be mainly through an inflammasome-dependent autophagy mechanism (Table 2, Fig. 2) [9–13, 33, 34].

Based on the results of previous experiments, several unconventional secretion mechanisms are proposed for the secretion of cytokines.

- 1. Secretion via secretory lysosome. IL-1 $\beta$  may be incorporated into secretory lysosomes that undergo [Ca<sup>2+</sup>]-dependent exocytosis with the release of the mature IL-1 $\beta$  with the help of the PLA2G5.
- 2. Secretory autophagy: IL-1 $\beta$ -autophagosome may fuse with the plasma membrane to release soluble IL-1 $\beta$ , which is inflammasome-dependent [11–13].
- 3. Secretion via exosomes: P2RX7 that is activated by ATP leads to the formation of multivesicular bodies (MVBs) containing exocytosis with IL-1 $\beta$ , caspase-1 and other inflammasome components. These MVBs undergo exocytosis with the release of IL-1 $\beta$  after the lysis of the exosome membrane.
- 4. Secretion via microvesicle shedding. P2RX7 induces immediate shedding of membrane-derived microvesicles containing IL-1 $\beta$  and inflammasome components. These cytokines are then released extracellularly after microvesicle lysis.
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5. Release by translocation through permeabilized plasma membrane, which occurs in cells undergoing pyroptosis, due to the sustained activation of the inflammasome.

IL-1 $\beta$  and IL-18 induce the expression of secondary mediators that attract immune cells to the site of the infection. Although IL-1 $\beta$  and IL-18 have a beneficial role in promoting inflammation and eliminating infectious pathogens, mutations that result in constitutive inflammasome activation and the over-production of IL-1 $\beta$  and IL-18 have been linked to auto-inflammatory and autoimmune disorders.

For these reasons, mechanisms that inhibit IL-1 $\beta$  and IL-18 signaling are of great interest to us. Decoy or soluble receptors that sequester IL-1 $\beta$  and IL-1 $\beta$  antagonists and disrupt IL-1 receptor heterodimerization are intrinsic pathways that inhibit IL-1 $\beta$  signaling. Similarly, naturally occurring IL-18 binding protein (IL-18 BP) can prevent IL-18 from binding to its receptor. Based on this bio-analysis of the secretion signaling pathway of IL, three approaches are proposed to increase the extracellular release of cytokines and high-throughput construction of the cytokine solution chip for HTS of molecular drugs.

- Knockdown of the *NLRP2* gene, along with highly expressed PLA2G5 for hydrolysis of the phospholipid monolayer to promote the local release of IL-1β, IL18, etc., in HBMECs.
- (2) Knockdown of the *NLRP2* gene and activation of probable G protein-coupled receptor 132 (*G2A*) gene expression to inhibit cell proliferation and promote inflammation to obtain a high quantity of IL-1β and IL-18.
- (3) Knockdown of both the *NLRP2* gene and cell adhesion molecule gene (*VCAM-1*) to promote cell suspension and local inflammation to obtain a high titer of IL-1β, IL-18 and other cytokines in HBMECs.

# High-throughput construction of new ultrasensitive solid cytokine chips for HTS of anti-inflammatory drugs

Because the binding of small-molecule drugs with cytokines will present different binding kinetics and electrochemical signaling, the cytokine solution chip can employ an electrochemical three-electrode system for detecting the electron transfer between the drugs and cytokines, and the effective lead drugs can be screened [20, 22–26, 34–36]. If the IL-1 $\beta$  and IL-18 ae expressed as chimera IL-1 $\beta$ -V5 tag and IL-18-V5 tag, the cytokines can be loaded on the solid glass slide through V5 antibody binding for HTS of molecular drugs based on the ELISA principle. That is, the lead drugs conjugated to the biotin can be screened out by binding the peroxidase-conjugated avidin, which can catalyze the TMB

chromogenic reagent to present a blue color for ultrasensitive identification of the lead drugs (Fig. 4a).

Remarkably, these cytokines can be tethered to the Nicoated chip via chelation of  $(2\times)$   $6 \times$  His tag: Ni, if they expressed as chimera IL-1 $\beta$ -6 × His tag  $(2\times)$  and IL-18-6 × His tag  $(2\times)$  (Fig. 4b). By applying an alternating electric field perpendicular to the Ni-coated surface, the variation of the amplitude can characterize the pharmacodynamic activity of the lead drugs (Fig. 4b).

## High-throughput construction of the liquid and microfluidic cytokine chips for HTS of anti-inflammatory drugs

If the cytokines were secreted as a cytokine storm, these massive cytokines can be purified individually (Tables 2 and 3). Because lysine is widely present in peptide chains of proteins, the active agent of bovine serum albumin (BSA) can react with these lysines to make them anchor on the BSA-coated glass slide, which exposes the corresponding active groups for HTS of molecular drugs (Fig. 5a).

On the contrary, if the candidate drugs were anchored on the chip surface through the avidin–biotin interaction, the microfluidic chip could be made to let the IL-1 $\beta$  cytokine solution flow through the fixed molecular drugs for HTS of candidate drugs, and identify the lead drugs by mass spectrometry (MS) detection, because a high molecular ion peak can be generated according to charge/mass (m/z) of highly ionized cytokines (Table 2, Fig. 5b). The candidate drugs can also be labeled by CY-5/CY-3 fluorescent dye for HTS of effective drugs by applying a microgravity field [26]. IL-1 $\beta$ /IL-18 cytokine drug screening has broad application for the diagnosis and treatment of inflammasome-related diseases, including auto-inflammatory disorders, Crohn's disease, vitiligo, gout, asbestosis, Alzheimer's disease and even tumors (Fig. 5c).

## Recombinant herpes virion liquid electrochemical sensor can be fabricated by high-throughput construction for HTS of anti-inflammatory drugs or clinical diagnosis and treatment of inflammatory diseases

Different molecular drug targets will generate variable electrochemical signaling for monitoring, due to the different structural properties of single transmembrane (IFNAR2) and multi-transmembrane protein (GPCR)

As just mentioned above, YojI binds IFNAR2 and triggers the intracellular anti-inflammatory signaling related to the synthesis and secretion of cytokines IL-1 $\beta$ /IL-18 [1, 2, 7, 8, 15–19]. Thus, the IFNAR2 receptor should retain the anti-inflammatory properties, which can be utilized for molecular drug docking and screening. A model can be used to explain the drug effectiveness: If the drug binds the single transmembrane protein IFNAR2, it mostly binds the extracellular chain



Fig. 4 Construction of new ultrasensitive cytokine and virion chips for HTS of anti-inflammatory drugs or clinical diagnosis and treatment of inflammatory diseases. a Chimera cytokine IL-1 $\beta$ -V5 tag and IL-18-V5 tag were highly expressed and were loaded on the solid chip through V5

antibody for HTS of molecular drugs based on the ELISA principle. **b** Chimera protein IL-1 $\beta$ -6 × His tag (**2**×) and IL-18-6 × His tag (**2**×) were expressed and loaded on the Ni-coated solid chip for HTS of molecular drugs of IL-1 $\beta$  cytokines



Fig. 5 High-throughput construction of the cytokine solution chip for HTS of anti-inflammatory drugs. The drug activity can be characterized either by the settlement of micro-fluid, microgravity condition, Cy-5/Cy-3 labeling or ELISA principle

(27-243aa) of IFNAR2 to exert its pharmacological activity. However, if the drug binds GPCR with seven-transmembrane domains, it can bind different loops of GPCR or the Nterminal of this protein receptor to perform its pharmacological activity (Fig. 6a). The different binding sites between the IFNAR2 and GPCR will generate different electrochemical signaling for monitoring, which is our primary interest.



Fig. 6 High-throughput construction of a liquid electrochemical sensor integrated with recombinant membrane protein virions for HTS of antiinfective and anti-inflammatory drugs (2019-novel coronavirus/human/ bacteria) and clinical diagnosis of inflammatory diseases. **a** Molecular drugs target different sites of transmembrane proteins: small drugs bind the extracellular domain of IFNAR2, N-terminal domain or different loops of seven-transmembrane domain protein receptor G2A (loop i, loop

ii, loop iii); the membrane protein interaction between Spike (trimer) and ACE2 (dimer). **b** The IFNAR2/ACE2/Spike and GPCR-recombinant herpes virion solution (IFNAR2/ACE2/Spike-HSV-1 and GPCR-HSV-1) can be applied to fabricate the new virion liquid electrochemical sensor for HTS of anti-inflammatory drugs or clinical diagnosis and treatment of inflammatory diseases, especially for the prevention and cure of 2019-novel coronavirus pneumonia

#### (IFNAR2/ACE2/Spike)-HSV-1 and (GPCR)-HSV-1 can be rapidly replicated for high-throughput construction of the recombinant virion liquid electrochemical sensor

Due to the similar single-transmembrane structure between Spike of 2019-nCoV, ACE2 of alveolar epithelial cells II (ATII) and IFNAR2 of HBMEC, the (IFNAR2/IL10RA/ACE2/Spike)-HSV-1 and (GPCR)-HSV-1 can be fabricated and rapidly replicated for highthroughput construction of the recombinant virion liquid sensor. Because the detected solution is a non-aqueous solution, and the IR [Deviation caused by I (current) and R (resistance) of the liquid] is much larger, such as blood and virion liquid, a three-electrode system should be applied, comprising a working electrode, an Ag wire as the quasi-reference electrode and a Pt coil as the counter electrode (Fig. 6b). The electrochemical signal will load on the working electrode and counter electrode which are linked to an electrical access loop for detection of the electron transition and electric current passing through [20, 22–25, 37, 38].

The high-throughput construction of a recombinant virion liquid electrochemical sensor is shown in (Fig. 6) [20, 22–25].

Because the single-transmembrane protein IFNAR2 presents a very long extracellular domain (27-243aa), which is similar to CD4 and IFNAR2, Spike (14-1195aa) of 2019nCoV and ACE2 (18-740aa) of alveolar epithelial cells II, they can be very easily incorporated into the envelope of HSV-1 under the glycoprotein B promotor of the HSV-1 genome to make the IFNAR2/ACE2/Spike-HSV-1 and GPCR recombinant virions (www.uniprot.org) [1, 2, 20, 21, 26]. Therefore, the IFNAR2/ACE2/Spike and GPCRrecombinant herpes virions (IFNAR2/ACE2/Spike-HSV-1 and GPCR-HSV-1) can be created through high-throughput construction to obtain a new virion solution electrochemical sensor for probing drug activity. The applied electrochemical sensor is very sensitive for monitoring electron transition of IFNAR2/IL10RA/ACE2/Spike/GPCR-molecular drug interaction in a small area of the virion solution, which can be compared with the electrical signal of the negative control virions (wild-type virions without IFNAR2/IL10RA/ACE2/ Spike or GPCR incorporation) to gain electrical signaling of clinical diagnosis and to identify effective drugs (Fig. 6b) [2, 9, 10, 21, 26-32].

In previous studies, recombinant purified virions were arrayed in a 384-well plate and spotted on FAST slides (Whatman) in a  $4 \times 4$  pattern to fabricate the virion display (VirD) array in order to profile functional membrane proteins using the Cy5-labeled antibody, ligand and lectin to detect immunofluorescence intensity for characterization of their structure and activity (Fig. 1) [2, 20, 21]. Next, virion nano-

oscillators were made to allow us to measure mobility and charge changes associated with molecular binding strength on each recombinant virion, characterized by the variation in binding kinetics and amplitude (Fig. 3). These novel analytical characterization approaches and real-time photoelectric measurement techniques provide us a new reference of theoretical analysis for the construction of the recombinant virion liquid electrochemical sensor. It can be applied not only in the detection of infectious diseases (patient blood sample that has been immunized with antibodies of bacteria/virus/2019nCoV), but also in the detection of diabetes, obesity, tissue injury, acute or chronic inflammation (pancreatitis or rheumatoid arthritis), cardiovascular disease and tumors (Fig. 6b). Subsequently, the screened drugs can be docked to the protein receptors on the virion envelope for designing highly effective anti-inflammatory drugs based on their crystal structure with computer-aided drug design. Because of the flexibility of the VirD platform, we can co-infect human cells with more than one type of VirD-GPCR to produce chimera, heterodimer or multimer VirD-GPCRs to establish the multifunctional virion liquid electrochemical platform for HTS of molecular drugs and the clinical diagnosis and treatment of infectious and inflammatory diseases.

# Anti-inflammatory and anti-2019-nCoV drugs can be developed

If we have identified the drugs with strong effectiveness by the virion and cytokine liquid chips, we can obtain the drug-recombinant virion crystal structures to design and modify the drugs based on the membrane protein receptor structure and their electronic binding properties [39–41]. Both the membrane protein receptors (Fig. 6a) and the transcription factor NF-kB and NLRP2 proteins can be designed as drugs for blocking the membrane protein interactions or blocking the protein aggregation and DNA binding properties to suppress inflammation in the human body [36, 42].

#### Conclusions

We have discovered that the YojI-IFNAR2 and YojI-IL10RA membrane protein interactions mediate antiinflammatory signaling cross-talk, which is related to the NLRP2 protein inhibition of inflammasome-mediated synthesis and secretion of IL-1 $\beta$  and IL-18. This novel unrevealed anti-inflammatory cellular signaling was applied to investigate the high-throughput construction of liquid cytokine chips and for the development of the single (IFNAR2/ IL10RA/ACE2/Spike) and multi-transmembrane protein (GPCR) recombinant herpes virion liquid electrochemical sensor (IFNAR2/IL10RA/ACE2/Spike-HSV-1 and GPCR- HSV-1) for HTS of anti-inflammatory drugs or clinical diagnosis and treatment of inflammatory diseases. Recently, the RNA coronavirus 2019-nCoV has posed a serious global public health threat, with death rates increasing every day. The infection occurs mainly through the membrane protein interaction of Spike with ACE2. Because the Spike protein is a trimer and ACE2 is a dimer (Fig. 6a), immune response is of tremendous importance for the production of synthetic antibodies to eliminate the 2019-nCoV pathogen. Thus, the molecular and protein drugs, vaccines (the singletransmembrane protein of Spike and ACE2 which removed the transmembrane domain) and antibodies can be developed to block the Spike and ACE2 membrane protein interaction to inhibit the 2019-nCoV infection (Fig. 1) [34-36, 39–41], which can be gained through recombinant herpes virion-immunized mice. Recently, angiotensin-converting enzyme 2 (ACE2) has received much attention. It has been identified as a functional receptor that is related to vasoconstriction, which plays a vital role in the development of hypertension, atherosclerosis and kidney disease, and which requires our further investigation (Fig. 6a). Interestingly, we can knock down the NLRP2 gene to monitor the dynamic conformation variation of ACE2 in response to blood rheology [39]. As a result, studies on the interactions between 2019-nCoV, ACE2 and the host innate immune system, and NLRP2 gene function are critical for infectious, cardiovascular and even tumor disease therapy, and to control serious epidemics. In the future, in addition to microfluidic and electrochemical devices, other advanced physical techniques such as field spectroscopy, quartz microbalance (QCM) and sonic waves can be applied for HTS of drugs. The newly constructed cytokine and virion liquid sensors have significant value for clinical diagnosis and treatment of infectious and inflammatory diseases (Figs. 5 and 6) [27, 36, 42].

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#### Compliance with ethical standards

No animal experiments were performed in this work. It is in compliance with ethical standards.

**Conflict of interest** The authors declare that they have no conflict of interest.

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