Electronic Supplementary Information

Information processing using an integrated DNA reaction network

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1. Oligonucleotide sequences used in this work.

| Name | Sequence from 5' to 3' |
|-------|--|
| A1 | AGC CAG CTG AGC CAA TTC ATG GAC CAG AAC AAC C |
| A1* | GGT CGT TCT GGT CCA TGA ATT GGC TCA GCT GGC T |
| CBc2 | TGA GCC AAT TCA TGG ACC AGA ACA ACC TGA CGC TCC TAG CTT CAC TGA CTC ACT CTA CT |
| Bc2 | CAG TAG AGT GAG TCA GTG AAG TCA GTG CTA GGA GCG TCA |
| WBc2 | TGA GTC AGT GAA GCT AGG AGC GTC AGG TTG T |
| CBa3' | TGA GCC AAT TCA TGG ACC AGA ACA ACC ATC AAA ATA TGT GAA TTT TTA CTA CCC AAG |
| | AA |
| Ba3' | CAT GGC CTT CTT GGG TAG TAA AAA TTT CAG CTC ACA TAT TTT GAT |
| WBa3' | GGT AGT AAA AAT TCA CAT ATT TTG ATG GTT GT |
| CBa3' | TGA GCC AAT TCA TGG ACC AGA ACA ACC ATC AAA ATA TGT GAA TTT TTA CTA CCC AAG |
| | AA |
| A2 | CCTTAGCAGAGCTGTGGAGTGTGACAATGGTGTT |
| A2* | AAC ACC ATT GTC ACA CTC CAC AGC TCT GCT AAG G |
| CBa1* | CCT TCT TGG GTA GTA AGA GAG TTG CCC TCC TTA GCA GAG CTG TGG AGT GTG ACA AT |
| WBa1* | GCT AAG GAG GGC AAC TCT CTT ACT ACC CAA G |
| Ba1* | AGG GCA ACT CTC CTT TTT TTT ACT ACC CAA GAA GGC C |
| CBf2' | CAG AGC TGT GGA GTG TGA CAA TGG TGT TGC AAC CGC AGC TAG GAT AGA AGT TGC CCA A |
| WBf2' | ACT TCT ATC CTA GCT GCG GTT GCA ACA CC |
| Bf2 | TTG GGC AAC TTC TAT CCT TTG GCA GCT GCG GTT GC |
| CBm2 | CAG AGC TGT GGA GTG TGA CAA TGG TGT TTG ACG CTC CTA GCT TCA CTG ACT CAC TCT AC |
| WBm2 | GAG TCA GTG AAG CTA GGA GCG TCA AAC ACC |
| A3 | TTG TTC CAC TTA TTT TAA TAG TTG TAG TTG TCG G |
| A3* | CCG ACA ACT ACA ACT ATT AAA ATA AGT GGA ACA A |
| CBc1 | CAC TTA TTT TAA TAG TTG TAG TTG TCG GGA GTG ACA CTG TAG TCA GTG CTA GGA GGT TC |
| Bc1 | TGA ACC TCC TAG CAC TGA CTT CAC TGA CAG TGT CAC TC |
| WBc1 | CCT AGC ACT GAC TAC AGT GTC ACT CCC GAC A |
| CBf1 | CAC TTA TTT TAA TAG TTG TAG TTG TCG GAG GGC AAC TCT CGC ATT TCA GCT CAA ATA TT |
| Bf1 | ATC AAA TAT TTG AGC TGA AAT GCC AAA GGA GAG TTG CCC T |
| WBf1 | TGA GCT GAA ATG CGA GAG TTG CCC TCC GAC A |
| OR1 | ACC TAC CAA AAC ACT CCT CCT AAA ACA CCT CAC CCA |
| COR1 | TGG GTG AGG TGT TTT AGG GTT TTG GTA GGT TTG TTC CAC TTA TTT TAA TAG TTG TAG |
| WOR1 | GAA CAA ACC TAC CAA AAC CCT AAA ACA CC |
| B1d | TGAACCTCCTAGCACTGGAGACTACGTCTCAG |
| B2d | CAGTAGAGTGAGTCAGTGAGTACTGACGTA |
| A4 | ACT CTT TGG CAA CGA CCC CTC GTC ACA ATA AAG A |
| | |

Table S1. The oligonucleotide sequences used in this work.

A4* TCT TTA TTG TGA CGA GGG GTC GTT GCC AAA GAG T CBc2 GCA ACG ACC CCT CGT CAC AAT AAA GAT GAC GCT CCT AGC TTC ACT GAC TCA CTC TAC T Bc2 CAG TAG AGT GAG TCA GTG AAG TCA GTG CTA GGA GCG TCA WBc2 GTG AGT CAG TGA AGC TAG GAG CGT CAT CTT TA GCA ACG ACC CCT CGT CAC AAT AAA GAA GGG CAA CTC TCG CAT TTC AGC TCA AAT ATT CBa1' TG ATC AAA TAT TTG AGC TGA AAT GCC AAA GGA GAG TTG CCC T Ba1' TTT GAG CTG AAA TGC GAG AGT TGC CCT TCT TTA WBa1' CBa2 GCA ACG ACC CCT CGT CAC AAT AAA GAG ACA TTG TCA GTT TTT TAA AGG AGA GTT GCC C Ba2 GGG CAA CTC TCC TTT AAA AAG GAG AGA CTG ACA ATG TC WBa2 TCT CCT TTA AAA AAC TGA CAA TGT CTC TTT A OR2 COR2 GCA ACG ACC CCT CGT CAC AAT AAA GAT TTA GGA GGA GTG GTA GGT GAT AGA TTG AG ATC ACC TAC CAC TCC TCC TAA ATC TTT A WOR2 MBBa1* ROX-GCA ACT CAA AAA AGG AGA GTT GC-BHQ2 Sa CAG GTG ACA TTG TCA GTC TCT CCA CTA CCC AAG AAG GC GGG TGG GTA GTG GAG AGA CTG ACA A Fa CAG GTG ACA TTG TCA GTC TCT CCA CTA CCC AAG AAG G Oaa GGG TGG GTA GTG GAG AGA CTG ACA A Saa MBBc1 ROX-TCA GTG CTA GGA GGT GCA CTG A-BHQ2 HBc2 AGT GAA GTG GGC AGA GGG GTG GGT TGG GCT CTG ACT TCA CTG ACT CAC HF1 CCA AAG GAG AGT GGG TTG GGT TGG GAA CTC TCC TTT GGC HF2 CAA AGG ATA GAA GTT GCC CAA GGG TGG GTG GGT GGG CAA CTT CT HF3 TTT CAG CTC AAA TAT TTG ATT ATT CCC TTC CCT TCC CAA TAA CCC AAA TAT TTG AG Fo Dabcyl-AGG AGT GTT TTG GTA GGT So CAA AAC ACT CCT-6-FAM CAG GTG ACA dT(BHQ2)TG TCA GTC TCT CCA CTA CCC AAG AAG GC L-Sa L-Fa GGG TGG GTA GTG GAG AGA CTG ACA A-ROX A3-3 TGCTCTTTCTGCATTCCTGG A3-5 TGGTGTTTCTGCATTCCTGG A3-8 TGGTCTTACTGCATTCCTGG A3-10 TGGTCTTTCTCCATTCCTGG

2. The scheme and PAGE results of the one-to-one translation.



Figure S1. The *A2*-to-*Ba1** translation. (A) The diagram of *A2*-to-*Ba1** in translation module. (B) The abstract description and scheme of *A2*-to-*Ba1** translation by cascading two-step DNA strand displacement reaction. (C) The PAGE results of the translation of *A2*-to-*Ba1**.



Figure S2. The *A2*-to-*Bf2* translation. (A) The diagram of *A2*-to-*Bf2* in translation module. (B) The abstract description and scheme of *A2*-to-*Bf2* translation by cascading two-step DNA strand displacement reaction. (C) The PAGE results of the translation of *A2*-to-*Bf2*.



Figure S3. The *A4*-to-*Ba1*' translation. (A) The diagram of *A4*-to-*Ba1*' in translation module. (B) The abstract description and scheme of *A4*-to-*Ba1*' translation by cascading two-step DNA strand displacement reaction. (C) The PAGE results of the translation of *A4*-to-*Ba1*'.



Figure S4. The *A4*-to-*Ba2* translation. (A) The diagram of *A4*-to-*Ba1'* in translation module. (B) The abstract description and scheme of *A4*-to-*Ba2* translation by cascading two-step DNA strand displacement reaction. (C) The PAGE results of the translation of *A4*-to-*Ba2*.



Figure S5. The *A3*-to-*B01* translation. (A) The diagram of *A3*-to-*B01* in translation module. (B) The abstract description and scheme of *A3*-to-*B01* translation by cascading two-step DNA strand displacement reaction. (C) The PAGE results of the translation of *A3*-to-*B01*.



Figure S6. The *A4*-to-*Bo2* translation. (A) The diagram of *A4*-to-*Bo2* in translation module. (B) The abstract description and scheme of *A4*-to-*Bo2* translation by cascading two-step DNA strand displacement reaction. (C) The PAGE results of the translation of *A4*-to-*Bo2*.

3. The scheme of one-to-many and many-to-one translation.



Figure S7. The scheme of one-to-many. (A) The diagram of *A3*-to-*Ba1/Bc1/Bf1* in translation module. (B) The abstract description and scheme of *A4*-to-*Ba1* translation. (C) The abstract description and scheme of *A4*-to-*Bc1* translation. (D) The abstract description and scheme of *A4*-to-*Bf1* translation.











Figure S8. The scheme of many-to-one. (A) The diagram of A1/A2/A4-to-Bc2 in translation module. (B) The abstract description and scheme of A1-to-Bc2 translation. (C) The abstract description and scheme of A2-to-Bc2 translation. (D) The abstract description and scheme of A4-to-Bc2 translation.

4. AND gate.



Figure S9. The original fluorescence spectrum and PAGE results of the AND gate, and the G-quadruplex fluorescent probe Thioflavin T (ThT) was employed to generate output signals.

5. The YES gate.



Figure S10. The YES gate. (A) The abstract diagram of YES gate. (B) The logic circuit and signal generation mechanism of YES gate. (C) Schematic and normalized fluorescence results of YES gate.

6. The Comparator gate.



Figure S11. The Comparator gate. (A) The abstract diagram of the Comparator gate. (B) The logic circuit and signal generation mechanism of the Comparator gate. (C) Schematic and normalized fluorescence results of the Comparator gate.



Figure S12. The original fluorescence spectrum (A) and PAGE results (B) of the Comparator gate.

7. The OR gate.



Figure S13. The OR gate. (A) The abstract diagram of the OR gate. (B) The logic circuit and signal generation mechanism of the OR gate. (C) Schematic and normalized fluorescence results of the OR gate.



Figure S14. The original fluorescence spectrum and PAGE results of the OR gate.

8. The Feynman gate.



Figure S15. The Feynman gate. (A) The abstract diagram of the Feynman gate. (B) The logic circuit and signal generation mechanism of the Feynman gate. (C) Schematic and normalized fluorescence results of the Feynman gate.



Figure S16. The original fluorescence spectrum (A) and PAGE results (B) of the Feynman gate.

9. The PAGE results and fluorescence spectrum of AND-AND gate.



Figure S17. The PAGE results (A) and original fluorescence spectrum (B) of the AND-AND gate.

10. The PAGE results and fluorescence spectrum of AND-AND-NOT gate.



Figure S18. The PAGE results (A) and original fluorescence spectrum (B) of the AND-AND-NOT gate.

11. The original fluorescence spectrum of cascaded AND and OR gate.



Figure S19. The original fluorescence spectrum of cascaded AND (A) and OR gate (B).

12. The influence of the translation process upon adding different targets.



Figure 20 The PAGE result of the parallel translation from A2 to B1d and A3 to B2d (A); lane 1: B2d, lane 2: A2+A2*, lane 3: A3+A3*, lane 4: CB2d+WB2d, lane 5: CB2d+B2d, lane 6: CB1d+WB1d, lane 7: CB1d+B1d, lane 8: A2*+CB1d+B1d+A3*+CB2d+B2d, lane 9: A2*+CB1d+B1d+A3*+CB2d+B2d+A2+A3. And the parallel translation from A3 to B01 and A4 to B02 (B); lane 1: A3+A3*, lane 2: A4+A4*, lane 3: CBo2+WBo2, lane 4: CBo1+WBo1, lane 5: Bo1, lane 6: Bo2, lane 7: A3*+CBo1+Bo1+A4*+CBo2+Bo2+WBo1+WBo2, lane 8: A3*+CBo1+Bo1+A4*+CBo2+Bo2+WBo1+WBo2+A3, lane 9: A3*+CBo1+Bo1+A4*+CBo2+Bo2+WBo1+WBo2+A4; lane 10: A3*+CBo1+Bo1+A4*+CBo2+Bo2+WBo1+WBo2+A3+A4. (C) The influence of single mutation at different location on the translation process; lane 1: A3*+B2c+CB2c+WB2c+A3-3, lane 2: A3*+B2c+CB2c+WB2c+A3-5, lane 3: A3*+B2c+CB2c+WB2c+A3-8, lane 4: A3*+B2c+CB2c+WB2c+A3-10. X of A3-x represents the mutation location. (D) The comparison of fluorescence results individual AND and OR (blue green) and AND-OR cascade in parallel (orange).