

FIG. 1

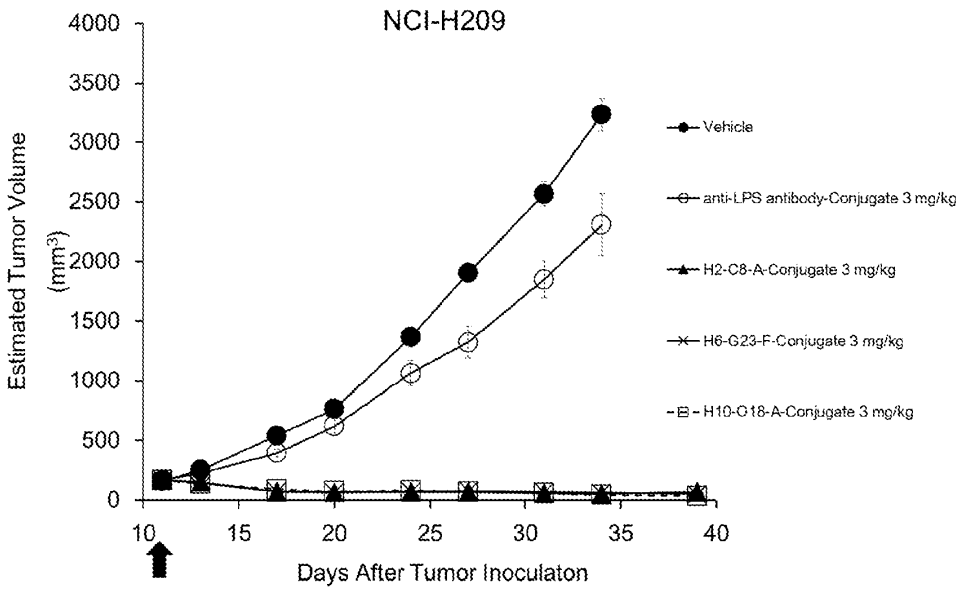


FIG. 2

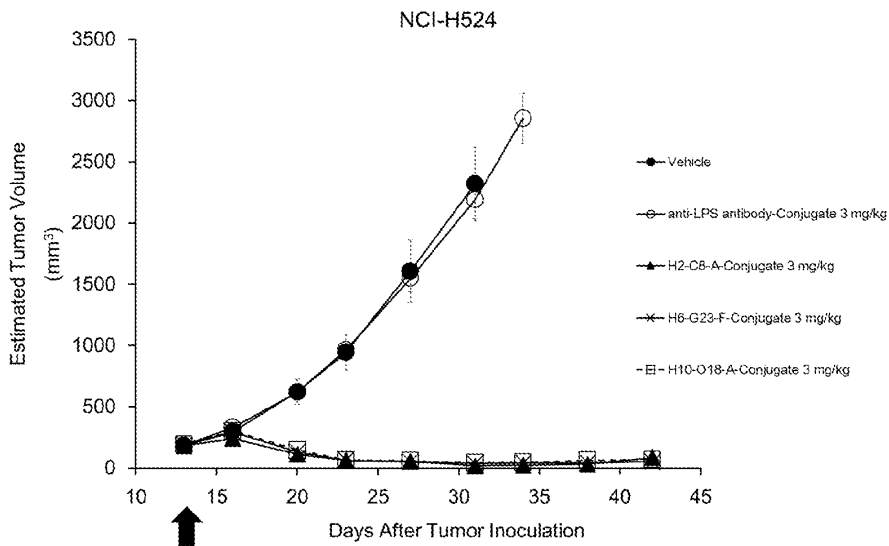


FIG. 3

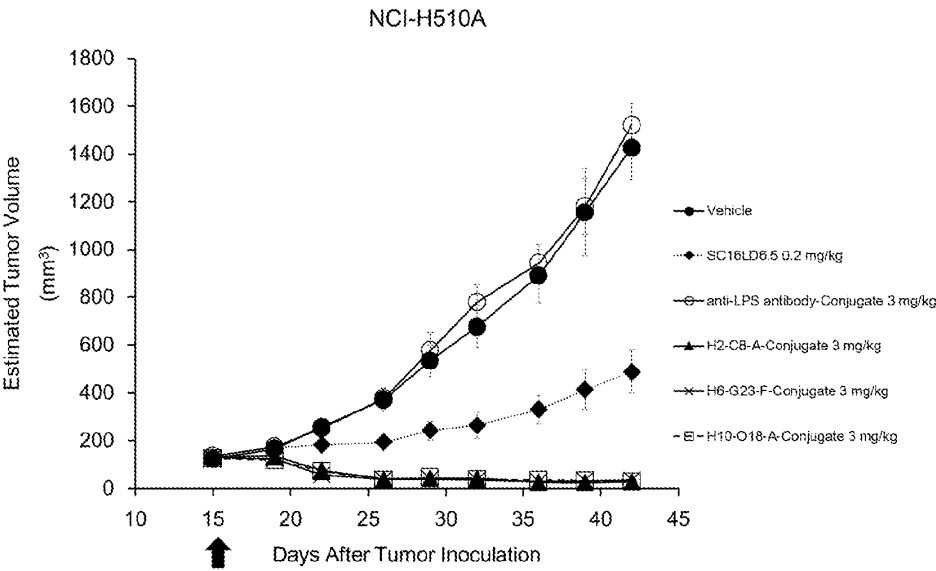


FIG. 4A

>NM_016941.4 Homo sapiens delta like canonical Notch ligand 3 (DLL3),
transcript variant 1, mRNA (SEQ ID NO: 55)

ACTCCCAGACCCCCACCAGAAGGCCATGGTCTCCCCACGGATGTCCGGGCTCCTCTCCAGACTGTG
ATCCTAGCGCTCATTTTCTCCCCAGACACGGCCCGCTGGCGTCTTCGAGCTGCAGATCCACTCTTTCC
GGCCGGGTCCAGGCCCTGGGGCCCCGCGGTCCCCCTGCAGCGCCCGGCTCCCCTGCCGCCTCTTCTTCAG
AGTCTGCCTGAAGCCTGGGCTCTCAGAGGAGGCCGCCGAGTCCCCGTGCGCCCTGGGCGCGGCGCTGAGT
GCGCGCGGACCGGTCTACACCGAGCAGCCCCGAGCGCCCGCGCCTGATCTCCACTGCCCGACGGCCTCT
TGCAGGTGCCCTTCCGGGACGCCTGGCCTGGCACCTTCTCTTTCATCATCGAAACCTGGAGAGAGGAGTT
AGGAGACCAGATTGGAGGGCCCGCCTGGAGCCTGCTGGCGCGCGTGGCTGGCAGGCGGCGCTTGGCAGCC
GGAGGCCCGTGGGCCCGGACATTACAGCGCGCAGGCGCCTGGGAGCTGCGCTTCTCGTACCGCGCGCGCT
GCGAGCCGCCTGCCGTGGGACCGGTGCACGCGCCTCTGCCGTCCGCGCAGCGCCCCCTCGCGGTGCGG
TCCGGGACTGCCCCCTGCGCACCGCTCGAGGACGAATGTGAGCGCCGCTGGTGTGCCGAGCAGGCTGC
AGCCCTGAGCATGGCTTCTGTGAACAGCCCGGTGAATGCCGATGCCTAGAGGGCTGGACTGGACCCCTCT
GCACGGTCCCTGTCTCCACCAGCAGCTGCCTCAGCCCCAGGGGCCCGTCTCTGCTACCACCGGATGCCT
TGTCCCTGGGCCTGGGCCCTGTGACGGGAACCCGTGTGCCAATGGAGGCAGCTGTAGTGAGACACCCAGG
TCCTTTGAATGCACCTGCCCGCGTGGGTTCTACGGGCTGCGGTGTGAGGTGAGCGGGGTGACATGTGCAG
ATGGACCCCTGCTTCAACGGCGGCTTGTGTGTCGGGGGTGCAGACCCTGACTCTGCCTACATCTGCCACTG
CCCACCCGGTTTCCAAGCTCCAACCTGTGAGAAGAGGGTGGACCGGTGCAGCCTGCAGCCATGCCGCAAT
GGCGGACTCTGCCTGGACCTGGGCCACGCCCTGCGCTGCCGCTGCCGCGCCGGCTTCGCGGGTCTCTGCT
GCGAGCACGACCTGGACGACTGCGCGGGCCGCGCCTGCGCTAACGGCGGCACGTGTGTGGAGGGCGGGCG
CGCGACCGCTGCTCCTGCGCGCTGGGCTTCGGCGGCCGCGACTGCCGCGAGCGCGCGGACCCGTGCGCC
GCGCGCCCCTGTGCTCACGGCGGCCGCTGCTACGCCCACTTCTCCGGCCTCGTCTGCGCTTGCCTCCCCG
GCTACATGGGAGCGCGGTGTGAGTTCCAGTGCACCCCGACGGCGCAAGCGCCTTGCCCGCGGCCCCGCC
GGGCCTCAGGCCCGGGGACCCCTCAGCGCTACCTTTTGCCCTCCGGCTCTGGGACTGCTCGTGGCCGCGGGC
GTGGCCGGCGCTGCGCTCTTGCTGGTCCACGTGCGCCGCCGTGGCCACTCCCAGGATGCTGGGTCTCGCT
TGCTGGCTGGGACCCCGGAGCCGTCACTCCACGCACTCCCGGATGCACTCAACAACCTAAGGACGCAGGA
GGGTTCCGGGGATGGTCCGAGCTCGTCCGTAGATTGGAATCGCCCTGAAGATGTAGACCCTCAAGGGATT

FIG. 4A CON'T

TATGTCATATCTGCTCCTTCCATCTACGCTCGGGAGGTAGCGACGCCCCCTTTCCCCCGCTACACACTG
GGCGCGCTGGGCAGAGGCAGCACCTGCTTTTCCCTACCCCTTCCTCGATTCTGTCCGTGAAATGAATTGG
GTAGAGTCTCTGGAAGGTTTAAAGCCCATTTTCAGTTCTAACTTACTTTTCATCCTATTTTGCATCCCTCT
TATCGTTTTGAGCTACCTGCCATCTTCTCTTGAAAAACCTATGGGCTTGAGGAGGTACGATGCCGACT
CCGCCAGAGCTTTTCCACTGATTGTACTCAGCGGGGAGGCAGGGGAGGCAGAGGGGCAGCCTCTCTAATG
CTTCCTACTCATTTTGTTCCTAGGCCTGACGCGTCTCCTCCATCCGCACCTGGAGTCAGAGCGTGGAATTT
TTGTATTTGCTCGGTGGTCCCCAGTCTCTGCCCCAGAGGCTTTGGAGTTCAATCTTGAAGGGGTGTCTGG
GGGAACCTTACTGTTGCAAGTTGTAAATAATGGTTATTTATATCCTATTTTTTCTACCCCATCTCTCTA
GAAACACCTATAAAGGCTATTATTGTGATCA

>NP_058637.1 delta-like protein 3 isoform 1 precursor [Homo sapiens] (SEQ ID
NO: 50)

MVSPRMSGLLSQTVILALIFLPQTRPAGVFELQIHSFGPGPGAPRSPCSARLPCRLFFRVCLKPGLSE
EAAESPCALGAALSARGPVYTEQPGAPAPDLPLPDGLLQVPFRDAWPGTFSFIETWREELGDQIGGPAW
SLLARVAGRRRLAAGGPWARDIQRAGAWELRFSYRARCEPPAVGTACTRLCRPRSAPSRCPGLRPCAPL
EDECEAPLVCRAGCSPEHGFCEQPGECRCLEGWTGPLCTVPVSTSSCLS PRGPSSATTGCLVPGPGPCDG
NPCANGGSCSETPRSFECTCPRGFYGLRCEVSGVTCADGPCFNGGLCVGGADPDSAYICHCPPGFQGSNC
EKRVDRCSLQPCRNGGLCLDLGHALRCRCRAGFAGPRCEHDLDDCAGRACANGGTCVEGGGAHRCSCALG
FGGRDCRERADPCAARPCAHHGRGYAHFSGLVACAPGYMGARCEFPVHPDGASALPAAPPGLRPGDPQR
YLLPPALGLLVAAGVAGAALLLVHVRRRGHSQDAGSRLLAGTPEPSVHALPDALNNLRTQEGSGDGPSSS
VDWNRPEDVDPQGIYVISAPSIYAREVATPLFPPLHTGRAGQRQHLLFPYPSSILSVK

FIG. 4B

>NM_203486.3 Homo sapiens delta like canonical Notch ligand 3 (DLL3),
transcript variant 2, mRNA (SEQ ID NO: 56)

ACTCCCAGACCCCCACCAGAAGGCCATGGTCTCCCCACGGATGTCCGGGCTCCTCTCCAGACTGTG
ATCCTAGCGCTCATTTTCTCCCCAGACACGGCCCGCTGGCGTCTTCGAGCTGCAGATCCACTCTTTCC
GGCCGGGTCCAGGCCCTGGGGCCCCGCGGTCCCCCTGCAGCGCCCGGCTCCCCTGCCGCCTCTTCTTCAG
AGTCTGCCTGAAGCCTGGGCTCTCAGAGGAGGCCGCCGAGTCCCCGTGCGCCCTGGGCGCGGCGCTGAGT
GCGCGCGGACCGGTCTACACCGAGCAGCCCCGAGCGCCCGCGCCTGATCTCCACTGCCCGACGGCCTCT
TGCAGGTGCCCTTCCGGGACGCCTGGCCTGGCACCTTCTCTTTCATCATCGAAACCTGGAGAGAGGAGTT
AGGAGACCAGATTGGAGGGCCCGCCTGGAGCCTGCTGGCGCGCGTGGCTGGCAGGCGGCGCTTGGCAGCC
GGAGGCCCGTGGGCCCGGACATTACAGCGCGCAGGCGCCTGGGAGCTGCGCTTCTCGTACCGCGCGCGCT
GCGAGCCGCCTGCCGTGGGACCGGTGCACGCGCCTCTGCCGTCCGCGCAGCGCCCCCTCGCGGTGCGG
TCCGGGACTGCCCCCTGCGCACCGCTCGAGGACGAATGTGAGCGCCGCTGGTGTGCCGAGCAGGCTGC
AGCCCTGAGCATGGCTTCTGTGAACAGCCCGGTGAATGCCGATGCCTAGAGGGCTGGACTGGACCCCTCT
GCACGGTCCCTGTCTCCACCAGCAGCTGCCTCAGCCCCAGGGGCCCGTCTCTGCTACCACCGGATGCCT
TGTCCCTGGGCCTGGGCCCTGTGACGGGAACCCGTGTGCCAATGGAGGCAGCTGTAGTGAGACACCCAGG
TCCTTTGAATGCACCTGCCCGCGTGGGTTCTACGGGCTGCGGTGTGAGGTGAGCGGGGTGACATGTGCAG
ATGGACCCCTGCTTCAACGGCGGCTTGTGTGTCGGGGGTGCAGACCCTGACTCTGCCTACATCTGCCACTG
CCCACCCGGTTTCCAAGCTCCAACCTGTGAGAAGAGGTTGGACCGGTGCAGCCTGCAGCCATGCCGCAAT
GGCGGACTCTGCCTGGACCTGGGCCACGCCCTGCGCTGCCGCTGCCGCGCCGGCTTCGCGGGTCTCTCGCT
GCGAGCACGACCTGGACGACTGCGCGGGCCGCGCCTGCGCTAACGGCGGCACGTGTGTGGAGGGCGGGCG
CGCGACCGCTGCTCCTGCGCGCTGGGCTTCGGCGGCCGCGACTGCCGCGAGCGCGCGGACCCGTGCGCC
GCGCGCCCCTGTGCTCACGGCGGCCGCTGCTACGCCCACTTCTCCGGCCTCGTCTGCGCTTGCCTCCCCG
GCTACATGGGAGCGCGGTGTGAGTTCCAGTGCACCCCGACGGCGCAAGCGCCTTGCCCGCGGCCCCGCC
GGGCCTCAGGCCCGGGGACCCCTCAGCGCTACCTTTTGCCCTCCGGCTCTGGGACTGCTCGTGGCCGCGGGC
GTGGCCGGCGCTGCGCTCTTGCTGGTCCACGTGCGCCGCCGTGGCCACTCCCAGGATGCTGGGTCTCGCT
TGCTGGCTGGGACCCCGGAGCCGTCACTCCACGCACTCCCGGATGCACTCAACAACCTAAGGACGCAGGA
GGGTTCCGGGGATGGTCCGAGCTCGTCCGTAGATTGGAATCGCCCTGAAGATGTAGACCCTCAAGGGATT

FIG. 4B CON'T

TATGTCATATCTGCTCCTTCCATCTACGCTCGGGAGGCCTGACGCGTCTCCTCCATCCGCACCTGGAGTC
AGAGCGTGGATTTTGTATTTGCTCGGTGGTGCCAGTCTCTGCCCCAGAGGCTTTGGAGTTCAATCTTG
AAGGGGTGTCTGGGGAACTTTACTGTTGCAAGTTGTAAATAATGGTTATTTATATCCTATTTTTTCTCA
CCCCATCTCTCTAGAAACACCTATAAAGGCTATTATTGTGATCA

>NP_982353.1 delta-like protein 3 isoform 2 precursor [Homo sapiens] (SEQ
ID NO: 51)
MVSPRMSGLLSQTVILALIFLPQTRPAGVFELQIHSFGPGPGPGAPRSPCSARLPCRLFFRVCLKPGLSE
EAAESPCALGAALSARGPVYTEQPGAPAPDLPLPDGLLQVPFRDAWPGTFSFIIETWREELGDQIGGPAW
SLLARVAGRRRLAAGGPWARDIQRAGAWELRFSYRARCEPPAVGTACTRLCRPRSAPSRGPGRLRPCAPL
EDECEAPLVCRAGCSPEHGFCEQPGECRCLEGWTGPLCTVPVSTSSCLS PRGPSSATTGCLVPGPGPCDG
NPCANGGSCSETPRSFECTCPRGFYGLRCEVSGVTCADGPCFNGGLCVGGADPDSAYICHCPPGFQGSNC
EKRVDRCSLQPCRNGGLCLDLGHALRCRCRAGFAGPRCEHDLDDCAGRACANGGTCVEGGGAHRCSCALG
FGGRDCRERADPCAARPCAHGGRCYAHFSGLVACAPGYMGARCEFPVHPDGASALPAAPPGLRPGDPQR
YLLPPALGLLVAAGVAGAALLLVHVRRRGHSQDAGSRLLAGTPEPSVHALPDALNNLRTQEGSGDGPSSS
VDWNRPEDVDPQGIYVISAPSIYAREA

FIG. 5A

Nucleotide Sequence of Heavy Chain Variable Domain (V_H) of 7-I1-B (SEQ ID NO: 1)

GAGGTGCAGCTGGTGGAGTCTGGGGGGGGCTTGGTAAAGCCTGGGGGGTCCCTT
AGACTCTCCTGTGCAGCCTCTGGATTCACTTTCAGTAACACCTGGATGAGCTGGG
TCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGGTTGGCCGTATTTAAAGCAAAT
CTGATGGTGGGACAACAGACTACGCTGCACCCGTGAAAGGCAGATTCACCATCT
CAAGAGATGATTCAAAAAACACGCTGTATCTGCAAATGAACAGCCTGAAAACCG
AGGACACAGCCGTGTATTACTGTACCCAGTATTATTGGAACCTCTTTGACTACTG
GGGCCAGGGAACCCTGGTCACCGTCTCCTCA

Amino Acid Sequence of Heavy Chain Variable Domain (V_H) of 7-I1-B (SEQ ID NO: 2)

EVQLVESGGGLVKPGGSLRLSCAAS**GF**TFSNTWMSWVRQAPGKGLEWVGRIK**SKS**
DGGTTDYAAPVKGRFTISRDDSKNTLYLQMNSLKTEDTAVYYCTIQYYWNSFDYWGQ
GTLVTVSS

FIG. 5B

Nucleotide Sequence of Light Chain Variable Domain (V_L) of 7-I1-B (SEQ ID NO: 6)

GACATCCAGATGACCCAGTCTCCATCCTCCCTGTCTGCATCTGTAGGAGACAGAG
TCACCATCACTTGCCAGGCGAGTCAGGACATTAGCAACTATTTAAATTGGTATCA
GCAGAAACCAGGGAAAGCCCCTAAGCTCCTGATCTACGATGCATCCAATTTGGA
AACAGGGGTCCCATCAAGGTTCAAGTGGAAAGTGGATCTGGGACAGATTTTACTTTC
ACCATCAGCAGCCTGCAGCCTGAAGATATTGCAACATATTACTGTCAACAGTATG
ATAATCTCCCGCTCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAA

Amino Acid Sequence of Light Chain Variable Domain (V_L) of 7-I1-B (SEQ ID NO: 7)

DIQMTQSPSSLSASVGDRVTITC**QASQ**DISNYLNWYQQKPGKAPKLLIYD**ASNLETG**
VPSRFSGSGSGTDFTFTISLQPEDATYYCQQYDNLPLTFGGGGTKVEIK

FIG. 6A

Nucleotide Sequence of Heavy Chain Variable Domain (V_H) of 2-C8-A (SEQ ID NO: 11).

GAGGTGCAGCTGGTGGAGTCTGGGGGAGGCTTGGTCCAGCCTGGGGGGTCCCAG
AGACTCTCCTGTGCAGCCTCTGGATTCACCTTTAGTAGCTATTGGATGAACTGGG
TCCGCCAGGCTCCAGGGAAGGGGCTGGAGTGGGTGGCCAACATAAAGGAAGAT
GGAAGTGAGAAATACTATGTGGACTCTGTGAAGGGCCGATTCACCATCTCCAGA
GACAACGCCAAGAACTCACTGTATCTGCAAATGAACAGCCTGAGAGCCGAGGAC
ACGGCTGTGTATTACTGTGCGAGAGATCCGGGCTGGGCTCCCTTTGACTACTGGG
GCCAGGGAACCCTGGTCACCGTCTCCTCA

Amino Acid Sequence of Heavy Chain Variable Domain (V_H) of 2-C8-A (SEQ ID NO: 12).

EVQLVESGGGLVQPGGSQRLSCAAS**GF****TFSSY**WMNWVRQAPGKGLEWVANIK**EDG**
SEKYYVDSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCARDDPGWAPFDYWGQ
GTLTVSS

FIG. 6B

Nucleotide Sequence of Light Chain Variable Domain (V_L) of 2-C8-A (SEQ ID NO: 16).

GACATCCAGATGTCCCAGTCTCCATCCTCACTGTCTGCATCTGTAGGAGACAGAG
TCACCATCACTTGTCGGGCGAGTCAGGGCATTAGCAATTATTTAGCCTGGTTTCA
GCAGAAACCAGGGAAAGCCCCTAAGTCCTGATCTATGCTGCATCCAGTTTGCA
AAGTGGGGTCCCATCAAAGTTCAGCGGCAGTGGATCTGGGACAGATTTCACTCTC
GCCATCAGCAGCCTGCAGCCTGAAGATTTTGCAACTTATTACTGCCAACAGTATA
ATAGTTTCCCGTACACTTTTGGCCAGGGGACCACGCTGGAGATCAAA

Amino Acid Sequence of Light Chain Variable Domain (V_L) of 2-C8-A (SEQ ID NO: 17).

DIQMSQSPSSLSASVGDRVTIT**C****RASQ****GISNY**LAWFQQKPGKAPKSLIYAASSLQSGV
PSKFSGSGSGTDFTLAISLQPEDFATYYCQQYNSPYTFGQGTTL**EIK**

FIG. 7A

Nucleotide Sequence of Heavy Chain Variable Domain (V_H) of 10-O18-A (SEQ ID NO: 21).

CAGGTGCAGCTGCAGGAGTCGGGCCCAGGACTGGTGAAGCCTTCGGAGACCCTG
TCCCTCACCTGCACTGTCTCTGGTGGCTCCATCAATAGTTACTACTGGAGCTGGA
TCCGGCAGCCCCAGGGAAGGGACTGGAGTGGATTGGGTATATCTTTTACAGTG
GGATCACCAACTACAACCCCTCCCTCAAGAGTCGAGTCACCATATCATTAGACAC
GTCCAAGAACCAGTTCTCCCTGAAGCTGAGCTCTGTGACCGCTGCGGACACGGCC
GTGTATTACTGTGCGAGAATCGGCGTGGCTGGTTTTTACTTTGACTACTGGGGCC
AGGGAACCCTGGTCACCGTCTCCTCA

Amino Acid Sequence of Heavy Chain Variable Domain (V_H) of 10-O18-A (SEQ ID NO: 22).

QVQLQESGPGLVKPSETLSLTCTVSGGSINSYYWSWIRQPPGKGLEWIGYIFYSGITN
YNPSLKSRTISLDTSKNQFSLKLSSVTAADTAVYYCARIGVAGFYFDYWGQGLVT
VSS

FIG. 7B

Nucleotide Sequence of Light Chain Variable Domain (V_L) of 10-O18-A (SEQ ID NO: 26).

GAAATTGTGTTGACGCAGTCTCCAGGCACCCTGTCTTTGTCTCCAGGGGAAAGAG
CCACCCTCTCCTGCAGGGCCAGTCAGAGTGTTAGCAGCAGCTACTTAGCCTGGTA
CCAGCAGAAACCTGGCCAGGCTCCCAGGCTCCTCATCTATGGTGCATCCAGCAG
GGCCACTGGCATCCCAGACAGGTTCAAGTGGCAGTGGGTCTGGGACAGACTTCAC
TCTACCATCAGCAGACTGGAGCCTGAAGATTTTGCAGTGTATTACTGTCAGCAG
TATGGTACCTCACCGCTCACTTTTCGGCGGAGGGACCAAGGTGGAGATCAAA

Amino Acid Sequence of Light Chain Variable Domain (V_L) of 10-O18-A (SEQ ID NO: 27).

EIVLTQSPGTLSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLLIYGASSRATG
IPDRFSGSGSGTDFTLTISRLEPEDFAVYYCQOYGTSPITFGGGTKVEIK

FIG. 8A

Nucleotide Sequence of Heavy Chain Variable Domain (V_H) of 6-G23-F (SEQ ID NO: 31).

CAGGTGCAGCTGGTGCAGTCTGGGGCTGAGGTGAAGAAGCCTGGGGCCTCAGTG
AAGGTTTCCTGCAAGGCATCTGGATACACCTTCACCAGCTACTATATACTGGG
TGCACAGGCCCTGGACAAGGGCTTGAGTGGATGGGAATAATCGACCCAAGTG
ATGGTAGCACAACTACGCACAGAAGTTCCAGGGCAGAGTCACCATGACCAGGG
ACACGTCCACGAGCACAGTCTACATGGAGCTGAGCAGCCTGAGATCTGAGGACA
CGGCCGTGTATTACTGTGCGAGAGATCGGGAATATACTACTACGGTTTGGACGT
CTGGGGCCAAGGGACCACGGTCACCGTCTCCTCA

Amino Acid Sequence of Heavy Chain Variable Domain (V_H) of 6-G23-F (SEQ ID NO: 32).

QVQLVQSGAEVKKPGASVKVSCKASGYTFTSYIHWRQAPGQGLEWMGIIDPSD
GSTNYAQKFQGRVTMTRDTSTSTVYMESSLRSEDTAVYYCARDREYNYYGLD/VWG
QGTTTVTVSS

FIG. 8B

Nucleotide Sequence of Light Chain Variable Domain (V_L) of 6-G23-F (SEQ ID NO: 36).

GATGTTGTGATGACTCAGTCTCCACTCTCCCTGCCCCGTCACCCTTGGACAGCCGG
CCTCCATCTCCTGCAGGTCTAGTCAAAGCCTCGTATACCGTGATGGAAACACCTA
CTTGAATTGGTTTCAGCAGAGGCCAGGCCAATCTCCAAGGCGCCTAATTTATAAG
GTTTCTAACCGGGACTCTGGGGTCCCAGACAGATTCCGCGGCAGTGGGTCAGGC
ACTGATTTCACTGAAAATCAGCCGGGTGGAGGCTGAGGATGTTGGGGTTTATT
ACTGCATGCAAGGTACACACTGGCCTCCGACGTTCCGCCAAGGGACCAAGGTGG
AAATCAAA

Amino Acid Sequence of Light Chain Variable Domain (V_L) of 6-G23-F (SEQ ID NO: 37).

DVVMTQSPSLPVTLGQPASISCRSSQSLVYRDGNTYLNWFQQRPGQSPRRLIYKVS
NRDSGVDPDRFRGSGSGTDFTLKISRVEADVGVYYCMOGTHWPPTFGQGTKVEIK

FIG. 9

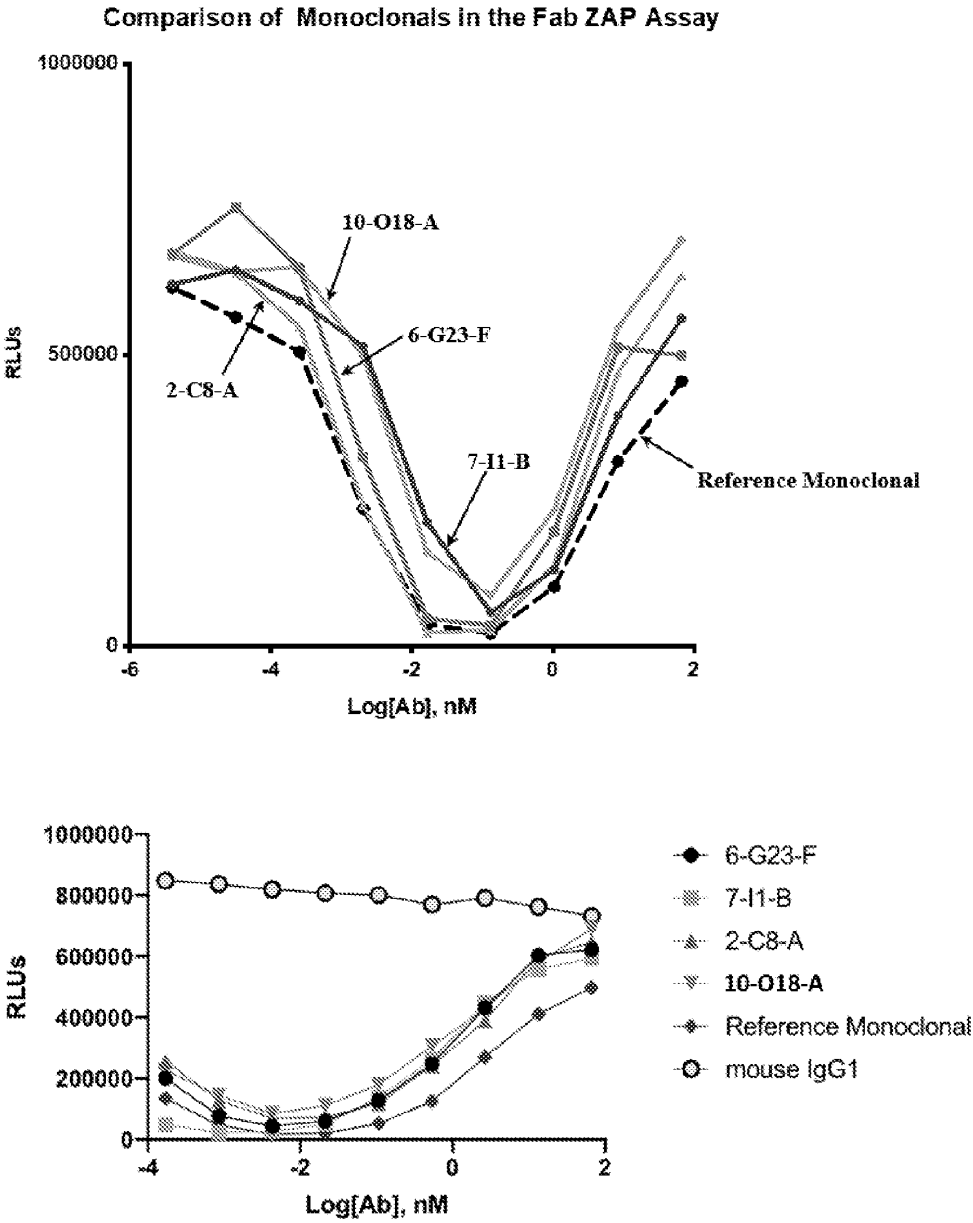


FIG. 10A

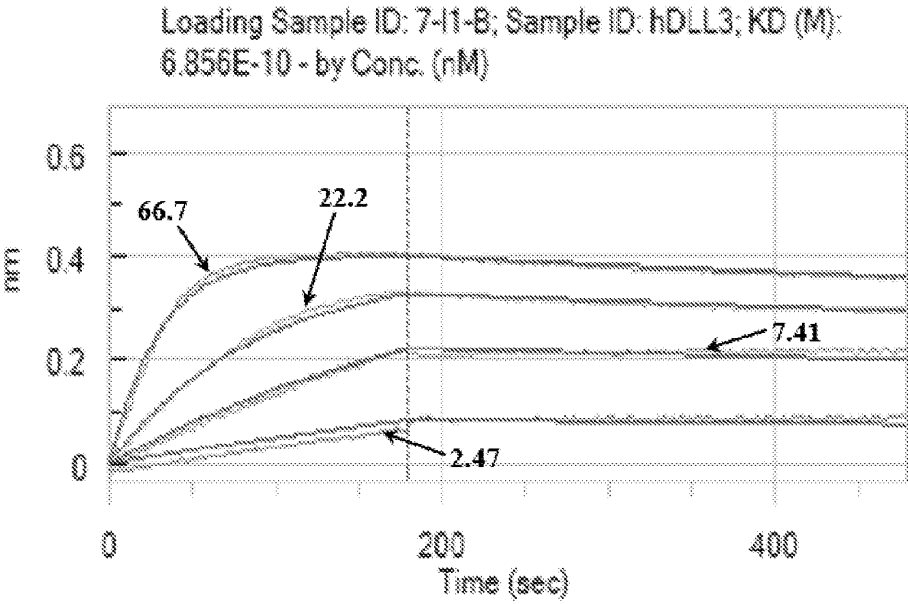


FIG. 10B

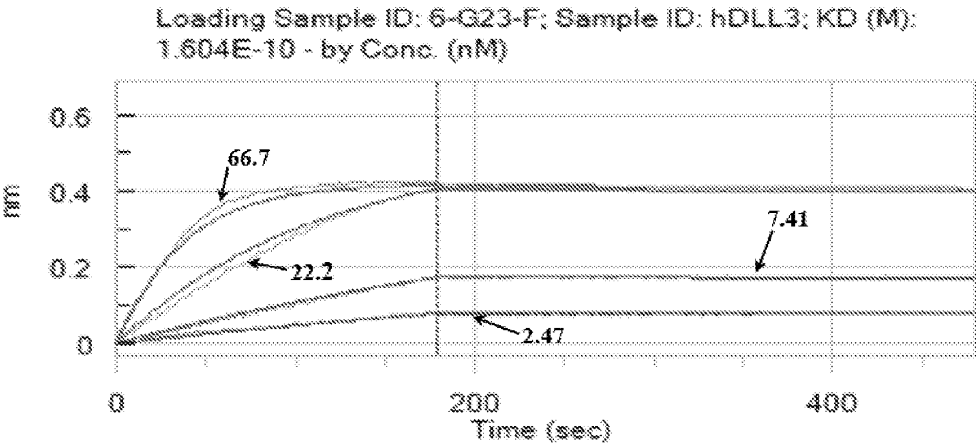


FIG. 10C

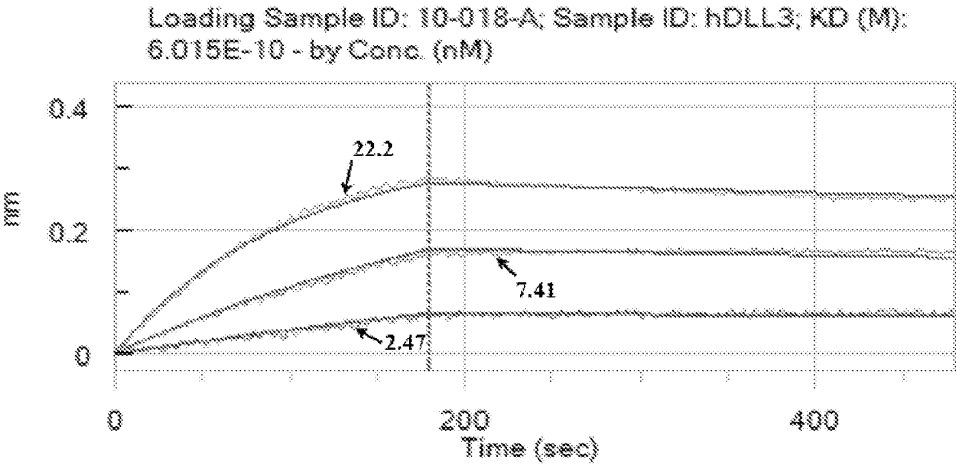


FIG. 10D

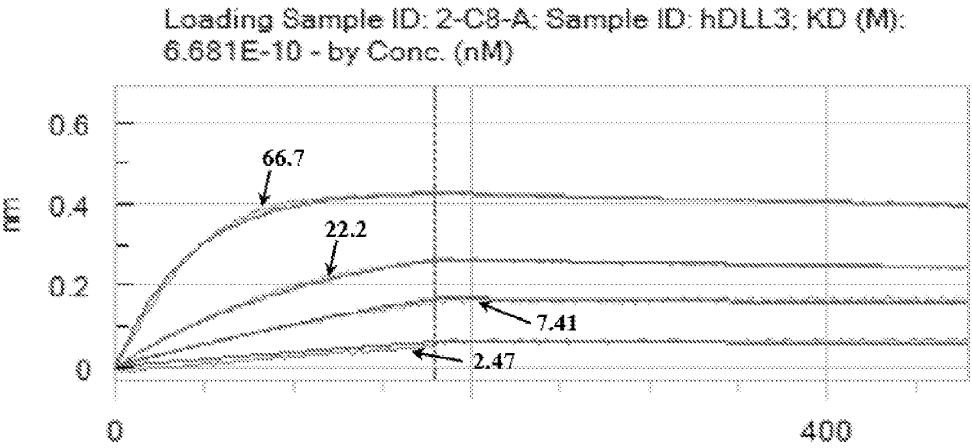


FIG. 10E

	Kd nM	Kon (1/Ms)	Kdis(1/s)	FullX^2	FullR^2
7-I1-B	0.68	5.02E+05	3.44E-04	0.168	0.99
10-O18-A	0.62	4.49E+05	2.67E-04	0.038	0.97
6-G23-F	0.16	4.05E+05	6.49E-05	0.15	0.97
2-C8-A	0.668	3.63E+05	2.42E-04	0.073	0.99

FIG. 11

Monoclonal	DLL3	DLL4	DLL1
10-O18-A	1.361	0.056	0.126
2-C8-A	1.359	0.07	0.098
6-G23-F	1.459	0.05	0.206
7-I1-B	1.302	0.96	0.127
Background (anti-mouse IgG HRP0)	0.085	0.085	0.095

FIG. 12

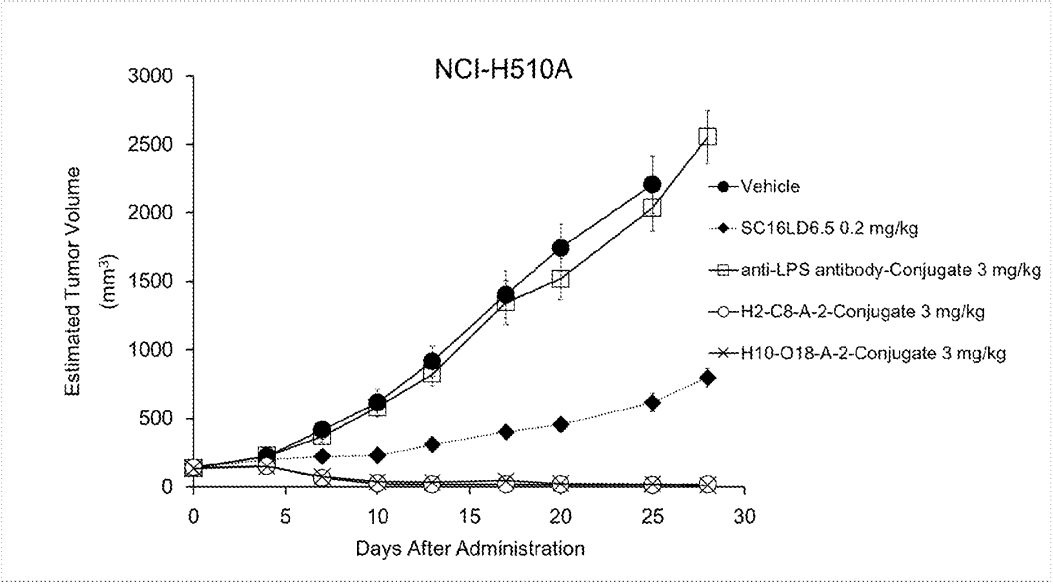


FIG. 13

