

**Fig. S1.**

WT <i>BCL-xL</i>	AGCAAGCGCTGAGGGAGGCAGGCGACGAGTTGAAC TGCGG-----
Octa/ <i>BCL-xL</i> KO <i>BCL-xL</i> a	AGCAAGCGCTGAGGGAGGCAGGCGACGAGTTGAAC TGCGG <ins>ACCGCCGCC</ins> TATGAAAGGTGGGCTCGGAATCGTTCCGGACGCCGGCTGGATGATCCT
Octa/ <i>BCL-xL</i> KO <i>BCL-xL</i> b	AGCAAGCGCTGAGGGAGGCAGGCGACGAGTTGAAC TGCGG <ins>AAGGCAA</ins> ATGCCGAAAAAGGGAATACGGGCACACGGAATGTTGAATACTCATACTCTCC
WT <i>BCL-xL</i>	-----
Octa/ <i>BCL-xL</i> KO <i>BCL-xL</i> a	CCAGCGCGGGATCTCATGCTGGAGTTCTCGCCCACCCAAC <ins>TGTTATTG</ins> CAGCTTATAATGTTACAAAT <ins>AA</ins> AGCAATAGCATCACAAATT <ins>TCACA</ins> AT <ins>AAA</ins>
Octa/ <i>BCL-xL</i> KO <i>BCL-xL</i> b	TTTTCAATATTATTGAAGCATTATCAGGGTTATTGTCTCATGAGCGGATACATATT <ins>TGA</ins> ATGTATTAGAAAATAACAAATAGGGTTCCGCGCACATTCC
WT <i>BCL-xL</i>	-----
Octa/ <i>BCL-xL</i> KO <i>BCL-xL</i> a	GCATTTTTCACTGCATTCTAGTTGTGGTTGTCAAAC <ins>CTCATCAATGTATCTTATCATGCTGTATACCGTCGACCTCTAGCTAGAGCTTGGCGTAA</ins> TATCATGGT
Octa/ <i>BCL-xL</i> KO <i>BCL-xL</i> b	CCGAAAAGTGCCACCTGACGTGACGGATC <ins>GGGAGATCTCCGATCCC</ins> TATGGTCACTCTCAGTACAATCTGCTCTGATGCCGATAGTTAAGCCAGTATCTGC
WT <i>BCL-xL</i>	----- <ins>TACCGGCGGGCATTCA</ins> GTGACCTGACATCCCAGCT
Octa/ <i>BCL-xL</i> KO <i>BCL-xL</i> a	CATAGCTGTTCCCTGTGTGAAATTGTTATCCGCTCA----- <ins>TACCGGCGGGCATTCA</ins> GTGACCTGACATCCCAGCT
Octa/ <i>BCL-xL</i> KO <i>BCL-xL</i> b	TCCCTGCTTGTGTGGAGGTCGCTGAGTAGTGCGCGAGC <ins>AAAATTAAAGCT</ins> TACCGGCGGGCATTCA <ins>GTGACCTGACATCCCAGCT</ins>

**Fig. S1 Mutations of the targeted *BCL-xL* locus in Octa/*BCL-xL* KO cells.**

Genomic sequence of *BCL-xL* around the CRISPR target sites aligned with the wild-type sequence. The CRISPR target sequence is underlined. Insertions are labeled in green and the stop codons are labeled in red.