

MBD4^{+/+} Leader

```

TTCTCCCCCTCTCCTCTCCCTCTCCAGGTTCCCTGGTGCAGGCAGCGCTGACTCAGCCGGCCTCGGTGTCAGCAAACCCAGGAGAA

```

MBD4^{Δ/Δ}

MBD4^{+/+} CDR1

```

ACCGTCAAGATCACCTGCTCCGGGGGTGGCAGCTATGCTGGAAGTTACTATTATGGCTGGTACCAAGCAGAAGTCTCCTGGCAGTGCC
C11.1-----T
C11.3-----*T

```

MBD4^{Δ/Δ}

MBD4^{+/+} CDR2

```

CCTGTCACTGTGATCTATGACAACGACAAGAGACCCCTCGGACATCCCTCACGATTCTCCGGTTCCAATCCGGCTCCACAGCCACA
C11.1-----T
C11.3-----T
C11.4 *T
C11.5-----T
C11.7-----T
C11.8-----T
C11.10-----T
C11.12-----T

```

MBD4^{Δ/Δ}

MBD4^{+/+} CDR3

```

TTAAACCATCACTGGGGTCCGAGCCGATGACGAGGCTGTCTATTCTGTGGGAGCTACGAAGACAAACAGTGGTGCTGCATTGGGGCC
C11.13-----T

```

MBD4^{Δ/Δ}

MBD4^{+/+}

```

GGGACAAACCTGACCGTCCTAGGTGAGTCGCTGACCTCGTCT

```

MBD4^{Δ/Δ}

Suppl. Figure 1. SHM in the IgL gene of MBD4^{Δ/Δ}.11 subclones. The leader and CDR1, CDR2 and CDR3 motifs in the reference sequence are boxed. Mutations are from MBD4^{Δ/Δ}.11 (n=8, green). AID hotspot motifs (RGYW, WRCY) are highlighted in bold.

MBD4^{+/+} Leader

TTCTCCCCCTCTCCTCTCCCT	TCCAGGTTCCCTGGTGCAGG	CAGCGCTGACTC	AGCCGGCCTCGGTGTCAGCAAACCCAGGAGAA
-----------------------	----------------------	--------------	---

MBD4^{ΔΔ}

MBD4^{+/+} CDR1

ACCGTCAAGATCACCTGCT	CCGGGGGTGGCAGCTATGCTGGAAGTTACTATTATGGC	TGGTACCAGCAGAAGTCTCCTGGCAGTGCC
---------------------	--	--------------------------------

MBD4^{ΔΔ}

C12.1	G
C12.2	G
C12.3	G
C12.4	G
C12.5	G
C12.6	G
C12.7	G
C12.8	G
C12.9	G
C12.10	G
C12.11	G
C12.12	G

MBD4^{+/+} CDR2

CCTGTCACTGTGATCTATGACAACGACAAGAGACCCCTCGG	ACATCCCTCACGATTCTCCGGTTCCAATCC	GGCTCCACAGCCACA
---	--------------------------------	------------------------

MBD4^{ΔΔ}

MBD4^{+/+} CDR3

TTAACCATCACTGGGGTCCGAGCCGATGACGAGGCTGTCTATTCTGTGGGAGCTACGAAGACAAACAGTGGTGCTGCATTTGGGGCC		
---	--	--

MBD4^{ΔΔ} C12.1 TG-A T A

MBD4^{+/+}

GGGACAAACCTGACCGTCCTAGGTGAGTCGCTGACCTCGTCT		
--	--	--

MBD4^{ΔΔ}

Suppl. Figure 2. SHMs occur in the IgL gene of MBD4^{ΔΔ}.12 subclones. The leader and CDR1, CDR2 and CDR3 motifs in the reference sequence are boxed. Mutations are from MBD4^{ΔΔ}.12 (n=12, blue) below the reference sequence. AID hotspot motifs (RGYW, WRCY) are highlighted in bold.

MBD4^{+/+}

Leader

TTCTCCCTCTCCTCTCCCTCTCCAGGTTCCCTGGTGCAGGCAGCGCTGACTC**AGCCGGCCTCGGTGTCAGCAAACCCAGGAGAA**

MBD4^{Δ/Δ}

MBD4^{+/+}

CDR1

ACCGTCAAGATCACCT**GCTCCGGGGTGGCAGCTATGCTGGAAGTTACTATTATGGC**TGGTACCAGCAGAAGTCTCCT**GGCAGTGCC**
C14.1-----A-----

MBD4^{Δ/Δ}

C14.2 A
C14.3 A
C14.4 A
C14.5 A
C14.6 A
C14.7 A
C14.8 A
C14.9 A
C14.10 A
C14.11 A
C14.12 A

MBD4^{+/+}

CDR2

CCTGTCACTGTGATCTAT**ACAACGACAAGAGACCCCTCGG**ACATCCCTTCACGATTCTCC**GGTTCCAATCCGGCTCCACAGCCACA**
C14.10-----C-----

MBD4^{Δ/Δ}

MBD4^{+/+}

CDR3

TTAACCATCACTGGGGTCCG**AGCCGATGACGAGGCTGTCTATTCTGTGGGAGCTACGAAGACAAACAGTGGTGCTGCATTTGGGGCC**
C14.1-----A-----

MBD4^{Δ/Δ}

C14.2 A
C14.4 A
C14.6 A
C14.8 A
C14.9 A
C14.11 A
C14.12 A

MBD4^{+/+}

GGGACA**ACCC**CTGACCGTCCTAGGTGAGTCGCTGACCTCGTCT

MBD4^{Δ/Δ}

Suppl. Figure 3. SHMs occur largely at G/C residues in the IgL gene of MBD4^{Δ/Δ} subclones.

The leader and CDR1, CDR2 and CDR3 motifs in the reference sequence are boxed. Mutations are from MBD4^{Δ/Δ}.14 (n=12, red) deletion clones, below the reference sequence. AID hotspot motifs (RGYW, WRCY) are highlighted in bold.

MBD4^{+/+} Leader

TTCTCCCCCTCTCCTCTCCCT	TCCAGGTTCCCTGGTGCAGG	CAGCGCTGACTC	AGCCGGCCTCGGTGTCAGCAAACCCAGGAGAA
-----------------------	----------------------	--------------	---

MBD4^{Δ/Δ}

WT.12	A
WT.11	A
WT.9	A
WT.8	A
WT.6	A
WT.5	A
WT.4	A
WT.3	A
MBD4^{+/+}	CDR1
WT.2	A
WT.1	A
ACCGTCAAGATCACCTGCTCGGGGGTGGCAGCTATGCTGGAAGTTACTATTATGGCTGGTACCAGCAGAAGTCTCCTGGCAGTGCC	

MBD4^{Δ/Δ}

MBD4^{+/+}

CCTGTCACTGTGATCTAT	ACAACGACAAGAGACCCCTCGG	ACATCCCTCACGATTCTCC	GGTTCCAATCCGGCTCCACAGCCACA
--------------------	-------------------------------	---------------------	-----------------------------------

MBD4^{Δ/Δ}

WT.12	T	
WT.11	T	
WT.9	T	A
WT.8		C
WT.6	T	A
WT.5	T	A
WT.4	T	
MBD4^{+/+}	CDR2	
WT.3	T	
WT.2	T	A
WT.1	T	
TTAACCATCACTGGGTCCGAGCCGATGACGAGGCTGTCTATTCTGTGGAGCTACGAAGACAACAGTGGTGCTGCA TTTGGGCC		

MBD4^{Δ/Δ}

MBD4^{+/+}

GGGACA	ACCC	CTGACCGTCCTAGGTGAGTCGCTGACCTCGTCT
--------	-------------	-----------------------------------

MBD4^{Δ/Δ}

Suppl. Figure 4. SHMs occur in the IgL gene of MBD4^{+/+} subclones. The leader and CDR1, CDR2 and CDR3 motifs in the reference sequence are boxed. Mutations are from the parental (n=11, black; above the reference sequence). AID hotspot motifs (RGYW, WRCY) are highlighted in bold.