

Supplementary Data

Supplementary Figures

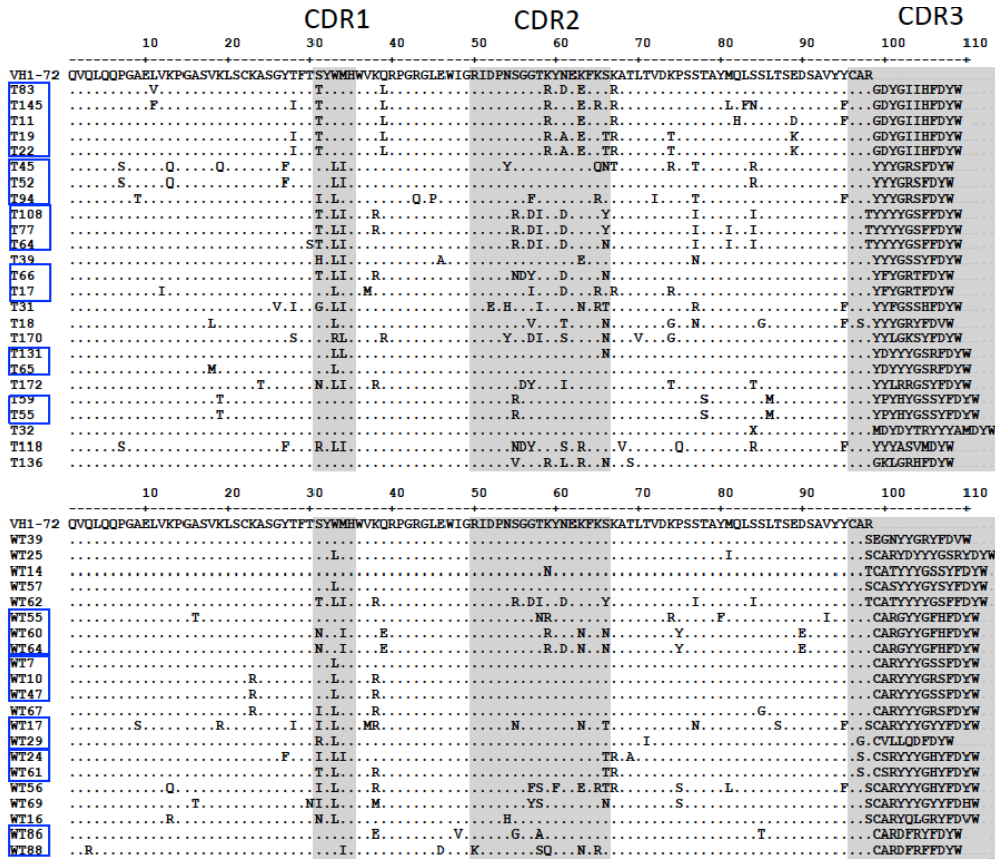


Figure 1A

Figure 1A: Localization of amino acid substitutions in distinct clones encoding VH1-72 heavy chains of anti-NP monoclonal antibodies obtained from wild type and TACI-deficient mice. Shown are independent VH sequences compared to the germline VH1-72 (indicated above each group). The boxed sequences correspond to the same clone, defined by the V(D)J junction. CDR1, CDR2 and CDR3, defined according to Kabat are shadowed. Sequences shown are all IgG1.

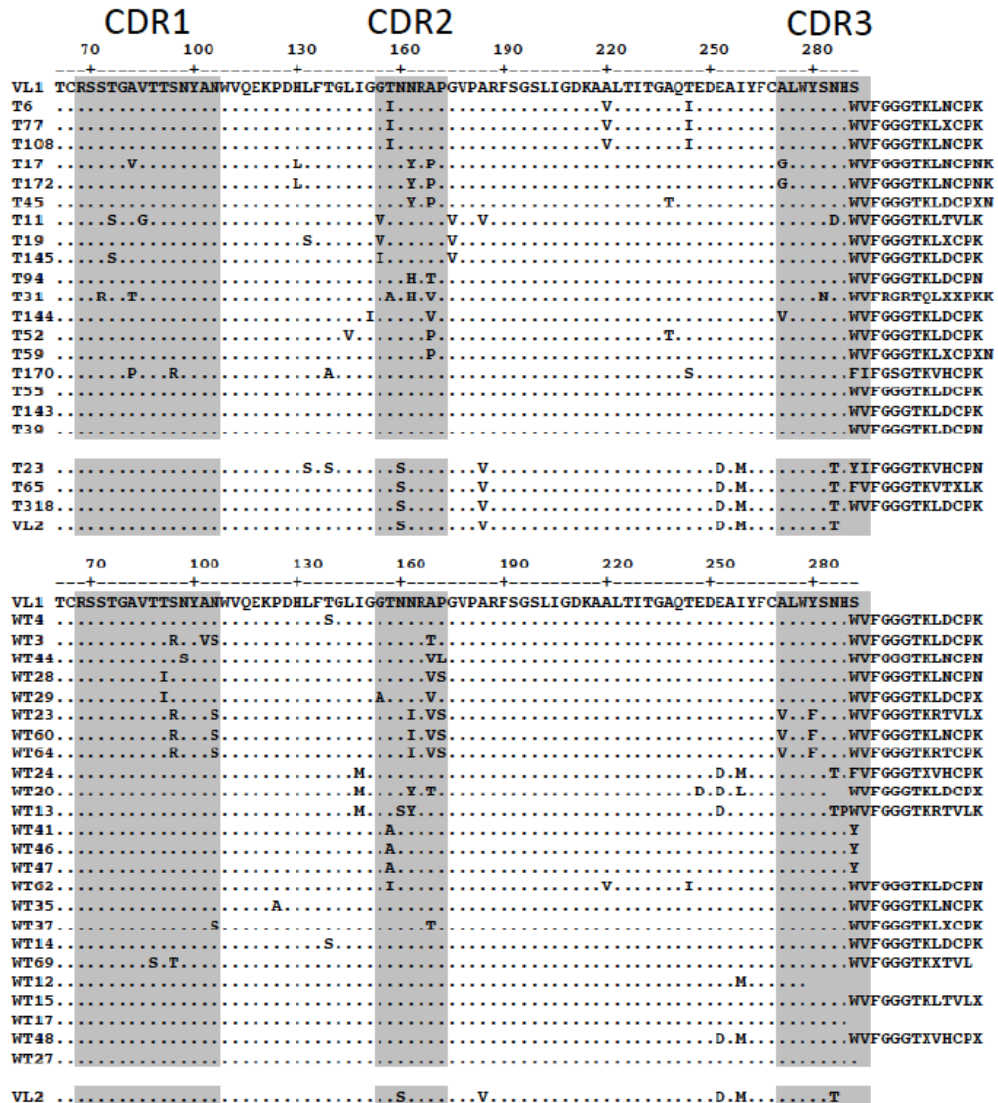


Figure 1B

Figure 1B: Localization of amino acid substitutions in distinct clones encoding Ig light chains of anti-NP monoclonal antibodies obtained from wild type and TACI-deficient mice. Shown are independent VL (V λ 1 or V λ 2) sequences compared to the germline (indicated above and below each group). CDR1, CDR2 and CDR3, defined according to Kabat are shadowed. The VL frequency of mutations was the same (1.7%) in clones derived from wt or from TACI-deficient mice.

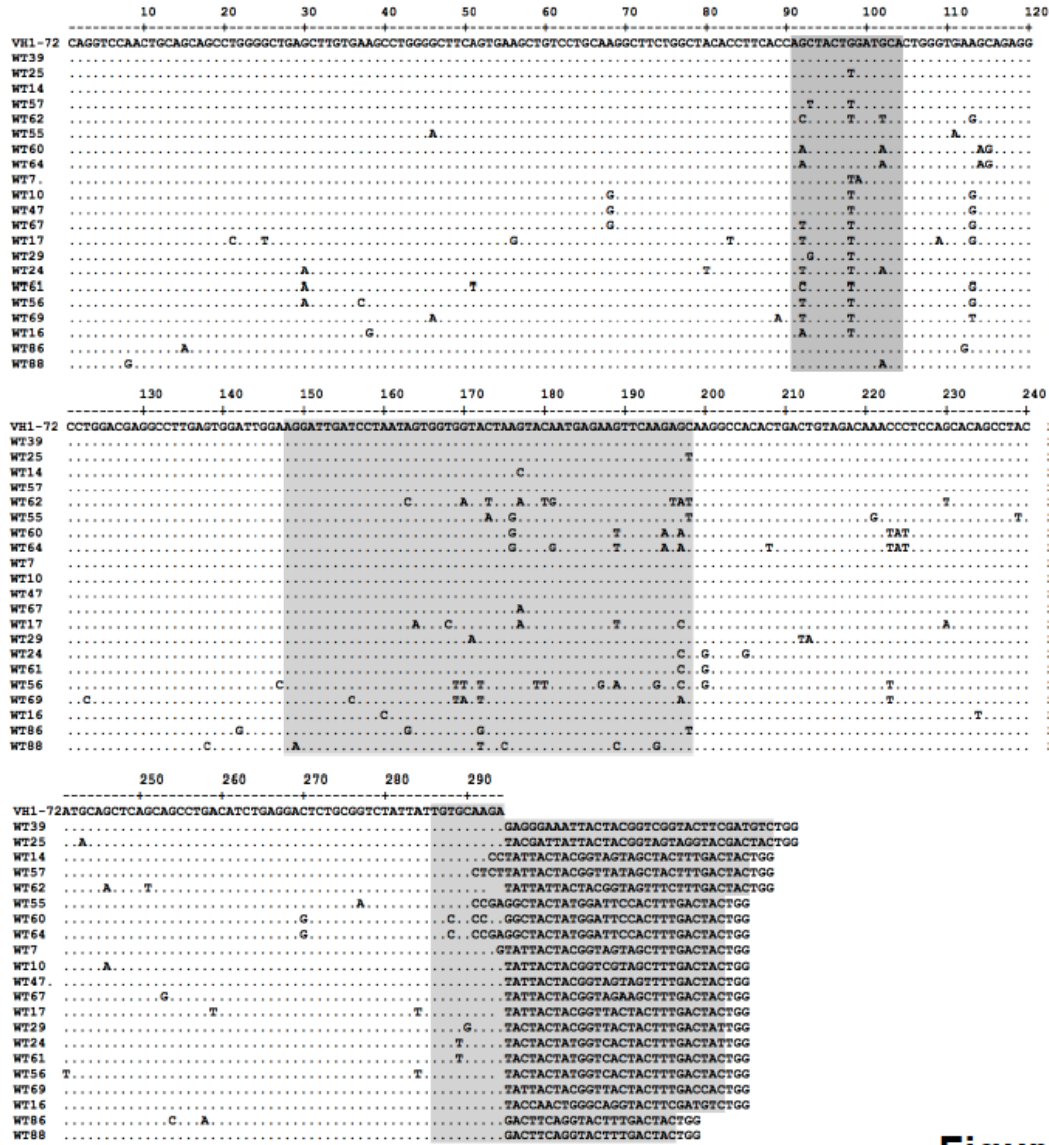


Figure 1C

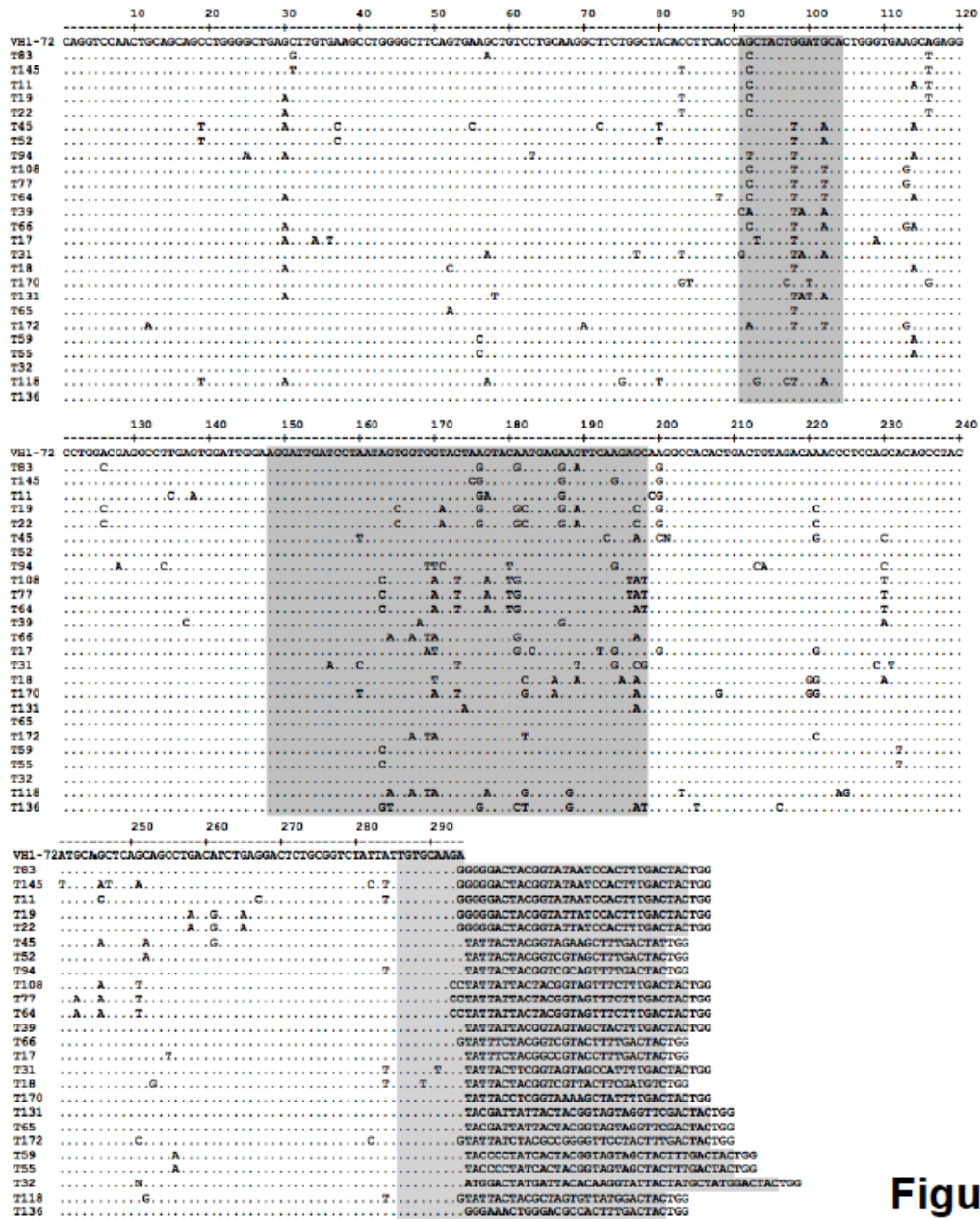


Figure 1D

Figures 1C and 1D: Nucleotide sequences of VH1-72 heavy chains of anti-NP monoclonal antibodies obtained from wild type and TAC1-deficient mice. Shown are independent VH sequences compared to the germline VH1-72 (indicated above each group, 1C, TAC1-ko, 1D, wt). CDR1, CDR2 and CDR3, defined according to Kabat are shadowed. Sequences shown correspond to the amino acid sequences shown in figure 1A.

	10	20	30	40	50	60	70	80	90	100	110	120		
VL1	ACTTGTGGCTCAAGTACTGGGGCTGTACAACTAGTAACTATGCCCACTGGGTCCAAGAAAACCAGATCATTTTTCACCTGGTCTAATAGGTGGTACCAACAACCGAGCTCCAGTGTPT													
WT4								G.....					
WT3			A.....	T..G.....							A.....		
WT44	A.....		G.....								T..T.....		
WT28		T.....									T..T.....		
WT29		T.....							C.....		T.....		
WT23		C.....		G.....						T.....	T..T.....		
WT60		C.....		G.....						T.....	T..T.....		
WT64		C.....		G.....						T.....	T..T.....		
WT24								G.....			T.....		
WT20								G.....		T.....	A.....		
WT10						G.....		G.....		G..T.....			
WT41										G.....			
WT46										G.....			
WT47										G.....			
WT62								T.....		T.....			
WT35					G.....								
WT37			G.....								A.....		
WT14						G.....							
WT09		T.....	C.....										
WT12													
WT15													
WT17													
WT40													
WT27													
VL2				T.....						G.....			
	130	140	150	160	170	180	190	200	210	220	230			
VL1	CCCTGCCAGATTCCTCAGGCTCCCTGATTGGAGACAAGGGCTGCCCTCACCATPCACAGGGGCLCAGACTGAGGATGAGGCAATATATTTCTGTGCTCTATGGGTACAGCAACCATTT													
WT4											C.....	GGGTGTTCCGGTGGAGGAACCAAA	
WT3										T.....	C.....	GGGTGTTCCGGTGGAGGAACCAAA	
WT44											C.....	GGGTGTTCCGGTGGAGGAACCAAA	
WT28												GGGTGTTCCGGTGGAGGAACCAAA	
WT29		T.....										GGGTGTTCCGGTGGAGGAACCAAA	
WT23										T.....	T.....	GGGTGTTCCGGTGGAGGAACCAAA	
WT60										T.....	T.....	GGGTGTTCCGGTGGAGGAACCAAA	
WT64										T.....	T.....	GGGTGTTCCGGTGGAGGAACCAAA	
WT24											C.....	TTCTTTCCGGTGGAGGAACCAAA	
WT20					C.....		T.....	T..G.....			C.....	GGGTGTTCCGGTGGAGGAACCAAA	
WT13							T.....	G.....			C..C.....	GGGTGTTCCGGTGGAGGAACCAAA	
WT41													
WT46													
WT47													
WT62					TG.....							C.....	GGGTGTTCCGGTGGAGGAACCAAA
WT35												C.....	GGGTGTTCCGGTGGAGGAACCAAA
WT37												C.....	GGGTGTTCCGGTGGAGGAACCAAA
WT14												C.....	GGGTGTTCCGGTGGAGGAACCAAA
WT60												C.....	GGGTGTTCCGGTGGAGGAACCAAA
WT12								G.....					
WT15									G.....				
WT17											T..G.....		
WT48												T..G.....	GG
WT27													G
VL2	T.....											C.....	

Figure 1E

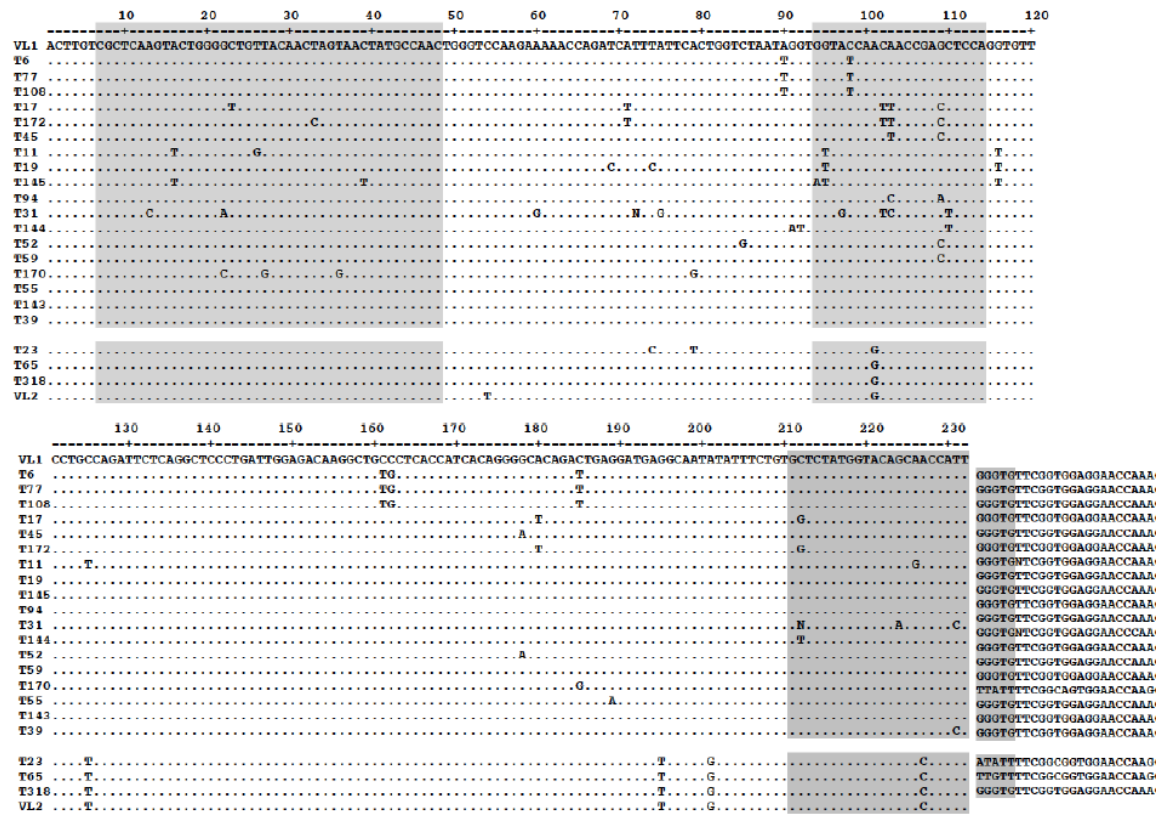


Figure 1F

Figures 1E and 1F: Nucleotide sequences of Light chains of anti-NP monoclonal antibodies obtained from wild type and TACI-deficient mice. The VL sequences compared to the germline V (indicated above each group, 1E, TACI-ko, 1F, wt). CDR1, CDR2 and CDR3, defined according to Kabat are shadowed. Sequences shown correspond to the aminoacid sequences shown in figure 1B.

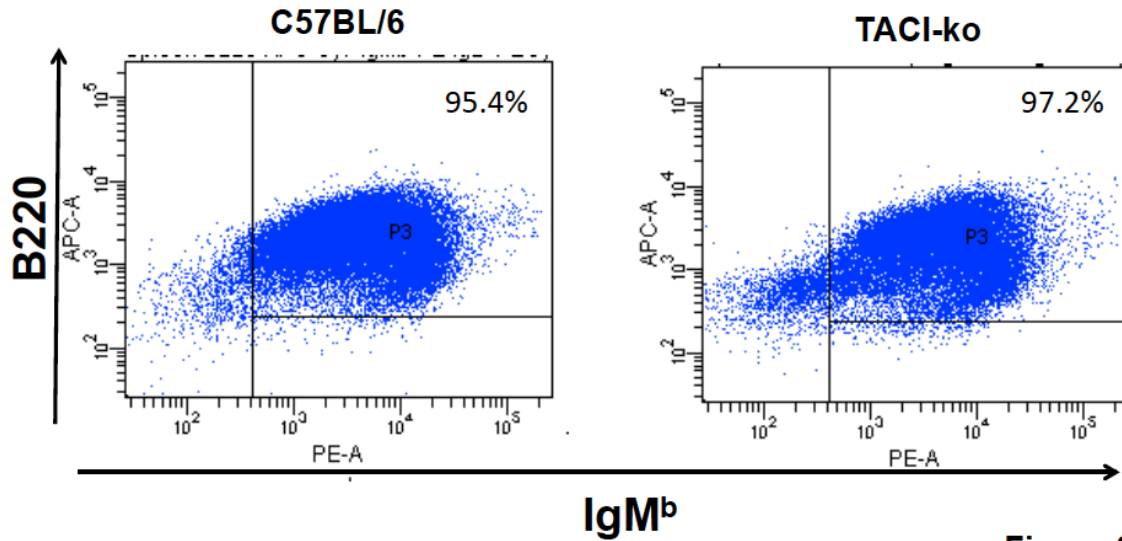


Figure 2

Figure 2: Flow Cytometer analysis of splenocytes obtained from wild type or TACI-deficient mice 10 days after re-infection with *C. rodentium*. Y-axis depicts staining with anti-B220 antibody; the X-axis depicts staining with anti-IgM^b antibody. Since IgM^b is the IgM allotype expressed by B cells of C57BL/6 mice, the figure indicates that B cells express the C57BL/6 IgM allotype.

Germinal center B cell clones from *C. rodentium* infected mice

C57BL/6				TACI-ko			
Clone	IgH chain	Strain	Junction	Clone	IgH chain	Strain	Junction
BL6_E08	Musmus IGHV1-18*01 F	C57BL/6	CARITYNSREFFYW	CI_D09	Musmus IGHV1-19*01 F	C57BL/6	CALGSSSFDFYW
BL6_H07	Musmus IGHV1-18*01 F	C57BL/6	CARWRNGYFDVW	CI_E02	Musmus IGHV1-19*01 F	C57BL/6	CARSFAYW
BL6_C02	Musmus IGHV1-31*01 F	C57BL/6	CARSITTAPFDYW	CI_A09	Musmus IGHV1-19*01 F	C57BL/6	CARDSSGYAYW
BL6_C07	Musmus IGHV1-54*01 F	C57BL/6	CARSMITKAMDYW	CI_D01	Musmus IGHV1-19*01 F	C57BL/6	CATPDGYYGYFDVW
BL6_D03	Musmus IGHV1-64*01 F	C57BL/6	CARRGLYGSSPYFDYW	CI_C01	Musmus IGHV1-20*01 F	C57BL/6	CARTGTTFDYW
BL6_F03	Musmus IGHV1-7*01 F	C57BL/6	CARGDGYPPYYAMDYW	CI_D04	Musmus IGHV1-39*01 F	C57BL/6	CARTLYYFDYW
BL6_G01	Musmus IGHV1-7*01 F	C57BL/6	CARGYYYGSSYGYFDVW	CI_H02	Musmus IGHV1-50*01 F	C57BL/6	CARYDGLGYFDYW
BL6_H09	Musmus IGHV1-72*01 F	C57BL/6	CTRGFYDGFSSWFAYW	CI_H12	Musmus IGHV1-52*01 F	C57BL/6	CARWGLRRSYAMDYW
BL6_E02	Musmus IGHV1-81*01 F	C57BL/6	CARVYYAMDYW	CI_C12	Musmus IGHV1-52*01 F	C57BL/6	CARFHYWYFDVW
BL6_E06	Musmus IGHV1-81*01 F	C57BL/6	CAREDGSSSWFAYW	CI_E11	Musmus IGHV1-53*01 F	C57BL/6	CARGGVITTVARYWYFDVW
BL6_A06	Musmus IGHV1-81*01 F	C57BL/6	CAREGDYSFAYW	CI_H10	Musmus IGHV1-53*01 F	C57BL/6	CARTAQTWGVFDYW
BL6_B02	Musmus IGHV1-81*01 F	C57BL/6	CARWTAQATCYW	CI_F03	Musmus IGHV1-64*01 F	C57BL/6	CARGYSIYWFVW
BL6_G11	Musmus IGHV1-81*01 F	C57BL/6	CARDSAGQAWFAYW	CI_F11	Musmus IGHV1-64*01 F	C57BL/6	CARKGLYRNYGVAYW
BL6_H08	Musmus IGHV1-82*01 F	C57BL/6	CARSGSSYYYAMDYW	CI_E04	Musmus IGHV1-64*01 F	C57BL/6	CARRWLLRSDYW
BL6_H02	Musmus IGHV1-82*01 F	C57BL/6	CARSIIYGIYAMDYW	CI_B02	Musmus IGHV1-69*01 F	C57BL/6	CARWYDFDYW
BL6_B09	Musmus IGHV1-9*01	C57BL/6	CARCSNLYYFDYW	CI_C10	Musmus IGHV1-72*01 F	C57BL/6	CVRWGYFDVW
BL6_F09	Musmus IGHV1-9*01 F	C57BL/6	CARGSNLYYFDYW	CI_E05	Musmus IGHV1-80*01 F	C57BL/6	CARFLTTVVATDYAMDYW
BL6_D09	Musmus IGHV1-9*01 F	C57BL/6	CARSTFDYW	CI_A06	Musmus IGHV1-81*01 F	C57BL/6	CTLDVPFAYW
BL6_E12	Musmus IGHV1-9*01 F	C57BL/6	CARRGGAYW	CI_D06	Musmus IGHV1-81*01 F	C57BL/6	CARYYYGSSFFDYW
BL6_C09	Musmus IGHV10-1*01 F	C57BL/6	CVRPSLYDGYLGGFAYW	CI_D02	Musmus IGHV1-85*01 F	C57BL/6	CARGGIYGITGAMDYW
BL6_G09	Musmus IGHV10-3*02 F	C57BL/6	CVRHYGSYYAMDYW	CI_G12	Musmus IGHV1-85*01 F	C57BL/6	CXXXGIXGIXGAXXX
BL6_B01	Musmus IGHV14-2*01 F	C57BL/6	CALYGDRAWFAYW	CI_F07	Musmus IGHV10-3*01 F	C57BL/6	CVRGNAMDYW
BL6_H11	Musmus IGHV14-4*01 F	C57BL/6	CTTSTTVASDYW	CI_B12	Musmus IGHV14-2*01 F	C57BL/6	CARYPEAYW
BL6_G02	Musmus IGHV3-6*01 F	C57BL/6	CARGRSGPFAYW	CI_D04	Musmus IGHV14-2*01 F	C57BL/6	CALEGGFAYW
BL6_H04	Musmus IGHV5-17*01 F	C57BL/6	CARSLIFSAYW	CI_C05	Musmus IGHV2-2*01 F	C57BL/6	CARKGDYYDAMDYW
				TACI_B09	Musmus IGHV3-6*01 F	C57BL/6	CAREGNWEAMDYW
				TACI_H07	Musmus IGHV3-6*01 F	C57BL/6	CAKGAARGFAYW
				TACI_G01	Musmus IGHV5-17*01 F	C57BL/6	CAMGGYSAWFAYW
				TACI_G02	Musmus IGHV5-17*01 F	C57BL/6	CARNWVDYW
				TACI_B10	Musmus IGHV5-4*01 F	C57BL/6	CARNDRCFYDAXDYXGQ
				TACI_B01	Musmus IGHV5-6*01 F	C57BL/6	CARLAYYSNYGWYFDVW

Figure 3

Figure 3: Identification of VH genes and V(D)J junctions of B cell clones isolated from the spleen of mice infected with *C. rodentium* 10 days earlier. Figure shows that all VH genes isolated match best to C57BL/6 germline alleles. Germinal center B cells (CD19+, GL7+ and Fas+) were isolated by single cell sorting from wt or TACI-deficient mice. CDN, amplification and sequencing were done according to (1).

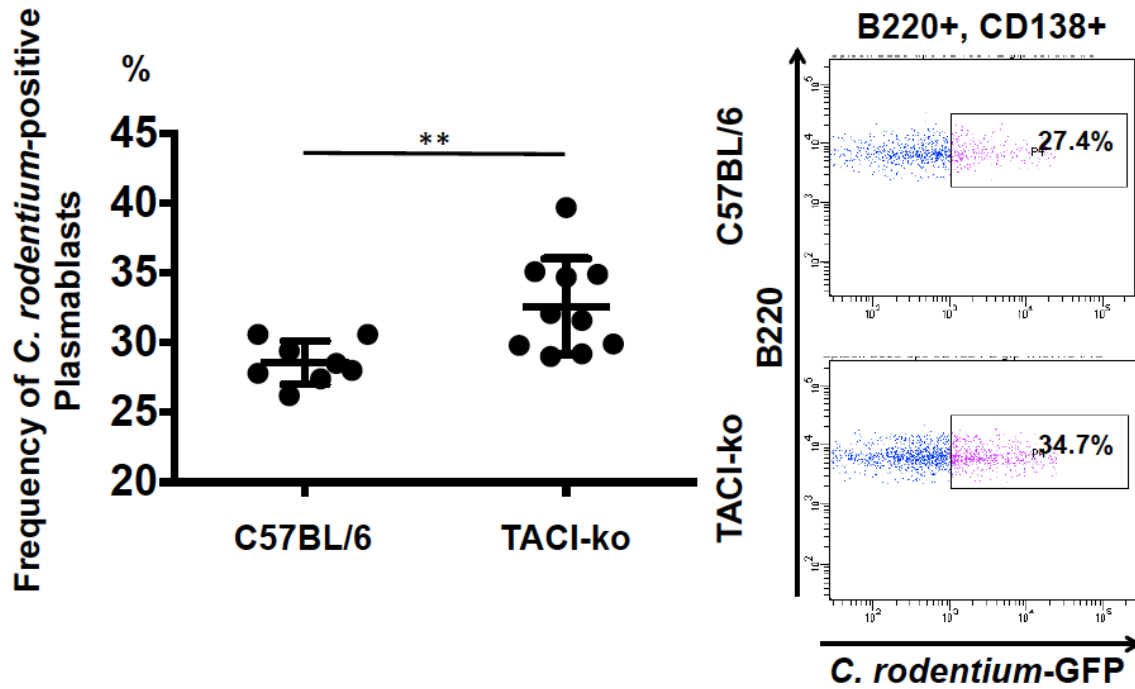


Figure 4

Figure 4: Identification of *C. rodentium* binding B cells. The figure shows flow cytometry analysis of splenocytes obtained from mice infected twice with *C. rodentium* with a month interval. Cells were analyzed 10 days after second infection. The left panel shows the frequency (Y-axis) of plasmablasts that bind to GFP-expressing *C. rodentium*. Panels on the right show representative dot plots (from N=10/strain) of splenocytes stained with anti-B220 antibody (Y-axis), anti-CD138 antibody and incubated with GFP-expressing *C. rodentium* (X-axis). Cells shown are both B220+ and CD138+ (plasmablasts). Rectangles depict the population that stains positively (above background) with the GFP-labeled *C. rodentium*.

Supplementary methods:

Primers for real-time PCR:

CTATTACGTGGATCGCAATGATGA (cIAP1-F),
TCTCCAGGGCCAAAATGCACCACT (cIAP1-R); CGGGACATCTTGACGGAC
(BCL6-F), CAGGGCTGATTTTCAGGATCTA (BCL6-R);
CATTCTGCCCAAAGATCAGT (TACI-F), TGGTGCCTTCCTGAGTTGTCT (TACI-
R); AGCCCATCACCATCTTCCAGGAG (GAPDH-F),
CCTCTCGCCACAGCTTTCCA (GAPDH-R)

Primers for sequencing:

Forward Primers:

V1-72 (IMGT) (2, 3)) specific forward primer

(CTTGACCCAGATGTCCCTTCTTCTCCAGCAGG);

Universal forward primers (AGGTSMARCTGCAGSAGTCWGG),

(GGGAATTCGAGGTGCAGCTGCAGGAGTCTGG) and

(SARGTNMAGCTGSAGSAGTC);

Reverse primers:

J4 reverse primer (TTCCTTGACCCCAGTAGTCCA).

Nested PCR:

The nested PCR was done using the same forward primer and one of several reverse primers located on J1-J2 intron (ATGGCCTGACATGGGGAGATCTG), J2-J3 intron

(GGGTCTAG AGGTGTCCCTAGTCCTTCATGACC) or J3-J4 intron (GGGTCTAG AGGTGTCCCTAGTCCTTCATGACC).

Amplification and sequencing of VH and VL exons from single sorted germinal center B cells isolated from mice infected with *C. rodentium* were done according to (1). Primers for amplifying and sequencing $\lambda 1$ and $\lambda 2$ light chain exons from hybridomas were according to (1)

References:

1. Tiller T, Busse CE, and Wardemann H. Cloning and expression of murine Ig genes from single B cells. *Journal of Immunological Methods*. 2009;350(1-2):183-93.
2. Giudicelli V, Chaume D, and Lefranc MP. IMGT/GENE-DB: a comprehensive database for human and mouse immunoglobulin and T cell receptor genes. *Nucleic Acids Res*. 2005;33(Database issue):D256-61.
3. Alamyar E, Giudicelli V, Li S, Duroux P, and Lefranc MP. IMGT/HighV-QUEST: the IMGT(R) web portal for immunoglobulin (IG) or antibody and T cell receptor (TR) analysis from NGS high throughput and deep sequencing. *Immunome Res*. 2012;8(1):26.