

LncRNAs downregulated in childhood acute lymphoblastic leukemia modulate apoptosis, cell migration, and DNA damage response

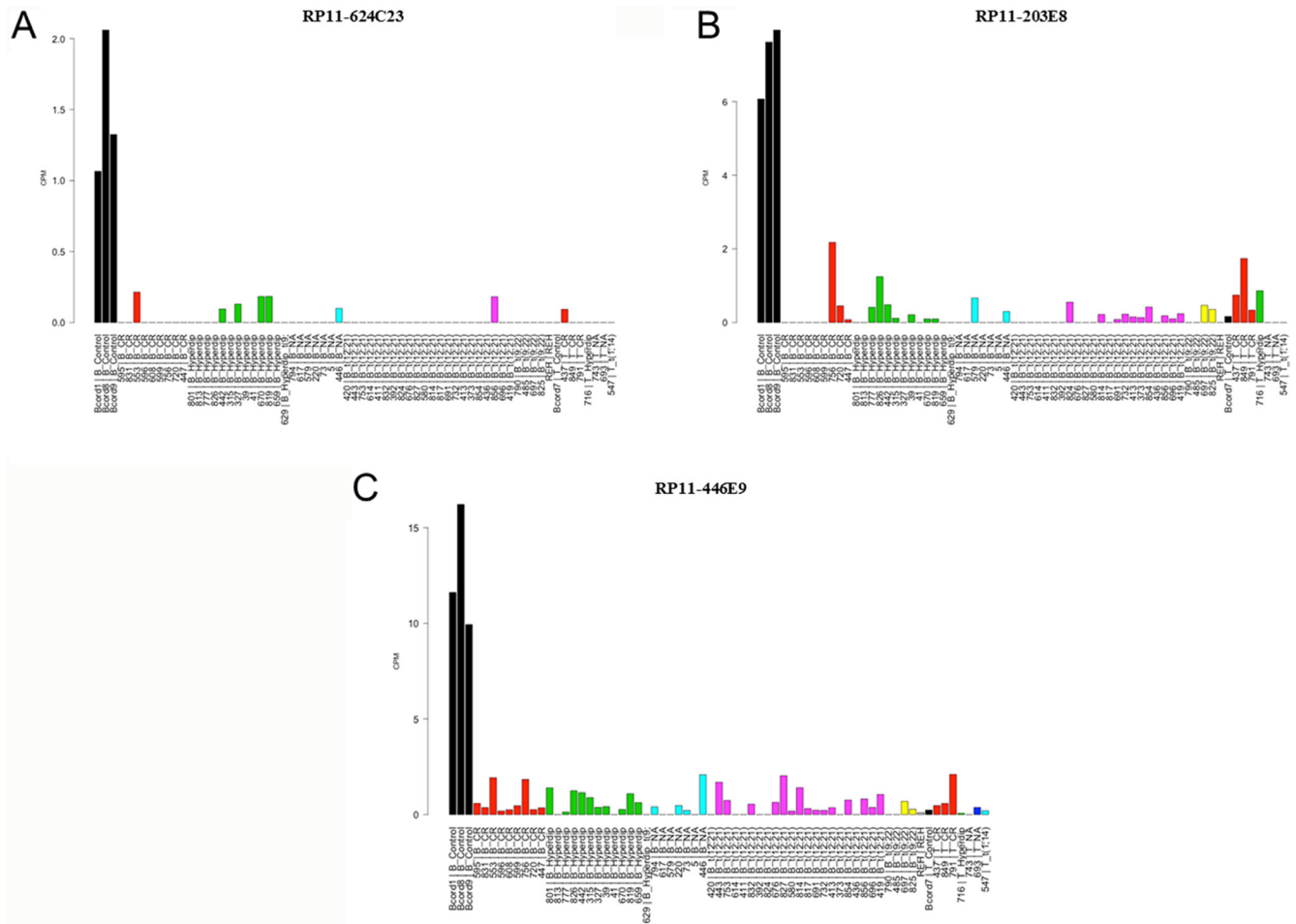
SUPPLEMENTARY MATERIALS

Supplementary Table 1: Characteristics of PCR amplimers

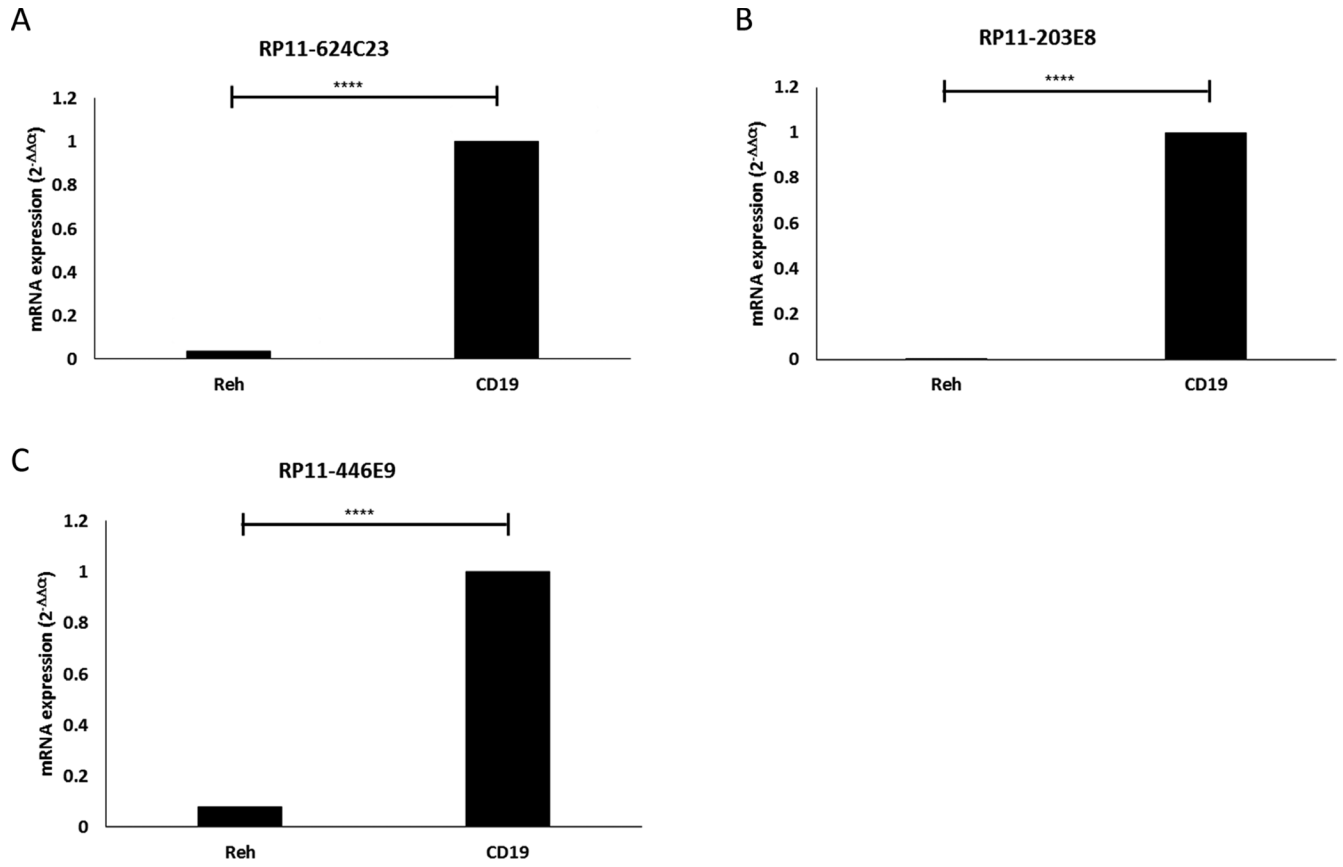
RP11-624C23.1	Forward	5'- GGGGACAAGTTTGTACAAAAAAGCAGGCTGAACAAGTGGGGAGAAATCTTA-3'
	Reverse	5'- GGGGACCACTTTGTACAAGAAAGCTGGGTGTTATGATTCCCATTTGGGTTG-3'
RP11-203E8	Forward	5'-GGGGACAAGTTTGTACAAAAAAGCAGGCTAAAACCTTATCAATTTCTTATATTTTAT-3'
	Reverse	5'- GGGGACCACTTTGTACAAGAAAGCTGGGTCTCAAATTATACCCTCTTTGTAG-3'
RP11-446E9.2	Forward	5'- GGGGACAAGTTTGTACAAAAAAGCAGGCTTCTCGAACTCCTGACCTCAGGC-3'
	Reverse	5'- GGGGACCACTTTGTACAAGAAAGCTGGGTGTGTTTCACTGGTTTTATTTCACA-3'

Supplementary Table 2: Characteristics of qPCR amplimers

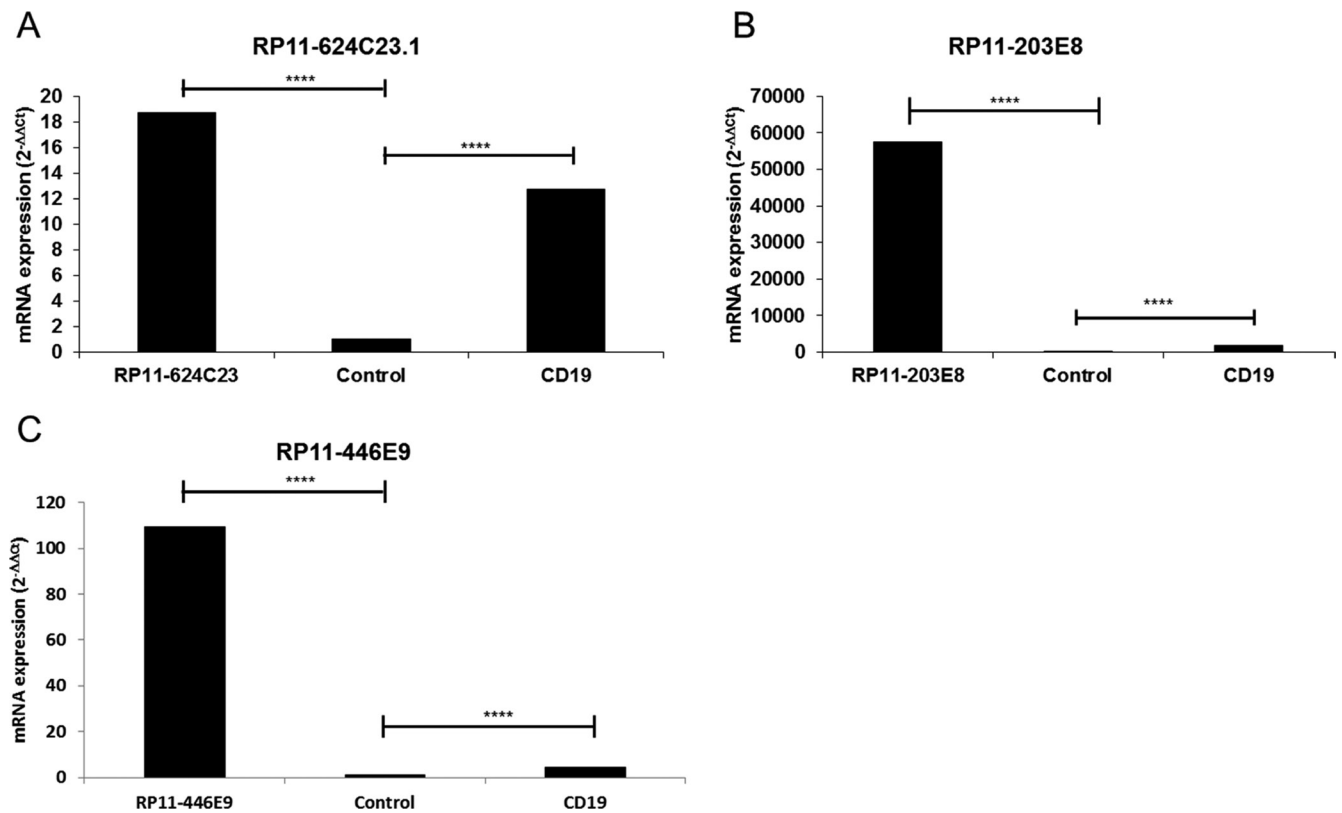
RP11-624C23.1	Forward	5'-CTGAGCCCTCTCTCCCAATA-3'
	Reverse	5'-ATCACTGGGCAGGTCAAGAG-3'
RP11-203E8	Forward	5'-CTCCCAATCCAAGAAACAG-3'
	Reverse	5'-GCCAAAATCCTCTCCATCA-3'
RP11-446E9.2	Forward	5'-CCCCGTCCTCTCACAACTA-3'
	Reverse	5'-GACAACTGGCTGTGGAACCT-3'
ADAM28	Forward	5'-AAGAACTCCCTGGGGTGAAG-3'
	Reverse	5'-AAAGCACTGCAATTTTCCA-3'
LYN	Forward	5'-CAGTACCAAGGTGGCTGTGA-3'
	Reverse	5'GGCCATGTACTCGGTGATG-3'
GAPDH	Forward	5'-TGCACCACCAACTGCTTAGC-3'
	Reverse	5'-GGCATGGACTGTGGTCATGAG-3'



Supplementary Figure 1: The expression of lncRNAs *RP11-624C23*, *RP11-203E8*, and *RP11-446E9* lncRNAs is downregulated in pre-B cALL patients. RNA sequencing was performing on 63 patients, 3 normal samples as control and Reh cell line as described in Ouimet et al. and Lajoie et al.

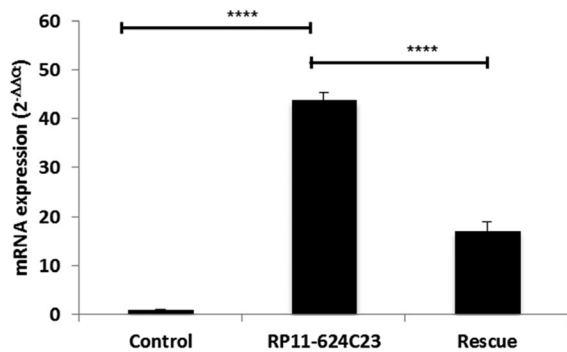


Supplementary Figure 2: LncRNAs *RP11-624C23.1*, *RP11-203E8*, and *RP11-446E9* are down-regulated in Reh cell line. QPCR of *RP11-624C23.1* (A), *RP11-203E8* (B), and *RP11-446E9* (C) lncRNAs were done on Reh using CD19⁺ primary cell (CD19) as control. **** $P \leq 0.0001$.

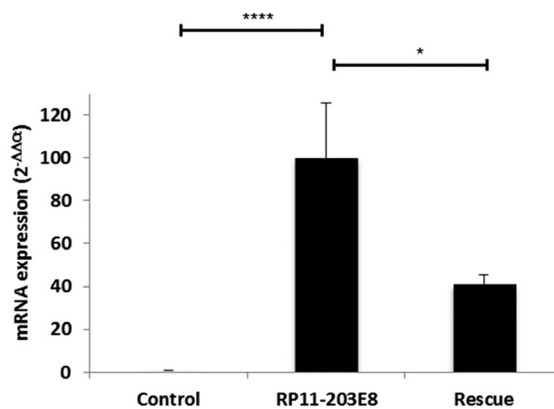


Supplementary Figure 3: LncRNAs *RP11-624C23.1*, *RP11-203E8*, and *RP11-446E9* are up-regulated in constructs overexpressing them. QPCR of *RP11-624C23.1* (A), *RP11-203E8* (B), and *RP11-446E9* (C) lncRNAs were done on Reh after infection with lentiviral expression vector. CD19⁺ primary cell (CD19) and Reh-pLenti (Control) were used as controls. Results were normalized on *GAPDH* expression. **** $P \leq 0.0001$.

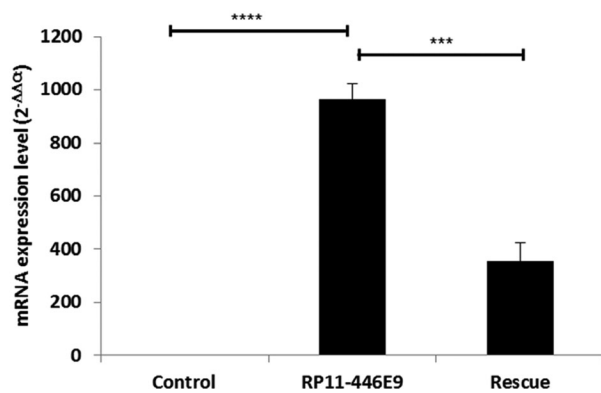
A



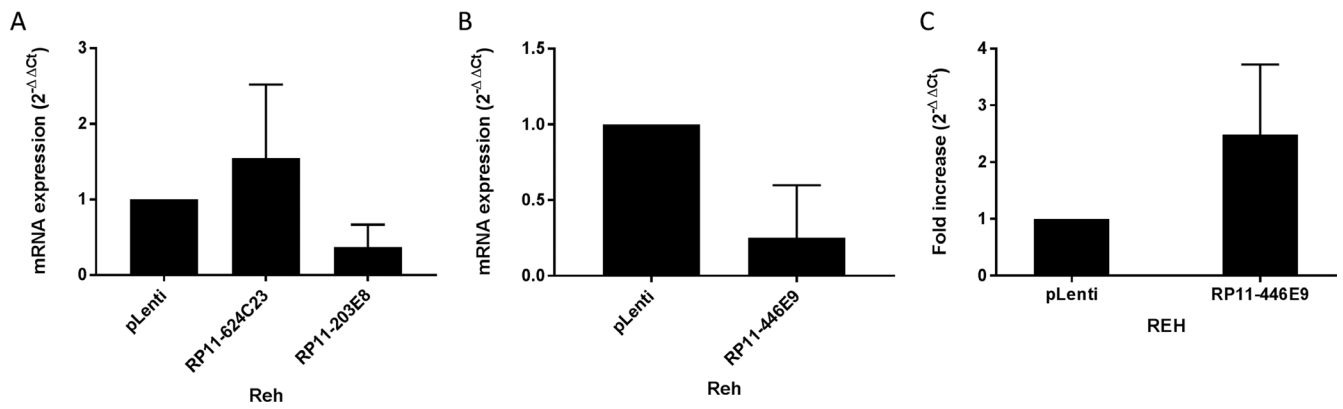
B



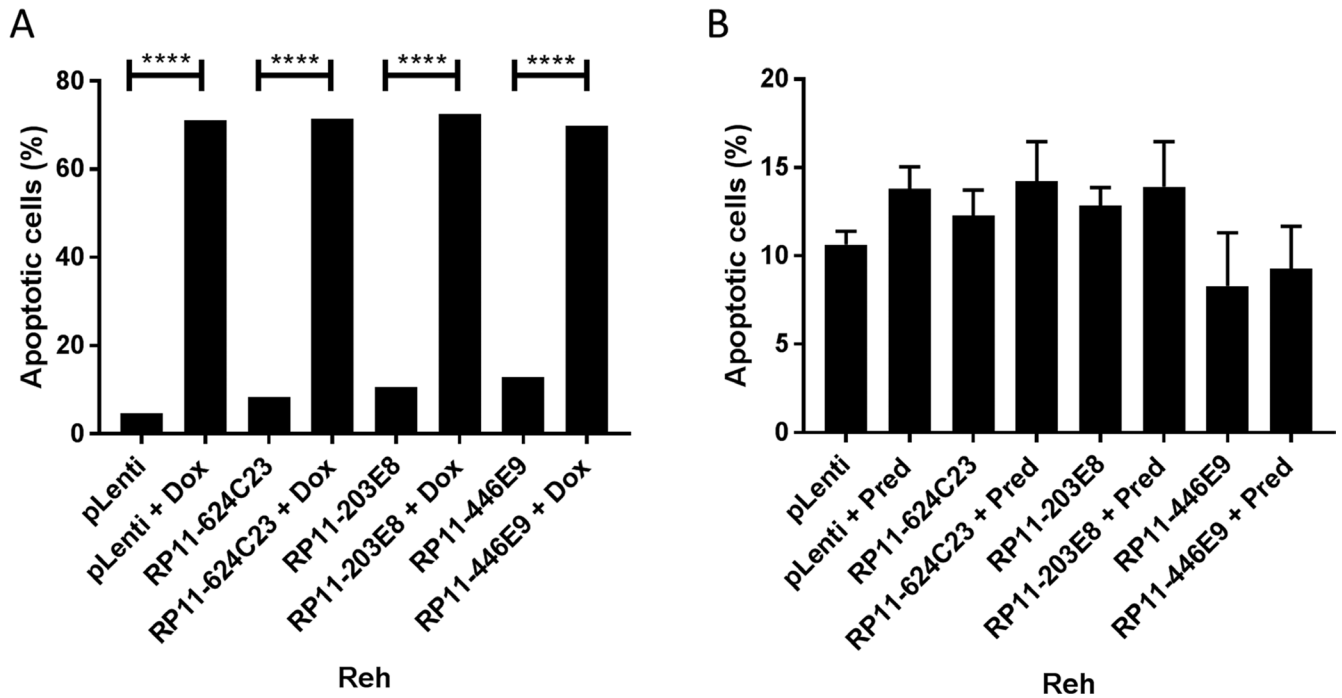
C



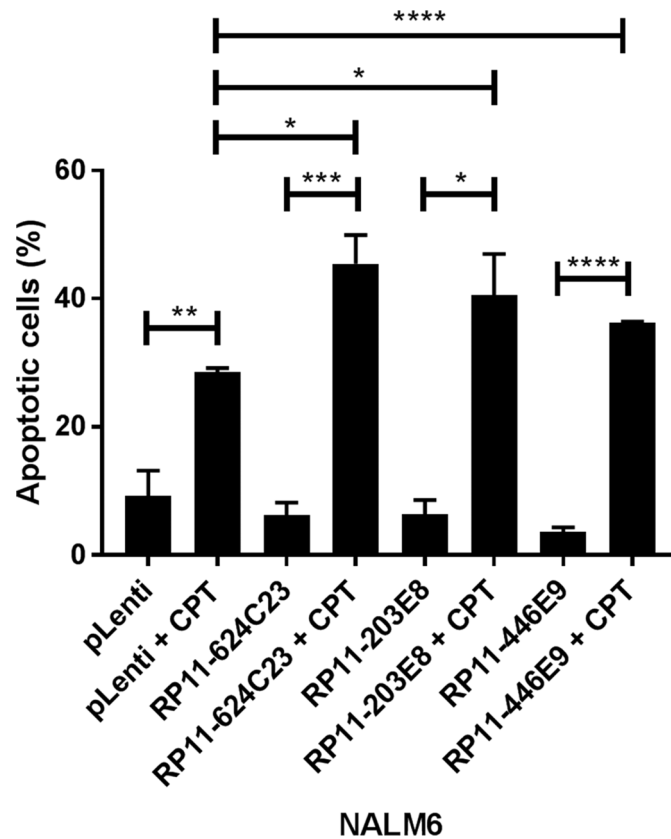
Supplementary Figure 4: Overexpression of *RP11-624C23.1*, *RP11-203E8*, and *RP11-446E9* is repressed by shRNA-mediated silencing. QPCR of *RP11-624C23.1* (A), *RP11-203E8* (B) and *RP11-446E9* (C) lncRNAs were done on Reh-*RP11-624C23.1* (A), Reh-*RP11-203E8* (B) and Reh-*RP11-446E9* (C) after shRNA-mediated silencing. Results were normalized on *GAPDH* expression. *, ***, and **** $P \leq 0.05$, ≤ 0.005 and ≤ 0.0001 , respectively.



Supplementary Figure 5: Expression levels of genes neighboring *RP11-624C23.1* and *RP11-203E8*. QPCR of *ADAM28* (A), *LYN* (B), and *RPS* (C) were done in control empty vector Reh cells (pLenti) and Reh cells overexpressing *RP11-624C23.1* or *RP11-203E8*. Results were normalized on each respective gene's expression in the pLenti control.



Supplementary Figure 6: Doxorubicine and prednisolone sensitivity are not affected by *RP11-624C23.1*, *RP11-203E8*, or *RP11-446E9* overexpression. Control (pLenti) Reh cells or Reh cells overexpressing *RP11-624C23.1*, *RP11-203E8*, or *RP11-446E9* were treated with doxorubicine (150 nM, 24 h) (A) or prednisolone (750 μ M, 24 h) (B). Apoptosis was measured using Annexin V/PI.



Supplementary Figure 7: NALM-6 pre-B cALL cells overexpressing *RP11-624C23.1*, *RP11-203E8*, or *RP11-446E9* are sensitive to camptothecin treatment. Control (pLenti) NALM-6 cells or NALM-6 cells overexpressing *RP11-624C23.1*, *RP11-203E8*, or *RP11-446E9* were treated with camptothecin (5 uM, 3 h). Apoptosis was measured using Annexin V/PI.

REFERENCES

1. Ouimet M, Drouin S, Lajoie M, Caron M, St-Onge P, Gioia R, Richer C, Sinnett D. A childhood acute lymphoblastic leukemia-specific lncRNA implicated in prednisolone resistance, cell proliferation, and migration. *Oncotarget*. 2017; 8:7477–88. <https://doi.org/10.18632/oncotarget.13936>.
2. Lajoie M, Drouin S, Caron M, St-Onge P, Ouimet M, Gioia R, Lafond MH, Vidal R, Richer C, Oualkacha K, Droit A, Sinnett D. Specific expression of novel long non-coding RNAs in high-hyperdiploid childhood acute lymphoblastic leukemia. *PLoS One*. 2017; 12:e0174124. <https://doi.org/10.1371/journal.pone.0174124>.