

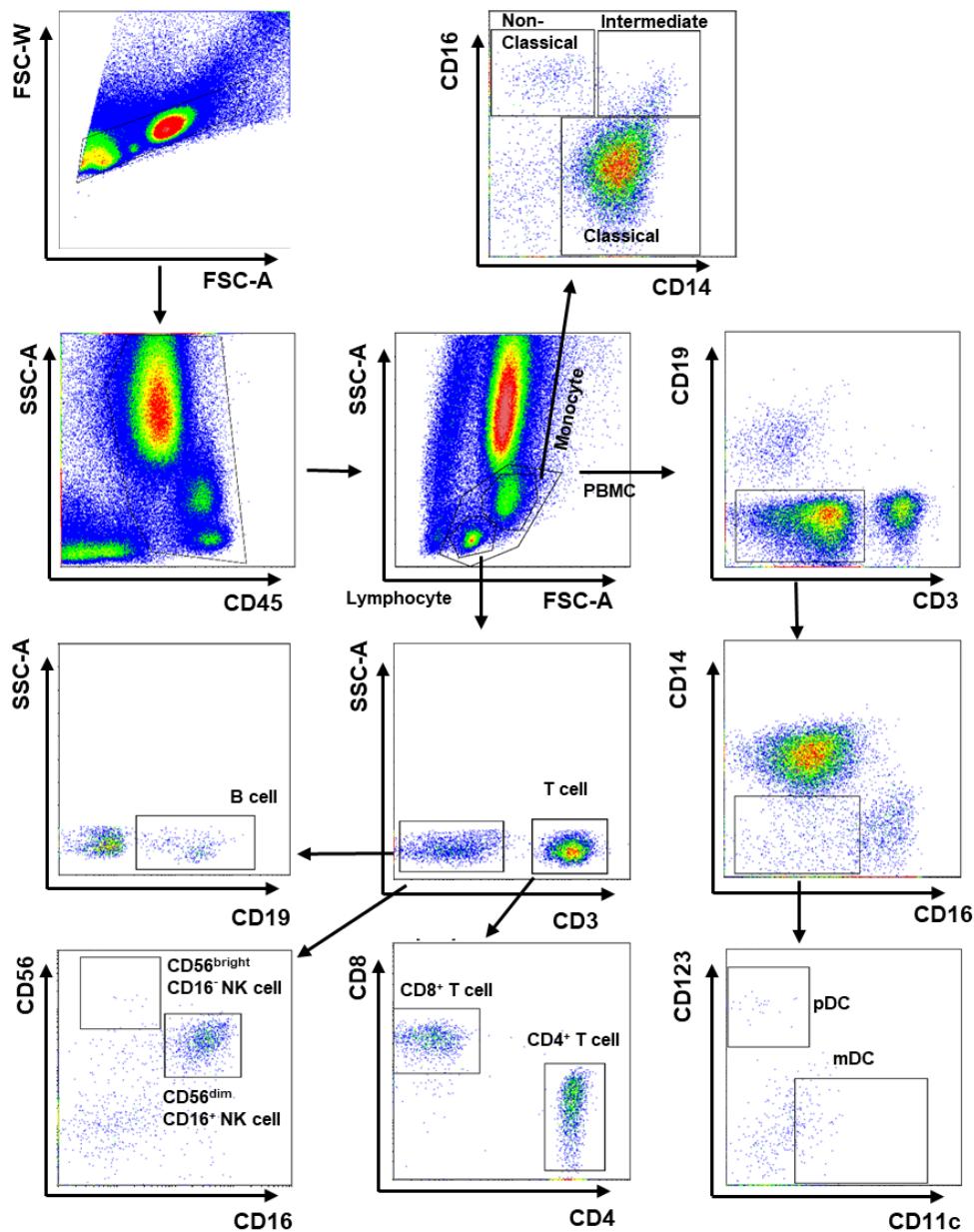
## SUPPLEMENTARY DATA

# Elevated CD4<sup>+</sup> T Cell Senescence Associates with Impaired Immune Responsiveness in Severe COVID-19

Jie Zhang<sup>1,2,3#</sup>, Chun Chang<sup>1#</sup>, Zhaoyuan Liang<sup>2</sup>, Tingting Hu<sup>1</sup>, Zhongnan Yin<sup>2,3</sup>, Ying Liang<sup>1</sup>, Ting Zhang<sup>2,3</sup>, Yanling Ding<sup>1</sup>, Xianlong Li<sup>2</sup>, Xiaoyan Gai<sup>1</sup>, Xiaoxue Yang<sup>2,3</sup>, Xin Li<sup>1</sup>, Xixuan Dong<sup>2,3</sup>, Jiaqi Ren<sup>1</sup>, Yafei Rao<sup>1</sup>, Jun Wang<sup>1</sup>, Jianling Yang<sup>2</sup>, Lixiang Xue<sup>2,3\*</sup>, Yongchang Sun<sup>1\*</sup>

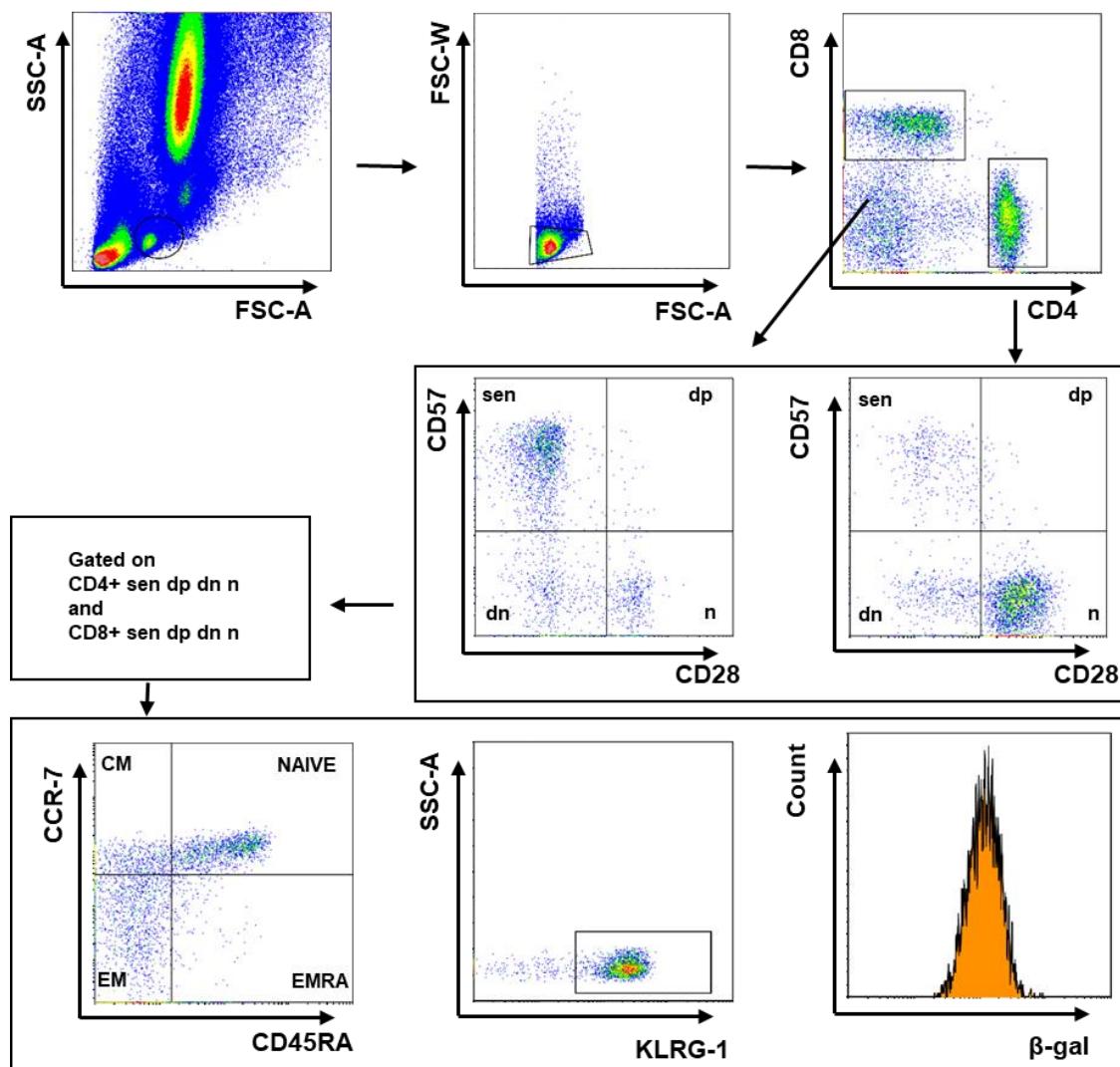
# SUPPLEMENTARY DATA

## Gating strategy of S1 Panel



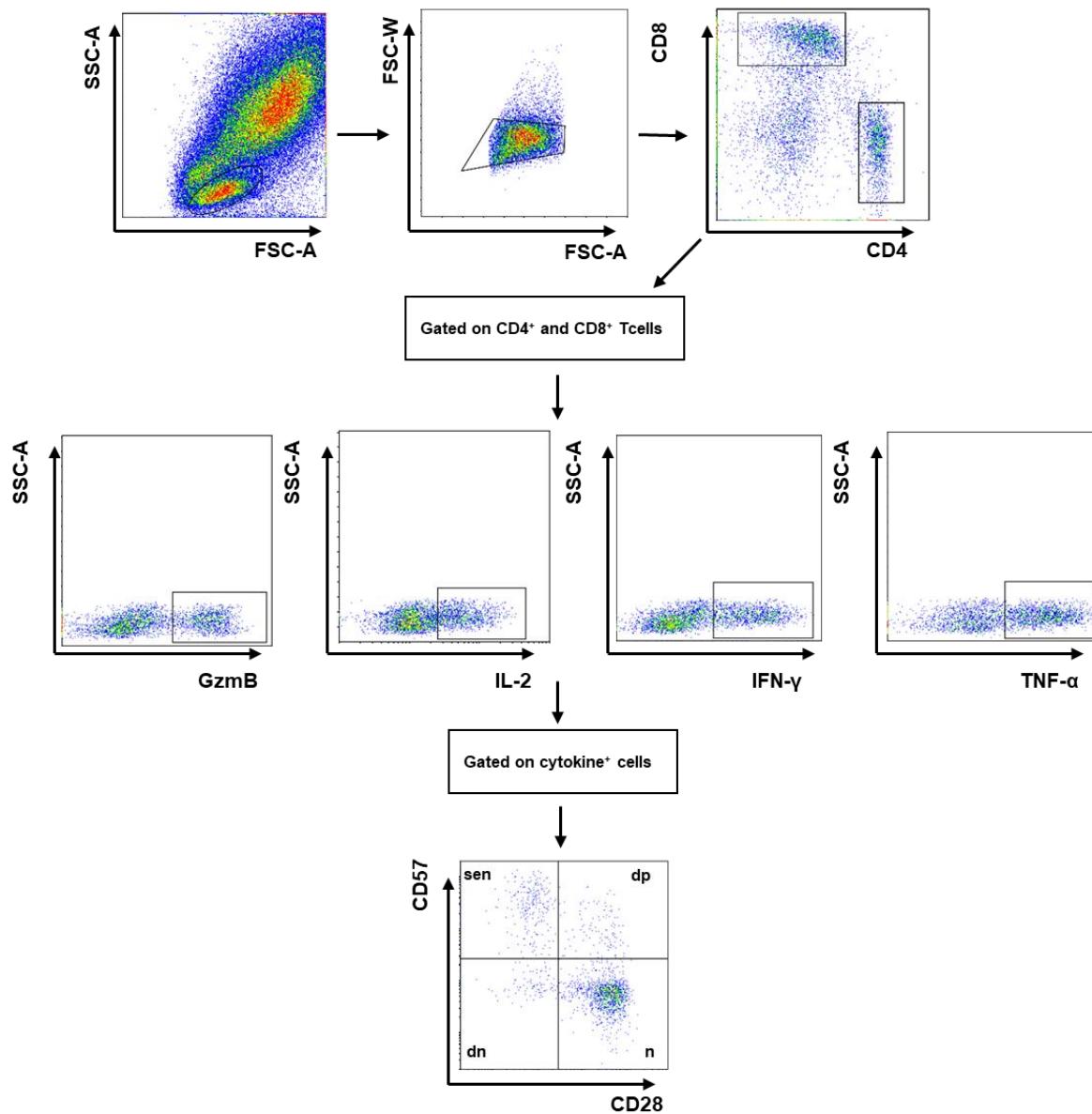
# SUPPLEMENTARY DATA

## Gating strategy of S2 Panel



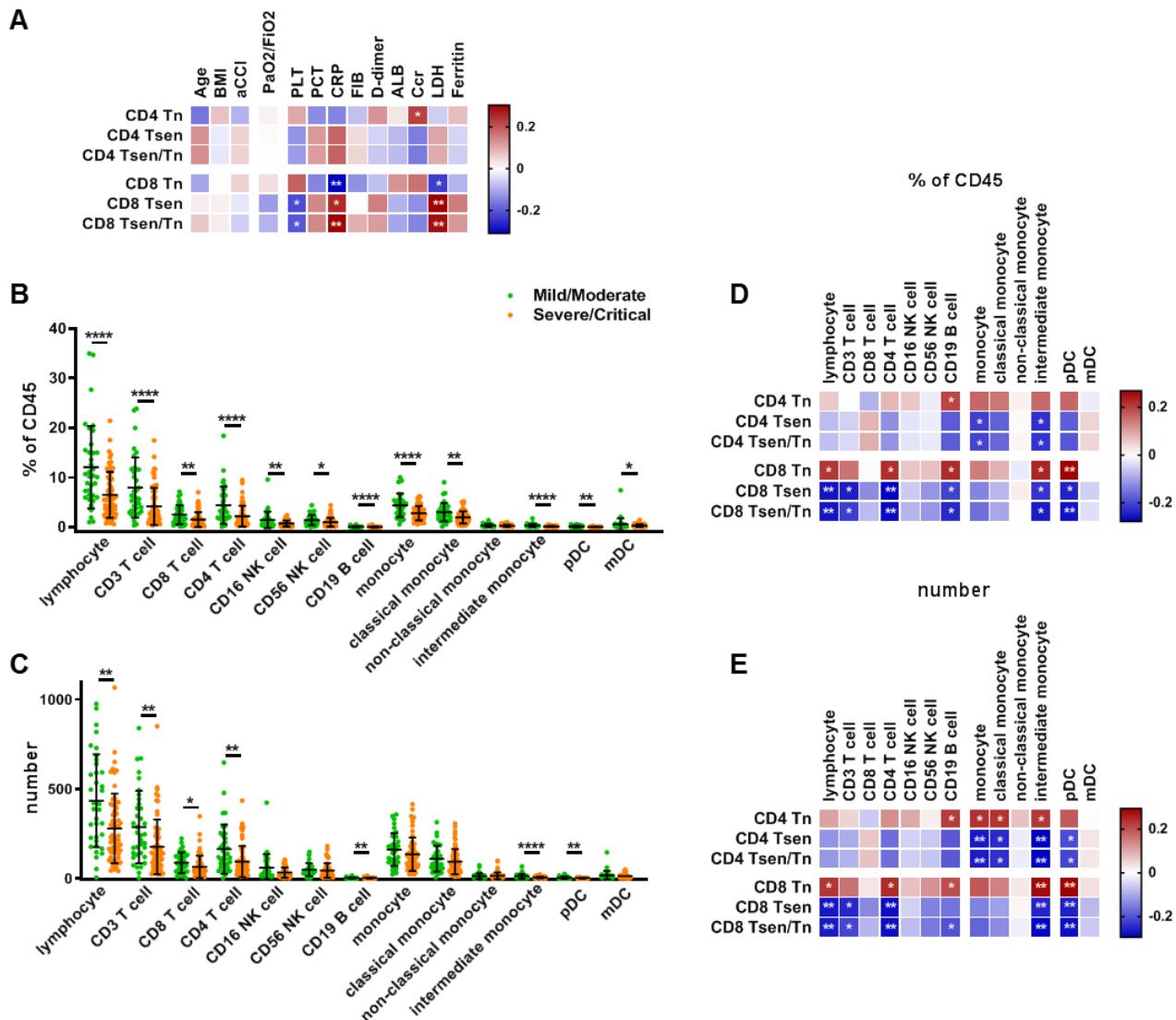
# SUPPLEMENTARY DATA

## Gating strategy of S3 Panel



**Supplementary Figure 1.** The gating strategy of S1, S2 and S3 Panel.

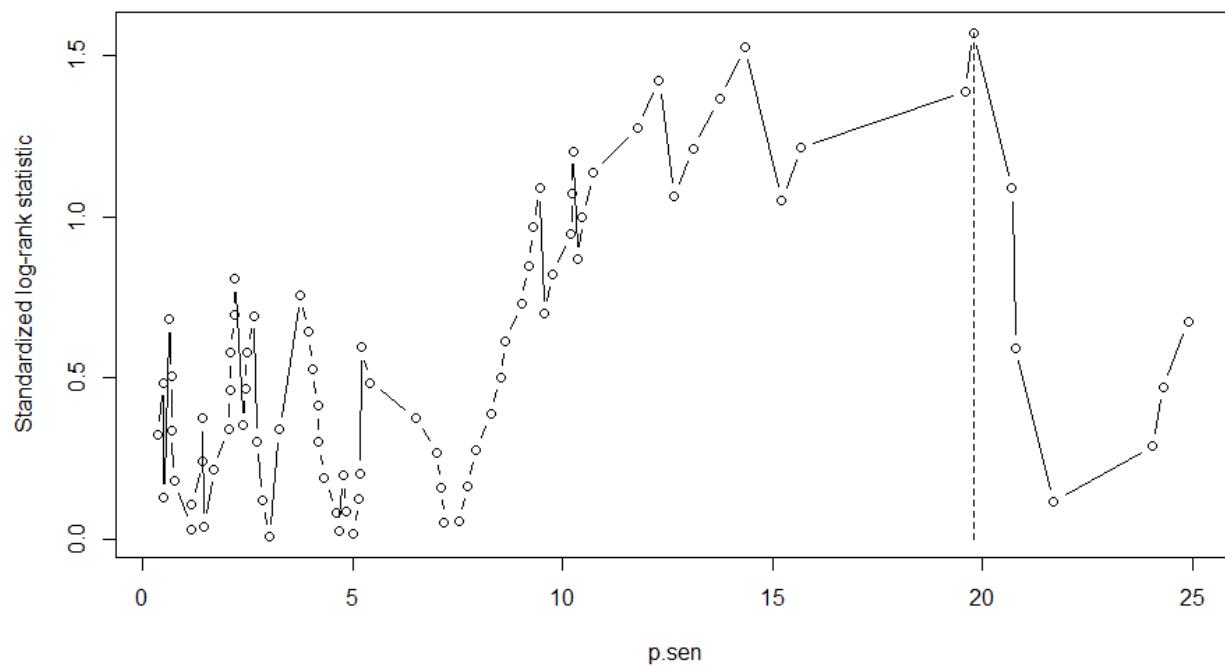
## SUPPLEMENTARY DATA



**Supplementary Figure 2.** (A) Correlations between the 3 subsets (Tn, Tsen, Tsen/Tn) of T cells and clinical phenotypes ( $n=100$ ). (B) The percentage of different immune cell subtypes in  $CD45^+$  white blood cells of mild/moderate ( $n=36$ ) or sever/critical ( $n=64$ ) patients. (C) The number of different immune cell subtypes in mild/moderate ( $n=36$ ) or mild/moderate ( $n=64$ ) patients. (D) Correlations between the 3 subsets (Tn, Tsen, Tsen/Tn) of T cells and the percentage of different immune cell subtypes in  $CD45^+$  white blood cells ( $n=100$ ). (E) Correlations between the 3 subsets (Tn, Tsen, Tsen/Tn) of T cells and the number of different immune cell subtypes ( $n=100$ )

\*,  $p < 0.05$ ; \*\*,  $p < 0.01$ ; \*\*\*,  $p < 0.001$ ; \*\*\*\*,  $p < 0.0001$ .

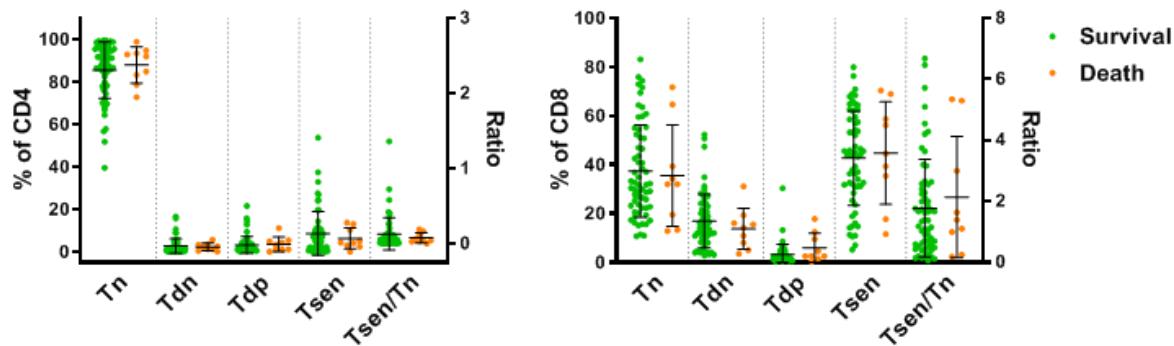
## SUPPLEMENTARY DATA



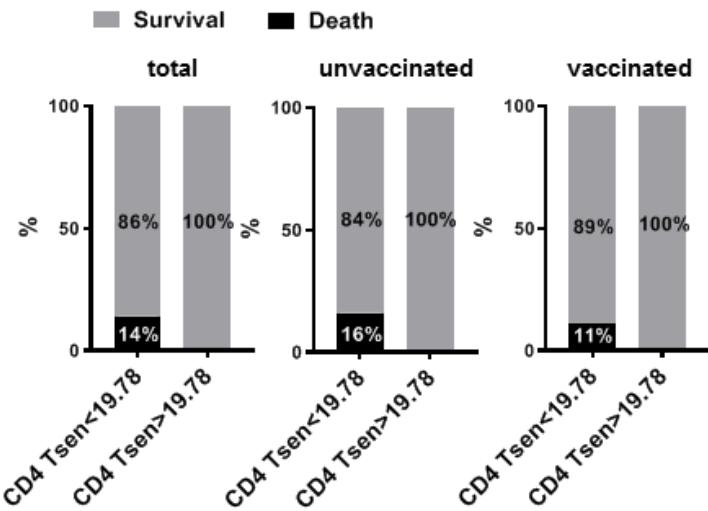
**Supplementary Figure 3. Optimization of CD4 Tsen cut-off according to maximization of log-likelihood ratio method.** Variation of death rate according to circulating senescent lymphocytes (% CD28<sup>-</sup>CD57<sup>+</sup> among CD4<sup>+</sup> T-cells)

## SUPPLEMENTARY DATA

**A**

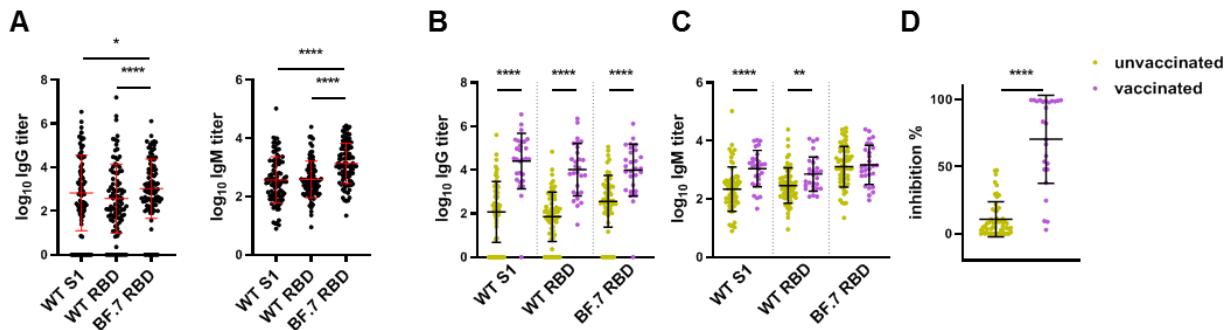


**B**



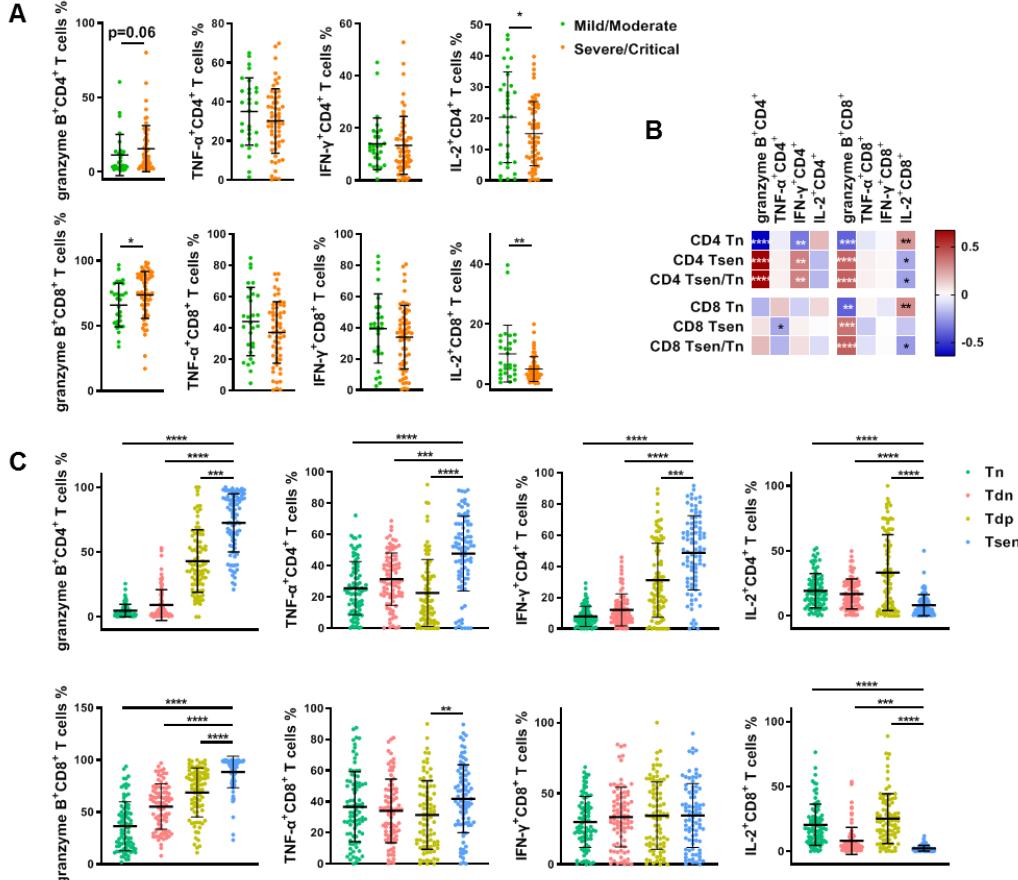
**Supplementary Figure 4. CD4<sup>+</sup> Tsens was not related to long-term outcomes of COVID-19.** (A) The percentage of CD4<sup>+</sup> and CD8<sup>+</sup> T cell subsets: CD28<sup>+</sup>CD57<sup>-</sup> (Tn), CD28<sup>-</sup>CD57<sup>-</sup> (Tdn), CD28<sup>+</sup>CD57<sup>+</sup> (Tdp), and CD28<sup>-</sup>CD57<sup>+</sup> (Tsen) and Tsen/Tn ratio in the survival (n=61) compared to the death (n=9). Groups were compared using Mann–Whitney U-test. Bars show mean with SD. (B) Long-term (One year) survival rates in patients with COVID-19 stratified by levels of CD4 Tsen (the entire cohort: CD4 Tsen≤19.78%, n=63 versus CD4 Tsen>19.78%, n=7; unvaccinated group: CD4 Tsen≤19.78%, n=45 versus CD4 Tsen>19.78%, n=5; vaccinated group: CD4 Tsen≤19.78%, n=18 versus CD4 Tsen>19.78%, n=2). P values for difference between survival rates were calculated using Fisher exact test.

# SUPPLEMENTARY DATA



**Supplementary Figure 5. The spike specific antibody titers and the inhibition rate of neutralization antibody in COVID-19 patients.** (A) The titer of IgG or IgM in plasma against the protein of WT S1, WT RBD and BF.7 RBD (n=97). (B-C) The titer of IgG or IgM against the protein of WT S1, WT RBD and BF.7 RBD in unvaccinated (n=55) or vaccinated patients (n=26). (D) The inhibition rate of neutralization antibody in unvaccinated (n=55) or vaccinated patients (n=26)

\*, p<0.05; \*\*, p <0.01; \*\*\*, p <0.001 ; \*\*\*\*, p<0.0001.



**Supplementary Figure 6. The percentage of T cells that release cytokines.** (A) The percentage of different cytokines (granzyme B, TNF- $\alpha$ , IFN- $\gamma$ , IL-2) released CD4 $^{+}$  or CD8 $^{+}$  T cells in mild/moderate (n=29) or severe/critical (n=58) patients . (B) Correlations between the percentage of 3 subsets of T cells (Tn, Tsen, Tsen/Tn) and the percentage of cytokine released T cells (n=87). (C) The percentage of different cytokines (granzyme B, TNF- $\alpha$ , IFN- $\gamma$ , IL-2) released subsets (Tn, Tdn, Tdp, Tsen) of CD4 $^{+}$  and CD8 $^{+}$  T cells (n=87). \*, p<0.05; \*\*, p <0.01; \*\*\*, p <0.001 ; \*\*\*\*, p<0.0001.

# SUPPLEMENTARY DATA

Supplementary Table 1. Reagents used in this research.

Reagent in Different Panels	Com	Catalog	Clone
<b>Panel S1</b>			
PE anti-human CD4	Biolegend	300508	RPA-T4
PerCP-cy5.5 anti-human CD8	Biolegend	344710	SK1
PE-Cy7 anti-human CD56	Biolegend	318318	HCD56
Brilliant Violet 605 anti-human CD16	Biolegend	360727	B73.1
PE-CF594 anti-human CD14	Biolegend	325634	HCD14
APC-Cy7 anti-human CD3	Biolegend	300426	UCHT1
APC anti-human CD123	Biolegend	306012	6H6
Brilliant Violet 421 anti-human CD19	Biolegend	302233	HIB19
Brilliant Violet 510 anti-human CD11c	Biolegend	371513	S-HCL-3
Brilliant Violet 650 anti-human CD45	Biolegend	304043	HI30
<b>Panel S2</b>			
APC-Cy7 anti-human CD4	Biolegend	357416	A161A1
PerCP-cy5.5 anti-human CD8	Biolegend	344710	SK1
PE-Cy7 anti-human CD28	Biolegend	302926	CD28.2
Brilliant Violet 421 anti-human CD57	Biolegend	359608	HNK-1
Brilliant Violet 650 anti-human CD45RA	Biolegend	304135	HI100
PE-CF594 anti-human CCR7	Biolegend	353236	G043H7
APC anti-human KLRG-1	Biolegend	367715	SA231A2
Cellular Senescence Detection Kit-SPiDER-β Gal		SG03	
<b>Panel S3</b>			
APC-Cy7 anti-human CD4	Biolegend	357416	A161A1
PerCP-cy5.5 anti-human CD8	Biolegend	344710	SK1
PE-Cy7 anti-human CD28	Biolegend	302926	CD28.2
Brilliant Violet 421 anti-human CD57	Biolegend	359608	HNK-1
FITC anti-human IFN-γ	Biolegend	359607	HNK-1
PE-CF594 anti-human TNF-α	Biolegend	502946	Mab11
APC anti-human GranzymeB	Biolegend	372204	QA16A02
Brilliant Violet 605 anti-human IL-2	Biolegend	500331	MQ1-17H12
<b>Panel S4</b>			
APC-Cy7 anti-human CD4	Biolegend	357416	A161A1
PerCP-cy5.5 anti-human CD8	Biolegend	344710	SK1
PE-Cy7 anti-human CD28	Biolegend	302926	CD28.2
Brilliant Violet 421 anti-human CD57	Biolegend	359608	HNK-1
APC anti-human CD27	Biolegend	302809	O323
<b>Panel S5</b>			

## SUPPLEMENTARY DATA

Brilliant Violet 510 anti-human CD4	Biolegend	357419	A161A1
Brilliant Violet 650 anti-human CD8	Biolegend	344729	SK1
PE-Cy7 anti-human CD28	Biolegend	302926	CD28.2
Brilliant Violet 421 anti-human CD57	Biolegend	359608	HNK-1
p21 Alexa Fluor® 488	Cell Signaling	5487	12D1
p16 PE	Cell Signaling	82548	D7C1M
<b>Panel S6</b>			
APC-Cy7 anti-human CD4	Biolegend	357416	A161A1
PerCP-cy5.5 anti-human CD8	Biolegend	344710	SK1
PE-Cy7 anti-human CD28	Biolegend	302926	CD28.2
Brilliant Violet 421 anti-human CD57	Biolegend	359608	HNK-1
APC-Cy7 anti-human CD3	Biolegend	300426	UCHT1
APC anti-human CD40L	Biolegend	310809	24-31
<b>Panel S7</b>			
CFSE	Biolegend	423801	
APC-Cy7 anti-human CD19	Biolegend	302233	HIB19
PE anti-human CD38	Biolegend	303505	HIT2
PE-Cy7 anti-human IgD	Biolegend	348209	IA6-2

**Supplementary Table 2.** Demographics, Characteristics, and Clinical Features of Patients With Coronavirus Disease 2019<sup>a</sup>.

Characteristics	All cases (n=100)	CD4 Tsen low (n=84)	CD4 Tsen high (n=16)	P-value <sup>b</sup>
Age, y(n)	80.10±9.89	79.98±9.89	80.69±10.22	0.797
Sex, male	64 (64%)	53 (63.1%)	11 (68.8%)	0.666
BMI, kg/m <sup>2</sup>	23.81±3.91 (96)	23.84±3.97	23.69±3.74	0.895
<18.5	7 (7.3%)	5 (6.2%)	2 (13.3%)	0.578
18.5-23.9	40 (41.7%)	35 (43.2%)	5 (33.3%)	
24.0-27.9	37 (38.5%)	30 (37.0%)	7 (46.7%)	
≥28.0	12 (12.5%)	11 (13.6%)	1 (6.7%)	
Smoking History, yes (n)	35 (35.0%)	27 (32.1%)	8 (50.0%)	0.170
Any comorbidity				
Diabetes	25 (25%)	22 (26.2%)	3 (18.8%)	0.753
Hypertension	52 (52.0%)	40 (47.6%)	12 (75.0%)	<b>0.045</b>
Cardiovascular diseases	24 (24.0%)	23 (27.4%)	1 (6.3%)	0.135
COPD	11 (11.0%)	8 (9.5%)	3 (18.8%)	0.519
Asthma	4 (4.0%)	3 (3.6%)	1 (6.3%)	0.508
aCCI	4.92±1.23	4.69±1.25	4.85±1.33	0.431
Signs and symptoms				
Fever	82 (82.0%)	68 (81.0%)	14 (87.5%)	0.787
Cough	85 (85.0%)	70 (83.3%)	15 (93.8%)	0.492
Sputum Production	80 (80.0%)	66 (78.6%)	14 (87.5%)	0.633

## SUPPLEMENTARY DATA

Dyspnea	60 (60.0%)	52 (61.9%)	8 (50.0%)	0.373
Medication				
Glucocorticoids	84 (84.0%)	71 (81.3%)	84 (84.0%)	0.743

BMI, body mass index; aCCI, age-adjusted Charlson Comorbidity Index; COPD, chronic obstructive pulmonary disease.

a. Continuous variables were presented as mean  $\pm$  SD (n); categorical variables are shown as n (%). Medication and respiratory support information was recorded during entire hospital stay; other information was recorded at admission.

b. P-values were from t-test for continuous data and from  $\chi^2$  test for categorical data.

**Supplementary Table 3.** Laboratory Characteristics on Admission for Severely and Critically Ill Patients With Coronavirus Disease 2019<sup>a</sup>.

Characteristics	All cases (n=100)	CD4 Tsen low (n=84)	CD4 Tsen high (n=16)	P-value <sup>b</sup>
Blood routine				
White blood cell count, 10 <sup>9</sup> /L	7.56 $\pm$ 2.9	7.42 $\pm$ 3.05	7.64 $\pm$ 2.77	0.797
<3.5	2 (2.0%)	2 (2.4%)	0 (0.0%)	0.429
3.5~9.5	75 (75.0%)	61 (72.6%)	14 (87.5%)	
>9.5	23 (23.0%)	21 (25.0%)	2 (12.5%)	
Neutrophil count, 10 <sup>9</sup> /L	6.39 $\pm$ 2.77	6.42 $\pm$ 2.84	6.20 $\pm$ 2.46	0.785
Lymphocyte count, 10 <sup>9</sup> /L	0.78 $\pm$ 0.48	0.78 $\pm$ 0.49	0.76 $\pm$ 0.38	0.686
Platelet count, 10 <sup>9</sup> /L	214.62 $\pm$ 81.04	216.29 $\pm$ 84.38	201.75 $\pm$ 69.30	0.492
Hemoglobin, g/L	121 $\pm$ 28.41	121.96 $\pm$ 32.37	114.69 $\pm$ 17.60	0.056
Inflammatory markers				
Procalcitonin, ng/mL	0.38 $\pm$ 1.10	0.29 $\pm$ 0.73	0.99 $\pm$ 2.30	0.710
<0.1	53 (54.1%)	36 (48.09%)	7 (50.0%)	0.883
0.1~0.3	30 (30.6%)	26 (34.7%)	4 (28.6%)	
>0.3	15 (15.3%)	13 (17.3%)	3 (21.4%)	
C-reactive protein, mg/L	24.76 $\pm$ 40.66	25.14 $\pm$ 37.66	44.04 $\pm$ 63.40	0.037
$\leq$ 8	41 (42.7%)	36 (44.4%)	45 (55.6%)	0.424
>8	55 (57.3%)	5 (33.3%)	10 (66.7%)	
Coagulation function				
D-dimer, ug/mL	2.77 $\pm$ 5.00	2.86 $\pm$ 4.77	1.08 $\pm$ 1.17	0.387
$\leq$ age/100	47 (49.0%)	38 (47.5%)	42 (52.5%)	0.523
> age/100	49 (51.0%)	9 (56.3%)	7 (43.8%)	
Serum biochemical indicators				
Serum albumin level, g/L	31.89 $\pm$ 4.87	31.89 $\pm$ 4.05	31.19 $\pm$ 4.32	0.531
Creatinine, $\mu$ mol/L	98.34 $\pm$ 92.85	96.75 $\pm$ 71.64	99.23 $\pm$ 103.42	0.413
Serum urea nitrogen, mmol/L	9.96 $\pm$ 8.53	10.29 $\pm$ 9.20	8.48 $\pm$ 4.52	0.472
Total bilirubin, $\mu$ mol/L	12.02 $\pm$ 5.66	12.52 $\pm$ 6.28	9.67 $\pm$ 2.96	0.227
Alanine Aminotransferase, U/L	37.49 $\pm$ 38.22	38.50 $\pm$ 38.75	31.85 $\pm$ 19.36	0.525
Aspartate Aminotransferase, U/L	42.47 $\pm$ 35.11	42.86 $\pm$ 27.99	41.31 $\pm$ 19.70	0.821
Creatine kinase, U/L	109.06 $\pm$ 182.15	111.80 $\pm$ 192.01	94.07 $\pm$ 117.18	0.151
Creatine kinase-MB, U/L	15.56 $\pm$ 31.62	17.11 $\pm$ 34.32	7.50 $\pm$ 3.03	0.024

a. Continuous variables were presented as median (interquartile range); categorical variables are shown as n (%).

b. P-values were from t-test for normally distributed continuous data and from Mann-Whitney U test for abnormally distributed continuous data. P-values were from  $\chi^2$  test for categorical data.

# SUPPLEMENTARY DATA

**Supplementary Table 4.** Plasma soluble factors in COVID-19 patients.

Cytokines	Mild/Moderate (n=19)	Severe/Critical (n=34)	P-value <sup>b-0</sup>
IL-2	63.46±15.86	58.34±9.993	0.3852
IL4	39.51±30.01	12.17±4.094	0.2385
IL-10	5.809±2.370	12.39±2.627	0.1013
IL-6	22.02±9.565	71.54±23.36	0.1300
IL-17a	5.221±1.582	3.519±0.8046	0.2913
TNF- $\alpha$	14.86±7.479	14.28±3.242	0.9345
sFas	544.0±70.80	701.1±63.76	0.1240
sFasL	18.27±3.782	13.22±1.946	0.1940
IFN- $\gamma$	123.9±48.29	180.7±49.86	0.4578
GranzymeA	171.2±95.14	102.5±19.70	0.3659
GranzymeB	467.8±133.9	473.7±101.8	0.9723
Perforin	1050±106.2	914.5±72.63	0.2859
Granulysin	2399±143.8	2423±195.5	0.9341
IL8	11.21±4.082	16.36±5.718	0.5351
IP10	522.2±182.0	1228±255.5	0.0612
Eotaxin	79.40±10.17	75.35±6.513	0.7271
TARC	40.55±7.107	38.25±6.708	0.8266
MCP1	164.4±37.54	278.2±68.7	0.2434
RANTES	953.0±99.30	931.9±119.4	0.9056
MIP1 $\alpha$	7.077±3.271	9.218±7.993	0.8466
MIG	492.9±111.6	758.0±141.0	0.2054
ENA78	19.69±4.668	23.79±5.070	0.5937
MIP3 $\alpha$	10.84±3.767	10.23±2.242	0.8816
GRO $\alpha$	23.08±6.923	17.57±3.826	0.4516
ITAC	30.68±5.317	61.00±14.72	0.1389
MIP1 $\beta$	2.773±0.5034	4.034±1.288	0.4791
TSLP	7.891±5.190	9.460±4.710	0.8331
IL1 $\alpha$	37.59±13.15	26.94±8.934	0.4941
IL1 $\beta$	11.53±3.404	12.43±2.684	0.8386
GM-CSF	3.230±2.045	6.398±2.057	0.3198
IFN $\alpha$ 2	5.942±1.260	4.956±0.9000	0.5221
IL23	53.35±13.81	46.99±6.493	0.6383
IL12p40	85.19±18.50	75.34±17.31	0.7169
IL12p70	2.748±1.025	4.177±0.7668	0.2697
IL15	152.8±39.19	115.3±22.73	0.3779
IL18	89.88±32.42	137.735.91	0.3796
IL11	108.9±33.62	140.5±28.84	0.4965
IL27	73.97±16.57	66.37±18.42	0.7843
IL33	63.12±27.80	43.12±13.72	0.4729
IL5	3.911±1.476	7.316±1.396	0.1231
IL13	7.809±4.382	16.50±4.347	0.1993
IL9	23.78±15.36	11.18±1.710	0.2836
IL17F	5.537±4.154	3.267±0.609	0.4798
IL22	6.358±2.188	4.644±0.8429	0.3904

a. Different cytokine concentrations (pg/mL) of mild/moderate or severe/critical patients in plasma were presented as mean ± SEM

b. P-values were from t-test for normally distributed continuous data and from Mann-Whitney U test for abnormally distributed continuous data.