

Table S3. IgM V_H amino-acid sequence of generated HBsAg-specific B cell clones.

CLONES	-----FR1-----	-----CDR1-----	-----FR2-----	-----CDR2-----	-----FR3-----	-----CDR3-----	-----FR4-----
VH3-73 6 β 2-D4-3	EVQLVESGGGLVQPGGSLKLSCAAS GGGLVQPGGSLKLSCAAS	GFTFSGSA GFTFSGSA	MHWVRQASGKGLEWGR MHWVRQASGKGLEWGR	IRSKANSYAT IRSKANSYAT	AYAASVK GRFTISRDDSNTAYLQMNSLKTEDTAVYYC AYAASVK GRFTISRDDSNTAYLQMNSLKTEDTAVYYC	TR TSRKSSSSDY	WGQGTLTVSS WGQGTLTVSS
6 β 1-E3-10	EVQLVESGGGLVQPGGSLKLSCAAS	GFTFSGSA	MHWVRQASGKGLEWGR	IRSKANSYAT	AYAA W K GRFTISRDDSNTAYLQMNSLKTEDTAVYYC	TR TSRKSSSSDY	WGQGTLTVSS
3 β 2-F2-2	EVQLVESGGGLVQPGGSLKLSCAAS	GFTFSGSA	MHWVRQASGKGLEWGR	IRSK P NYSYAT	AYAASVK GRFTISRDDSNTAYLQMNSLKTEDTAVYYC	TR TSLSGRGVDY	WGQGTLTVSS
6 β 2-G11-6	EVQLVESGGGLVQPGGSLKLSCAAS	GFTFSGSA	MHWVRQASGKGLEWGR	IRSKANS Y AT	AYAASVK GRFTISRDDSNTAYLQMNSLKTEDTAVYYC	TR TKSSSSDY	WGQGTLTVSS
5 β 2-D5-5 γ 5-8	EVQLVESGGGLVQPGGSLKLSCAAS EVQLVESGGGLVQPGGSLKLSCAAS	GFTFSGSA GFTFSGSA	MHWVRQASGKGLEWGR MHWVRQASGKGLEWGR	IRSKANSYAT IRSKANSYAT	AYAASVK GRFTISRDDSNTAYLQMNSLKTEDTAVYYC AYAASVK GRFTISRDDSNTAYLQMNSLKTEDTAVYYC	TR TR TYSSSWYFDY TR TYSSSWYFDY	WGQGTLTVSS WGQGTLTVSS
γ 6-7 5 β 1-E3-5	EVQLVESGGGLVQPGGSLKLSCAAS EVQLVESGGGLVQPGGSLKLSCAAS	GFTFSGSA GFTFSGSA	MHWVRQASGKGLEWGR MHWVRQASGKGLEWGR	IRSKANSYAT IRSKANSYAT	AYAASVK GRFTISRDDSNTAYLQMNSLKTEDTAVYYC AYAASVK GRFTISRDDSNTAYLQMNSLKTEDTAVYYC	TR RGYYGSGSYYGDY TR RGYYGSGSYYCDY	WGQGTLTVSS WGQGTLTVSS
VH4-34 2 β 2-D6-11	QVQLQQWGAGLLKPSETLSLCAVY QVQLQQWGAGLLKPSETLSLCAVY	GGSF S YY GGSF S YY	WSWIROPPGKGLEWIGE WSWIROPPGKGLEWIGE	INHSGST INHSGST	YNPNSLK SRVTISVDTSKNQFSLKLSVTAADTAVYYC YNPNSLK SRVTISVDTSKNQFSLKLSVTAADTAVYYC	AR AR GFY	WGQGTLTVSS WGQGTLTVSS
VH3-30 3 β 2-G1-7	QVQLVESGGGVVQPGRSRLSCAAS QVQLVESGGGVVQPGRSRLSCAAS	GFTFSSYG GFTFSSYG	MHWVRQAPGKGLEWAV MHWVRQAPGKGLEWAV	ISYDGSNK ISYDGSNK	YYADSVK GRFTISRDNSKNTLYLQMNSLRAEDTAVYYC YYADSVK GRFTISRDNSKNTLYLQMNSLRAEDTAVYYC	AR AKAVVDRARDGYNLY	WGQGTLTVSS WGQGTLTVSS
γ 6-14	PGRSLRLSCAAS	GFTF S Y A	MHWVRQAPGKGLEWAV	ISYDGSNK	YYADSVK GRFTISRDNSKNTLYLQMNSLRAEDTAVYYC	AR CTYYGSGIGFDY	WGQGTLTVSS
VH3-15 2 β 2-H8-2 γ 2-15	EVQLVESGGGLVKPGGSLRLSCAAS EVQLVESGGGLVKPGGSLRLSCAAS GSLRLSCAAS	GFTFSNAW GFTFSNAW GFTFSNAW	MSWVRQAPGKGLEWGR MSWVRQAPGKGLEWGR MSWVRQAPGKGLEWGR	IKSKTDGGTT IKSKTDGGTT IKSKTDGGTT	DYAAPVK GRFTISRDDSNTLYLQMNSLKTEDTAVYYC DYAAPVK GRFTISRDDSNTLYLQMNSLKTEDTAVYYC DYAAPVK GRFTISRDDSNTLYLQMNSLKTEDTAVYYC	TT LINWGIRD LINWGIRD	WGQGTLTVSS WGQGTLTVSS WGQGTLTVSS
VH5-51 γ 1-4	EVQLVQSGAEVKKPGESLKISCKGS EVQLVQSGAEVKKPGESLKISCKGS	GYSFTSYW GYSFT T YW	IGWVRQMPGKGLEWMGI IGWVRQMPGKGLEWMGI	IYPGDS T IYPGDS T	RYSPSFQ GQVTISADKSISTAYLQWSSLKASDTAMYYC RYSPSFQ GQVTISADKSISTAYLQWSSLKASDTAMYYC	AR AR HSEYYYDSSGYYLDY	WGQGTLTVSS WGQGTLTVSS
VH5-A γ 6-21	EVQLVQSGAEVKKPGESLRISCKGS EVQLVQSGAEVKKPGESLRISCKGS	GYSFTSYW GYSFTSYW	ISWVRQMPGKGLEWMGR ISWVRQMPGKGLEWMGR	IDPSDSYT IDPSDSYT	NYSPSFQ GHVTISADKSISTAYLQWSSLKASDTAMYYC KYSPSFQ GHV T ADKSISTAYLQWSSLKASDTAMYYC	AR AR HLREAVADFFPMDV	WGQGTLTVSS WG

The germ-line sequence is given for each V_H family, with indication of framework regions (FR) and complementary determining regions (CDR). Highlighted amino-acids correspond to N-additions (in the CDR3 region) and somatic hyper-mutation events, whether it results in a silent mutation (green) or not (red). Clones with identical BCR sequences are grouped together.