Supplementary information

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The names of the authors: Helin Wang¹, Xiuhua Hu¹, Mingfang Zhang¹, Lin Yang¹, Yueying Xu¹, Xiaoxu Gu¹, Junjun Jiang¹*, Weiwei Hu¹*

The affiliations and addresses of the authors:

¹ Microbial and Viral Platforms (MVP), WuXi Biologics, 291 Fucheng Road, Hangzhou 311106, China

Corresponding author:

Dr. Weiwei Hu

E-mail: hu weiwei0501@wuxibiologics.com

Dr. Junjun Jiang

E-mail: jiang junjun@wuxibiologics.com

Telephone and fax number: +86-571-60954091

Supplementary figures:

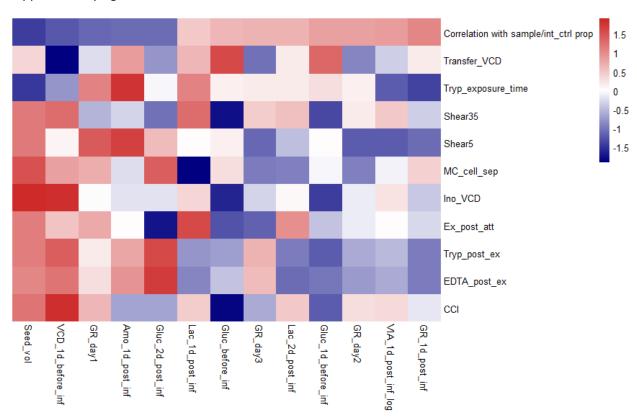


Figure S1 Correlation factor matrix of secondary correlation analysis

The first row of the matrix shows the correlation between the performance parameters and the sample/internal control proportion (normalized titer). The other rows show the correlation between the performance parameters and the process parameters.

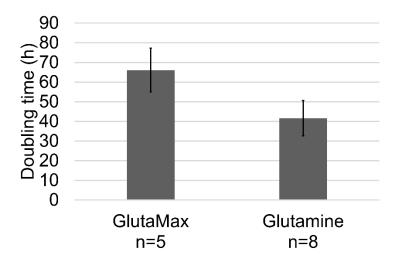


Figure S2 Doubling time of Vero cells cultured with GlutaMax or glutamine as supplements

Vero cells were cultured with VP-SFM medium in T flasks, and GlutaMax (n=5) or glutamine (n=8) were used as source of glutamine. Doubling times of the cells were shown in the figure.

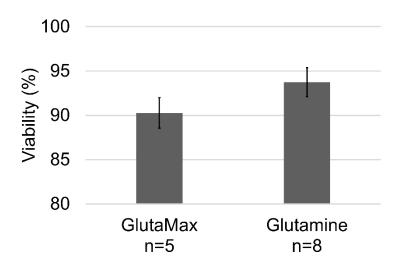


Figure S3 Viability of Vero cells cultured with GlutaMax or glutamine as supplements

Vero cells were cultured with VP-SFM medium in T flasks, and GlutaMax (n=5) or glutamine (n=8) were used as source of glutamine. Viabilities of the cells were shown in the figure.

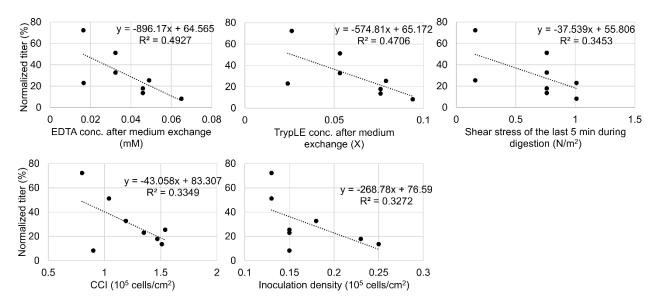


Figure S4 Linear regression analysis between screened operation parameters and normalized titer. Correlation between normalized titer and the five operation parameters (EDTA and TrypLE concentration after medium exchange, shear stress of the last 5 min during digestion, CCI and inoculation density) were indicated by correlation analysis and PCA. And linear regression was performed to select the optimized level of these parameters.

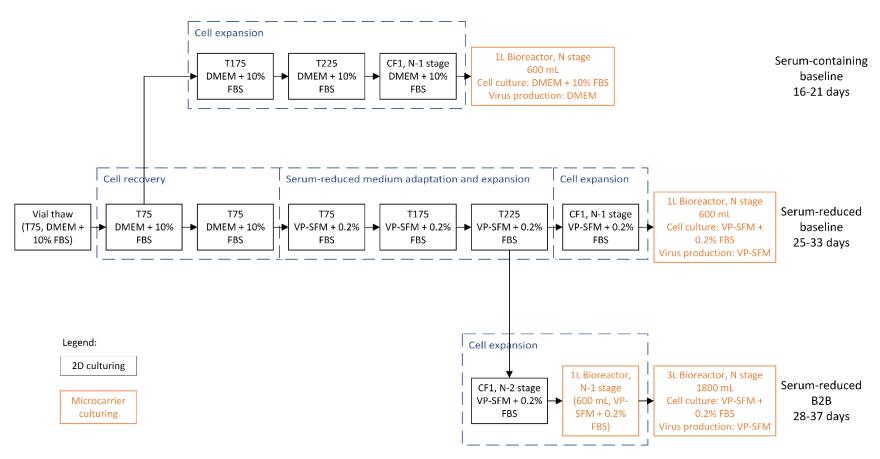


Figure S5 Expansion flow chart for Vero cell culture and HSV-1 production. For the serum-containing baseline process, the cells were recovered and amplified with 2D culturing in DMEM + 10% FBS medium, and finally inoculated into a 1L bioreactor for virus production. For the serum-reduced baseline process, the cells were vial thawed and passaged twice in T75 flask in DMEM + 10% FBS medium for cell recovery, and then directly adapted to VP-SFM + 0.2% FBS medium and amplified simultaneously in 2D culture vessels, then inoculated into a 1L bioreactor for virus production. For the serum reduced bead-to-bead transfer process, the cells cultured in CF1 were inoculated into a 1L bioreactor as N-1 stage. Then bead-to-bead transfer was performed and

the cells were inoculated into a 3L bioreactor for N-stage virus production. Serum was only used during cell expansion, and serum-free medium was used during virus production in N-stage.

Supplementary tables:

Table \$1 Experimental design

Category	Evaluated Parameter	Culture Vessel	Evaluated Performance
Medium composition	Basal medium type T flask		Doubling time, viability, virus titer
optimization	Serum concentration		
	Glutamine supplement type and concentration		
Baseline process transfer	Agitation speed	Bioreactor	Microcarrier uniformity
from spinner flask to	Test baseline process in bioreactors	Bioreactor	Cell growth rate, viability, metabolite levels,
bioreactors			and virus titer
Preliminary bead-to-bead	TrypLE concentration	Spinner flask	Detaching percentage, overall recovery
transfer process	Agitation during digestion		proportion, and viability drop after
development	Residue volume during washing procedure	_	digestion
	Transfer operation reduction		Cell growth rate, viability, metabolite levels,
	Cell and microcarrier separation	_	and virus titer in subculture
	Medium exchange in subculture		
Virus production	47 parameters from historical experiments	N/A	Correlation with virus titer
optimization of the bead-	Residue EDTA and TrypLE concentration in subculture	Spinner flask	Detaching percentage, overall recovery
to-bead transfer process	Shear stress during agitation		proportion, and viability drop after
	Cell densities of inoculation	_	digestion
	CCI		Cell growth rate, viability, metabolite levels,
			and virus titer in subculture
Bead-to-bead transfer	Test optimized bead-to-bead transfer, and subsequent	1L bioreactor for	Detaching percentage, overall recovery
process in bioreactors	cell culture and virus production processes in	N-1 stage	proportion, and viability drop after
	bioreactors	1L and 3L	digestion
		bioreactors for N-	Cell growth rate, viability, metabolite levels,
		stage	and virus titer in subculture

Table S2 Performance and fluid dynamics parameters measured or calculated during the whole procedure

Process stage	Measured or calculated performance parameters
Vial thaw and cell amplification with 2D culturing	Measured: VIA and VCD during cell passage
	Calculated: specific growth rate
N-1 cell amplification with microcarrier	Measured: daily sampling and measure of VIA, VCD, osmolality, pH, pCO ₂ , pO ₂ ,
	concentrations of Na ⁺ , K ⁺ , glucose, ammonium, lactose, glutamine, and glutamic acid.
	Calculated: specific growth rate
Bead-to-bead transfer	Measured: VIA and VCD before the microcarrier sedimentation of wash ("before wash"
	for short), before digestion, and before inoculation
	Calculated: viability drop, overall recovery proportion, and calibrated overall recovery
	proportion.
	$Viability\ drop = VIA_{before\ wash} - VIA_{before\ inoculation}$
	$Overall\ recovery\ proportion = \frac{VCD_{inoculum} \times Vol_{inoculum}}{VCD_{before\ wash} \times Vol_{before\ wash}} \times 100\%$
	$VCD_{before\ wash} \times Vol_{before\ wash}$
	$Overall\ recovery\ proportion_{Calibrated} = \frac{{\it VCD}_{inoculum} \times {\it Vol}_{inoculum}}{{\it VCD}_{before\ wash} \times {\it Vol}_{before\ wash} - {\it Sampling\ loss}} \times $
	100% *
	Sampling loss was calculated as followed:
	Sampling loss = $VCD_{Sample\ 1} \times Vol_{Sample\ 1} + VCD_{Sample\ 2} \times Vol_{Sample\ 2} + \cdots$ + $VCD_{Sample\ n} \times Vol_{Sample\ n}$
	* Calibrated overall recovery proportions were only used in experiments conducted with
	spinner flasks. The overall recovery proportions in bioreactors were not calibrated as
	sampling loss was neglectable.
N-stage cell culturing	Measured: daily sampling and measurement of VIA, VCD, osmolality, pH, pCO ₂ , pO ₂ ,
	concentrations of Na ⁺ , K ⁺ , glucose, ammonium, lactose, glutamine, and glutamic acid.
	Calculated: specific growth rate
Virus infection and amplification	Measured: daily sampling and measurement of VIA, VCD, osmolality, pH, pCO ₂ , pO ₂ ,
	concentrations of Na ⁺ , K ⁺ , glucose, ammonium, lactose, glutamine, and glutamic acid;
	sampling and measurement of volumetric virus titers at different HPIs.
	Calculated: specific growth rate, cell-specific virus production, normalized volumetric titer
Fluid dynamics parameters	Power input per volume:
	$P/V = N_p n^3 \rho D_i^5 / V$

Process stage	Measured or calculated performance parameters	
	Kolmogorov eddy length:	
	$\eta = \left[v^3 / \left(N_p n^3 D_i^5 / V \right) \right]^{\frac{1}{4}}$	
	Maximum shear stress:	
	$\tau_{max} = 5.33 \rho (v N_p n^3 D_i^5 / V)^{\frac{1}{2}}$	
	η is Kolmogorov eddy size (m), v is kinematic viscosity (m ² /s), V is volume (m ³), N _p is	
	impeller's Newton number or impeller power number, n is agitation speed (r/s), ρ is	
	culture medium density (kg/m³), D _i is impeller diameter (m).	

 Table S3 Analyzed operation and performance parameters

Process stage	Parameter	Short name	Туре	Tested range	Unit
N-1 culturing	Transfer VCD - N-1 stage	Transfer_VCD	Process parameter	1.56 - 1.93	10 ⁵ cells/cm ²
N-1 culturing	VIA before digestion - N-1	N-1_VIA_before_digestion	Performance	99.1 - 99.2	%
	stage		parameter		
Digestion	TrypLE exposure time	Tryp_exposure_time	Process parameter	54 - 76	min
Digestion	Shear stress of the beginning	Shear35	Process parameter	0.16 - 0.19	Pa
	30/35 min during digestion				
Digestion	Shear stress of the last 5 min	Shear5	Process parameter	0.16 - 1.01	Pa
	during digestion				
Digestion	Cell/microcarrier separation	MC_cell_sep	Process parameter	Yes/no	N/A
Digestion	VIA after digestion - N-1	VIA_post_digestion	Performance	95.5 - 98.2	%
	stage		parameter		
Termination	TrypLE concentration after	Tryp_post_ter	Mixed	0.74 - 1.58	X
	termination				
Termination	EDTA concentration after	EDTA_post_ter	Mixed	0.51 - 0.96	mM
	termination				
Termination	Medium proportion after	Med_prop_post_ter	Mixed	0 - 39.4	%
	termination				
N-stage inoculation	Inoculation VCD - N-stage	Ino_VCD	Process parameter	0.13 - 0.25	10 ⁵ cells/cm ²

Process stage	Parameter	Short name	Туре	Tested range	Unit
N-stage inoculation	Inoculum volume - N-stage	Seed_vol	Mixed	2.37 - 15.5	mL
N-stage inoculation	Calculated trypLE concentration after inoculation	Try_post_ino	Process parameter	0.026 - 0.30	X
N-stage inoculation	Calculated EDTA concentration after inoculation	EDTA_post_ino	Process parameter	0.016 - 0.18	mM
N-stage inoculation	VIA upon inoculation	VIA_upon_ino	Performance parameter	96.8 - 100	%
N-stage cell culturing	Medium exchange after attaching period	Ex_post_att	Process parameter	Yes/no	N/A
N-stage cell culturing	TrypLE concentration after medium exchange	Tryp_post_ex	Process parameter	0.024 - 0.094	X
N-stage cell culturing	EDTA concentration after medium exchange	EDTA_post_ex	Process parameter	0.016 - 0.065	mM
N-stage cell culturing	VCD one day before infection	VCD_1d_before_inf	Performance parameter	0.45 - 0.94	10 ⁵ cells/cm ²
N-stage cell culturing	VIA one day before infection	VIA_1d_before_inf	Performance parameter	96.7 - 99.7	%
N-stage cell culturing	VIA right before infection	VIA_upon_inf	Performance parameter	99.3 - 100	%
N-stage cell culturing	Cell specific growth rate of day 1 - N stage	GR_day1	Performance parameter	0.18 - 1.35	day ⁻¹
N-stage cell culturing	Cell specific growth rate of day 2 - N stage	GR_day2	Performance parameter	0.62 - 1.21	day ⁻¹
N-stage cell culturing	Cell specific growth rate of day 3 - N stage	GR_day3	Performance parameter	0.16 - 0.68	day ⁻¹
N-stage cell culturing	Glucose concentration one day before infection	Gluc_1d_before_inf	Performance parameter	2.4 - 3.5	g/L
N-stage cell culturing	Glucose concentration right before infection	Gluc_before_inf	Mixed	1.25 - 2.03	g/L

Process stage	Parameter	Short name	Туре	Tested range	Unit
Virus production	CCI	CCI	Process parameter	0.80 - 1.54	10 ⁵ cells/cm ²
Virus production	VCD one day after infection	VCD_1d_post_inf	Performance	0.49 - 1.71	10 ⁵ cells/cm ²
			parameter		
Virus production	VCD two days after infection	VCD_2d_post_inf	Performance	0.27 - 0.86	10 ⁵ cells/cm ²
\	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	parameter	74.5.00.6	0/
Virus production	VIA one day after infection	VIA_1d_post_inf	Performance parameter	74.5 - 99.6	%
Virus production	VIA two days after infection	VIA_2d_post_inf	Performance	59.0 - 82.4	%
			parameter		
Virus production	Cell specific growth rate of	GR_1d_post_inf	Performance	-0.58 - 0.61	day ⁻¹
	the first day after infection -		parameter		
	N stage				
Virus production	Cell specific growth rate of	GR_2d_post_inf	Performance	-1.220.59	day ⁻¹
	the second day after		parameter		
	infection - N stage				
Virus production	Glucose concentration one	Gluc_1d_post_inf	Mixed	2.59 - 3.73	g/L
	day after infection				
Virus production	Glucose concentration two	Gluc_2d_post_inf	Mixed	1.81 - 3.44	g/L
	days after infection				
Virus production	Lactate concentration one	Lac_1d_post_inf	Performance	0.81 - 1.73	g/L
	day after infection		parameter		
Virus production	Lactate concentration two	Lac_2d_post_inf	Performance	1.07 - 2.53	g/L
	days after infection		parameter		
Virus production	Ammonia concentration one	Amo_1d_post_inf	Performance	1.73 - 2.64	g/L
	day after infection		parameter		
Virus production	Ammonia concentration two	Amo_2d_post_inf	Performance	2.35 - 3.00	mM
	days after infection		parameter		
Virus production	Glutamine concentration	Gln_1d_post_inf	Performance	1.22 - 2.13	mM
	one day after infection		parameter		
Virus production	Glutamine concentration	Gln_2d_post_inf	Performance	0.81 - 1.21	mM
	two days after infection		parameter		

Process stage	Parameter	Short name	Туре	Tested range	Unit
Virus production	Normalized titer: ratio of	Titer_sample_vs_intctrl	Performance	8.30 - 72.3	%
	sample/internal control		parameter		
	infectivity titer				

Table S4 Marks of secondary correlation analysis

Parameters	Negative mark	Positive mark	Total mark
Transfer_VCD	0	3	3
Tryp_exposure_time	0	0	0
Shear35	-4	1	-3
Shear5	-7	0	-7
MC_cell_sep	-1	0	-1
Ino_VCD	-4	0	-4
Ex_post_att	-2	3	1
Tryp_post_ex	-10	0	-10
EDTA_post_ex	-10	0	-10
CCI	-4	0	-4

 Table S5 Levels of selected operation parameters in original and optimized processes

Parameter	Original level*1	Optimized level*2
EDTA concentration after medium exchange (mM)	<0.64	<0.16
TrypLE concentration after medium exchange (X)	<0.08	<0.02
Maximum shear stress of last 5 min during digestion (N/m²)	1.02	0.38
CCI (10 ⁵ cells/cm ²)	1.5 ± 0.2	1.1±0.2
Inoculation density (10 ⁵ cells/cm ²)	0.15±0.3	0.12 ± 0.3

^{*1} Average level of historical data.

Table S6 Abbreviations

Abbreviation	Full name	
B2B	Bead-to-bead transfer	
BM	Basal medium	
BR	Bioreactor	
CCI	Cell concentration at the point of infection	
CF	Cell factory	
DMEM	Dulbecco's modified eagle medium	
DO	Dissolved oxygen	
DPBS	Dulbecco's phosphate-buffered saline	
FBS	Fetal bovine serum	
HPI	Hour post-infection	
HSV-1	Herpes simplex virus type-1	
JS	Just suspended	
OV	Oncolytic virus	
PCA	Principal component analysis	
rpm	Revolutions per minute	
SCM	Serum-containing medium	
SP	Spinner flask	
SRM	Serum-reduced medium	
STI	Soybean trypsin inhibitor	
VCD	Viable cell density	
VIA	Viability	

^{*2} Theoretical level for process design. Applied level may fluctuate around 20% due to operation deviations.