Table S1. List of Antibodies Used in the Present Study

Primary antibody	S Used in the Present Student Catalog number	Company					
FLAG-tag	MA1-91878	Thermo Fisher Scientific					
hTM4SF4	NBP1-69647	Novus Biologicals					
Sox2	#3579	Cell Signaling Technology (CST)					
Oct4	#2750	CST					
β-catenin	#8480	CST					
ALDH1A1	ab6192	Abcam					
ALDH1A3	ab129815	Abcam					
CD44	#3570	CST					
β-actin	sc-47778	Santa Cruz, Biotechnology					
E-cadherin	ab15148	Abcam					
N-cadherin	#610920	BD Transduction					
Vimentin	MA5-14564	Invitrogen					
Snail	sc-10432	Santa Cruz					
Twist	sc-15393	Santa Cruz					
Slug	sc-166476	Santa Cruz					
Zeb1	sc-25388	Santa Cruz					
Phospho-IGF1R	sc-81499	Santa Cruz					
IGF1Rβ	sc-3027	Santa Cruz					
IL-1β	AF-401-NA	R&D system					
Osteopontin	AF1433	R&D system					
IGF1	AF-291-NA	Bio-Techne					
Phospho-PI3K	sc-293115	Santa Cruz					
PI3K	#3011	CST					
Phospho-AKT	#9271	CST					
AKT	#9272	CST					
Phospho-GSK3β	#9322	CST					
GSK3β	#9315	CST					
Phospho-JAK	#3711	CST					
JAK	#3230	CST					
Phospho-FAK	#8556	CST					
FAK	#3285	CST					
Phospho-STAT3	#9145	CST					
STAT3	sc-8019	Santa Cruz					
PD-L1	#13684	CST					
B7-H4	12080-1-AP	Proteintech					
CD63	ab8219	Abcam					
CD63-biotin	ab134331	Abcam					
Normal mouse IgG1	sc-3877	Santa Cruz					

Table S2. Primers Used for Recombinant PCR of the Chimeric 2B7 Antibody Gene Cloning

Gene		Primer Sequences (5'→3')
Heavy chain	Forward	GAC GAA TTC ACT CTA ACC ATG GAA TGG A
signal peptide	Reverse	CTT CAC CTC GGA GTG GAC ACC TGT AGT TA
Heavy chain	Forward	GTC CAC TCC GAG GTG AAG CTG GAG GAG TC
variable region	Reverse	TTG GGC CCT TGG TGG AGG CTG CAG AGA CAG TGA CCA G
Light chain	Forward	CTG CAA AGC TTC GGC ACG AGC A
signal peptide	Reverse	CAC AAT ATC TCC TTC AAC ACC AGA CAA CC
Light chain	Forward	GTT GAA GGA GAT ATT GTG ATG ACC CAG TCT
variable region	Reverse	CCA CCG TAC GTT TGA TTT CCA GCT T

Table S3. Primers Used Recombinant PCR to Produce a Gene Sequence Encoding the Heavy Chain Variable Region, Referred to as Hz2B7-1.0

Primer	<b>Sequence (5</b> `→ 3`)
2B7-S-H-5`	GAC GAA TTC ACT CTA ACC ATG GAA TGG AGC TGG GTC TTT CTC TTC CTG TCA GTA ACT ACA GGT GTC CAC TCC GAG ATC ACC CTG AA
2B7-S-H-3`	CTC AGG CTG AAG CCG CTG AAG GTG CAG GTC AGG GTC AGG GTC TGG GTG GGC TTC ACC AGG GTG GGG CCG CTC TCC TTC AGG GTG ATC TCG GAG
2B7-HC-1-5`	CAG CGG CTT CAG CCT GAG CAC TTA TGG TAT AGG AGT AAG CTG GAT CAG GCA GCC CCC CGG CAA GGC CCT GGA GTG GCT GGC CCA CAT TTG
2B7-HC-1-3`	CCT GGT TCT TGC TGG TGT CCT TGG TGA TGG TCA GCC TGC TCT TCA GGG CTG TGT TAT AGT ACT TAT CAT TCC ACC AAA TGT GGG CCA GCC AC
2B7-HC-2-5`	ACA CCA GCA AGA ACC AGG TGG TGC TGA CCA TGA CCA ACA TGG ACC CCG TGG ACA CCG CCA CCT ACT ACT GCG CCA GGA AGG AGG GCA GCT
2B7-HC-2-3`	GGG CCC TTG GTG GAG GCG CTG CTC ACG GTC ACC AGG GTG CCC TGG CCC CAG TAA GCA AAG GGG GCC GAG CTG CCC TCC TTC CTG G
Ch57-HC-5`	GAC GAA TTC ACT CTA ACC AT
2B7-WH-3`	TTG GGC CCT TGG TGG AGG CGC TGC T

Table S4. Primers Used for Humanized 2B7 Antibody Affinity Maturation

Gene	Primer	Sequence $(5 \rightarrow 3)$						
	h2B7-H.C-W55S-5`	TTT GGT CGA ATG ATA ATA AGT A						
	h2B7-H.C-W55S-3`	TCA TTC GAC CAA ATG TGG GCC A						
II a a a ser Charles	h2B7-H.C-W55Y-5`	TTT GGT ATA ATG ATA ATA AGT A						
Heavy Chain	h2B7-H.C-W55Y-3`	TCA TTA TAC CAA ATG TGG GCC A						
	h2B7-H.C-SLIC-5`	GCC AGT GTG CTG GAA TTC ACT CTA ACC						
	h2B7-H.C-SLIC-3`	AAG ACC GAT GGG CCC TTG GTG GAG						
	h2B7-L.C-N31F-5`	TTT TAT TCA GTA GCA ATC AA						
	h2B7-L.C-N31F-3`	CTA CTG AAT AAA AGG CTC TGA C						
Tiple Chain	h2B7-L.C-N31V-5`	TTT TAG TAA GTA GCA ATC AA						
Light Chain	h2B7-L.C-N31V-3`	CTA CTT ACT AAA AGG CTC TGA C						
	h2B7-L.C-SLIC 5`	ATA GGG AGA CCC AAG CTT CGG CAC GAG CAG A						
	h2B7-L.C-SLIC 3`	TGG TGC AGC CAC CGT ACG CTT GAT CTC CA						

Table S5. Isotypes of Mouse Anti-hTM4SF4 mAbs Generated in This Study

Clones	ELISA*	Ig Isoty	pe
	(OD450)	НС	LC
<b>2B</b> 7	3.654	IgG1	Kappa
4C1	3.578	IgG2a	Kappa
8E2	3.686	IgG1	Kappa
8E5	3.521	IgG2a	Kappa
12A8	3.685	IgG2b	Kappa

<sup>\*</sup>Coating antigen: hTM4SF4 (126-140) peptide-BSA

Table S6. Summary of SPR analysis for Mouse Anti-TM4SF4 Antibodies

mAb	2B7	4C1	12A8
k <sub>a</sub> (1/Ms) #	4.85E+04	1.69E+04	2.47E+04
$k_{\rm d}$ (1/s)	1.29E-04	1.09E-03	2.06E <b>-</b> 04
<b>K</b> <sub>D</sub> ( <b>M</b> ) §	2.66E-09	6.45E-08	8.34E-09
Rmax (RU)	103.4	129.5	131.6
$\chi^2 (RU^2)^{\$}$	1.47	7.42	0.254
$U$ -value $^{\xi}$	12	7	1

 $<sup>^{\#}</sup>$ Sensor Chip: Biotin-hTM4SF4 (T126–E140) peptide captured SA sensor  $^{\$}$   $k_a/k_d$  model: 1:1 binding mass transfer.  $^{\$}$  $\chi^2$ :< 10% of Rmax is significant for the experimental data

Table S7. Summary of the SPR analysis for Humanized 2B7 Antibodies

mAb	Chi-2B7	Hz2B7-1.1	Hz2B7-1.2	Hz2B7-4.3
k <sub>a</sub> (1/Ms) #	5.077E+03	7.523E+03	1.259E+04	9.585E+03
$k_{\rm d}$ (1/s)	3.084E-04	1.837E-04	7.590E-05	7.827E-04
$K_{D}(M)^{\S}$	6.074E-08	2.442E-08	6.030E-09	8.165E-08
Rmax (RU)	26.53	76.30	81.32	63.53
$\chi^2 (RU^2)^{\$}$	0.0809	0.252	1.78	0.352
U-value	3	2	12	1

<sup>&</sup>lt;sup>E</sup>The U-value: an estimate of the uniqueness of the calculated values for rate constants and Rmax. If the U-value is below about 15 the parameter values are not significantly correlated.

 $<sup>^{\#}</sup>$ Sensor Chip: Biotin-TM4SF4<sub>(126-140)</sub> peptide captured SA sensor  $^{\S}$   $k_a/k_d$  model: 1:1 binding mass transfer.  $^{\S}$   $\chi^2$  :< 10% of Rmax is significant for the experimental data  $^{\S}$ The U-value: an estimate of the uniqueness of the calculated values for rate constants and Rmax. If the U-value is below about 15 the parameter values are not significantly correlated.

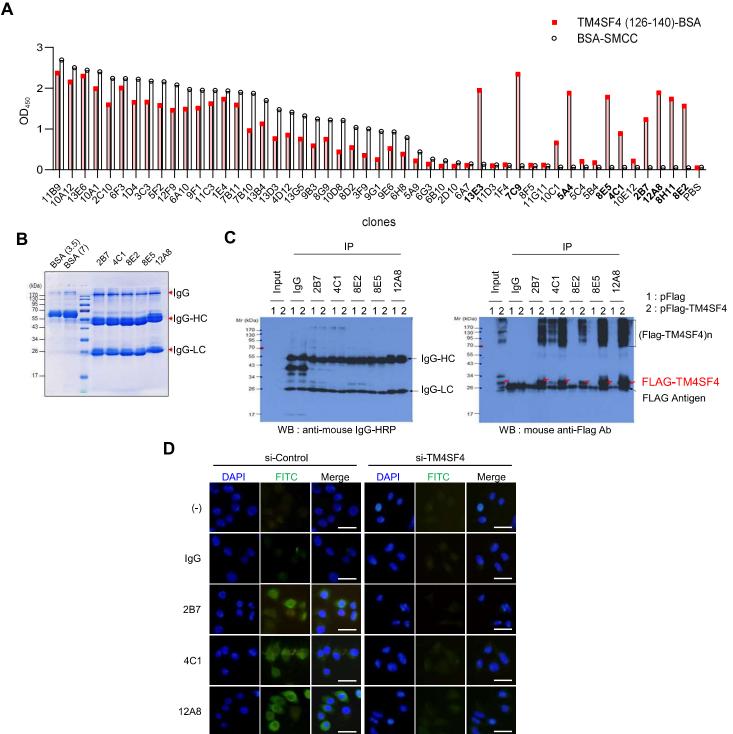


Figure S1. Generation of anti-hTM4SF4 mAbs using a 15-mer peptide derived from the large extracellular loop of hTM4SF4 (A) BALB/c mice were immunized with an hTM4SF4 (126–140) peptide-BSA conjugate for B cell hybridoma generation, which yielded fifty B cell hybridoma clones. The reactivity of antibodies produced by these B cell hybridomas against antigenic peptide were determined by ELISA using the hTM4SF4(126–140) peptide conjugated to BSA. BSA modified with SMCC was used as a control antigen. (B) Five antibodies against hTM4SF4(126–140) were purified and analysed by 10% reducing SDS-PAGE. (C) Five novel anti-hTM4SF4 antibodies immunoprecipitated the FLAG tagged TM4SF4 expressed in HEK293T cells (original data of Figure 1D). (D) A549 cells were stained with anti-hTM4SF4 mAbs (2B7, 4C1 and 12A8) and an FITC-conjugated secondary reagent, whereas TM4SF4-knockdown suppressed FITC staining with these antibodies.

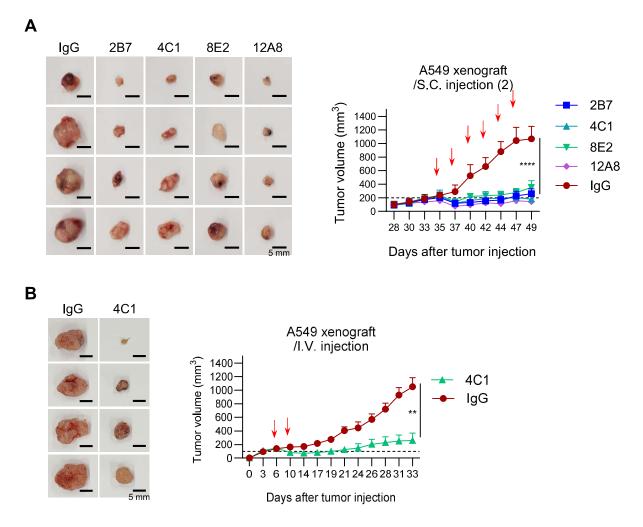


Figure S2. Anti-hTM4SF4 mAbs inhibit NSCLC growth in vivo mouse xenograft model (A) Tumor growth inhibition in an A549 NSCLC xenograft mouse model by subcutaneous injection of anti-hTM4SF4 antibodies. Antibodies (2B7, 4C1, 8E2, and 12A8) were administered when the tumor size of xenograft model reached approximately 200 mm³. Injections were administered six times at 2–3 day intervals (indicated by red arrows on the plot), with an injection dose of 10 μg per mouse, totaling 60 μg per mouse. (B) Tumor growth inhibition in an A549 NSCLC xenograft mouse model by intravenous injection of anti-hTM4SF4 antibodies. Antibody, 4C1 was administered intravenously to the mice five days after the subcutaneous injection of tumor cells. Injections were administered two times at 4-day intervals (indicated by red arrows on the plot).

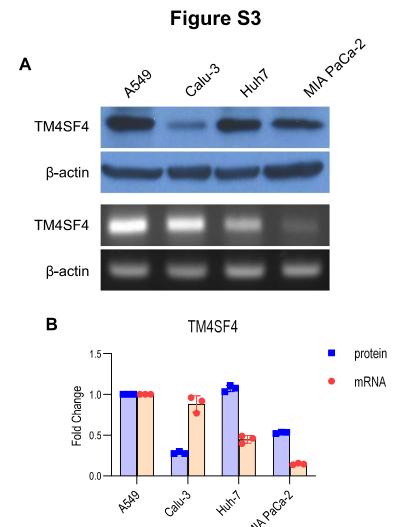


Figure S3. TM4SF4 expression in various tumor cell lines. (A) TM4SF4 expression in various tumor cell lines was analyzed by Western blotting and RT-PCR in A549 and Calu-3 (NSCLC cell lines), Huh-7 (hepatocellular carcinoma), and MIA PaCa-2 (pancreatic carcinoma) cells. Western blot and RT-PCR analysis revealed that TM4SF4 was highly expressed in A549, Huh-7 and MIA PaCa-2 cells, whereas its expression was relatively low in Calu-3 cells. (B) All experiments were performed in triplicate, and band intensities were quantified by ImageJ software and normalized to β-actin expression.

#### A. Mouse 2B7 Heavy chain variable region: Subgroup (IIIC)

GAG E	GTG V	AAG K	CTG L	GAG E	GAG E	TCT S	GGC G	CCT P	GGG G	ATA I	TTG L	CAG O	CCC P	TCC S	CAG Q	ACC T	CTC L	54 18
						V-2.2.	-					· •			CDR-	-н1		
AGT	CTG	ACT	TGT	TCT	TTC	TCT	GGG	TTT	TCA	CTG	AGC	ACT	TAT	GGT	ATA	GGA	GTA	108
S	L	T	C	S	F	S	G	F	S	L	S	T	Y	G	I	G	Λ	36
AGC	TGG	ATT	CGT	CAG	CCT	TCT	GGG	AAG	GGT	CTG	GAG	TGG	CTG	GCA	CAC	ATT	TGG	162
S	W	I	R	Q	P	S	G	K	G	L	E	W	L	A	H	I	W	54
N/A				CDR	-H2							7.2						
TGG	AAT	GAT	AAT	AAG	TAC	TAT	AAC	ACA	GCC	CTG	AAG	AGC	CGG	CTC	ACA	ATC	TCC	216
W	N	D	N	K	Y	Y	N	T	A	L	K	S	R	L	T	I	S	72
AAG	GAT	ACC	TCC	AAC	AAC	CAG	GTA	TTC	CTC	AAG	ATC	GCC	AGT	GTG	GAC	ACT	GCA	270
K	D	T	S	N	N	Q	Λ	F	L	K	I	A	S	Λ	D	$\mathbf{T}$	Α	90
													CDR-	-н3				
GAT	ACT	GCC	ACA	TAC	TAC	TGT	GCT	CGA	AAG	GAG	GGC	AGC	TCG	GCC	CCC	TTT	GCT	324
D	T	A	T	Y	Y	С	A	R	K	E	G	T	S	A	P	F	A	108
TAC	TGG	GGC	CAA	GGG	ACT	CTG	GTC	ACT	GTC	TCT	GCA	360						
F	W	G	Q	G	Т	L	Λ	T	V	S	Α	120						

#### B. Mouse 2B7 Light chain variable region: Subgroup (IIIC)

GAT	ATT	GTG	ATG	ACC	CAG	TCT	CCA	TCC	TCC	CTG	GCT	ATG	TCA	GTA	GGA	CAG	AAG	54
D	I	Λ	M	T	Q	S	P	S	S	L	A	M	S	Λ	G	Q	K	18
												CDR	-L1					
GTC	ACT	ATG	AGC	TGC	AAG	TCC	AGT	CAG	AGC	CTT	TTA	AAT	AGT	AGC	AAT	CAA	AAG	108
Λ	T	M	S	C	K	S	S	Q	S	L	L	N	S	S	N	Q	K	36
AAC	TAT	TTG	GCC	TGG	TAC	CAG	CAG	AAA	CCA	GGA	CAG	TCT	CCT	AAA	CTT	CTG	ATA	162
N	Y	L	A	W	Y	Q	A	E	D	L	A	D	Y	F	C	Q	Q	54
	25		CD	R-L2	2	55036												
TAC	TTT	GCA	TCC	ACT	AGG	GAA	TCT	GGG	GTC	CCT	GAT	CGC	TTC	ATA	GGC	AGT	GGA	216
H	Y	Α	S	T	R	E	S	G	Λ	P	D	R	F	Ι	G	S	G	72
TCT	GGG	ACA	GAT	TTC	ACT	CTT	ACC	ATC	AGC	AGT	ATG	CAG	GCT	GAA	GAC	CTG	GCA	270
S	G	T	D	F	T	L	T	I	S	S	M	Q	A	E	D	L	A	90
				2.0			CDR	-L3				400						
GAT	TAC	TTC	TGT	CAG	CAA	CAT	TAT	AGA	ACT	CCT	CCG	ACG	TTC	GGT	GGA	GGC	ACC	324
D	Y	F	C	Q	Q	H	Y	R	T	P	P	Т	F	G	G	G	Т	108
AAG	CTG	GAA	ATC	AAA	339													
K	L	E	I	K	113													

**Figure S4 Sequence analysis of heavy and light chain variable regions of mouse 2B7 mAb (A)** Nucleotide and amino acid sequences of heavy chain of 2B7. CDRs 1, 2 and 3 are also shown. **(B)** Nucleotide and amino acid sequences of light chain of the 2B7. CDRs 1, 2 and 3 are also shown.

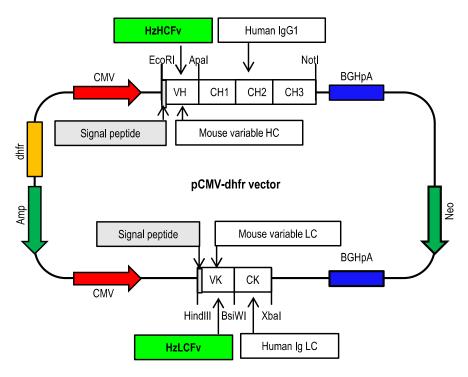
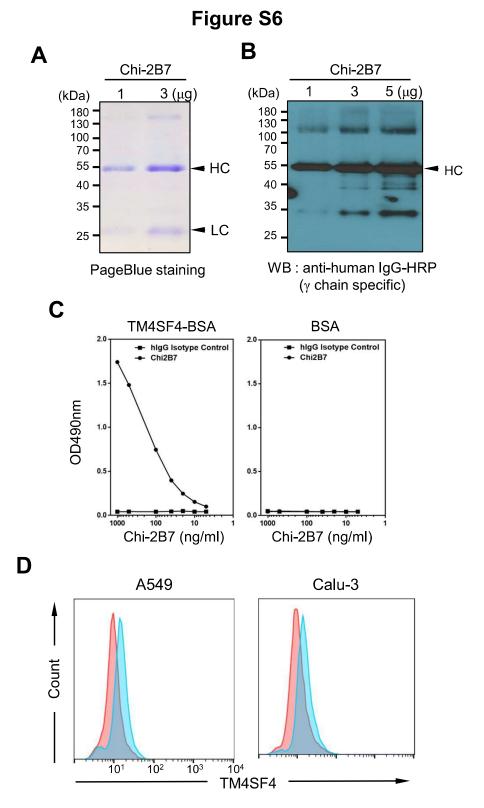


Figure S5. Schematic representation of the monoclonal antibody expression vectors for the chimeric and humanized versions of the 2B7 antibody. For the construction of the chimeric antibody expression vector (pdCMV-dhfr-Chi-2B7), the variable region gene of the 2B7 murine antibody and the human IgG1 constant region gene were inserted into the pCMV-dhfr vector. To develop the humanized version of the 2B7 antibody (Hz2B7-1.1), the CDRs of the 2B7 antibody were grafted onto the framework regions of the most homologous human antibody, 3QRG (described in detail in the Materials and Methods section), and inserted into the pCMV-dhfr vector, resulting in the pdCMV-dhfr-Hz2B7-1.1 vector.



**Figure S6.** Characterization of Chimeric 2B7 Antibody (A) 10% reducing SDS PAGE analysis of Chi-2B7, with heavy chain (HC) and light chain (LC) indicated. (B) Western blot of Chi-2B7 probed using α-human IgG-γ chain specific-HRP. (C) Binding affinity of Chi-2B7 to TM4SF4 peptide conjugated BSA and BSA (1 ug/ml) antigen by indirect ELISA. (D) Flow cytometry analysis of Chi-2B7 binding to NSCLC cell lines (A549, Calu-3), comparing Chi-2B7 staining (blue) to FITC-conjugated secondary antibody staining as a control (red).

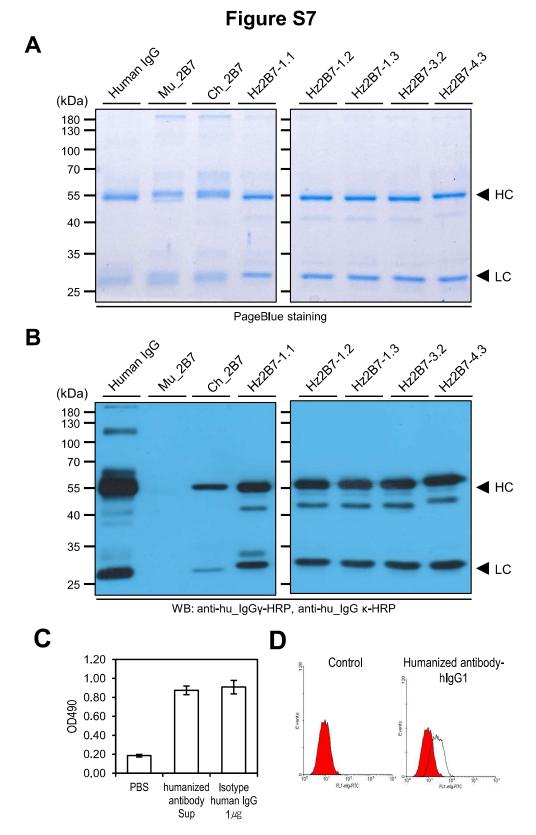


Figure S7. Characterization of Humanized 2B7 Antibodies (A) 10% reducing SDS PAGE analysis of various version of humanized 2B7 antibodies, with heavy chain (HC) and light chain (LC) indicated. (B) Western blot humanized 2B7 antibodies probed using α-human IgG-γ chain specific-HRP. (C) Detection of humanized antibody in cell cultured supernatant by indirect ELISA. Human IgG1 was used as control. (D) Flow cytometry analysis of humanized antibody binding to NSCLC A549 cell line. Hz2B7 antibody staining (unfilled) to FITC-conjugated secondary antibody staining as a control (red).

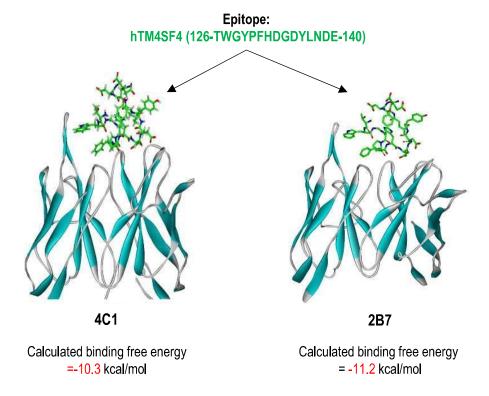


Figure S8. Docked pose of the epitope within the CDRs of the 4C1 and 2B7 mouse antibodies. The CDRs of antibodies are displayed in a ribbon representation, while the peptide epitope is depicted as a stick model. Calculated binding free energies were also indicated.

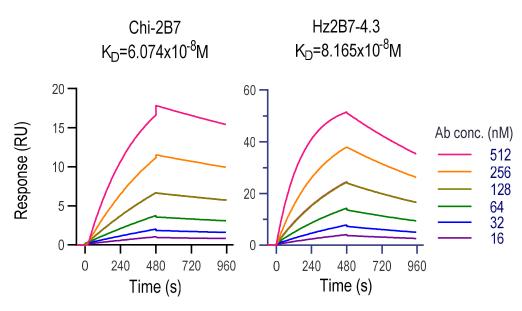


Figure S9. SPR sensorgrams showing the concentration-dependent binding kinetics of chimeric or humanized 2B7 antibody (Chi2B7, Hz2B7-4.3) to the hTM4SF4 peptide. Biotinylated hTM4SF4126–140 peptide was immobilized on an SA sensor chip, and the antibodies were applied at increasing concentrations. KD values are presented on each sensorgram, with detailed kinetic parameters of the antigen-antibody interactions provided in Table S7.

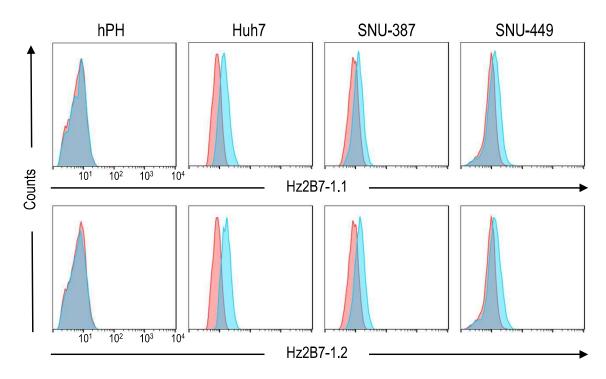


Figure S10. Binding activity of humanized 2B7 antibodies to hepatocellular carcinoma cell lines. The binding activity of Hz2B7-1.1 and Hz2B7-1.2 to primary hepatocytes (hPH) and hepatocellular carcinoma cell lines (Huh7, SNU-387, SNU 449) was measured by Flow cytometry.

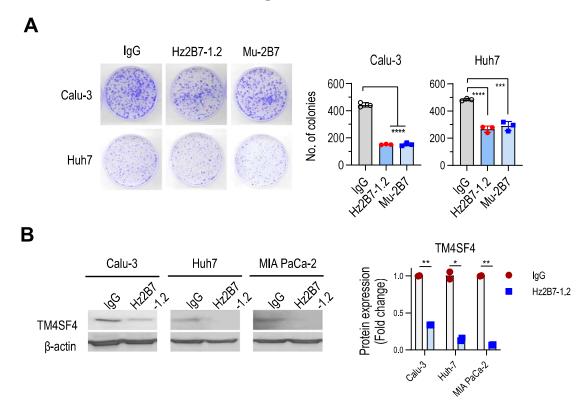


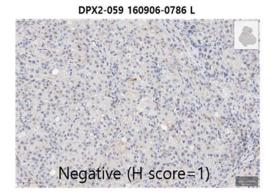
Figure S11. Hz2B7-1.2 antibody treatment suppressed tumor cell growth and reduced TM4SF4 expression (A) Colony-forming assay. Calu-3 (non–small cell lung cancer, NSCLC) and Huh-7 (hepatocellular carcinoma, HCC) cells were treated with Hz2B7-1.2 antibody (5  $\mu$ g/mL) and cultured for 10 days for colony formation. (B) TM4SF4 expression analysis. Various tumor cell lines were treated with Hz2B7-1.2 antibody (5  $\mu$ g/mL) for 48 h, washed, harvested, and analyzed for TM4SF4 expression by Western blot.

#### Figure S12 Α siCtl siTM4SF4 (#1) siTM4SF4 (#2) s 8.07 12.9 43.7 BrdU-FITC 103 103 G2.M G2.M G2.M 2.28 2.11 G0.G1 G0.G1 G0.G1 101 = 53.3 89.1 78.8 B C A549 TM4SF4 1.4 siCtl 100 siTM4SF4 #1 Relative mRNA expression 2.0 - 0.1 population (%) siTM4SF4 #2 60 40 <u></u>8 20 ns 0.0 silhasfa #1 SITMASFAHR 0 G0/G1 G2/M D Ε hlgG isotype Hz2B7-1.2 A549 hlgG 44.0 49.8 60 Hz2B7-1.2 population (%) ns **BrdU-FITC** 103 102 G2.M G2.M = 20 3.55 G0.G1 G0.G1 ns 51.4 45.9 0 100K G0/G1 G2/M

Figure S12. TM4SF4 inhibition suppress NSCLC growth by blocking G1 to S phase transition (A) Cell cycle analysis in siTM4SF4 or control siRNA-transfected A549 cells. Cells were treated with 30µM BrdU post 48h of transfection and analyzed in flow cytometry by using anti-BrdU-FITC and PI. (B) Relative cell cycle population at each cell cycle phase in TM4SF4-knockdown A549 cells. (C) TM4SF4 siRNA treatment reduced TM4SF4 mRNA expression by approximately 50%, as determined by RT-qPCR analysis. The siRNA sequences targeting human TM4SF4 were as follows: #7104-1 (Bioneer, Daejeon, Korea), antisense: 5'-CACCUUUCCCAAGAGAUCUtt-3', #7101-2 (Bioneer, Daejeon, Korea), antisense: 5'-CACCUUUCCCAAGAGAUCUtt-3'. (D) Cell cycle analysis in Hz2B7-1,2 or control isotype antibody-treated A549 cells. Cells were treated with BrdU 72 h after antibody treatment (5 µg/ml) and analyzed by flow cytometry using anti-BrdU-FITC and Pl. (E) Relative cell cycle population at each cell cycle phase in Hz2B7-1.2-treated A549 cells (ns. not significant, \*, p<0.05, \*\*, p<0.1, \*\*\*, p<0.005).

S

Lung cancer PDX		Image		TM4SF4 gene expression	1+,2+,3+ %	2+,3+ %	3+ %	Neg %	H score	IHC result	staining pattern
	DPX0-059	160412-0604	R	8.07	8	0	0	92	8	0	
candidate #	DPX2-059	160906-0786	L		1	0	0	99	1	0	
1.	DPX2-059	160906-0787	R		2	0	0	98	2	0	
	SPX1-095	150819-0095	L	1.25	0	0	0	100	0	0	
candidate #	SPX2-095	151012-0356	L	1.87	18	2	0	82	20	1+	membranous + cytoplasmic
2	SPX2-095	151012-0360	L		8	0	0	92	8	0	



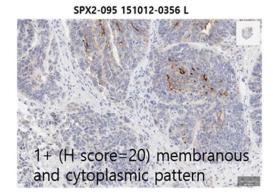
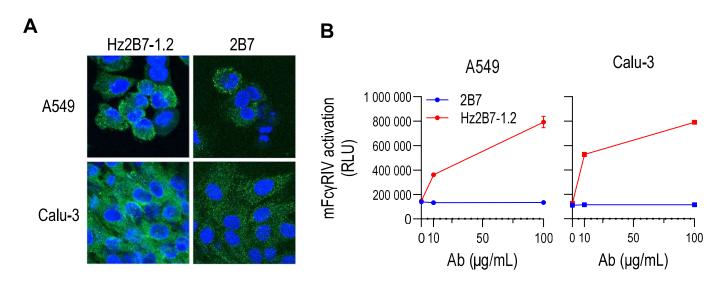


Figure S13. PDX model for lung adenocarcinoma expressing TM4SF4 To establish a PDX model for TM4SF4-expressing lung adenocarcinoma, the expression of TM4SF4 in patient tissues was analyzed by DEG analysis and IHC staining. DEG analysis showed that DPX0-059 had very high TM4SF4 expression; however, as shown in the table, no expression was detected by IHC. On the other hand, TM4SF4 expression was detected in passage 2 for SPX2-095. Therefore, we established a PDX model using patient tissue SPX2-095.



**Figure S14. Mouse FcγRIV ADCC Assay for Humanized Antibody Hz2B7-1.2** (A) Binding of anti-TM4SF4 antibodies to NSCLC cell lines. A549 and Calu-3 non–small cell lung cancer (NSCLC) cells cultured on coverslips were fixed, permeabilized, and stained with either humanized anti-TM4SF4 antibody Hz2B7-1.2 or mouse antibody 2B7, followed by FITC-conjugated secondary antibodies. Both antibodies bound to the cell surface of A549 and Calu-3 cells. **(B)** Mouse FcγRIV ADCC assay. Antibody-dependent cellular cytotoxicity (ADCC) activity was assessed against A549 and Calu-3 cells using mouse FcγRIV/NFAT–Jurkat effector cells (BPS Bioscience). The mouse IgG1 antibody 2B7 did not elicit detectable ADCC, whereas the humanized IgG1 antibody Hz2B7-1.2 induced robust levels of ADCC in both cell lines.