Appendix: Modeling the CD8+ T cell immune

2 response to influenza infection in adult and aged

₃ mice

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11 Appendix A

12 A.1 Model Equations

Model	Expressions
MA1	$dV/dt = pV(1-V/k_V) - c_V V T$, $dT/dt = s_T + rTV/(V+k_T) - d_T^{\text{half life}} T$
MA2	$dV/dt = pV(1 - V/k_V) - c_V V T$, $dT/dt = s_T + rTV/(V + k_T) - c_T/(1 + V^2)T - d_T^{\text{half life}}T$
MA3	$dV/dt = pV(1 - V/k_V) - c_V V T$, $dT/dt = s_T + rTV/(V + k_T) - (d_T^{\text{half life}} + d_T^{\text{downregulation}})T$
MA4	$dV/dt = pV(1 - V/k_V) - c_V VT, dT/dt = s_T + rTV/(V + k_T) - c_T/(1 + V^2)T - (d_T^{\text{half life}} + d_T^{\text{downregulation}})T$
MB1	$dU/dt = -\beta UV, dI/dt = \beta UV - d_1TI, dV/dt = pI - cV, dT/dt = s_T + rTV/(V + k_T) - d_T^{half life}T$
MB2	$dU/dt = -\beta UV, dI/dt = \beta UV - d_I TI, dV/dt = pI - cV, dT/dt = s_T + rTV/(V + k_T) - c_T/(1 + V^2)T - d_T^{\text{half life}}T$
MB3	$dU/dt = -\beta UV$, $dI/dt = \beta UV - d_ITI$, $dV/dt = pI - cV$, $dT/dt = s_T + rTV/(V + k_T) - (d_T^{half life} + d_T^{downregulation})T$
MB4	
MC1	$dV/dt = pV(1 - V/k_V) - c_V VT, dT/dt = s_T + rTV - d_T^{\text{half life}}T$
MC2	$dV/dt = pV(1 - V/k_V) - c_V V T$, $dT/dt = s_T + rTV - c_T/(1 + V^2)T - d_T^{\text{half life}}T$
MC3	$dV/dt = pV(1 - V/k_V) - c_V VT$, $dT/dt = s_T + rTV - (d_T^{\text{half life}} + d_T^{\text{downregulation}})T$
MC4	$dV/dt = pV(1 - V/k_V) - c_V VT, dT/dt = s_T + rTV - c_T/(1 + V^2)T - (d_T^{\text{half life}} + d_T^{\text{downregulation}})T$
MD1	$dU/dt = -\beta UV$, $dI/dt = \beta UV - d_I TI$, $dV/dt = pI - cV$, $dT/dt = s_T + rTV - d_T^{half\ life}T$
MD2	$dU/dt = -\beta UV, dI/dt = \beta UV - d_ITI, dV/dt = pI - cV, dT/dt = s_T + rTV - c_T/(1 + V^2)T - d_T^{\text{half life}}T$
MD3	$dU/dt = -\beta UV$, $dI/dt = \beta UV - d_I TI$, $dV/dt = pI - cV$, $dT/dt = s_T + rTV - (d_T^{\text{half life}} + d_T^{\text{downregulation}})T$
MD4	$dU/dt = -\beta UV, dI/dt = \beta UV - d_ITI, dV/dt = pI - cV, dT/dt = s_T + rTV - c_T/(1 + V^2)T - (d_T^{\text{half life}} + d_T^{\text{downregulation}})T$

Table A.1.1. Mathematical form of the 16 models considered.

A.2 Structural Identifiability Analysis

MA1		MA2		N	/IA3	MA4		
Parameters	Identifiability	Parameters	Identifiability	Parameters	Identifiability	Parameters	Identifiability	
k_T	✓			s_T	✓	d_T	✓	
c_V	\checkmark	k_T	\checkmark	k_V	\checkmark	c_T	\checkmark	
		c_V	\checkmark	T(t)	\checkmark	c_V	\checkmark	
k_V	\checkmark	p	\checkmark	r	\checkmark	k_V	\checkmark	
p	\checkmark	r	\checkmark	c_V	\checkmark	p	\checkmark	
T(t)	\checkmark	s_T	\checkmark	d_T	\checkmark	T(t)	\checkmark	
V(t)	\checkmark	k_V	\checkmark	V(t)	\checkmark	V(t)	\checkmark	
r	\checkmark	T(t)	\checkmark	p	\checkmark	r	\checkmark	
s_T	\checkmark	V(t)	\checkmark	k_T	\checkmark	s_T	\checkmark	
		c_T	\checkmark			k_T	\checkmark	

Table A.2.1. Model group A structural identifiabilities. Model group A uses a logistic form for viral dynamics and Michaelis-Menten form for CD8+ T cell proliferation.

N	ИВ1	N	/IB2	N	ИВ3	MB4		
Parameters	Identifiability	Parameters	Identifiability	Parameters	Identifiability	Parameters	Identifiability	
d_I	√	c_T	√	S_T	√	c_T	✓	
V(t)	\checkmark	p		c	\checkmark	V(t)	\checkmark	
r	\checkmark	T(t)	\checkmark	I(t)		r	\checkmark	
		k_T	\checkmark	T(t)	\checkmark	d_I	\checkmark	
c	\checkmark	β	\checkmark	β	\checkmark	c	\checkmark	
k_T	\checkmark	V(t)	\checkmark	d_I	\checkmark	d_T	\checkmark	
U(t)		d_I	\checkmark	U(t)		U(t)		
β	\checkmark	r	\checkmark	r	\checkmark	β	\checkmark	
T(t)	\checkmark	ST	\checkmark	V(t)	\checkmark	T(t)	\checkmark	
p		U(t)		p		p		
s_T	\checkmark	c	\checkmark	d_T	\checkmark	s_T	\checkmark	
I(t)				k_T	\checkmark	I(t)		
		I(t)				k_T	\checkmark	

Table A.2.2. Group B structural identifiabilities. Model group B uses a target cell limited model for viral dynamics and Michaelis-Menten form for CD8+ T cell proliferation.

N	MC1		MC2		ИС3	MC4		
Parameters	arameters Identifiability Parameters Ident		Parameters Identifiability		Parameters Identifiability		Identifiability	
				s_T	✓	d_T	✓	
c_V	\checkmark			k_V	\checkmark	c_T	\checkmark	
		c_V	\checkmark	T(t)	\checkmark	c_V	\checkmark	
k_V	\checkmark	p	\checkmark	r	\checkmark	k_V	\checkmark	
p	\checkmark	r	\checkmark	c_V	\checkmark	p	\checkmark	
T(t)	\checkmark	s_T	\checkmark	d_T	\checkmark	T(t)	\checkmark	
V(t)	\checkmark	k_V	\checkmark	V(t)	\checkmark	V(t)	\checkmark	
\dot{r}	\checkmark	T(t)	\checkmark	p	\checkmark	r	\checkmark	
s_T	\checkmark	V(t)	\checkmark			s_T	\checkmark	
		c_T	\checkmark					

Table A.2.3. Group C structural identifiabilities. Model group C uses a logistic form for viral dynamics and bilinear form for CD8+ T cell proliferation.

N	MD1	N	/ID2	N	/ID3	N	/ID4
Parameters	Identifiability	Parameters	Identifiability	Parameters	Identifiability	Parameters	Identifiability
d_I	√	c_T	✓	s_T	✓	c_T	✓
V(t)	\checkmark	p		c	\checkmark	V(t)	\checkmark
r	\checkmark	T(t)	\checkmark	I(t)		r	\checkmark
				T(t)	\checkmark	d_I	\checkmark
c	\checkmark	β	\checkmark	$\hat{\boldsymbol{\beta}}$	\checkmark	c	\checkmark
		V(t)	\checkmark	d_I	\checkmark	d_T	\checkmark
U(t)		d_I	\checkmark	U(t)		U(t)	
$\hat{\boldsymbol{\beta}}$	\checkmark	r	\checkmark	r	\checkmark	β	\checkmark
T(t)	\checkmark	ST	\checkmark	V(t)	\checkmark	T(t)	\checkmark
p		U(t)		p		p	
S_T	\checkmark	c	\checkmark	d_T	\checkmark	ST	\checkmark
I(t)						I(t)	
. /		I(t)				()	

Table A.2.4. Group D structural identifiabilities. Model group D uses a target cell limited model for viral dynamics and bilinear form for CD8+ T cell proliferation.

14 A.3 All Model Parameter Fits

15 A.3.1 Model Group A

		Model M	IA1 Parame	eters			
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes
p	Viral replication rate	d^{-1}	6.05	2.57	[1e-2,1e2]	1	-
k_V	Carrying capacity of virus	(PFU/ml)	1.2	1.2	-	-	Determined from data
c_V	Viral clearance rate	$d^{-1}cell^{-1}$	2.84e-6	1.78e-6	[1e-8,1e-4]	2	
s_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$d_T T(0)$	$d_T T(0)$	-	-	From steady state
r	CD8+ T cell proliferation rate	d^{-1}	2.21e-1	99.9	[1e-4,1e2]	1,3	
k_T	CD8+ T cell half saturation constant	(PFU/ml)	5.11e+3	2.35e+7	[1e1, 1e8]	4	
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	0.011	0.012	-	6	

Table A.3.1. Summary of MA1 parameters. The mathematical expression for MA1 is presented in A.1.1.

		Mod	el MA2 Paramete	rs			
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes
p	Viral replication rate	d^{-1}	5.93	3.07	[1e-2,1e2]	1	-
k_V	Carrying capacity of virus	(PFU/ml)	1.2	1.2	-	-	Determined from data
c_V	Viral clearance rate	$d^{-1}cell^{-1}$	2.61e-6	1.98e-6	[1e-8,1e-4]	2	
s_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$(d_T+c_T)T(0)$	$(d_T+c_T)T(0)$	-	-	From steady state
r	CD8+ T cell proliferation rate	d^{-1}	2.05e-1	1.68e-1	[1e-4,1e2]	1,3	
k_T	CD8+ T cell half saturation constant	(PFU/ml)	3.77e+3	1e1	[1e1, 1e8]	4	
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	0.011	0.012	-	6	
c_T	nonlinear CD8+ T cell downregulation rate	$(PFU/ml)d^{-1}$	2.49e-2	4.89e-2	[0,1e0]	-	Assumed similar to d_T

Table A.3.2. Summary of MA2 parameters. The mathematical expression for MA2 is presented in A.1.1.

-		Model	MA3 Parameters				
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes
P	Viral replication rate	$[d^{-1}]$	5.07 (4.52,5.68)	2.57	[1e-2,1e2]	1	-
k_V	Carrying capacity of virus	[PFU/ml][10 ⁶]	1.2	1.2	-	1	Determined from data
c_V	Viral clearance rate	$[d^{-1}cell^{-1}][10^{-6}]$	2.21 (1.92,2.62)	1.78	[1e-2,1e2]	2	
s_T	Homeostatic CD8+ T cell proliferation rate	$[cell \cdot d^{-1}]$	$d_T T(0)$		-	-	From steady state
r	CD8+ T cell proliferation rate	$[d^{-1}]$	2.74e-1 (2.36e-1,3.16e-1)	99.9	[1e-4,1e2]	1,3	
k_T	CD8+ T cell half saturation constant	[PFU/ml][10 ³]	2.48e3 (1.58e3,3.98e3)	2.35e7	[1e1, 1e7]	4	
$d_T^{\text{half life}}$	CD8+ T cell half life	$[d^{-1}]$	1.1e-2	1.2e-2	-	6	
$d_T^{\text{downregulation}}$	CD8+ T linear downregulation	$[d^{-1}]$	6.65-2 (4.43e-2,8.50e-2)	1.0e-4	[1e-4,1e0]	4,5	

Table A.3.3. Parameters for MA3, the selected adult model. The mathematical expression for MA3 is presented in A.1.1.

-		Model I	MA4 Parameters				
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes
p	Viral replication rate	$[d^{-1}]$	5.45	3.07	[1e-2,1e2]	1	=
k_V	Carrying capacity of virus	[PFU/ml][10 ⁶]	1.2	1.2	-	1	Determined from data
c_V	Viral clearance rate	$[d^{-1}cell^{-1}][10^{-6}]$	2.16	1.97	[1e-2,1e2]	2	
s_T	Homeostatic CD8+ T cell proliferation rate	$[cell \cdot d^{-1}]$	$(d_T+c_T)T(0)$	$(d_T+c_T)T(0)$	-	-	From steady state
r	CD8+ T cell proliferation rate	$[d^{-1}]$	2.73e-1	1.69e-1	[1e-4,1e2]	1,3	
k_T	CD8+ T cell half saturation constant	[PFU/ml][10 ³]	2.20e3	1e1	[1e1, 1e7]	4	
$d_T^{\text{half life}}$	CD8+ T cell half life	$[d^{-1}]$	1.1e-2	1.2e-2	-	6	
$d_T^{\text{downregulation}}$	CD8+ T linear downregulation	$[d^{-1}]$	6.47e-2	1.0e-4	[1e-4,1e0]	4,5	
c_T	nonlinear CD8+ T cell downregulation rate	$(PFU/ml)d^{-1}$	2.32e-9	4.86e-2	[0,1e0]	-	Assumed similar to d_T

Table A.3.4. Parameters for MA4, the selected adult model. The mathematical expression for MA4 is presented in A.1.1.

A.3.2 Model Group B

	Model MB1 Parameters										
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes				
β	Infection rate of uninfected cells	$[PFU/ml]^{-1}[d^{-1}][10^{-5}]$	2.23	4.63 (1.69,6.80)	[1e-3,1e1]	2	-				
d_I	Clearance rate of infected cells	$[cells^{-2}][10^{-7}]$	6.81	4.68 (3.85,5.32)	[1e-1,1e3]	2					
p	Production rate of virus	$[d^{-1}]$	1	1	-	2	Fixed for structural identifiability.				
c	Clearance rate of virus	$[d^{-1}][10^1]$	2.13e-1	1.23 (1.33e-1,1.88)	[1e-3,1e1]	2					
ST	Homeostatic CD8+ T cell proliferation rate	$[celld^{-1}]$		$d_T T(0)$	-	-	From steady state				
r	CD8+ T cell proliferation rate	$[d^{-1}][10^{-1}]$	4.45	2.15 (1.70, 2.71)	[1e-3,1e3]	1,3					
k_T	CD8+ T cell half saturation constant	[PFU/ml][10 ¹]	4.45e3	1.15 (1.0,13.1)	[1e0, 1e7]	4					
$d_T^{\text{half life}}$	CD8+ T half life	$[d^{-1}][10^{-2}]$	1.1	1.2	-	6					

Table A.3.5. Parameters for model MB1, the selected aged model. The mathematical expression for MB1 is presented in A.1.1.

	Model MB2 Parameters									
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes			
β	Infection rate of uninfected cells	$[PFU/m1]^{-1}[d^{-1}][10^{-5}]$	2.23	4.63	[1e-3,1e1]	2	-			
d_I	Clearance rate of infected cells	$[cells^{-2}][10^{-7}]$	6.79	4.68	[1e-1,1e3]	2				
p	Production rate of virus	$[d^{-1}]$	1	1	-	2	Fixed for structural identifiability.			
c	Clearance rate of virus	$[d^{-1}][10^1]$	2.14e-1	1.23	[1e-3,1e1]	2				
s_T	Homeostatic CD8+ T cell proliferation rate	$[celld^{-1}]$		$d_TT(0)$	-	-	From steady state			
r	CD8+ T cell proliferation rate	$[d^{-1}][10^{-1}]$	4.38	2.14	[1e-3,1e3]	1,3				
k_T	CD8+ T cell half saturation constant	[PFU/ml][10 ¹]	4.34e3	1.16	[1e0, 1e7]	4				
$d_T^{\text{half life}}$	CD8+ T half life	$[d^{-1}][10^{-2}]$	1.1	1.2	-	6				
c_T	CD8+ T cell nonlinear downregulation rate	$(PFU/ml)d^{-1}$	2.99e-3	2.08e-9	[0,1e0]	-	Assumed similar to d_T			

Table A.3.6. Parameters for model MB2. The mathematical expression for MB2 is presented in A.1.1.

		N	Model MB3 Paran	neters			
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes
β	Infection rate of uninfected cells	$(PFU/ml)^{-1}d^{-1}$	2.21e-5	1.77e-5	[1e-8,1e-4]	2	-
d_I	Clearance rate of infected cells	cells ^{−2}	5.55e-7	4.33e-6	[1e-8,1e-4]	2	
p	Production rate of virus	d^{-1}			1e0	2	Fixed for structural identifiability.
c	Clearance rate of virus	d^{-1}	2.19	1.42	[1e-2,1e2]	2	
s_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$(d_T + c_T)T(0)$	$(d_T + c_T)T(0)$	-	-	From steady state
r	CD8+ T cell proliferation rate	d^{-1}	4.18e-1	2.03e-1	[1e-4,1e2]	1,3	
k_T	CD8+ T cell half saturation constant	(PFU/ml)	1.05e4	1e1	[1e1, 1e8]	4	
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	1.1e-2	1.2e-2	-	6	
$d_T^{\text{downregulation}}$	CD8+ T cell linear downregulation rate	d^{-1}	8.37e-2	1.00e-4	[1e-4,1e0]	4,5	

Table A.3.7. Summary of MB3 parameters. The mathematical expression for MB3 is presented in A.1.1.

	Model MB4 Parameters											
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes					
β	Infection rate of uninfected cells	$(PFU/ml)^{-1}d^{-1}$	2.21e-5	4.63e-5	[1e-8,1e-4]	2	-					
d_I	Clearance rate of infected cells	cells ⁻²	5.55e-7	4.68e-7	[1e-8,1e-4]	2						
p	Production rate of virus	d^{-1}			1e0	2	Fixed for structural identifiability.					
c	Clearance rate of virus	d^{-1}	2.19	12.3	[1e-2,1e2]	2						
s_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$(d_T + c_T)T(0)$	$(d_T+c_T)T(0)$	-	-	From steady state					
r	CD8+ T cell proliferation rate	d^{-1}	4.18e-1	2.15e-1	[1e-4,1e2]	1,3						
k_T	CD8+ T cell half saturation constant	(PFU/ml)	1.05e4	1.14e1	[1e1, 1e8]	4						
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	1.1e-2	1.2e-2	[1e-4,1e0]	6						
$d_T^{\text{downregulation}}$	CD8+ T cell linear downregulation rate	d^{-1}	8.37e-2	1.00e-4	[1e-4,1e0]	4,5						
c_T	CD8+ T cell nonlinear downregulation rate	$(PFU/ml)d^{-1}$	5.98e-14	3.65e-8	[0,1e0]	-	Assumed similar to d_T					

Table A.3.8. Summary of MB4 parameters. The mathematical expression for MB4 is presented in A.1.1.

17 A.3.3 Model Group C

	Model MC1 Parameters											
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes					
p	Viral replication rate	d^{-1}	4.61	2.71	[1e-2,1e2]	1	-					
k_V	Carrying capacity of virus	(PFU/ml)			1.2e6 (fixed)	-	Determined from data					
c_V	Viral clearance rate	$d^{-1}cell^{-1}$	1.84e-6	2.06e-6	[1e-8,1e-4]	2						
S_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$d_T T(0)$	$d_TT(0)$	-	-	From steady state					
r	CD8+ T cell proliferation rate	$d^{-1}(PFU/ml)^{-1}$	2.08e-6	4.22e-6	$[1e-4/k_V, 1e2/k_V]$	1,3	Bounds are scaled.					
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	0.011	0.012	-	6						

Table A.3.9. Summary of MC1 parameters. The mathematical expression for MC1 is presented in A.1.1.

-	Model MC2 Parameters												
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes						
p	Viral replication rate	d^{-1}	5.42	3.16	[1e-2,1e2]	1	-						
k_V	Carrying capacity of virus	(PFU/ml)			1.2e6 (fixed)	-	Determined from data						
c_V	Viral clearance rate	$d^{-1}cell^{-1}$	2.13e-6	2.08e-6	[1e-8,1e-4]	2							
ST	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$(d_T+c_T)T(0)$	$(d_T+c_T)T(0)$	-	-	From steady state						
r	CD8+ T cell proliferation rate	$d^{-1}(PFU/ml)^{-1}$	1.31e-6	3.84e-6	$[1e-4/k_V, 1e2/k_V]$	1,3	Bounds are scaled.						
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	1.1e-2	1.2e-2	-	6							
c_T	CD8+ T cell nonlinear downregulation rate	$(PFU/ml)d^{-1}$	1.30e-1	2.27e-1	[0,1e0]	-	Assumed similar to d_T						

Table A.3.10. Summary of MC2 parameters. The mathematical expression for MC2 is presented in A.1.1.

	Model MC3 Parameters											
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes					
p	Viral replication rate	d^{-1}	4.60	2.71	[1e-2,1e2]	1	-					
k_V	Carrying capacity of virus	(PFU/ml)			1.2e6 (fixed)	-	Determined from data					
c_V	Viral clearance rate	$d^{-1}cell^{-1}$	1.84e-6	2.06e-6	[1e-8,1e-4]	2						
s_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$d_T T(0)$	$d_T T(0)$	-	-	From steady state					
r	CD8+ T cell proliferation rate	$d^{-1}(PFU/ml)^{-1}$	2.08e-6	4.28e-6	$[1e-4/k_V, 1e2/k_V]$	1,3	Bounds are scaled.					
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	0.011	0.012	-	6						
$d_T^{ m downregulation}$	CD8+ T cell linear downregulation rate	d^{-1}	1e-4	3.4e-3	[1e-4,0]	4,5						

Table A.3.11. Summary of MC3 parameters. The mathematical expression for MC3 is presented in A.1.1.

	Model MC4 Parameters												
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes						
p	Viral replication rate	d^{-1}	5.42	3.16	[1e-2,1e2]	1	-						
k_V	Carrying capacity of virus	(PFU/ml)			1.2e6 (fixed)	-	Determined from data						
c_V	Viral clearance rate	$d^{-1}cell^{-1}$	2.13e-6	2.08e-6	[1e-8,1e-4]	2							
ST	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$(d_T+c_T)T(0)$	$(d_T+c_T)T(0)$	-	-	From steady state						
r	CD8+ T cell proliferation rate	$d^{-1}(PFU/ml)^{-1}$	1.31e-6	3.84e-6	$[1e-4/k_V, 1e2/k_V]$	1,3	Bounds are scaled.						
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	1.1e-2	1.2e-2	-	6							
$d_T^{\text{downregulation}}$	CD8+ T cell linear downregulation rate	d^{-1}	1e-4	1e-4	[1e-4,0]	4,5							
c_T	CD8+ T cell nonlinear downregulation rate	$(PFU/ml)d^{-1}$	1.30e-1	2.27e-1	[0,1e0]	-	Assumed similar to d_T						

Table A.3.12. Summary of MC4 parameters. The mathematical expression for MC4 is presented in A.1.1.

A.3.4 Model Group D

	Model MD1 Parameters												
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes						
β	Infection rate of uninfected cells	$(PFU/ml)^{-1}d^{-1}$	2.15e-5	4.07e-5	[1e-8,1e-4]	2	-						
d_I	Clearance rate of infected cells	$cells^{-2}$	6.72e-7	4.28e-7	[1e-8,1e-4]	2							
p	Production rate of virus	d^{-1}			1e0	2	Fixed for structural identifiability.						
c	Clearance rate of virus	d^{-1}	2.06	11.3	[1e-2,1e2]	2							
s_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$d_TT(0)$	$d_TT(0)$	-	-	From steady state						
r	CD8+ T cell proliferation rate	$d^{-1}(PFU/ml)^{-1}$	2.61e-6	1.17e-5	$[1e-4/k_V, 1e2/k_V]$	1,3	Bounds are scaled.						
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	1.1e-2	1.2e-2	-	6							

Table A.3.13. Summary of MD1 parameters. The mathematical expression for MD1 is presented in A.1.1.

	Model MD2 Parameters										
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes				
β	Infection rate of uninfected cells	$(PFU/ml)^{-1}d^{-1}$	2.15e-5	1.67e-5	[1e-8,1e-4]	2	-				
d_I	Clearance rate of infected cells	$cells^{-2}$	6.72e-7	4.43e-7	[1e-8,1e-4]	2					
p	Production rate of virus	d^{-1}			1e0	2	Fixed for structural identifiability.				
c	Clearance rate of virus	d^{-1}	2.06	13.1	[1e-2,1e2]	2					
s_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$(d_T+c_T)T(0)$	$(d_T+c_T)T(0)$	-	-	From steady state				
r	CD8+ T cell proliferation rate	$d^{-1}(PFU/ml)^{-1}$	2.61e-6	7.78e-6	$[1e-4/k_V, 1e2/k_V]$	1,3	Bounds are scaled.				
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	0.0648	1.00e-4	[1e-4,1e0]	6					
c_T	CD8+ T cell nonlinear downregulation rate	$(PFU/ml)d^{-1}$	4.28e-12	1.72e-1	[0,1e0]	-	Assumed similar to d_T				

Table A.3.14. Summary of MD2 parameters. The mathematical expression for MD2 is presented in A.1.1.

	Model MD3 Parameters												
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes						
β	Infection rate of uninfected cells	$(PFU/ml)^{-1}d^{-1}$	2.08e-5	1.71e-5	[1e-8,1e-4]	2	-						
d_I	Clearance rate of infected cells	cells ⁻²	5.76e-7	6.00e-6	[1e-8,1e-4]	2							
p	Production rate of virus	d^{-1}			1e0	2	Fixed for structural identifiability.						
c	Clearance rate of virus	d^{-1}	2.05	1.25	[1e-2,1e2]	2							
s_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$d_T T(0)$	$d_T T(0)$	-	-	From steady state						
r	CD8+ T cell proliferation rate	$d^{-1}(PFU/ml)^{-1}$	3.13e-6	1.24e-5	$[1e-4/k_V, 1e2/k_V]$	1,3	Bounds are scaled.						
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	1.1e-2	1.2e-2	[1e-4,1e0]	6							
$d_T^{\text{downregulation}}$	CD8+ T cell linear downregulation rate	d^{-1}	5.42e-2	1.00e-4	[1e-4,1e0]	4,5							

Table A.3.15. Summary of MD3 parameters. The mathematical expression for MD3 is presented in A.1.1.

	Model MD4 Parameters											
Parameter	Description	Units	Adult	Aged	Search Bounds	Reference	Notes					
β	Infection rate of uninfected cells	$(PFU/ml)^{-1}d^{-1}$	2.08e-5	1.67e-5	[1e-8,1e-4]	2	-					
d_I	Clearance rate of infected cells	cells ⁻²	5.76e-7	4.43e-6	[1e-8,1e-4]	2						
p	Production rate of virus	d^{-1}			1e0	2	Fixed for structural identifiability.					
c	Clearance rate of virus	d^{-1}	2.05	1.31	[1e-2,1e2]	2						
s_T	Homeostatic CD8+ T cell proliferation rate	$cell \cdot d^{-1}$	$(d_T+c_T)T(0)$	$(d_T+c_T)T(0)$	-	-	From steady state					
r	CD8+ T cell proliferation rate	$d^{-1}(PFU/ml)^{-1}$	3.13e-6	7.78e-6	$[1e-4/k_V, 1e2/k_V]$	1,3	Bounds are scaled.					
$d_T^{\text{half life}}$	CD8+ T cell half life	d^{-1}	1.1e-2	1.2e-2	[1e-4,1e0]	6						
$d_T^{\text{downregulation}}$	CD8+ T cell linear downregulation rate	d^{-1}	5.42e-2	1.00e-4	[1e-4,1e0]	4,5						
c_T	CD8+ T cell nonlinear downregulation rate	$(PFU/ml)d^{-1}$	1.02e-11	1.72e-1	[0,1e0]	-	Assumed similar to d_T					

Table A.3.16. Summary of MD4 parameters. The mathematical expression for MD4 is presented in A.1.1.

A.4 Fit Model Plots

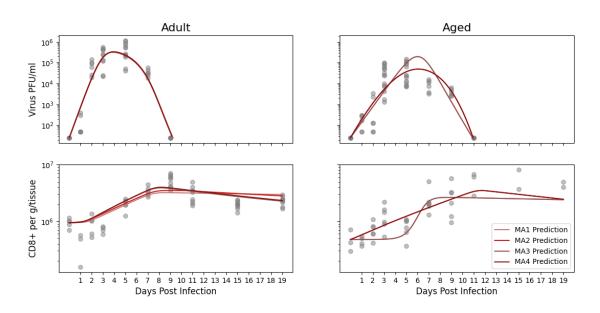


Figure A.4.1. Group A fitted model trajectories alongside the data from Toapanta et al.⁷.

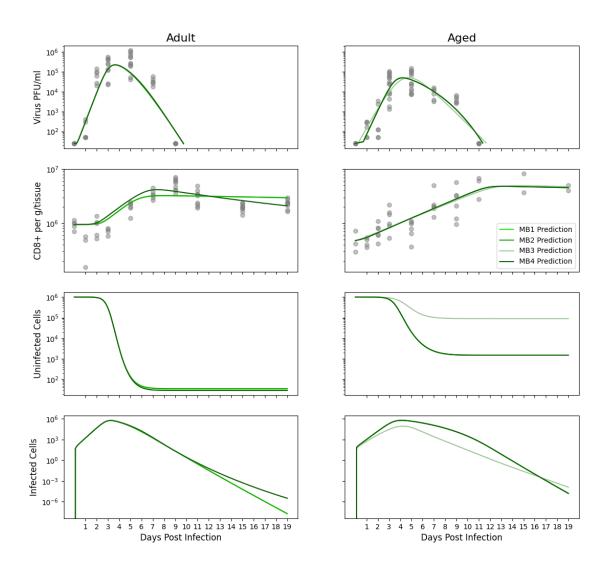


Figure A.4.2. Group B fitted model trajectories alongside the data from Toapanta et al.⁷.

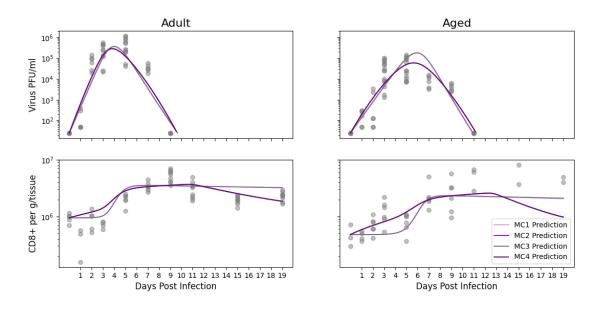


Figure A.4.3. Group C fitted model trajectories alongside the data from Toapanta et al.⁷.

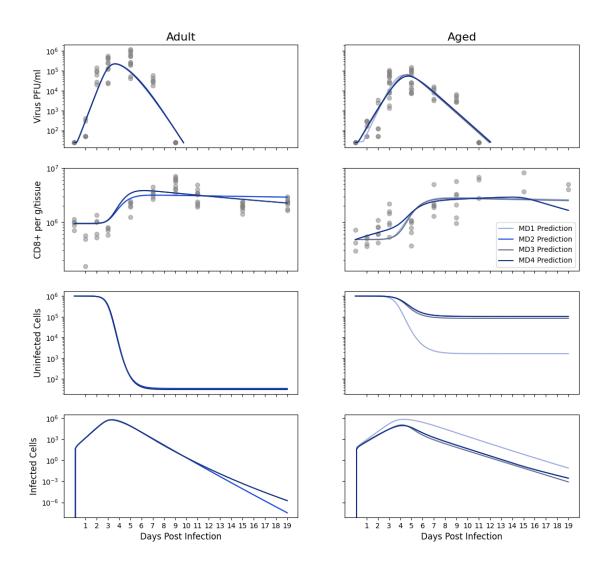


Figure A.4.4. Group D fitted model trajectories alongside the data from Toapanta et al.⁷.

20 A.5 Bootstrapped Predictions for the Selected Models

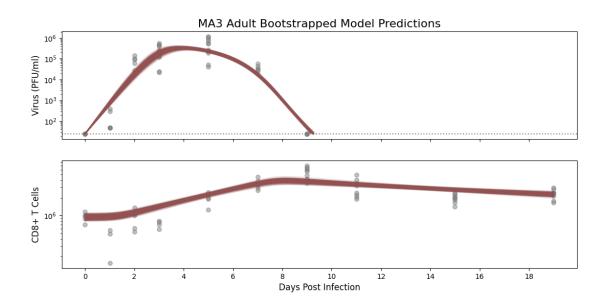


Figure A.5.1. Adult model (MA3) bootstrapped trajectories alongside the data from Toapanta et al.⁷.

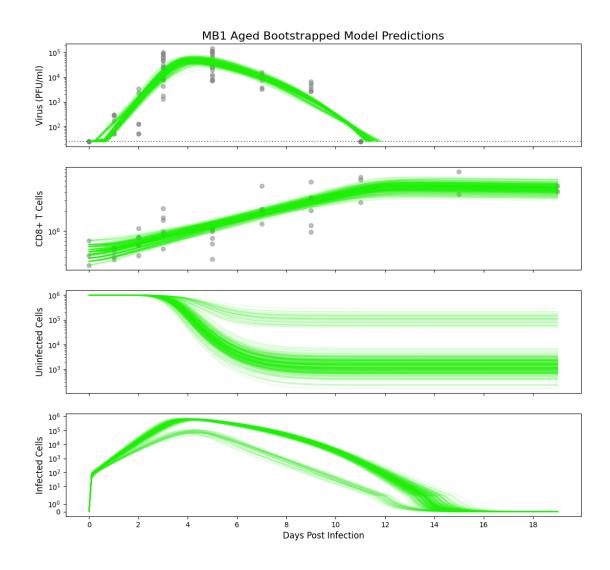


Figure A.5.2. Aged model (MB1) bootstrapped trajectories alongside the data from Toapanta et al.⁷.

21 References

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