

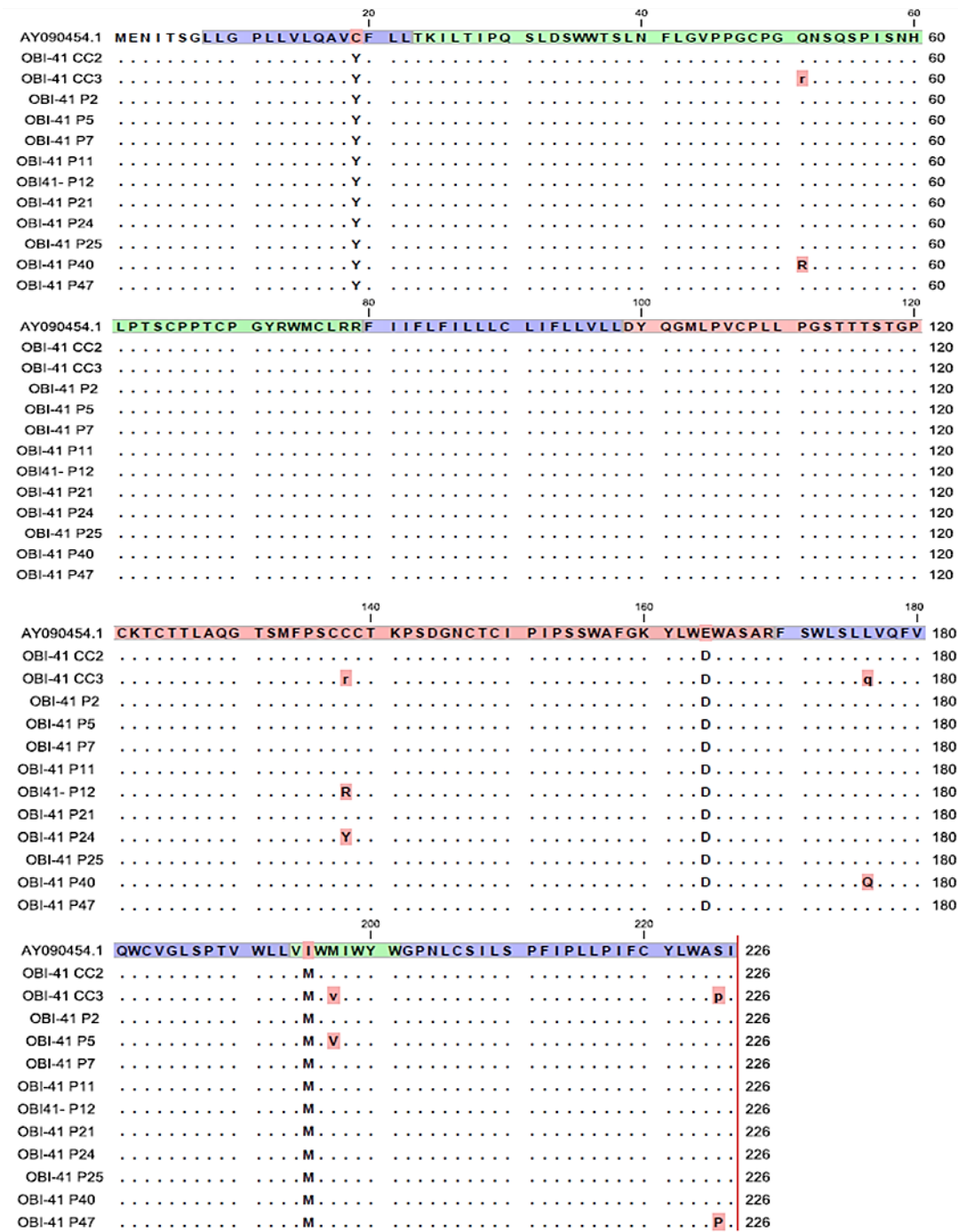
Supplementary Figure 1 legend

**Supplementary Figure 1. Deduced amino acid sequence analysis of the HBsAg.** Alignments of the deduced amino acid sequences obtained from ten S gen sequences of OBI patients. The alignment was performed using one reference HBV sequence genotype H (AY0954.1) with the CLCbio software. The colors shadow in the first line indicate the HBsAg domain. Residues identical to the genotype H reference consensus are indicated by dots. Mutations in red boxes.

Alignment of amino acid sequences deduced from OBI-39 sample clones

		20		40		60	
AY090454.1	MENITSGLLG	PLLVLQAV	CF	LLTKILTIPQ	SLDSWWTSLN	FLGVPPGCPG	QNSQSPI SNH 60
OBI-39 CC2	.....	.....	Y.	.....	.....	.....	60
OBI-39 P1 aa	.....	.....	Y.	.....	.....	.....	60
OBI-39 P3 aa	.....	.....	Y.	.....	.....	.....	60
OBI-39 P5 aa	.....	.....	Y.	.....	.....	.....	60
OBI-39 P7 aa	.....	.....	Y.	.....	.....	.....	60
OBI-39 P8 aa	.....	.....	Y.	.....	.....	.....	60
OBI-39 P9 aa	.....	.....	Y.	.....	P.	.....	60
OBI-39 P10 aa	.....	.....	Y.	.....	.....	.....	60
OBI-39 P12 aa	.....	.....	Y.	.....	.....	.....	60
OBI-39 P13 aa	.....	.....	Y.	.....	.....	.....	60
OBI-39 P14 aa	.....	.....	Y.	.....	.....	.....	60
		80		100		120	
AY090454.1	LPTSCPPTCP	GYRWMCLRRF	ITFLFILLLC	LIFLLVLLDY	QGMLPVCPLL	PGSTTTSTGP	120
OBI-39 CC2	.....	.....	.....	.....	.....	.....	120
OBI-39 P1 aa	.....	R.	.....	.....	H.	.....	120
OBI-39 P3 aa	.....	.....	.....	.....	.....	.....	120
OBI-39 P5 aa	.....	.....	.....	.....	.....	.....	120
OBI-39 P7 aa	.....	.....	.....	.....	.....	.....	120
OBI-39 P8 aa	.....	.....	.....	.....	.....	.....	120
OBI-39 P9 aa	.....	.....	R.	.....	.....	.....	120
OBI-39 P10 aa	.....	.....	.....	.....	.....	.....	120
OBI-39 P12 aa	.....	.....	.....	.....	.....	.....	120
OBI-39 P13 aa	.....	.....	.....	.....	.....	A.	120
OBI-39 P14 aa	.....	.....	.....	.....	.....	.....	120
		140		160		180	
AY090454.1	CKTCTTLAQG	TSMFPSCCCT	KPSDGNCTCI	PIPSWAFGK	YLWEWASARF	SWLSLLVQEV	180
OBI-39 CC2	.....	.....	.....	.....	.....	.....	180
OBI-39 P1 aa	.....	.....	.....	.....	.....	.....	180
OBI-39 P3 aa	.....	.....	.....	.....	.....	.....	180
OBI-39 P5 aa	.....	.....	.....	.....	.....	.....	180
OBI-39 P7 aa	.....	.....	P.	.....	.....	.....	180
OBI-39 P8 aa	.....	.....	.....	.....	.....	.....	180
OBI-39 P9 aa	.....	.....	.....	.....	.....	.....	180
OBI-39 P10 aa	.....	.....	.....	.....	.....	.....	180
OBI-39 P12 aa	.....	.....	.....	.....	.....	.....	180
OBI-39 P13 aa	.....	.....	I.	.....	.....	.....	180
OBI-39 P14 aa	.....	.....	.....	.....	.....	.....	180
		200		220			
AY090454.1	QWCVGLSPTV	WLLVIWMIWY	WGPNLCSILS	PFIPLLPIFC	YLWASI		226
OBI-39 CC2	.....	.....	.....	.....	.....		226
OBI-39 P1 aa	.....	.....	.....	I.	.....		226
OBI-39 P3 aa	.....	.....	.....	I.	.....		226
OBI-39 P5 aa	.....	.....	.....	.....	.....		226
OBI-39 P7 aa	.....	.....	.....	.....	.....		226
OBI-39 P8 aa	.....	.....	.....	I.	.....		226
OBI-39 P9 aa	.....	.....	.....	.....	.....		226
OBI-39 P10 aa	.....	S.	.....	.....	.....		226
OBI-39 P12 aa	.....	.....	.....	.....	.....		226
OBI-39 P13 aa	.....	.....	.....	.....	.....		226
OBI-39 P14 aa	.....	.....	.....	.....	.....		226

Alignment of amino acid sequences deduced from OBI-41 sample clones



Supplementary Figure 1. Alignment of amino acid sequences deduced from sample clones OBI-43. Residues identical to the genotype H reference consensus are indicated by dots. Mutations in red boxes.

Alignment of amino acid sequences deduced from OBI-43 sample clones

		20		40		60	
AY090454.1	MENITSGLLG	PLLVLQAVCF	LLTKILTIPQ	SLDSWWTSLN	FLGVPPGCPG	QNSQSPISNH	60
OBI-43	.....	Y.	.....	.....	.....	.....	60
OBI-43 P1	.....	Y.	.....	.....	.....	.....	60
OBI-43 P5	.....	Y.	.....	.....	.....	.....	60
OBI-43 P17	.....	Y.	.....	.....	.....	.....	60
OBI-43 P19	.....	Y.	.....	.....	.....	.....	60
OBI-43 P34	.....	Y.	.....	.....	.....	.....	60
OBI-43 P36	.....	Y.	.....	.....	.....	D.	60
OBI-43 P37	.....	Y.	.....	.....	.....	.....	60
OBI-43 P40	.....	Y.	.....	.....	.....	.....	60
OBI-43 P42	.....	Y.	.....	.....	.....	.....	60
OBI-43 P43	.....	Y.	.....	.....	.....	.....	60
		80		100		120	
AY090454.1	LPTSCPPTCP	GYRWMCLRRF	IIFLFILLLC	LIFLLVLLDY	QGMLPVCPLL	PGSTTTSTGP	120
OBI-43	.....	.....	.....	.....	.....	.....	120
OBI-43 P1	.....	.....	.....	.....	.....	.....	120
OBI-43 P5	.....	.....	V.	.....	.....	.....	120
OBI-43 P17	.....	.....	.....	.....	Y.	.....	120
OBI-43 P19	.....	.....	.....	.....	Y.	.....	120
OBI-43 P34	.....	.....	.....	.....	.....	.....	120
OBI-43 P36	.....	.....	.....	.....	.....	.....	120
OBI-43 P37	C.	.....	.....	.....	Y.	.....	120
OBI-43 P40	.....	.....	.....	.....	.....	.....	120
OBI-43 P42	.....	.....	.....	.....	.....	.....	120
OBI-43 P43	.....	.....	.....	.....	.....	.....	120
		140		160		180	
AY090454.1	CKTCTTLAQG	TSMFPSCCCT	KPSDGNCTCI	PIPSSWAFGK	YLWEWASARF	SWLSLLVQFV	180
OBI-43	.....	.....	.....	.....	.....	.....	180
OBI-43 P1	.....	.....	.....	.....	.....	.....	180
OBI-43 P5	.....	.....	.....	.....	.....	.....	180
OBI-43 P17	.....	.....	.....	.....	.....	.....	180
OBI-43 P19	.....	.....	.....	.....	.....	.....	180
OBI-43 P34	.....	.....	.....	.....	C.	.....	180
OBI-43 P36	.....	.....	.....	.....	.....	.....	180
OBI-43 P37	.....	.....	.....	.....	.....	.....	180
OBI-43 P40	.....	.....	.....	.....	.....	.....	180
OBI-43 P42	.....	.....	.....	.....	.....	.....	180
OBI-43 P43	.....	.....	.....	.....	.....	.....	180
		200		220			
AY090454.1	QWCVGLSPTV	WLLVIWMIWY	WGNLCSILS	PFIPLLPIFC	YLWAS	I	226
OBI-43	.....	M.	.....	.....	.....	.....	226
OBI-43 P1	T.	.....	.....	.....	.....	.....	226
OBI-43 P5	.....	.....	.....	.....	.....	.....	226
OBI-43 P17	.....	.....	.....	.....	.....	.....	226
OBI-43 P19	T.	.....	.....	.....	.....	.....	226
OBI-43 P34	.....	.....	.....	.....	.....	.....	226
OBI-43 P36	.....	.....	.....	.....	.....	.....	226
OBI-43 P37	.....	.....	.....	.....	.....	.....	226
OBI-43 P40	.....	.....	.....	.....	.....	.....	226
OBI-43 P42	.....	.....	.....	.....	.....	.....	226
OBI-43 P43	.....	.....	.....	.....	.....	.....	226

Supplementary Figure 1. Alignment of amino acid sequences deduced from sample clones OBI-43. Residues identical to the genotype H reference consensus are indicated by dots. Mutations in red boxes.

Alignment of amino acid sequences deduced from OBI-44 sample clones

		20		40		60	
AY090454.1	MENITSGLLG	PLLVLQAVCF	LLTKILTIPQ	SLDSWWTSLN	FLGVPPGCPG	QNSQSPISNH	60
OBI-44	.....	.....Y.	.....	.....	.....	.....	60
OBI-44 P8	.....	..... <span style="border: 1px solid red;">A</span> .	.....	.....	.....	.....	60
OBI-44 P11	.....	.....Y.	.....	.....	.....	.....	60
OBI-44 P12	.....	.....Y.	.....	.....	.....	.....	60
OBI-44 P14	.....	.....Y.	.....	.....	.....	.....	60
OBI-44 P17	.....	.....Y.	.....	.....	.....	.....	60
OBI-44 P19	.....	.....Y.	.....	.....	.....	.....	60
OBI-44 P20	.....	..... <span style="border: 1px solid red;">A</span> .	.....	.....	.....	.....	60
OBI-44 P22	.....	..... <span style="border: 1px solid red;">A</span> .	.....	.....	.....	.....	60
OBI-44 P26	.....	.....Y.	.....	.....	.....	.....	60
OBI-44 P30	.....	.....Y.	.....	.....	.....	.....	60
		80		100		120	
AY090454.1	LPTSCPPTCP	GYRWMCLRRF	IIFLFILLIC	LIFLLVLLDY	QGMLPVCPLL	PGSTTTSTGP	120
OBI-44	.....	.....	.....	.....	.....	.....	120
OBI-44 P8	.....	.....	.....	.....	.....	.....	120
OBI-44 P11	.....	.....	.....	.....	.....	.....	120
OBI-44 P12	.....	.....	.....	.....	.....	.....	120
OBI-44 P14	.....	.....	.....	.....	.....	.....	120
OBI-44 P17	.....	.....	.....	.....	.....	.....	120
OBI-44 P19	.....	.....	.....	.....	.....	.....	120
OBI-44 P20	.....	.....	.....	.....	.....	.....	120
OBI-44 P22	.....	.....	.....	.....	.....	.....	120
OBI-44 P26	.....	.....	.....	.....	.....	.....	120
OBI-44 P30	.....	.....	.....	.....	.....	.....	120
		140		160		180	
AY090454.1	CKTCTTLAQG	TSMFPSCCCT	KPSDGNCTCI	PIPSWAFGK	YLWEWASARE	SWLSLLVQEV	180
OBI-44	.....	.....	.....	.....	.....	.....	180
OBI-44 P8	.....	.....	.....	..... <span style="border: 1px solid red;">S</span> .....	.....	.....	180
OBI-44 P11	.....	.....	.....	.....	.....	.....	180
OBI-44 P12	.....	.....	.....	.....	.....	.....	180
OBI-44 P14	.....	.....	.....	.....	.....	.....	180
OBI-44 P17	.....	.....	.....	.....	.....	.....	180
OBI-44 P19	.....	.....	.....	.....	.....	.....	180
OBI-44 P20	.....	.....	.....	..... <span style="border: 1px solid red;">S</span> .....	.....	.....	180
OBI-44 P22	.....	..... <span style="border: 1px solid red;">R</span> .....	.....	.....	.....	.....	180
OBI-44 P26	.....	.....	.....	.....	.....	.....	180
OBI-44 P30	.....	.....	.....	.....	.....	.....	180
		200		220			
AY090454.1	QWCVGLSPTV	WLLVIWMIWY	WGPNLCSILS	PFIPLLPIFC	YLWASI		226
OBI-44	.....	.....	.....	.....	.....		226
OBI-44 P8	.....	.....	.....	.....	.....		226
OBI-44 P11	.....	.....	.....	.....	.....		226
OBI-44 P12	.....	.....	.....	.....	.....		226
OBI-44 P14	.....	.....	.....	.....	.....		226
OBI-44 P17	.....	.....	.....	.....	.....		226
OBI-44 P19	.....	.....	.....	.....	.....		226
OBI-44 P20	.....	.....	.....	.....	.....		226
OBI-44 P22	.....	.....	.....	.....	.....		226
OBI-44 P26	.....	.....	.....	.....	.....		226
OBI-44 P30	.....	.....	.....	.....	.....		226

Supplementary Figure 1. Alignment of amino acid sequences deduced from sample clones OBI-44. Residues identical to the genotype H reference consensus are indicated by dots. Mutations in red boxes.

Alignment of amino acid sequences deduced from OBI-50 sample clones

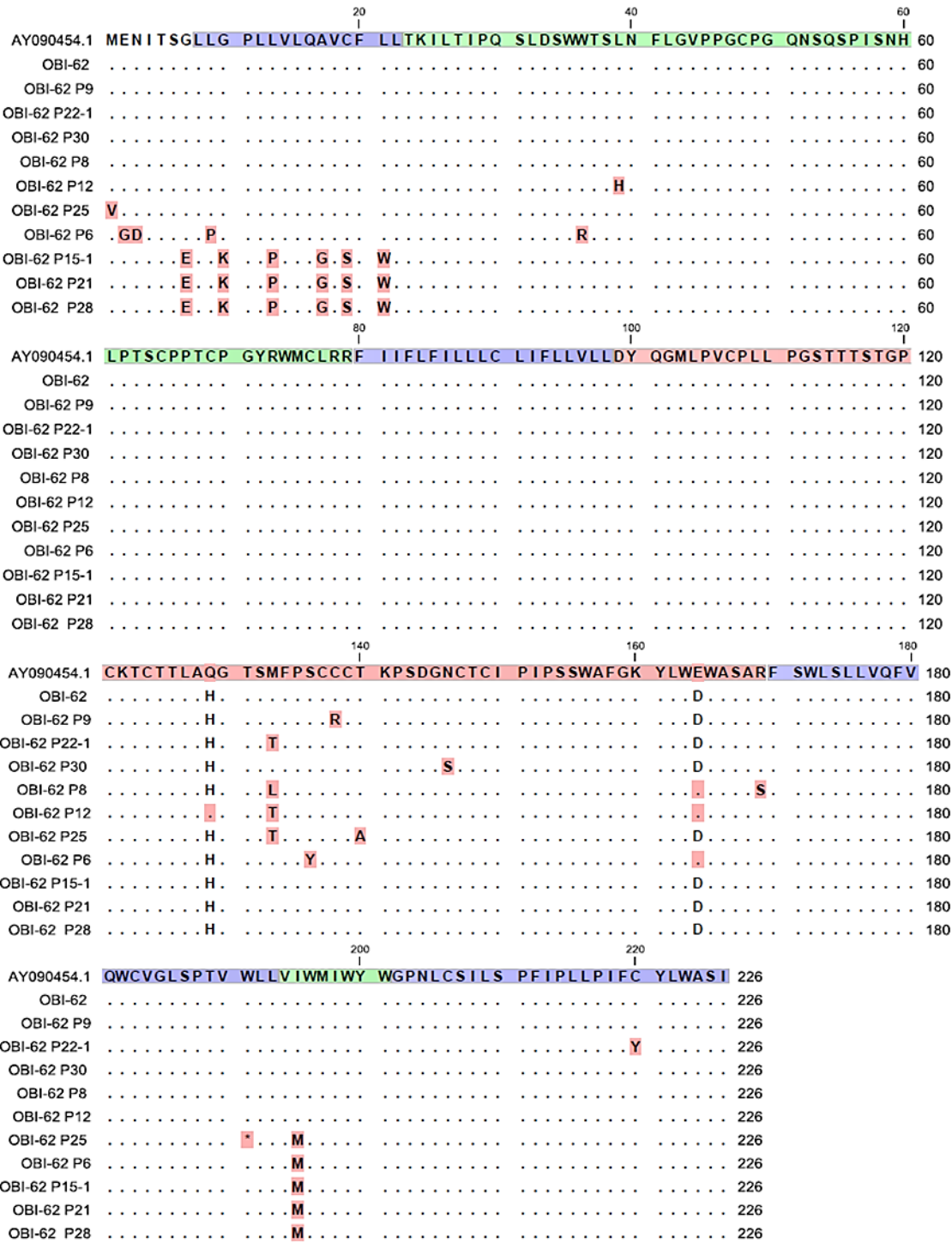
AY090454.1	MENITSGLLG	PLLVLQAVCF	LLTKILTIPQ	SLDSWWTSLN	FLGVPPGCPG	QNSQSPI	SNH	60
OBI-50	.....	.....	.....	.....	.....	.....	.....	60
OBI-50 P1	.....	A	.....	H	E	.....	.....	60
OBI-50 P3	.....	.....	.....	.....	.....	.....	.....	60
OBI-50 P6	.....	.....	RK	H	E	.....	.....	60
OBI-50 P13	.....	.....	.....	.....	.....	.....	.....	60
OBI-50 P15-1	.....	.....	R	K	.....	.....	.....	60
OBI-50 P17	P	.....	RK	.....	.....	.....	.....	60
OBI-50 P22	.....	.....	.....	.....	.....	.....	.....	60
OBI-50 P24	.....	.....	.....	.....	.....	.....	.....	60
OBI-50 P28-1	P	.....	.....	.....	.....	.....	.....	60
OBI-50 P32	.....	I	.....	.....	.....	.....	.....	60
AY090454.1	LPTSCPPTCP	GYRWMCLRRF	IIFLFILLLC	LIFLLVLLDY	QGMLPVCPLL	PGSTTTSTGP		120
OBI-50	.....	.....	.....	.....	.....	.....	.....	120
OBI-50 P1	.....	.....	.....	.....	.....	.....	.....	120
OBI-50 P3	.....	.....	.....	.....	.....	.....	.....	120
OBI-50 P6	.....	.....	.....	.....	.....	.....	.....	120
OBI-50 P13	.....	.....	.....	.....	.....	.....	.....	120
OBI-50 P15-1	.....	.....	.....	.....	.....	.....	.....	120
OBI-50 P17	.....	.....	.....	.....	.....	.....	.....	120
OBI-50 P22	.....	.....	.....	.....	.....	.....	.....	120
OBI-50 P24	.....	.....	.....	.....	.....	.....	.....	120
OBI-50 P28-1	.....	.....	.....	.....	.....	.....	*	120
OBI-50 P32	.....	.....	.....	.....	.....	.....	.....	120
AY090454.1	CKTCTTLAQG	TSMFPSCCCT	KPSDGNCTCI	PIPSSWAFGK	YLWEWASARF	SWLSLLVQFV		180
OBI-50	.....	H	.....	.....	*	.....	.....	180
OBI-50 P1	AF	*	.....	.....	*	.....	.....	180
OBI-50 P3	.....	H	.....	.....	D	.....	.....	180
OBI-50 P6	.....	H	.....	.....	D	.....	.....	180
OBI-50 P13	.....	H	L	G	D	.....	.....	180
OBI-50 P15-1	.....	H	.....	.....	D	S	P	180
OBI-50 P17	.....	*	.....	.....	*	.....	.....	180
OBI-50 P22	.....	H	L	G	D	.....	.....	180
OBI-50 P24	.....	H	.....	.....	D	.....	.....	180
OBI-50 P28-1	R	*	.....	.....	*	*	.....	180
OBI-50 P32	.....	*	.....	.....	*	.....	.....	180
AY090454.1	QWCVGLSPTV	WLLVIWMIWY	WGNLCSILS	PEIPLLPIFC	YLWASI			226
OBI-50	.....	*	.....	.....	.....	.....	.....	226
OBI-50 P1	.....	*	.....	.....	.....	.....	.....	226
OBI-50 P3	.....	M	V	.....	Y	.....	.....	226
OBI-50 P6	.....	R	M	.....	.....	.....	.....	226
OBI-50 P13	.....	M	.....	.....	.....	.....	.....	226
OBI-50 P15-1	.....	*	.....	.....	.....	.....	.....	226
OBI-50 P17	.....	*	.....	.....	.....	.....	.....	226
OBI-50 P22	.....	M	.....	.....	.....	.....	.....	226
OBI-50 P24	.....	M	V	.....	.....	.....	.....	226
OBI-50 P28-1	.....	M	.....	.....	.....	.....	.....	226
OBI-50 P32	.....	*	.....	.....	.....	.....	.....	226

Supplementary Figure 1. Alignment of amino acid sequences deduced from sample clones OBI-50. Residues identical to the genotype H reference consensus are indicated by dots. Mutations in red boxes.



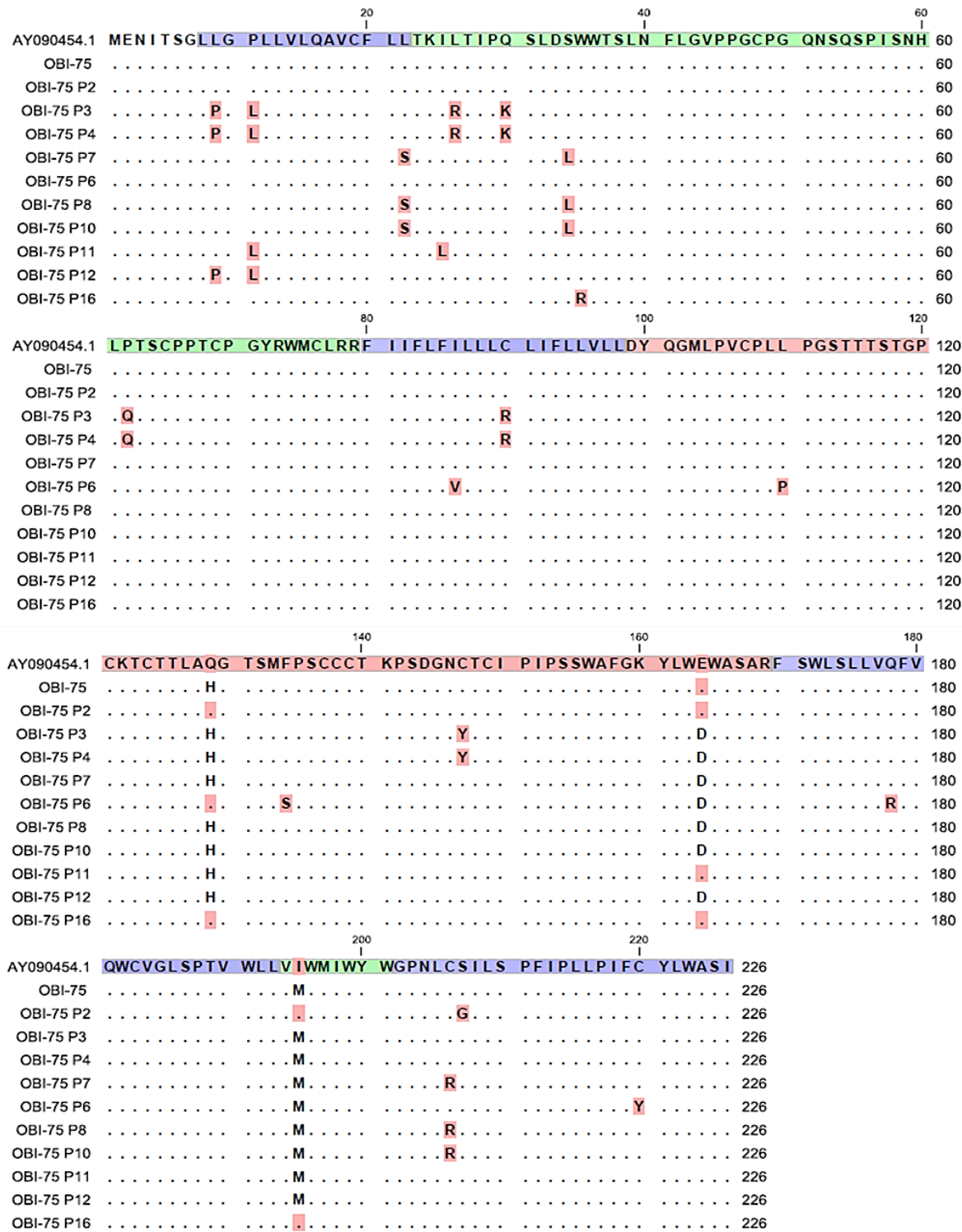


Alignment of amino acid sequences deduced from OBI-62 sample clones



Supplementary Figure 1. Alignment of amino acid sequences deduced from sample clones OBI-62. Residues identical to the genotype H reference consensus are indicated by dots. Mutations in red boxes.

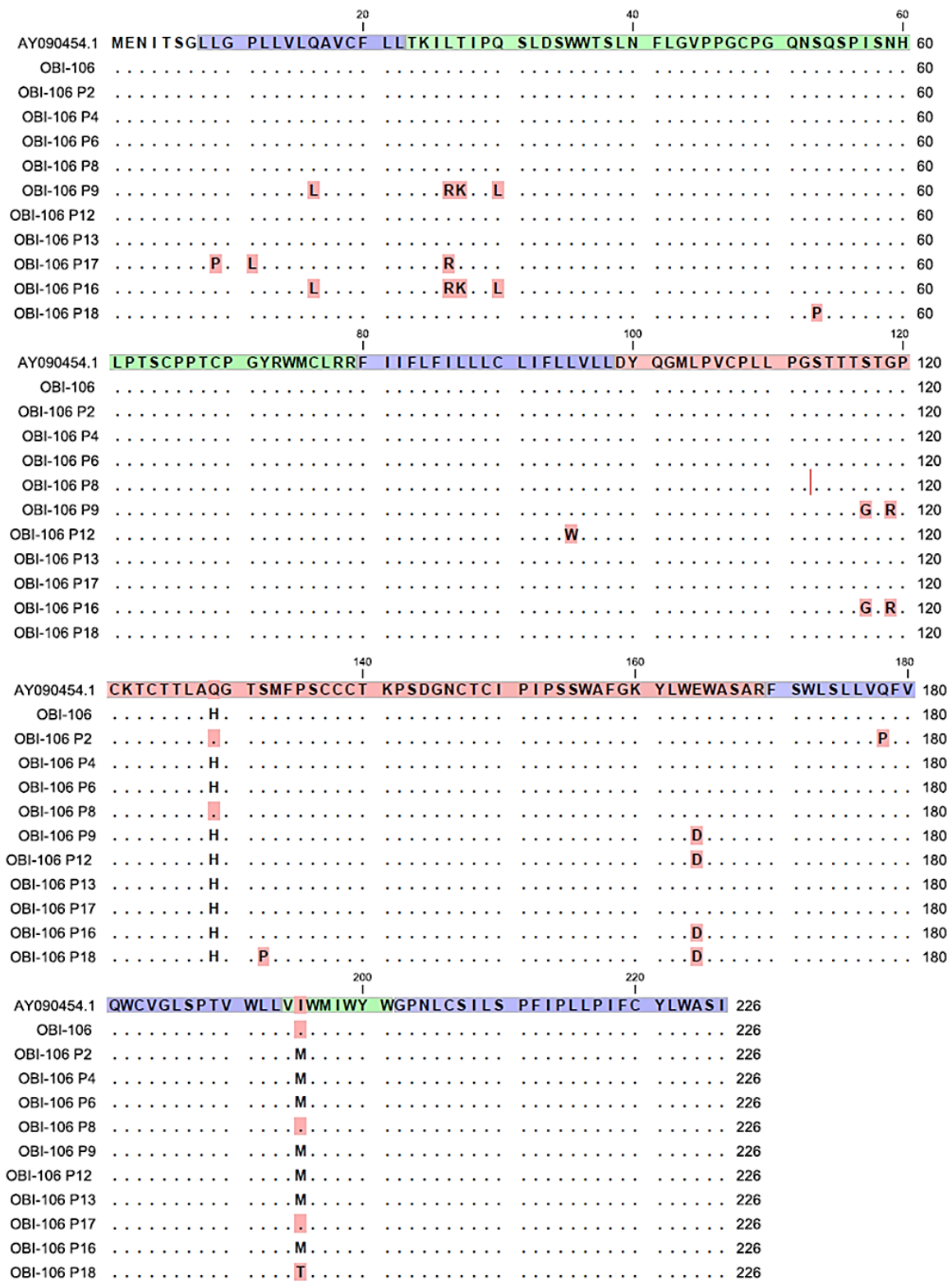
Alignment of amino acid sequences deduced from OBI-75 sample clones



Supplementary Figure 1. Alignment of amino acid sequences deduced from sample clones OBI-75. Residues identical to the genotype H reference consensus are indicated by dots. Mutations in red boxes.



Alignment of amino acid sequences deduced from OBI-106 sample clones



**Supplementary Figure 1.** Alignment of amino acid sequences deduced from sample clones OBI-106. Residues identical to the genotype H reference consensus are indicated by dots. Mutations in red boxes.

Alignment of amino acid sequences deduced from OBI-112 sample clones

		20		40		60	
AY090454.1	MENITSGLLG	PLLVLQAVCF	LLTKILTIPQ	SLDSWWTSLN	FLGVPPGCPG	QNSQSPI SNH	60
OBI-112	.....	.....Y.	.....	.....	.....	.....	60
OBI-112 P3	.....	.....Y.	.....	.....	.....	.....	60
OBI-112 P10	.....	.....Y.	.....	.....	.....	.....	60
OBI-112 P11	.....	.....Y.	.....	.....	.....	.....	60
OBI-112 P12	.....	.....Y.	.....	.....	.....	.....	60
OBI-112 P13	.....	.....Y.	.....	.....	.....	.....	60
OBI-112 P4	.....	.....Y.	.....	.....	.....	.....	60
OBI-112 P9	.....	......	.....	.....	.....	.....	60
OBI-112 P15	.....	.....Y.	.....	.....	.....	.....	60
		80		100		120	
AY090454.1	LPTSCPPTCP	GYRWMCLRRF	IIFLFILLLC	LIFLLVLLDY	QGMLPVCPLL	PGSTTTSTGP	120
OBI-112	.....	.....	.....	.....	.....	.....	120
OBI-112 P3	.....	.....	.....	.....	.....	.....	120
OBI-112 P10	.....	.....	.....	.....	.....	.....	120
OBI-112 P11	.....	.....	.....	.....	.....	.....	120
OBI-112 P12	.....	.....	.....	.....	.....	.....	120
OBI-112 P13	.....	.....	.....	.....	.....	.....	120
OBI-112 P4	.....	.....	.....	.....	.....	.....	120
OBI-112 P9	.....	.....	.....	.....	.....	.....	120
OBI-112 P15	.....	.....R.	.....	.....	.....	.....	120
		140		160		180	
AY090454.1	CKTCTTLAQG	TSMFPSCCCT	KPSDGNCTCI	PIPSSWAFGK	YLWEWASARF	SWLSLLVQFV	180
OBI-112	.....	.....	.....	.....	.....	.....	180
OBI-112 P3	.....	.....	.....	.....	.....	.....	180
OBI-112 P10	.....	.....	.....	.....	.....	.....	180
OBI-112 P11	.....	.....	.....	.....	.....	.....	180
OBI-112 P12	.....	.....	.....	.....	.....	.....	180
OBI-112 P13	.....	.....	.....	.....	.....	.....	180
OBI-112 P4	.....	.....	.....	.....	.....	.....	180
OBI-112 P9	.....	.....	.....	.....	.....	.....	180
OBI-112 P15	.....	.....	.....	.....	.....	.....	180
		200		220			
AY090454.1	QWCVGLSPTV	WLLVITWMIWY	WGPNLCSILS	PFIPLLPIFC	YLWASI		226
OBI-112	.....	......	.....	.....	.....	.....	226
OBI-112 P3	.....	.....M.	.....	.....	.....	.....	226
OBI-112 P10	.....	.....M.	.....	.....	.....	.....	226
OBI-112 P11	.....	.....M.	.....	.....	.....	.....	226
OBI-112 P12	.....	.....M.	.....	.....	.....	.....	226
OBI-112 P13	.....	.....M.	.....	.....	.....	.....	226
OBI-112 P4	.....	.....M.	.....	.....	.....	.....	226
OBI-112 P9	.....A.	.....M.	.....	.....	.....	.....	226
OBI-112 P15	.....	.....M.	.....	.....	.....	.....	226

Supplementary Figure 1. Alignment of amino acid sequences deduced from sample clones OBI-112. Residues identical to the genotype H reference consensus are indicated by dots. Mutations in red boxes.