



Supplementary Information for

**Lifelong CMV infection improves immune defense in old mice by
broadening the mobilized TCR repertoire against third-party infection**

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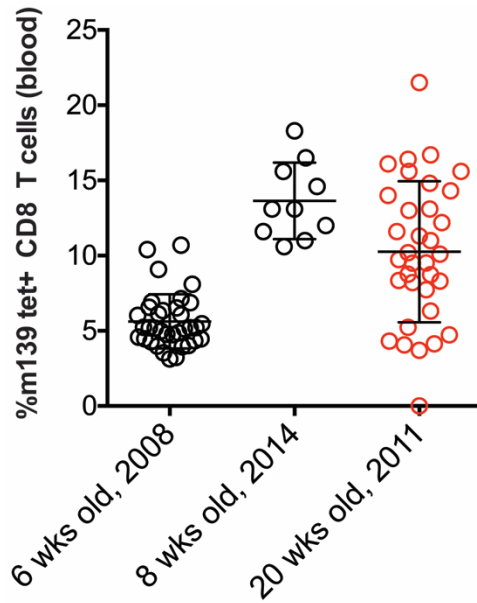


Fig. S1. Age of MCMV infection does not determine the magnitude of memory inflation over the lifespan. Steady state memory inflation in 17 month old mice was monitored in groups that had been infected with MCMV at 6, 8, or 20 weeks of age (n=35, n=10, n=34 respectively). Animals in the red dataset were used for this manuscript.

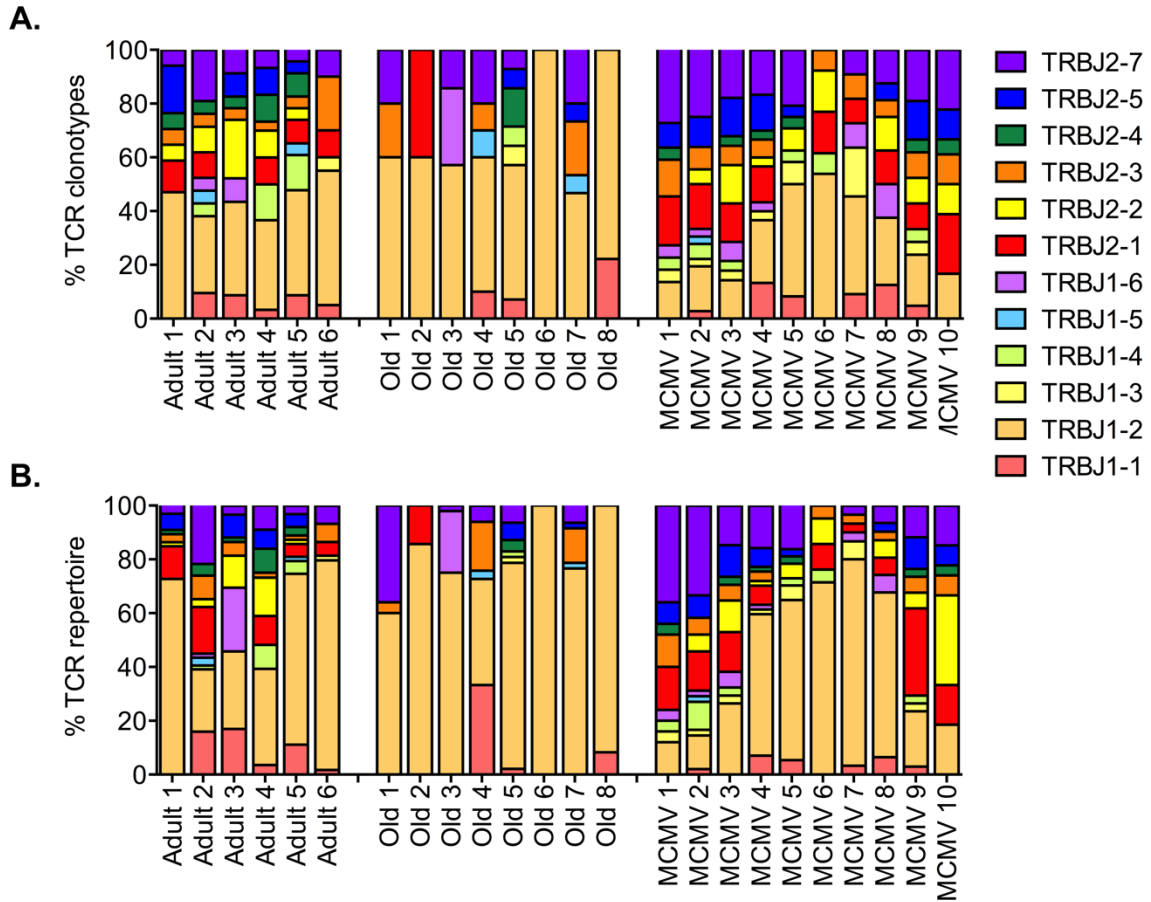


Fig. S2. MCMV+ mice show increased variability in their TRBJ gene choice.

Utilization of different TRBJ genes for the unique clonotypes (A) and the total repertoire (B) for each individual adult, old, and MCMV+ mouse are shown.

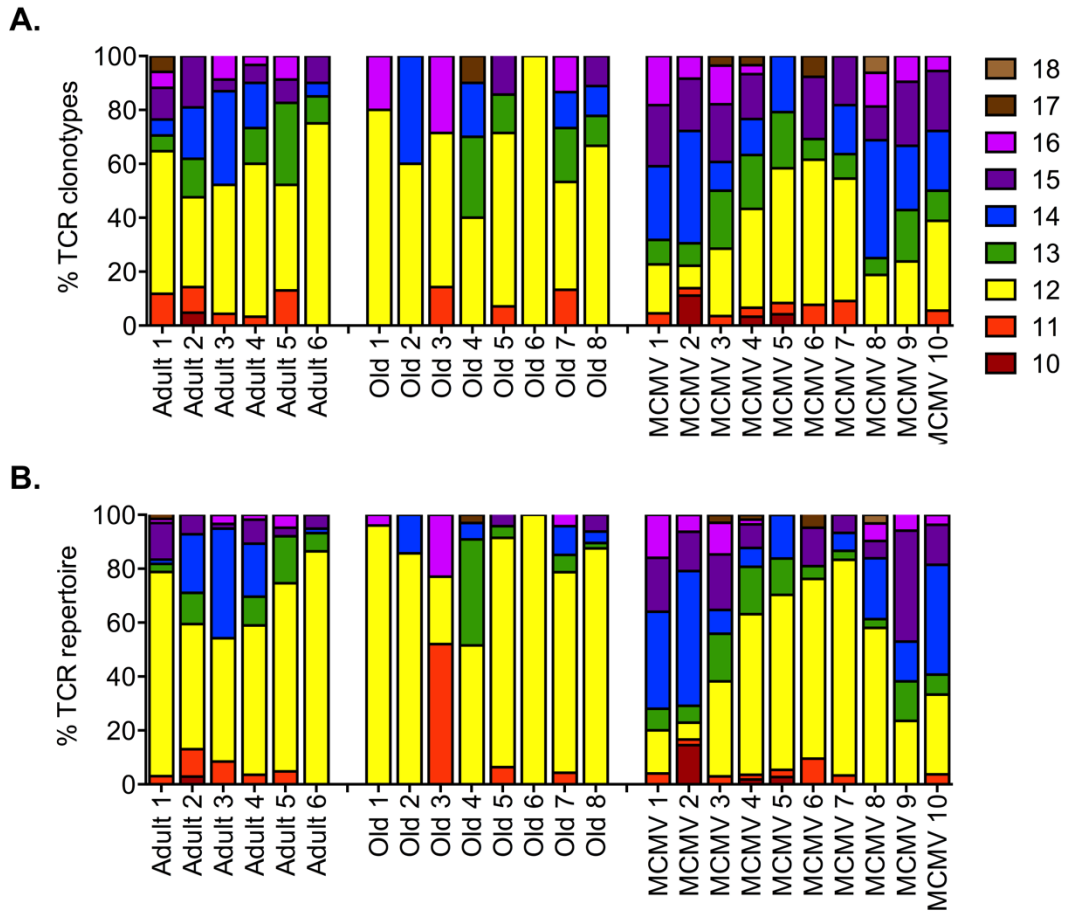


Fig. S3. MCMV+ mice show high variability in their CDR3 lengths in the OVA-specific TRBV12+ repertoire. The CDR3 amino acid length for unique clonotypes (A) and the total repertoire (B) for each individual adult, old, and MCMV+ mouse is shown.

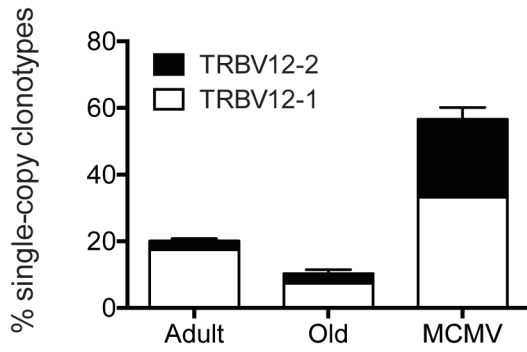


Fig. S4. Aging with MCMV infection results in the recruitment of many “single-copy” clonotypes into the OVA-specific repertoire. The frequency of clonotypes present as “single-copy” within the TRBV12-1+ or TRBV12-2+ OVA-specific CD8 repertoires of adult old, and MCMV+ old mice.

Table S1. Summary of the data used to compare TRBV12+ TCR β clonotypes (amino acid sequences) for the OVA-specific CD8 T cell response to Lm-OVA infection in adult, old, and old mice with lifelong-MCMV infection (“MCMV”).

Parameter	Adult (12 wk old)	Old (20 mo old)	MCMV+ (20 mo old, 15 w/ MCMV)
Number of mice	6	8	10
Total number of TCR β sequences across all mice	372	325	344
Range of number of TCR β sequences per mouse	56-69	25-48	21-57
Mean number of TCR β sequences per mouse	62	40.4	34.4

Table S2. Examples of alignments of TCR sequences with the TCR genes, showing the extent of germline encoding of the PR motif for several shared clonotypes, including some using genes other than TRBJ1-2. Blue =TRBV encoded, green = TRBJ encoded, red = TRBD encoded, black = non-template additions.

												No. of n- additions	No. of mice
C	A	S	S	P	R	A	S	D	Y	T	F		10
tgt	gcc	agc	tcc	ccc	agg	gcc	tcc	gac	tac	acc	ttc	4	4
tgt	gcc	agc	tct	ccc	agg	gcc	tcc	gac	tac	acc	ttc	2	3
tgt	gcc	agc	tct	cca	cgg	gcc	tcc	gac	tac	acc	ttc	3	2
tgt	gcc	agc	tcc	ccc	cgg	gcc	tcc	gac	tac	acc	ttc	5	2
tgt	gcc	agc	tct	ccc	agg	gcc	tcc	gac	tac	acc	ttc	4	1
C	A	S	S	P	R	A	T	D	Y	T	F		9
tgt	gcc	agc	tct	ccg	cgg	gct	acc	gac	tac	acc	ttc	5	9
tgt	gcc	agc	tct	ccg	cgg	gcg	acc	gac	tac	acc	ttc	5	1
C	A	S	S	P	R	D	S	D	Y	T	F		9
tgt	gcc	agc	tct	ccc	agg	gac	tcc	gac	tac	acc	ttc	1	4
tgt	gcc	agc	tct	cca	agg	gac	tcc	gac	tac	acc	ttc	2	3
tgt	gcc	agc	tcc	ccc	agg	gac	tcc	gac	tac	acc	ttc	3	3
tgt	gcc	agc	tct	ccc	agg	gat	tcc	gac	tac	acc	ttc	3	1
tgt	gcc	agc	tcc	ccc	cgg	gac	tcc	gac	tac	acc	ttc	5	1
tgt	gcc	agc	tcc	ccc	cgg	gac	tcc	gac	tac	acc	ttc	6	1
C	A	S	S	P	R	G	P	D	Y	T	F		7
tgt	gcc	agc	tcc	ccc	aga	ggg	ccc	gac	tac	acc	ttc	7	2
tgt	gcc	agc	tct	ccc	agg	gga	ccc	gac	tac	acc	ttc	3	1
tgt	gcc	agc	tct	ccc	cgg	ggg	ccc	gac	tac	acc	ttc	3	1
tgt	gcc	agc	tct	ccc	cgg	gga	ccc	gac	tac	acc	ttc	4	1
tgt	gcc	agc	tct	cct	agg	gga	ccc	gac	tac	acc	ttc	4	1
tgt	gcc	agc	tcc	ccc	agg	gga	ccc	gac	tac	acc	ttc	5	1
tgt	gcc	agc	tct	ccc	cgc	ggg	ccc	gac	tac	acc	ttc	5	1
C	A	S	S	P	R	G	T	D	Y	T	F		5
tgt	gcc	agc	tct	ccc	cgg	ggg	acc	gac	tac	acc	ttc	4	4
tgt	gcc	agc	tct	ccg	cgg	ggt	acc	gac	tac	acc	ttc	5	2
tgt	gcc	agc	tct	ccc	cgg	ggc	acc	gac	tac	acc	ttc	4	1
tgt	gcc	agc	tct	ccc	cgg	gga	acc	gac	tac	acc	ttc	5	1
tgt	gcc	agc	tct	ccc	cgg	ggt	acc	gac	tac	acc	ttc	5	1
C	A	S	S	P	R	N	S	D	Y	T	F		5
tgt	gcc	agc	tct	ccc	agg	aac	tcc	gac	tac	acc	ttc	1	3
tgt	gcc	agc	tcc	ccc	aga	aac	tcc	gac	tac	acc	ttc	3	1
tgt	gcc	agc	tcg	cca	aga	aac	tcc	gac	tac	acc	ttc	4	1
tgt	gcc	agc	tct	cca	aga	aac	tcc	gac	tac	acc	ttc	2	1
C	A	S	S	P	R	K	T	D	Y	T	F		5
tgt	gcc	agc	tct	ccc	cga	aag	acc	gac	tac	acc	ttc	7	5
TRBV12-1	tgt	gcc	agc	tct	ctc								
TRBD1: gggacagggggc TRBD2: gggactgggggggc													
ca aac tcc gac tac acc ttc												TRBJ1-2	
C	A	S	S	P	R	N	T	E	V	T	F		4
tgt	gcc	agc	tct	cca	cga	aac	aca	gaa	gtc	ttc	ttt	2	3
tgt	gcc	agc	tcc	ccg	aga	aac	aca	gaa	gtc	ttc	ttt	4	1
TRBV12-1	tgt	gcc	agc	tct	ctc								
TRBD1: gggacagggggc TRBD2: gggactgggggggc													
ca aac aca gaa gtc ttc ttt												TRBJ1-1	
C	A	S	S	P	R	E	D	T	Q	Y	F		3
tgt	gcc	agc	tct	ccc	cgg	gaa	gac	acc	cag	tac	ttt	3	1
tgt	gcc	agc	tcc	ccc	agg	gaa	gac	acc	cag	tac	ttt	3	1
tgt	gcc	agc	tcc	ccg	cga	gaa	gac	acc	cag	tac	ttt	4	1
TRBV12-1	tgt	gcc	agc	tct	ctc								
TRBD1: gggacagggggc TRBD2: gggactgggggggc													
aac caa gac acc cag tac ttt												TRBJ2-5	

Dataset S1. OVA-specific CDR3 sequences recovered from all animals.

