Panel	Short Description	Folder	File Path	Notes					
Figure 1:	Study Design and a Multi-parameter Disease Severity Metric	(DSM)		·					
А	Patient overview	A3	overview/patient.and.sample.overview.Rmd						
В	Experiment overview	NA	NA						
С	DSM parameter hm	A3	dsm/citeseq.patient.dsm.Rmd	Produces the DSM end point file					
D	DSM parameter model coef	A3	dsm/citeseq.patient.dsm.Rmd						
E	DSM validation box plot	A3	dsm/brescia.dsm.validation.Rmd, dsm/citeseq.patient.dsm.Rmd						
Figure 2:	Figure 2: A Multimodal Single Immune Cell Atlas of COVID-19 Reveals Cell Populations Associated with Disease and Disease Severity (DSM)								
Α	CITE hm	A2	3.UMAPandProteinVis.Figure2.R						
В	UMAP	A2	3.UMAPandProteinVis.Figure2.R						
С	CITE hm	A2	3.UMAPandProteinVis.Figure2.R						
D-I	Mono UMAP	A2	3.UMAPandProteinVis.Figure2.R						
Figure 3:	Figure 3: Cell-type-specific Gene Expression Signatures of COVID-19 Disease Status and Severity								
Α	COVIDvsHC bubble	A4	figures/gsea_bubble_plots.R						
В	DSM bubble	A4	figures/gsea_bubble_plots.R						
С	Mono IFN hm	A1	DEplot.subject.hm.Rmd						
D	Mono IFN trajecotry	A1	modulescore.TSO.Rmd						
E	TSO IFN coef	A1	NES.PC1groupTSO.interaction.Rmd						
F	pDC apopt hm	A1	DEplot.subject.hm.Rmd						
G	corr hm and scatters	A4	figures/fig3_heatmap_and_scatterplots.R						
Figure 4:	Cell-type-specific Gene Expression Profiles of Select Signatur	es and D	isease Severity Network	·					
А	severity network	A3	severity.network/severity.network.Rmd						
В	NK FA IL15 corr	A1	modulescore.corr.Rmd						
С	IL15 DSM corr	A1	modulescore.corr.Rmd						
D	NK FA hm	A1	DEplot.subject.hm.Rmd						
E	NK NFkB hm	A1	DEplot.subject.hm.Rmd						
F	NK IL1 hm	A1	DEplot.subject.hm.Rmd						
G	NK FA inflam corr	A1	modulescore.corr.Rmd						
Н	NK IFNG	A1	DEplot.subject.hm.Rmd						
I	NK FA mTOR, IFNG corr	A1	modulescore.corr.Rmd						
J	NK FA trajectory	A1	modulescore.TSO.Rmd						
K	NK inflam trajectory	A1	modulescore.TSO.Rmd						
L	IL15 trajectory	A1	modulescore.corr.Rmd						
Figure 5: Single Cell and Clonal Expansion Analysis in CD4+ and CD8+ T cells and Exhaustion Assessment in Clonal CD8+ T cells									
Α	CD8 cluster hm	A2	4.TcellReclustering.R						
В	CD8 CITE hm	A2	4.TcellReclustering.R						

C CD8 cluster FC A4 figures/fig4_plot_cd4_8_fig4_cluster_volcano.R D CD8 cluster overlap w celltypes A2 4.TcellReclustering.R E clonality DSM A4 figures/clonality/plot_pcgroup_onset_assoc_for_paper.R F CD8 exhaustion marker FC A4 figures/tcell_exhaustion/surface_marker_coefficients_with_pval.R G exhaustion gene GSEA curve A4 figures/tcell_exhaustion/exhaustion_fgsea_plots.R							
F CD8 exhaustion marker FC A4 figures/tcell_exhaustion/surface_marker_coefficients_with_pval.R							
G exhaustion gene GSEA curve A4 figures/tcell_exhaustion/exhaustion_fgsea_plots.R							
Figure 6: Analyses of Longitudinal Data and Timing Effects Suggest an Immune Response Juncture Whose Immune Profile is Predictive of Deceased-or-Recovery Outcomes							
A Mono cell trajectory A1 cell.freq.plot.Rmd							
B Mono Neut count A3 critical.juncture/dsm.high.v.low.cytokine.Rmd							
C NES TSO group A1 NES.PC1onsetgroup.interaction.Rmd							
D juncture bubble A4 figures/gsea_bubble_plots.R							
E NK, mono juncture trajectory A1 modulescore.TSO.Rmd							
F cytokine trajectory DSM A3 critical.juncture/dsm.high.v.low.cytokine.Rmd							
G scheme NA NA							
H Effect sizes of cytokine delta of delta A3 critical.juncture/deceased.v.recovered.cytokine.Rmd							
I cytokine trajectory decease recover A3 critical.juncture/deceased.v.recovered.cytokine.Rmd							
Supplementary Figures							
Figure S1: Subject treatment timeline							
A subject reatment timeline A3 overview/patient.and.sample.overview.Rmd							
B CITESeq patient DSM violin plot by WHO Scores A3 dsm/citeseq.patient.dsm.Rmd							
Figure S2: cell freq description							
A flow CITE corr A1 cell.freq.corr.Rmd							
B cell freq DSM A1 cell.freq.plot.Rmd							
C cell freq hm A1 cell.freq.plot.Rmd							
Figure S3: Cell-type-specific Gene Expression Signatures of COVID-19 Disease Status and Severity							
A DSM-low TSO bubble A4 figures/gsea_bubble_plots.R							
B DSM-high TSO bubble A4 figures/gsea_bubble_plots.R							
C mono translation hm A1 DEplot.subject.hm.Rmd							
D mono translation trajectory A1 modulescore.TSO.Rmd							
E German data pDC apopt hm1 A1 Schulte-Schrepping/cohort1.DEsubject.hm.Rmd							
F German data pDC apopt hm2 A1 Schulte-Schrepping/cohort2.DEsubject.hm.Rmd							
G German data pDC apopt GSEA curve A4 figures/validations/pdc_apoptosis/*							
Figure S4: Cell-type-specific Gene Expression Signatures of COVID-19 Disease Status and Severity							
A pDC apopt v. oxidative stress corr A3 cytokine.geneset.correlation/citeseq.T0.cytokine.pathway.corr							
B NK FA IL15 residual corr A3 cytokine.geneset.correlation/citeseq.T0.cytokine.pathway.corr							
C German data NK FA hm1 A1 Schulte-Schrepping/cohort1.DEsubject.hm.Rmd							

D	German data NK FA hm2	A1	Schulte-Schrepping/cohort2.DEsubject.hm.Rmd	
E	German data NK FA GSEA curve	A4	figures/validations/NK fattyacid gsea curves cohorts 1 2.R	
F	German data FA NFkB score corr	A1	Schulte-Schrepping/NK.modulescore.corr.Rmd	
G-H	Mono inflam hm	A1	DEplot.subject.hm.Rmd	
Figure S5	Exogeneous corticosteroid effect			
Α	glucocorticoid GSEA curve	A4	dsm_steroid_enrichment.R	
В	related scores and glucocorticoid score corr	A3	cytokine.geneset.correlation/citeseq.T0.cytokine.pathway.corr	
С	related scores and TSC22D3 corr	A3	cytokine.geneset.correlation/citeseq.T0.cytokine.pathway.corr	
D	TSC22D3 boxplot	A1	steroiduse.TSC22D3.Rmd	
Figure S6	: CD4 and CD8 exhaustion validation	•		
Α	CD4 cluster hm	A2	4.TcellReclustering.R	
В	CD4 CITE hm	A2	4.TcellReclustering.R	
С	CD4 cluster FC	A4	figures/fig4_plot_cd4_8_fig4_cluster_volcano.R	
D	CD4 cluster overlap w celltypes	A2	4.TcellReclustering.R	
E	exhaustion marker RNA	A4	figures/tcell_exhaustion/RNA_surface_marker_coeff_with_pval.R	
F	CD39+PD1+ freq boxplot	A4	figures/tcell_exhaustion/2020_12_09_cd39_pd1.R	
G	exhaustion cutoff boxplot	A4	figures/tcell_exhaustion/cutoff_markers_grid.R	
Н	exhaustion LE enrichment	A4	figures/tcell_exhaustion/wherry_exhaustion_enrichment.R	
I	German data exhaustion bubble	A4	figures/validations/exhaustion_plots.R	
J	German data exhaustion GSEA curve	A4	figures/validations/exhaustion_plots.R	
Figure S7	: Supporting data for critical juncture analysis			
Α	difference of difference bubble	A4	figures/gsea_bubble_plots.R	
В	Deceased v. recovered neutrophils and monocytes	A3	critical.juncture/deceased.v.recovered.cytokine.Rmd	
С	Outcome and severity comparison ES corr	A3	critical.juncture/deceased.v.recovered.cytokine.Rmd	
D	Yale cohort validation ES corr	A3	critical.juncture/cytokine.critical.juncture.validation.Rmd	
E	Deceased v. recovered titers	A3	critical.juncture/deceased.v.recovered.cytokine.Rmd	