One-Bit Full Adder-Full Subtractor Logical Operation Based on DNA Strand Displacement

Yanfeng Wang, Xing Li, Chun Huang, Guangzhao Cui, and Junwei $\text{Sun}^{(\boxtimes)}$

College of Electrical and Electronic Engineering, Zhengzhou University of Light Industry, No. 5 Dongfeng Road, Zhengzhou 450002, China junweisun@yeah.net

Abstract. DNA nanotechnology has become a reliable, programmable control method, which can realize complex reaction networks nanostructures due to the accuracy and predictability complementary DNA base pairing. In this paper, one-bit full adder-full subtractor is constructed to achieve two kinds of functions which are full adder function and full subtractor function, respectively. Based on the cascaded DNA strand displacement reaction, the digital logic circuit is further translated into its corresponding dual-rail logic circuit and seesaw cascade logic circuit. The simulation results prove the feasibility and effectiveness of the designed circuit.

Keywords: DNA strand displacement \cdot Full adder-full subtractor \cdot Dual-rail circuit \cdot Seesaw circuit

1 Introduction

DNA computing is a new field which combines computer science and molecularbiology subject. DNA acts as the computing tool, which has solved many problems, such as solving Hamition path, maximal clique problem [\[1](#page-7-0)[–3](#page-7-1)]. DNA computing has merged a lot of molecule operation technology, such as self-assembly $[4-6]$ $[4-6]$, fluorescence labeling $[7-9]$ $[7-9]$ strand displacement $[10-12]$ $[10-12]$ and probe machine [\[13](#page-8-4)]. DNA self-assembly technology and DNA stand displacement technology are two important technical support of DNA nanotechnology. DNA strand displacement technology is developed on the basis of DNA self-assembly technology. In recent years, DNA strand displacement [\[14](#page-8-5)[–16](#page-8-6)] is a new method in the biocomputing, and has become a common method in DNA self-assembly. Based on the strand displacement cascade reaction [\[17](#page-8-7)], the dynamical connection adjacent logic modules [\[18](#page-8-8)[–20\]](#page-8-9) have been achieved, which makes it possible for the researcher to construct large-scale, complicated logic circuits [\[21](#page-8-10)]. Moreover, with the advantage of high-capacity information accumulation, high performance parallel computing, programming and simulating, it had acquired an in-depth study in the field of molecular computing, nano-machine, diagnosis and remedy of the disease. DNA strand displacement technology has the gigantic proficiency in

-c Springer Nature Singapore Pte Ltd. 2016

M. Gong et al. (Eds.): BIC-TA 2016, Part I, CCIS 681, pp. 30–38, 2016.

DOI: 10.1007/978-981-10-3611-8 4

solving the math problem [\[22](#page-8-11)[–24](#page-8-12)], managing the nano-machine and discussing the life course. Based on DNA strand displacement, the construction of the biochemistry logic circuits has significant research means by mastering the design procedures.

Compared with the previous work $[25,26]$ $[25,26]$ $[25,26]$, the one-bit full adder-full subtractor logical operation is constructed for the first time in this paper. Based on the DNA strand displacement, there is the control terminal which concerns the function of the full adder-full subtractor. There are two kinds of function in the circuit, which are the full adder and the full subtroctor, respectively. The circuit could be applied to the construction of the biological computer in the future.

This paper is organized as follows. Firstly, the introduction is described in the Sect. [1.](#page-0-0) Then the strand displacement reaction mechanism is shown in the Sect. [2.](#page-1-0) The digit circuit and dual-rail circuit of the full adder-full subtractor is designed in the Sect. [3.](#page-2-0) The seesaw circuit and the simulation of the full adderfull subtractor are shown in the Sect. [4.](#page-4-0) Finally, the conclusion is given for the full adder-full subtractor.

2 The Background of DNA Strand Displacement

DNA strand displacement technology acts as an important technology of modern biological computing, which has been proved that it is a kind of nanoscale technologies to overcome circuit component miniaturization problem. DNA strand displacement response is a dynamic process and has the following three advantages. (1) DNA strand displacement response doesn't need special temperature requirements and can proceed in room temperature without annealing. (2) It is the spontaneous reaction without adding enzyme. (3) A dynamic cascade system can be constituted due to dynamic characteristics of DNA strand displacement response. The characteristics of DNA strand displacement technology provide a good way for building the nanoscale large-scale circuit.

In order to perform mathematical logic, DNA strand displacement cascade technology has been widely applied to the configuration of the basic DNA logic gates (AND gate, OR gate and NOT gate). DNA strand displacement reaction mechanism is shown in Fig. [1.](#page-2-1) In the Fig. [1,](#page-2-1) the domain t represents a short toehold and the domain t* is the complementary pairing of the domain t. The DNA single strand $\langle a \cdot b \rangle$ and strand $\langle b \cdot c \rangle$ represent input signal and output signal, respectively. The strand \langle b t \rangle and strand $\{t^*\}$ [b t] \langle c \rangle are recognition domain and the double strand complex, respectively. Firstly, the short toehold domain t of the strand *<*atb*>* and toehold domain t* of the double strand complex have a DNA complementary pairing. Then the domain b of the DNA single strand $\langle a, b \rangle$ and domain b^* of the double strand complex also conduct a DNA complementary pairing. Eventually, the output strand *<*btc*>* falls off from the double strand complex and releases the molecule complex *<*a*>*[t b]:*<*b*>* [t]*<*c*>* The whole reaction process can be considered that a DNA single strand $\langle a \cdot b \rangle$ replace of the DNA single strand $\langle b \cdot c \rangle$.

Fig. 1. The DNA strand displacement reaction process. The DNA single strand *<*a t b*>* and strand *<*btc*>* represent input signal and output signal. The strand *<*b t*>* is recognition domain.

3 The Digit Circuit and Dual-rail Circuit

The full adder-full subtractor also is a combinational logic circuit which performs simple logic operations of four binary digits. The truth table of the full adderfull subtractor is given in Table [1.](#page-2-2) In the Table [1,](#page-2-2) there are sixteen kinds of conditions which achieve two kinds of functions. Boolean functions of logic circuit is constructed by "AND", "OR" and "NOT" gates. Based on the function of the truth table one-bit the full adder-full subtractor combinational logic circuit is shown in Fig. [2.](#page-3-0) The full adder-full subtractor logic circuit haves four inputs which are x_0, x_1, x_2 and y_0 in the left side of the logic circuit and two outputs which are y_1 and y_2 in the right side of the logic circuit, respectively. In the

Logical function	Input x_0 Input x_1		Input x_2 Input y_0		Output y_1	Output y_2
Full adder	$\overline{0}$	θ	$\overline{0}$	$\overline{0}$	$\boldsymbol{0}$	$\overline{0}$
	$\overline{0}$	$\overline{0}$	$\boldsymbol{0}$	$\mathbf{1}$	$\mathbf{1}$	$\overline{0}$
	$\overline{0}$	$\overline{0}$	$\mathbf{1}$	$\overline{0}$	$\mathbf{1}$	$\overline{0}$
	$\overline{0}$	θ	$\mathbf{1}$	$\mathbf{1}$	$\overline{0}$	$\mathbf{1}$
	$\overline{0}$	$\mathbf{1}$	$\overline{0}$	$\overline{0}$	1	$\overline{0}$
	θ	$\mathbf{1}$	$\overline{0}$	1	$\overline{0}$	1
	$\overline{0}$	$\mathbf{1}$	$\mathbf{1}$	$\overline{0}$	$\overline{0}$	$\mathbf{1}$
	$\overline{0}$	$\mathbf{1}$	$\mathbf{1}$	$\mathbf{1}$	1	$\mathbf{1}$
Full subtractor	$\mathbf{1}$	$\overline{0}$	$\overline{0}$	$\overline{0}$	$\overline{0}$	$\overline{0}$
	$\mathbf{1}$	$\overline{0}$	$\overline{0}$	$\mathbf{1}$	$\mathbf{1}$	$\mathbf{1}$
	1	θ	$\mathbf{1}$	$\overline{0}$	1	1
	1	$\overline{0}$	$\mathbf{1}$	$\mathbf{1}$	$\overline{0}$	$\mathbf 1$
	$\mathbf{1}$	$\mathbf{1}$	$\overline{0}$	$\boldsymbol{0}$	1	$\overline{0}$
	$\mathbf{1}$	$\mathbf{1}$	$\overline{0}$	$\mathbf{1}$	$\overline{0}$	$\overline{0}$
	1	$\mathbf{1}$	$\mathbf{1}$	$\overline{0}$	$\overline{0}$	$\overline{0}$
	1	$\mathbf{1}$	$\mathbf{1}$	$\mathbf{1}$	$\mathbf{1}$	$\mathbf{1}$

Table 1. Truth table of the full adder-full subtractor.

Fig. 2. Digital logic circuit of one-bit full adder-full subtractor. x_0 , x_1 , x_2 and y_0 are input signal, y_1 and y_2 are output signal. x_0 is the control terminal which concerns the function of the circuit. y_0 is the low level carry-bit or low level borrow-bit.

logic circuit, the input x_0 is the control terminal and concerns the function of the circuit. If the value of x_0 is "0", the logic circuit implements the function of full adder in which x_1 and x_2 are two addends, y_0 , y_1 and y_2 indicate low level carry-bit the sum-bit and the high level carry-bit, respectively. On the other hand, the logic circuit achieves the function of full subtractor if the value of x_0 is "1". Under the circumstances, x_1 and x_2 are minuend and subtrahend, y_0, y_1 and y_2 indicate low level borrow-bit the difference-bit and the high level borrow-bit, respectively.

Fig. 3. The dual-rail logic circuits. (a) The dual-rail logic circuit of AND gate. (b) The dual-rail logic circuit of OR gate. (c) The dual-rail logic circuit of NAND gate. (d) The dual-rail logic circuit of NOR gate. (e) The dual-rail logic circuit of full adder-full subtractor.

On the basis of the given principle, one-bit the full adder-full subtractor combinational logic circuit should be translated into the corresponding dual-rail circuit to avoid generating error output signal. The logic gate consists of a pair of "AND" gate and "OR" gate. The dual-rail circuits of "AND" gate, "OR" gate, "NAND" gate and "NOR" gate are shown in Fig. $3(a)$ $3(a)$ –(d), respectively. The input signal state is represented by logic "ON" and "OFF". Taking the input x_0 as an example, if the input x_0 participates in the reaction, then the states of the x_0^0 and x_0^1 represent logic "OFF" and "ON", respectively. On the contrary, if the input x_0 can't participate in the reaction, then the states of the x_0^0 and x_0^1 represent logic "ON" and "OFF" respectively. According to the corresponding logical relationship, the dual-rail circuit of the full adder-full subtractor is constructed, as shown in Fig. $3(e)$ $3(e)$.

4 Seesaw Circuit and Simulation with Visual DSD

Based on DNA stand displacement response, DNA seesaw logic gates could be designed. The logic gates are divided into left side and right side, and are con-nected by a node, as shown in Fig. [4.](#page-4-1) In Fig. $4(a)$ $4(a)$, the input acts as the input single strand $\langle a \cdot b \rangle$, the strand $\{t^*\}$ [b t] $\langle c \rangle$ is the gate note. The output 1 acts as the output strand *<*btc*>*. And output 2 is other output strand. The red digit 0.6 is the threshold value. The value of fuel is two times of the total output value. The gates usually consist of amplifying gate which can produce multi-path outputs and integration gate which could receive multi-path inputs, respectively. One-input-four-output and one-input-five-output amplifying gates are shown in Fig. $4(b)$ $4(b)$ –(c), respectively. Two-input-one-output integration gate is shown in Fig. $4(d)$ $4(d)$. The seesaw motifs of "OR" gate and "AND" gate are shown in Fig. $4(e)$ $4(e)$ –(f).

Fig. 4. The seesaw motif of basic gates. (a) The amplifying gate of one-input-twooutput. (b) The amplifying gate of one-input-four-output. (c) The amplifying gate of one-input-five-output. (d) The integration gate of two-input-one-output. (e) OR gate. (f) AND gate. (Color figure online)

According to the seesaw circuit and the dual-rail logic circuit of full adderfull subtractor, the seesaw circuit of full adder-full subtractor is shown in Fig. [5,](#page-5-0) which could be simulated in the Visual DSD. There will be sixteen outputs which are produced along with the sixteen inputs. The simulation results of the full adder-full subtractor are shown in Fig. [6.](#page-6-0) In the Fig. [6,](#page-6-0) the blue curve and yellow curve represent the value of y_1^0 and y_1^1 , respectively; the red line and green line separately indicate the value of y_2^0 and y_2^1 , respectively In this paper, the total concentration of the reaction is 1000 nm. When the range is 0–100 nm, it expresses the logic "0". On the other hand, it expresses the logic "1" if the range is 900–1000 nm.

If the control terminal x_0 is "0", then the function of logic circuit is full adder whose simulation results are shown in Fig. $6(a)$ $6(a)$ –(h). Under this cases, if the input signal $x_1x_2y_0$ is "000", then the output signal y_1y_2 is "00" in Fig. [6\(](#page-6-0)a). If the input signal $x_1x_2y_0$ is "001", then the output signal y_1y_2 is "10" in Fig. [6\(](#page-6-0)b). If the input signal $x_1x_2y_0$ is "010", then the output signal y_1y_2 is "10", as shown in Fig. $6(c)$ $6(c)$. If the input signal $x_1x_2y_0$ is "011", then the output signal y_1y_2 is "01", as shown in Fig. $6(d)$ $6(d)$. If the input signal $x_1x_2y_0$ is "100", then the output

Fig. 5. The seesaw circuit of one-bit full adder-full subtractor.

signal y_1y_2 is "10" in Fig. $6(e)$ $6(e)$. If the input signal $x_1x_2y_0$ is "101", then the output signal y_1y_2 is "01" in Fig. [6\(](#page-6-0)f). If the input signal $x_1x_2y_0$ is "110", then the output signal y_1y_2 is "01", as shown in Fig. $6(g)$ $6(g)$. If the input signal $x_1x_2y_0$ is "111", then the output signal y_1y_2 is "11", as shown in Fig. $6(h)$ $6(h)$.

If the control terminal x_0 is "1", then the function of the logic circuit is full subtractor whose simulation results are shown in Fig. $6(i)$ $6(i)$ –(p). If the input signal $x_1x_2y_0$ is "000", then the output signal y_1y_2 is "00", as shown in Fig. [6\(](#page-6-0)i). If the input signal $x_1x_2y_0$ is "001", then the output signal y_1y_2 is "11" in Fig. [6\(](#page-6-0)j). If the input signal $x_1x_2y_0$ is "010", then the output signal y_1y_2 is "11", as shown in Fig. $6(k)$ $6(k)$. If the input signal $x_1x_2y_0$ is "011", then the output signal y_1y_2 is "01", as shown in Fig. $6(1)$ $6(1)$. When the input signal $x_1x_2y_0$ is "100", the output signal

Fig. 6. The simulation in Visual DSD. (Color figure online)

 y_1y_2 is "00", as shown in Fig. $6(m)$ $6(m)$. If the input signal $x_1x_2y_0$ is "101", then the output signal y_1y_2 is "11" in Fig. [6\(](#page-6-0)n). If the input signal $x_1x_2y_0$ is "110", then the output signal y_1y_2 is "11", as shown in Fig. [6\(](#page-6-0)o). If the input signal $x_1x_2y_0$ is "111", then the output signal y_1y_2 is "01", as shown in Fig. $6(p)$ $6(p)$.

According to the simulation results, the following conclusions can be obtained. The logic "ON" and logic "OFF" curves both enter into the stable area, which express the logic "1" and logic "0" correctly, respectively. The simulation results have a high validity.

5 Conclusion

In this paper, one-bit full adder-full subtractor logic circuit has been constructed by DNA strand displacement. Then the full adder-full subtractor combinational logic circuit has been converted to the corresponding the dual-rail logic circuit and the biochemical logic circuit. Finally, the seesaw logic circuit of the full adder-full subtractor is simulated in the visual DSD. The correctness of simulation results proves that the DNA strand displacement technique is a feasible method in the study of biochemical circuit. As a result of the limitation of the current scientific research platform and technology, the biochemical experiment also needs to continue to be explored, which will be the focus of the next research direction.

Acknowledgments. The work is supported by the State Key Program of the National Natural Science Foundation of China (Grant No. 61632002), the National Natural Science Foundation of China (Grant Nos. 61472371, 61472372, 61572446, 61602424 and 61603348), China Postdoctoral Science Foundation funded project (Grant Nos. 2015M570641 and 2016T90687), Basic and Frontier Technology Research Program of Henan Province (Grant No. 162300410220), Key Program of Higher Education of China Henan Province (Grant No. 17A120005) and the Science Foundation of for Doctorate Research of Zhengzhou University of Light Industry (Grant No. 2014BSJJ044).

References

- 1. Adleman, L.M.: Molecular computation of solutions to combinatorial problems. Science **226**, 1021–1024 (1994)
- 2. Carlson, R.: The changing economics of DNA synthesis. Nat. Biotechnol. **27**, 1091– 1094 (2009)
- 3. Turberfield, A.J., Mitchell, J.C., Yurke, B., Mills Jr., A.P., Blakey, M.I., Simmel, F.C.: DNA fuel for free-running nanomachines. Phys. Rev. Lett. **90**, 118102-1– 118102-4 (2003)
- 4. Yin, P., Choi, H.M.T., Calvert, C.R., Pierce, N.A.: Programming biomolecularselfassembly pathways. Nature **451**, 318–322 (2008)
- 5. Gothelf, K.V., LaBean, T.H.: DNA-programmed assembly of nanostructures. Org. Biomol. Chem. **3**, 4023–4037 (2005)
- 6. Shi, X.L., Lu, W., Wang, Z.Y., Pan, L.Q., Cui, G.Z., Xu, J., La Bean, T.H.: Programmable DNA tile self-assembly using a hierarchical sub-tile strategy. Nanotechnology **25**(7), 075602 (2014)
- 7. Zhang, D.Y., Turberfield, A.J., Yurke, B., Winfree, E.: Engineering entropydrivenreactions and networks catalyzed by DNA. Science **318**, 1121–1125 (2007)
- 8. Seelig, G., Soloveichik, D., Zhang, D.Y., Winfree, E.: Enzyme-free nucleic acid logiccircuits. Science **314**, 1585–1588 (2006)
- 9. Chiniforooshan, E., Doty, D., Kari, L., Seki, S.: Scalable, time-responsive, digital, energy-efficient molecular circuits using DNA strand displacement. In: Sakakibara, Y., Mi, Y. (eds.) DNA 2010. LNCS, vol. 6518, pp. 25–36. Springer, Heidelberg (2011). doi[:10.1007/978-3-642-18305-8](http://dx.doi.org/10.1007/978-3-642-18305-8_3) 3
- 10. Srinivas, N., Ouldridge, T.E., Sulc, P., Schaeffer, J.M., Yurke, B., Louis, A.A., Doye, J.P.K., Winfree, E.: On the biophysics and kinetics of toehold-mediated DNA strand displacement. Nucleic Acids Res. **41**(22), 10641–10658 (2013)
- 11. Shi, X.L., Wang, Z.Y., Deng, C.Y., Song, T., Pan, L.Q., Chen, Z.H.: A Novel Bio-Sensor Based on DNA Strand Displacement. PLoS ONE **9**(10), e108856 (2014). doi[:10.1371/journal.pone.0108856](http://dx.doi.org/10.1371/journal.pone.0108856)
- 12. Yang, J., Dong, C., Dong, Y.F., Liu, S., Pan, L.Q., Zhang, C.: Logic nanoparticle beacon triggered by the binding-induced effect of multiple inputs. ACS Appl. Mater. Interfaces **6**(16), 14486–14492 (2014)
- 13. Xu, J.: Probe machine. IEEE Trans. Neural Netw. Learn. Syst. **27**(7), 1405–1416 (2016)
- 14. Zhang, D.Y., Winfree, E.: J. Am. Chem. Soc. **131**, 17303–17314 (2009)
- 15. Soloveichik, D., Seelig, G., Winfree, E.: Proc. Nat. Acad. Sci. U.S.A. **107**, 5393– 53989 (2010)
- 16. Phillips, A., Cardelli, L.: J. R. Soc. Interface **6** (2009). doi[:10.1098/rsif.2009.0072.](http://dx.doi.org/10.1098/rsif.2009.0072.focus) [focus](http://dx.doi.org/10.1098/rsif.2009.0072.focus)
- 17. Eckhoff, G., Codrea, V., Ellington, A.D., Chen, X.: J. Syst. Chem. **1**, 13 (2010). doi[:10.1186/1759-2208-1-13](http://dx.doi.org/10.1186/1759-2208-1-13)
- 18. Qian, L., Winfree, E.: A simple DNA gate motif for synthesizing large-scale circuits. J. R. Soc. Interface **1**, 13 (2011). doi[:10.1098/rsif.2010.0729](http://dx.doi.org/10.1098/rsif.2010.0729)
- 19. Lund, K., Manzo, A.J., Dabby, N., Michelotti, N., Johnson-Buck, A., Nangreave, J., Taylor, J.S., Pei, R., Stojanovic, M.N., Walter, N.G., Winfree, E., Yan, H.: Nature **465**, 206–210 (2010)
- 20. Qian, L., Soloveichik, D., Winfree, E.: Efficient turing-universal computation with DNA polymers. In: Sakakibara, Y., Mi, Y. (eds.) DNA 2010. LNCS, vol. 6518, pp. 123–140. Springer, Heidelberg (2011). doi[:10.1007/978-3-642-18305-8](http://dx.doi.org/10.1007/978-3-642-18305-8_12) 12
- 21. Gaber, R., Lebar, T., Majerle, A., Ster, B., Dobnikar, A., Bencina, M., Jerala, R.: Designable DNA-binding domain enable construction of logic circuit in mammaliancells. Nat. Chem. Biol. **10**, 203–208 (2014)
- 22. Zhang, Z., Li, J., Pan, L., Ye, Y., Zeng, X., Song, T., Zhang, X., Wang, E.K.: Anovel visualization of DNA sequences, reflecting GC-content. MATCH Commun. Math. Comput. Chem **72**, 533–550 (2014)
- 23. Zeng, X., Xu, L., Liu, X., Pan, L.: On languages generated by spiking neural P systems with weights. Inform. Sci. **278**, 423–433 (2014)
- 24. Zhang, X., Liu, Y., Luo, B., Pan, L.: Computational power of tissue P systems for generating control languages. Inform. Sci. **278**, 285–297 (2014)
- 25. Wang, Y., Tian, G., Hou, H., et al.: Simple logic computation based on the DNA strand displacement. J. Comput. Theor. Nanosci. **11**, 1975–1982 (2014)
- 26. Cui, G., Zhang, J., Cui, Y., et al.: DNA strand-displacement digital logic circuit with fluorescence resonance energy transfer detection. J. Comput. Theor. Nanosci. **12**, 2095–2100 (2015)