

Programmable and scalable assembly of a flexible hexagonal DNA origami

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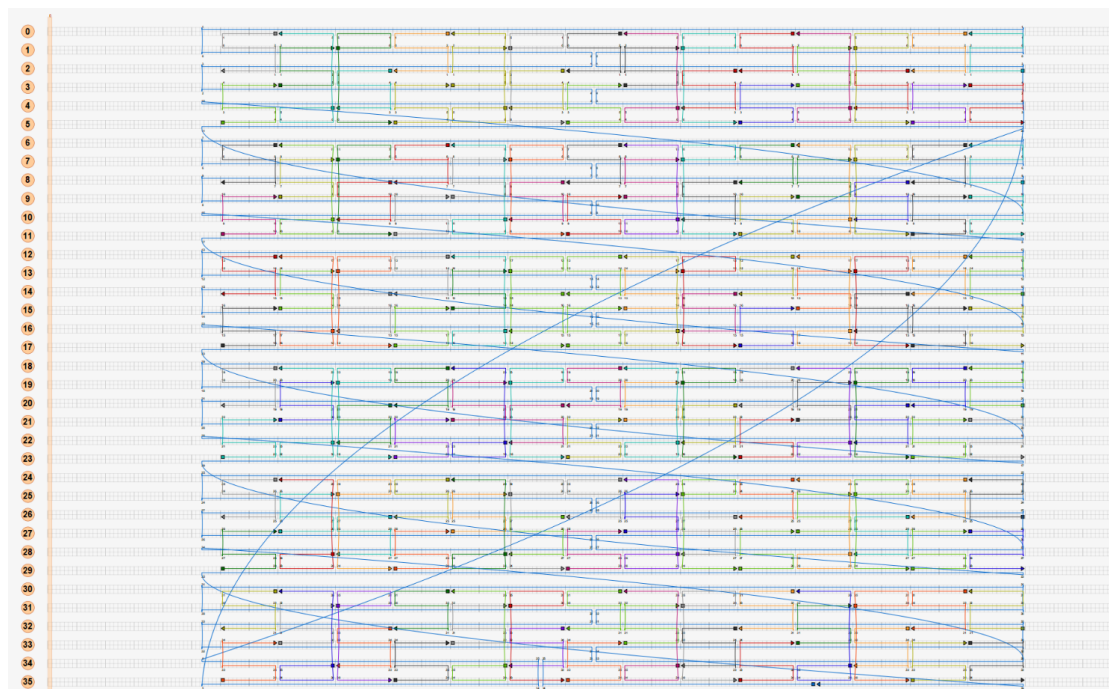


Figure 1. The caDNAno design figure of hexagonal origami.

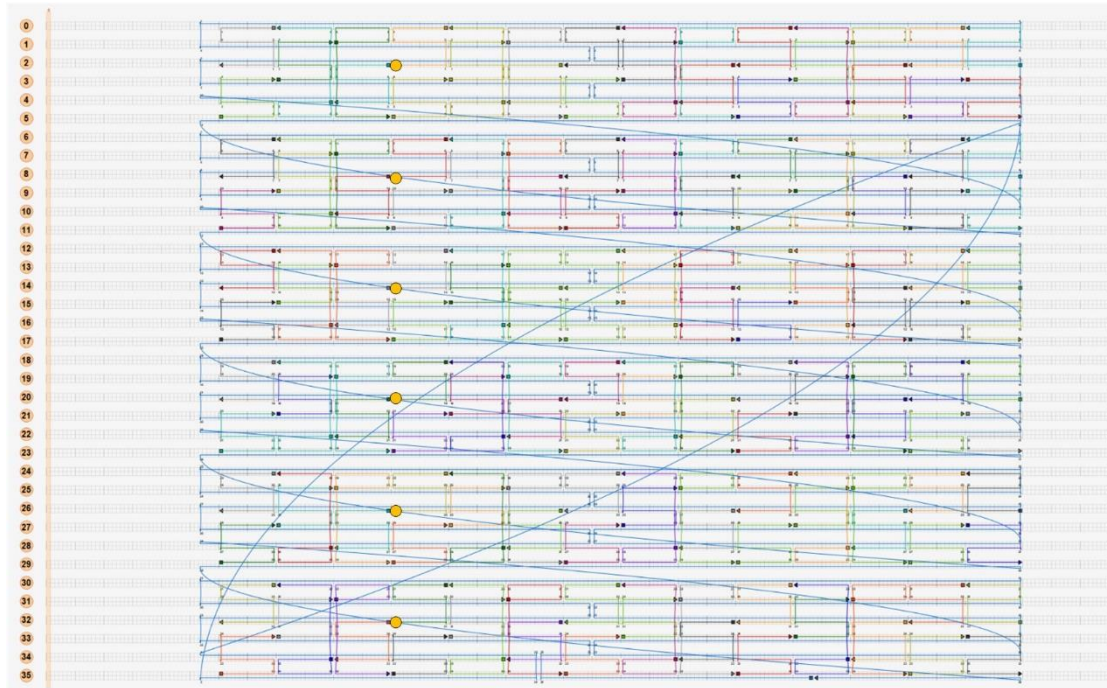


Figure 2. The yellow dots are the locations of biotin modified staple strands. They are illustrated in Table1.

Table1 Biotin modified strands.

| Location | Sequence |
|--------------|---|
| 6[97]8[84] | AATAGCAATAGCTATGGCATGATTAAGAGCAAAGACACCACGTTTTT |
| 0[97]2[84] | TGCTGATGCAAATCTAAATAAGGCGTTACTTACCAGTATAAATTTTT |
| 18[97]20[84] | ATCAACGTAACAAATTACCTTATGCGATGGAACAACATTATTTTTTT |
| 24[97]26[84] | AGCATAAAGCTAAAATGCAATGCCTGAGCAACCGTTCTAGCTTTTTT |
| 12[97]14[84] | GATAAGTGCCGTCGCAGAGCCACCACCCCTACAACGCCTGTATTTTT |
| 30[97]32[84] | ACACAACATACGAGGTGCCAGCTGCATTGTGAGACGGGCAACTTTTT |

We extended four poly-T at the 3' end for biotin modification.

Purified Results

The concentration of purification samples via gel extraction is relatively low. All the view fields of supplemental AFM images were 2 μ m. Because biotin-streptavidin dots are not recognizable at a higher view, and we can not get enough samples at a lower view.

We marked all the potential biotin-streptavidin dots via green circles. The binding efficiency of streptavidin attachment, origami shape-changing are statistically calculated at the end of this section.

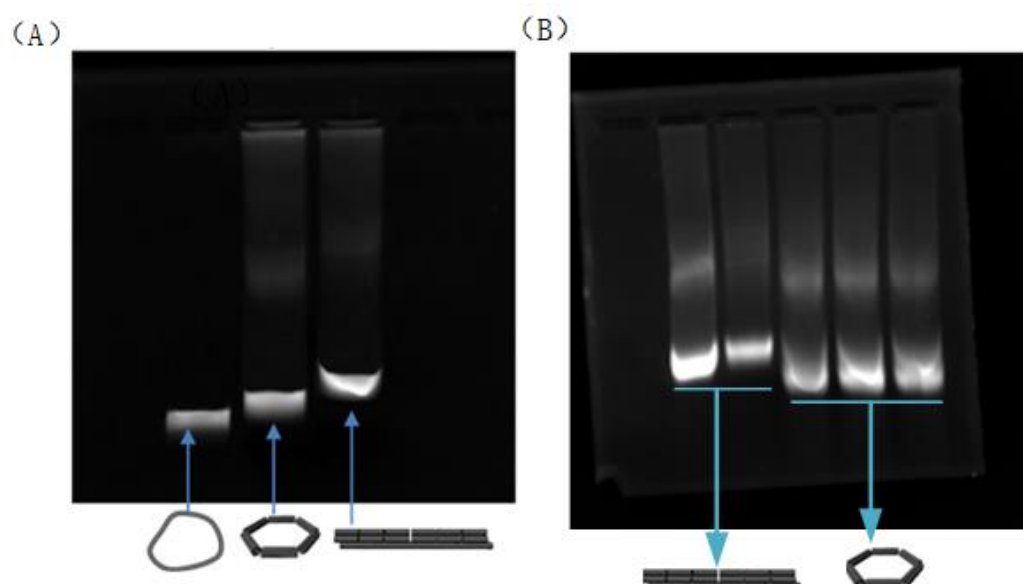


Figure 3. The Gel electrophoresis results. (A) The three lanes from left to right: M13; hexagonal origami; linear origami. (B) The two on the left are the linear origami. The three on the right are the hexagonal origami.

The molecular weight of linear origami is the same as that of hexagonal origami, but the linear origami has a larger morphological expansion and occupies a wider range of horizontal space. Therefore, it moves slower than the hexagonal structure.

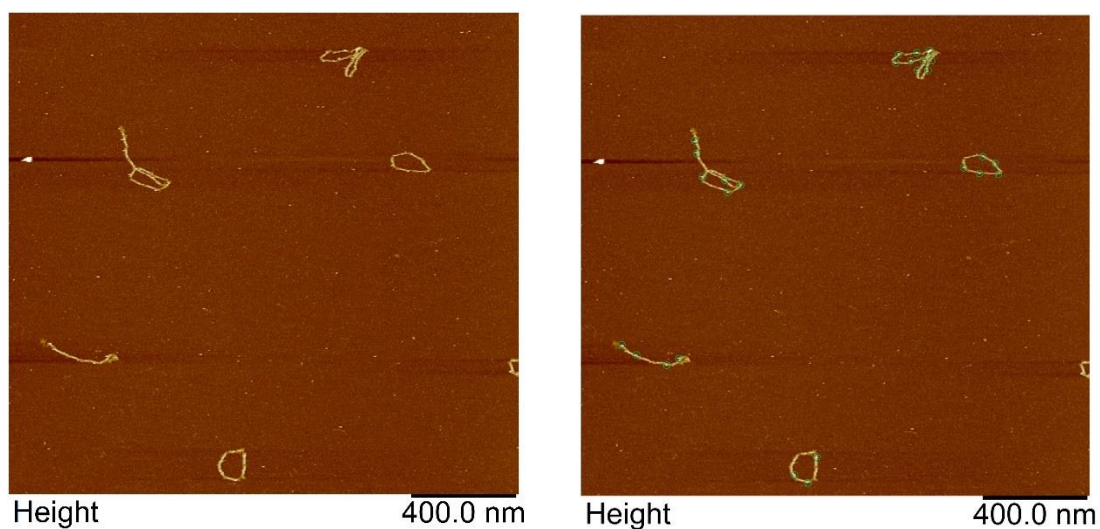


Figure 4. The hexagonal origami with gel purified results. There are seven origamis, two of them are shape changed. The possible biotin-streptavidins are marked with green dots. We counted all the possible dots on the seven origamis, manually.

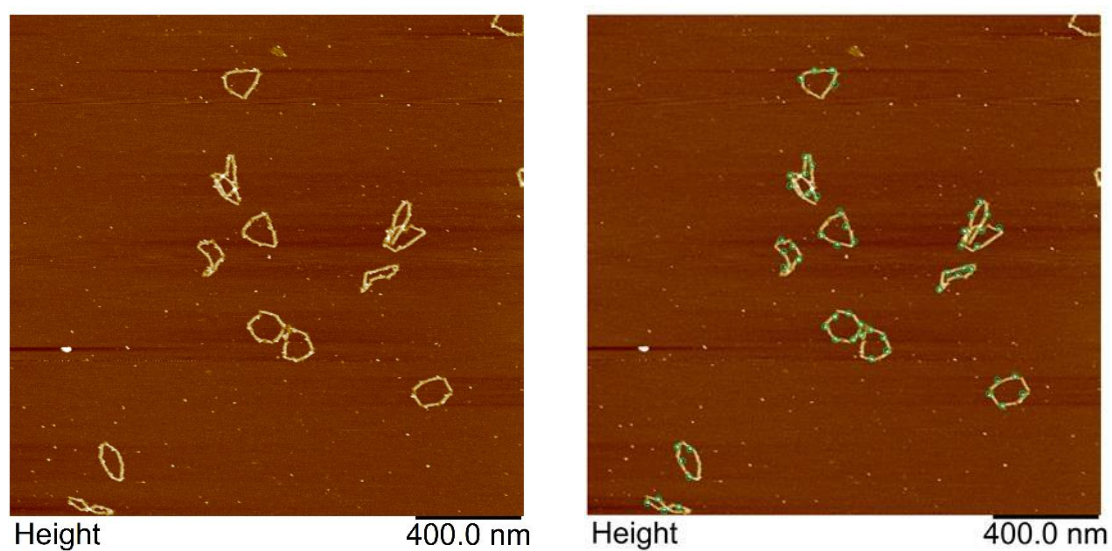
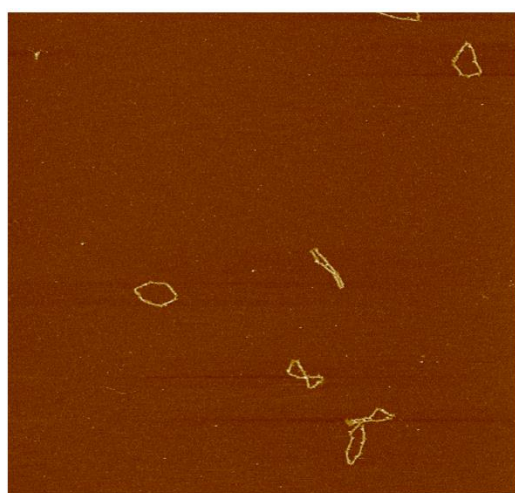
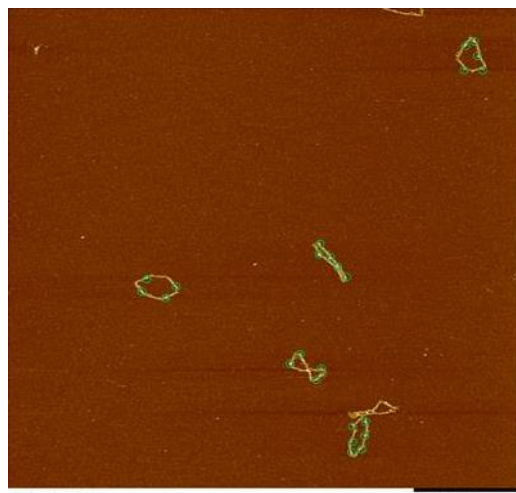


Figure 5. Purified results. There are 13 origamis, and possible biotin-streptavidins were marked with green dots.

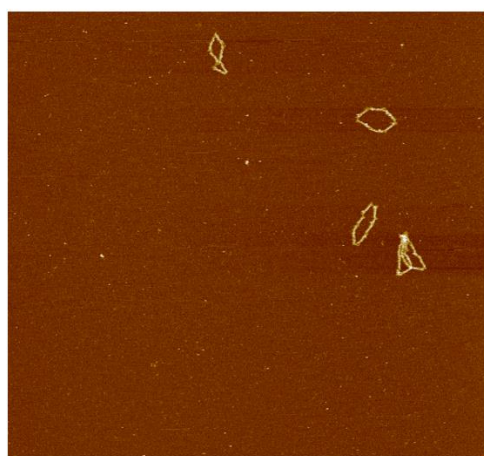


Height 400.0 nm

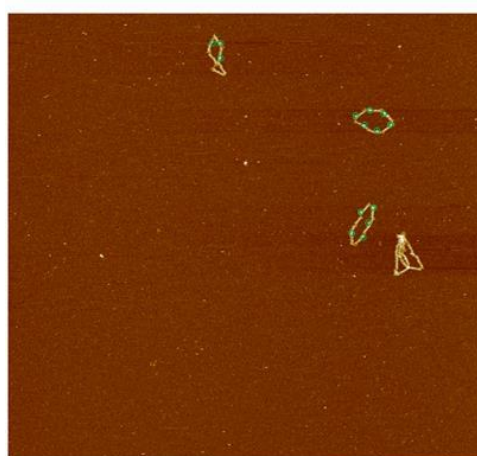


Height 400.0 nm

Figure 6. Purified results. There are six origamis, possible biotin-streptavidins are marked with green dots.

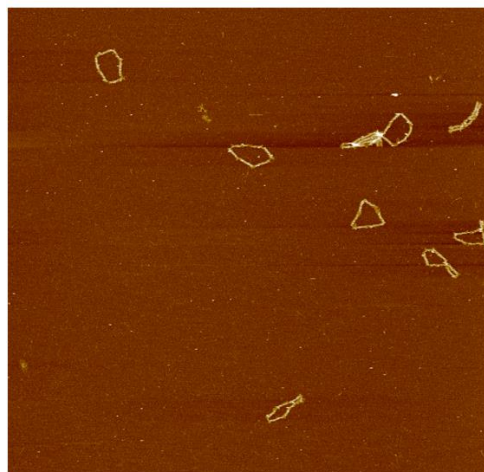


Height 400.0 nm

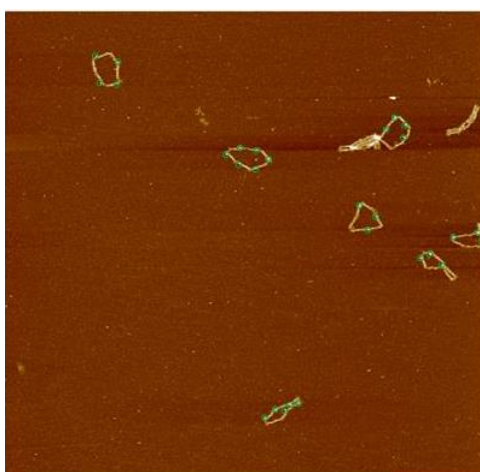


Height 400.0 nm

Figure 7. Purified results. There are five origamis, and we marked three of them.



Height 400.0 nm



Height 400.0 nm

Figure 8. Purified results. There are ten origamis, we marked seven of them.

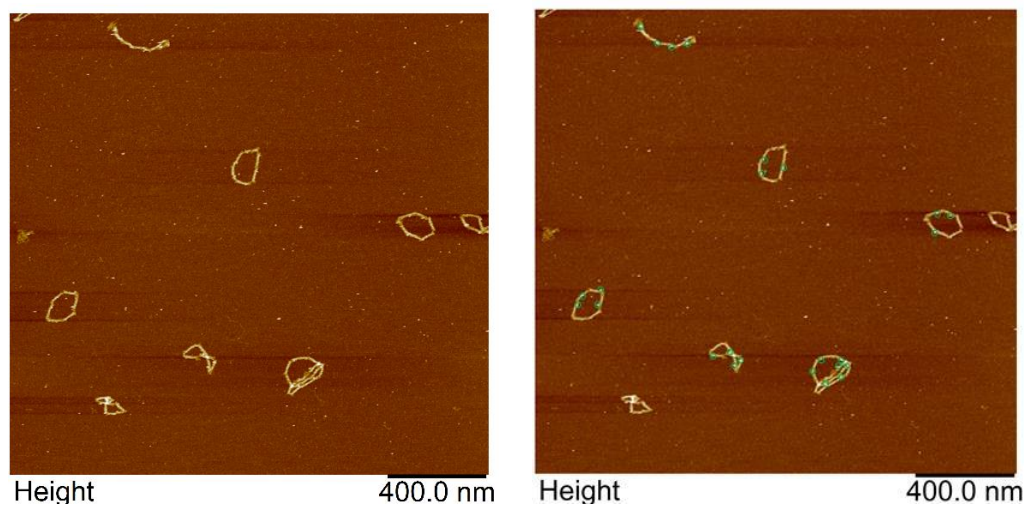


Figure 9. Purified results. There are nine origamis, one of them is shape changed.

The samples had attached labels in previous step. We suspect the labels could fall off from the structures. To identify the linking efficiency of biotin-streptavidin labels, we added excess streptavidin solution to the purified samples. The following results are the gel purified results with extra streptavidin solution.

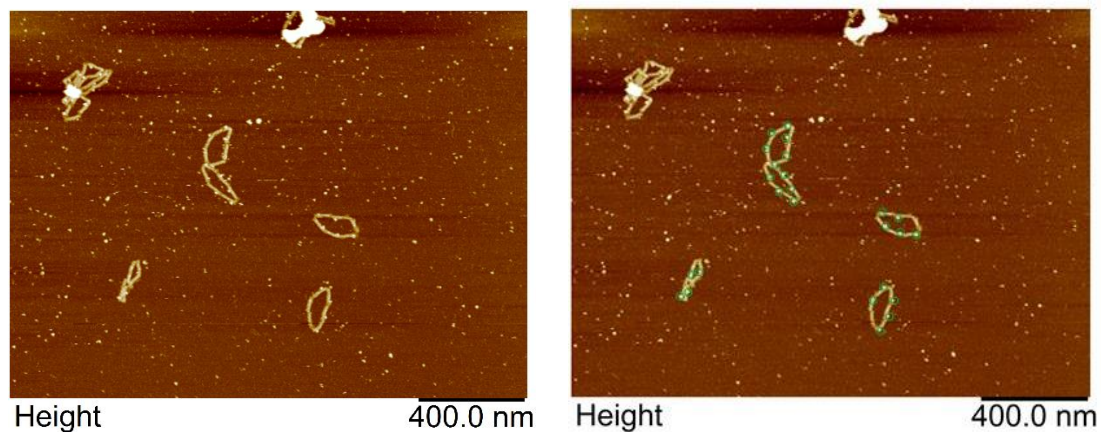


Figure 10. Purified results with extra streptavidin. Some origamis stacked together. We suspect they were stacked through the blunt end of these edges. These high white areas are the stacking positions.

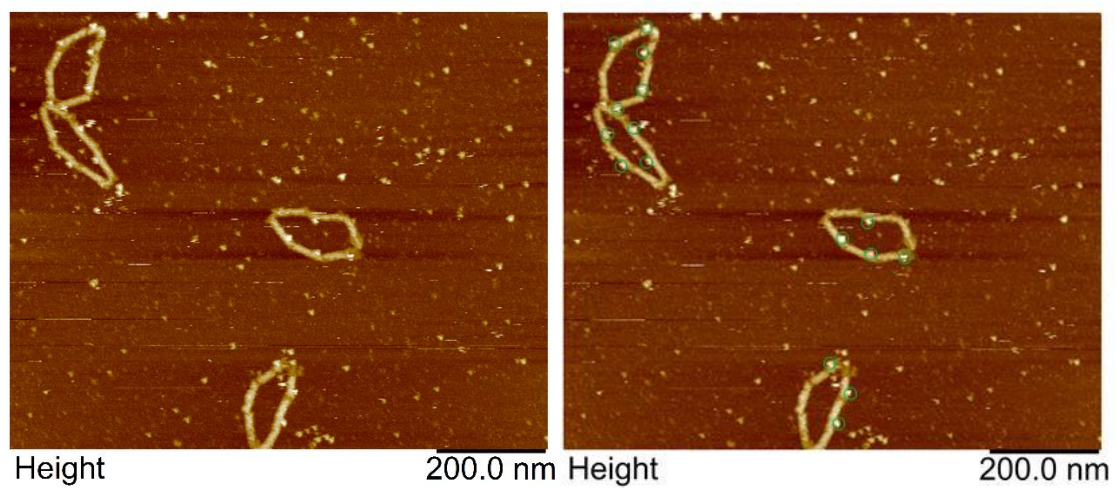


Figure 11. Purified results with excess streptavidin.

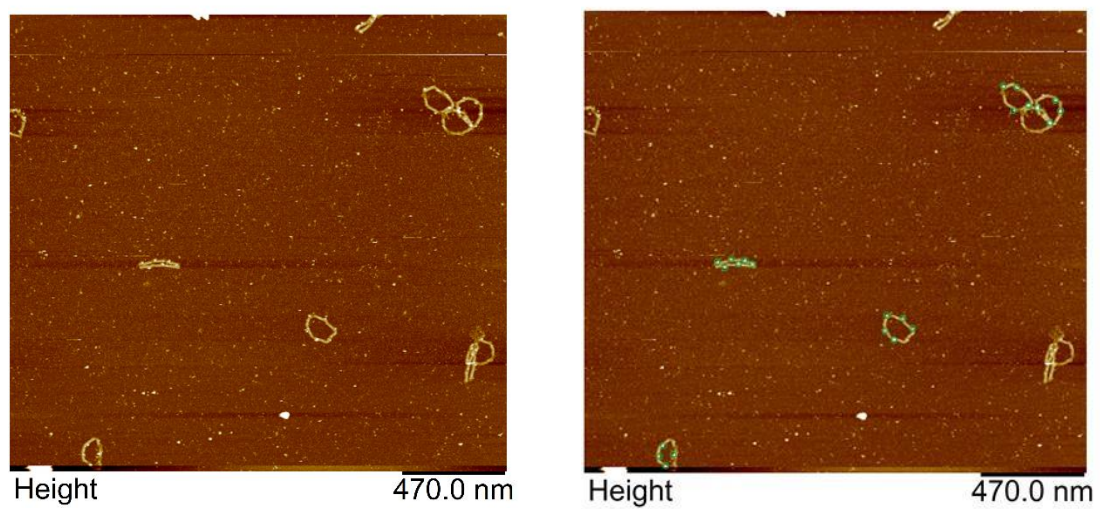


Figure 12. Purified results with excess streptavidin. There are nine origamis, one of them is shape changes. We marked 5 of them.

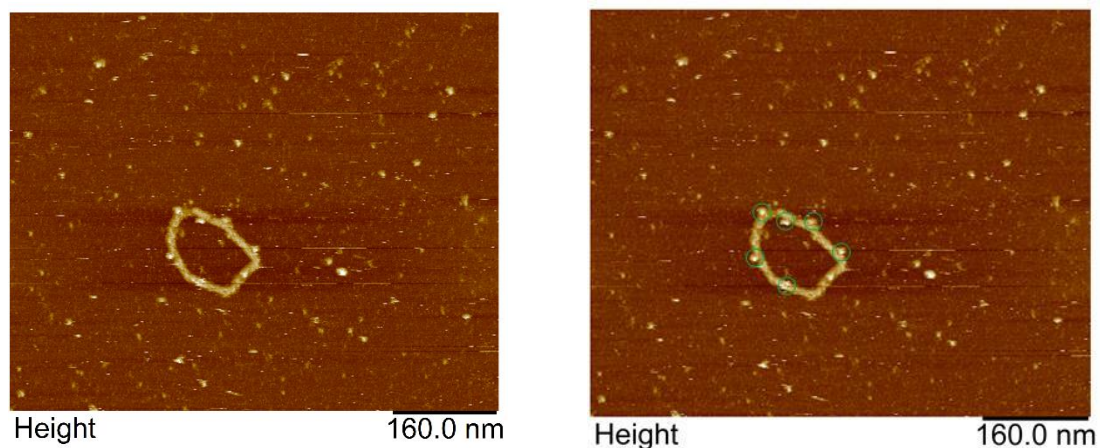


Figure 13. A typical hexagonal origami with biotin-streptavidin labels.

Yield statistics:

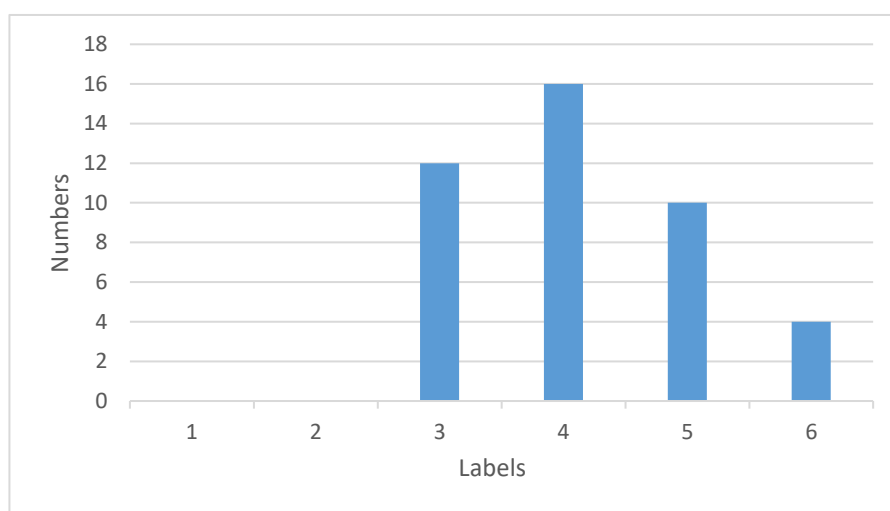


Figure 14. The statistical analysis of purified samples of biotin-streptavidin linking efficiency.

From the above Figures 3 to 8, we marked 42 origamis. There are four hexagonal origamis with six labels; Ten origamis with five labels; 16 origamis with four labels, 12 origamis with three labels. The total linking efficiency is approximately 70%.

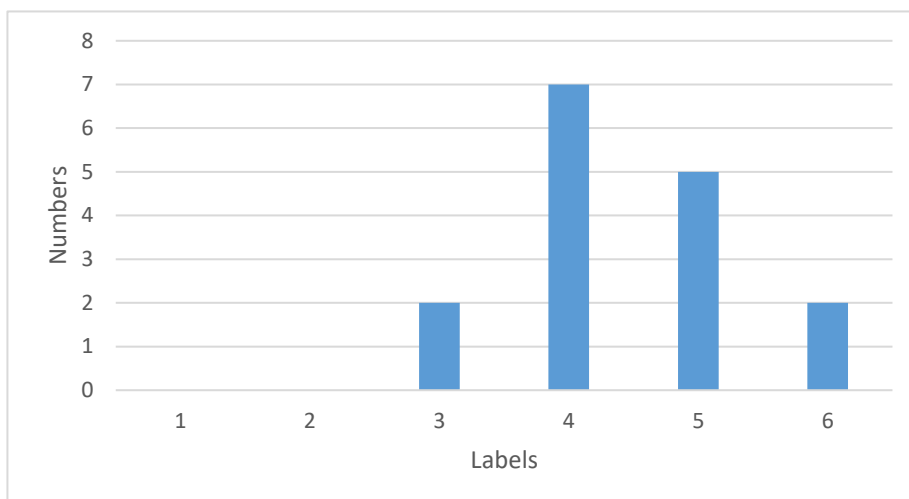


Figure 15. The statistical analysis of purified samples with extra streptavidin.

We marked 15 origamis in Figures 9 to 12. There are two hexagonal origamis with six labels; Five origamis with five labels; Seven origamis with four labels; One origami with Three labels. The total linking efficiency is approximately 75%.

Three defect structures were in linear shape. Some of the structures were randomly twisted together. The total number of the above-scanned original is 49; three among them are defects. Therefore, the ratio is 6%.

In conclusion, the linking efficiency is around 70%. There could be a slight deviation, as some labels are misidentified. The statistical results are reliable, because of the greater number of purification results; more images are not provided in this file, as limited by the length of supplementary files.

Lines with statistical analysis:

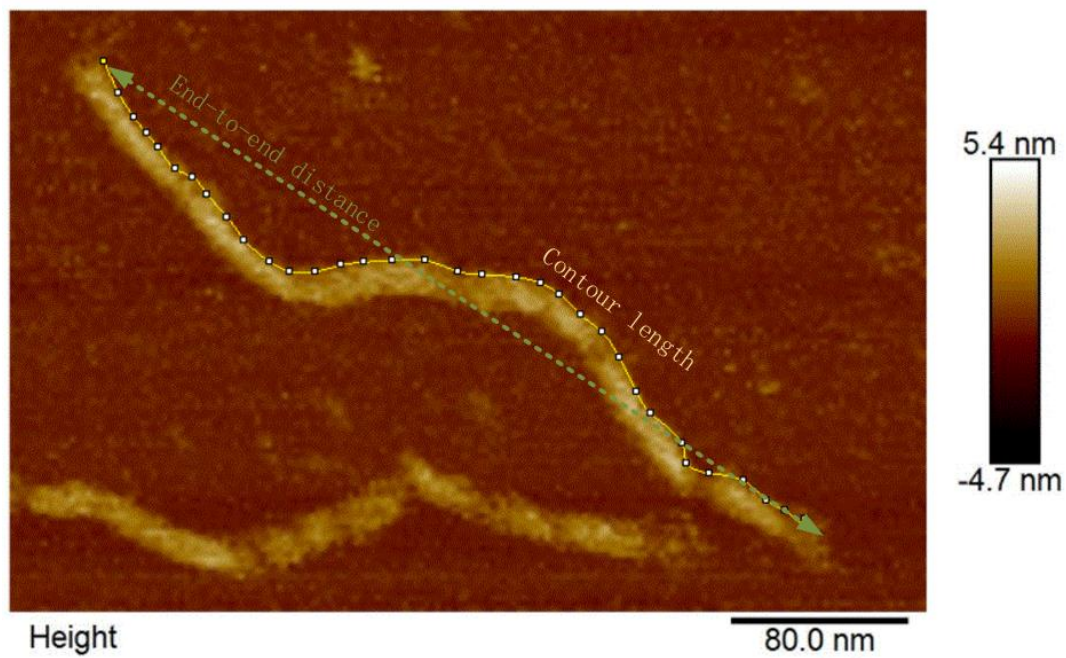


Figure 16. Analysis of a linear origami. The contour length and end-to-end distance were measured via ImageJ software. The contour length was 428.4nm, the end-to-end distance was 392nm.

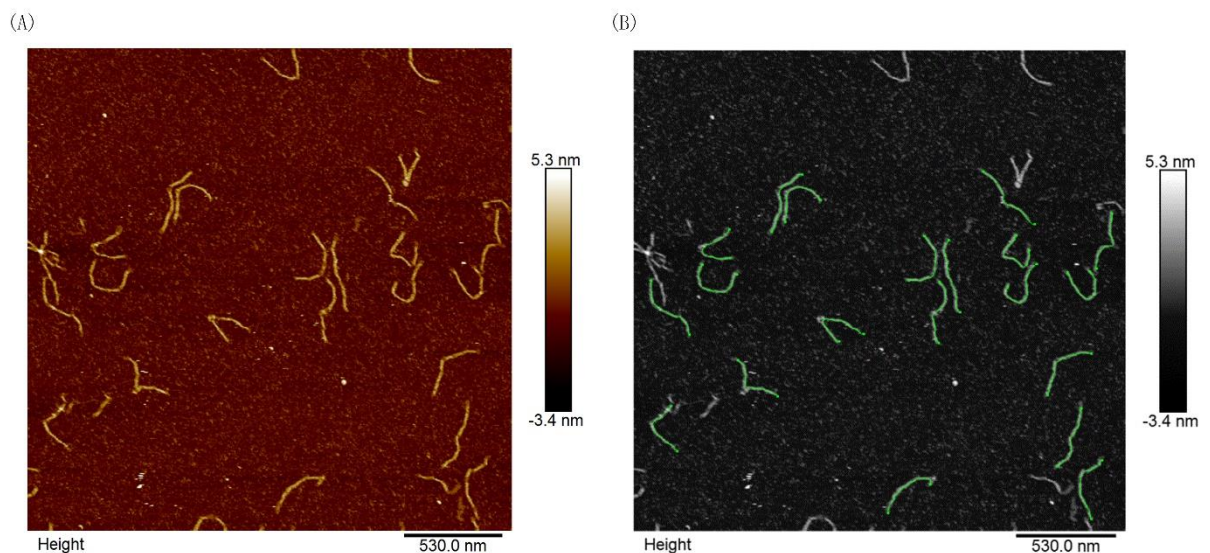


Figure 17. The linear origami analysis. (A) Original AFM image. (B) The green lines are the contour lines of selected origamis. The length of these green lines was calculated via ImageJ software.

We selected 20 origamis in figure 17(B) and measured their contour length and end-to-end distance via software. The results are illustrated in Table 2.

Table2 The contour length and end-to-end distance measurements.

| PathID | PrimaryPath | PathLength | End-to-End Length | LengthUnits | Contour Length | End-to-End Length | LengthUnits |
|--------|-------------|------------|-------------------|-------------|----------------|-------------------|-------------|
| 1 | TRUE | 2.8004 | 1.65645 | inches | 380.5672 | 225.10731 | nm |
| 2 | TRUE | 3.0081 | 2.68397941 | inches | 408.7931 | 364.74592 | nm |
| 3 | TRUE | 2.7632 | 1.12029128 | inches | 375.5118 | 152.24471 | nm |
| 4 | TRUE | 2.8935 | 1.39750329 | inches | 393.2192 | 189.91711 | nm |
| 5 | TRUE | 2.9944 | 1.07731306 | inches | 406.9313 | 146.40408 | nm |
| 6 | TRUE | 2.6272 | 2.20964671 | inches | 357.0297 | 300.28532 | nm |
| 7 | TRUE | 3.0999 | 2.60927957 | inches | 421.2685 | 354.5944 | nm |
| 8 | TRUE | 2.8618 | 2.56097834 | inches | 388.9113 | 348.03039 | nm |
| 9 | TRUE | 2.8355 | 1.89786004 | inches | 385.3372 | 257.91431 | nm |
| 10 | TRUE | 3.2439 | 2.91417492 | inches | 440.8377 | 396.0289 | nm |
| 11 | TRUE | 2.6998 | 2.19328547 | inches | 366.8959 | 298.06187 | nm |
| 12 | TRUE | 2.9839 | 2.37717019 | inches | 405.5044 | 323.05133 | nm |
| 13 | TRUE | 2.9481 | 1.44979062 | inches | 400.6392 | 197.02283 | nm |
| 14 | TRUE | 2.7418 | 2.45469609 | inches | 372.6036 | 333.5869 | nm |
| 15 | TRUE | 2.8046 | 0.84117055 | inches | 381.1379 | 114.31292 | nm |
| 16 | TRUE | 3.247 | 2.99891247 | inches | 441.259 | 407.54452 | nm |
| 17 | TRUE | 2.6503 | 1.81154803 | inches | 360.169 | 246.18473 | nm |
| 18 | TRUE | 2.7246 | 0.76439767 | inches | 370.2662 | 103.87968 | nm |
| 19 | TRUE | 3.055 | 2.50664455 | inches | 415.1667 | 340.64657 | nm |
| 20 | TRUE | 2.7092 | 2.34837547 | inches | 368.1733 | 319.13821 | nm |

We measured the selected 20 origamis. The mean square end-to-end distance of these linear origamis is 81826.74. The 2 dimensions equilibrium, formulation (1) was adopted to calculate the persistence length.[1]

$$\langle R^2 \rangle_{2D} = 4L_p L \left(1 - \frac{2L_p}{L} \left(1 - e^{\frac{-L}{2L_p}}\right)\right) \quad (1)$$

The $\langle R^2 \rangle_{2D}$ is mean-square end-to-end distance that obtained from table2, L is the contour length, L_p is the persistence length.

We brought these twenty values of L and the constant value $\langle R^2 \rangle_{2D}$ into formula (1) and calculated 20 times via MATLAB, respectively. And we got twenty

results. The average value of the twenty results (L_p) is 100, standard deviation is 18. Therefore, the persistence length is $100 \pm 18\text{nm}$.

Annealing Process:

1. Folding the origami structures

95°C to 85°C: -0.1°C Every 8 sec
85°C to 65°C: -0.1°C Every 30 sec
65°C to 45°C: -0.1°C Every 40 sec
45°C to 25°C: -0.1°C Every 2 min
25°C to 4°C: -0.1°C Every 15 sec

2. Linking biotin strands

45°C to 25°C: -1°C Per min
25°C to 4°C: -0.1°C Every 8 sec

Reference:

- [1] Mantelli S, Muller P, Harlepp S and Maaloum M 2011 Conformational analysis and estimation of the persistence length of DNA using atomic force microscopy in solution Soft Matter 7 3412

The DNA sequences:

The hexagonal origami:

| | |
|----------------|--|
| 35[210]33[223] | ATCACCCAAATCAATAACCAATAGGAACCCGTGCATCTGCCA |
| 20[125]22[112] | AAAATCTACGTTAAGATAAAAACCAAAATGCAAAAGAAGTTT |
| 11[42]9[55] | AGGGACATTCTGGCAAACCATCGATAGCACAAAAGGGCGACA |
| 17[126]15[139] | CAGCCATTGCAACATTGCCATAATTGCGCTAAAGGCTTTTCA |
| 7[154]11[167] | CCATATTATTTATCGAGCGCTAATATCACCTAAAACATCGCC |

23[84]21[97] TTAGAATCAGAGCGATAGTAAATGTTTCTATCATAACCCTC
 31[196]35[209] CTGCAAGGCGATTACAAGCTTGCATGCCCCACTACGTGAACC
 25[70]29[83] CTCATATATTTAATCGGTTGTACCAAACTAAAGGGAGCCCC
 30[55]32[42] TGCCTAATGAGTGATGCGTTGCGCTCACGAGAGAGTTGCAGC
 18[181]20[168] ATGAACGGTGTACAGAGATTTGTATCATTAAACGAAAGAGG
 13[70]17[83] AGAACCGCCACCCTAGAGGGTTGATATAAGTAGAAGAACTCA
 14[209]16[196] CCAGCATTGACAGGACCCTCAGAACCGCAGTTTGCCTTTAGC
 4[111]0[98] GAACGCGCCTGTTTCTGAACCTCAAATATACTATATGTAAA
 16[69]12[56] TAATTGTATCGGTTACATCACTTGCCTGAGTATAGCCCGGAA
 5[126]3[139] AATATCTGGTCAGTCGTAAAAACAACATACAATAACAGAAAT
 25[196]29[209] AGTAGATTTAGTTTGGGGCGCGAGCTGACCACCACACCCGCC
 1[112]5[125] CCGGAATCATAATTTAGGTTGGGTTATATCAAACCCTCAATC
 23[168]21[181] TTATAATCAGTGAGCCATCGCCACGCACAGCGAAAGACAGC
 27[140]25[153] AAAAGATTAAGAGGCTCAACATGTTTGTAGTCGGTGTCTGGA
 14[83]16[70] GCATTCCACAGACAGGATTTTGCTAAACCCAAAAGAGCCTT
 31[70]35[83] TCGGGAACCTGTCCCGGAAGCATAAAGAAACAATTCGACAA
 27[56]25[69] CTGGAGCAAACAAGGGAGAGGGTAGCTAAAATTTTTAGAACC
 35[84]33[97] CTCGTATTAAATCCATTTTGCGGAACAAAATCGGCAAAATCC
 24[181]26[168] TTCTACTAATAGTATAACAGTTGATTCCAGCTTAATTGCTGA
 4[153]0[140] CAAAATTATTTGCATGGCAAATCAACAGTGAGAGACTACCTT
 29[126]27[139] AAGGAAGGGAAGAAAGCGGATGAAAAGCATAATCATGCATCA
 34[153]30[140] ATGTGAGCGAGTAATTAAAGAACGTGGATCGTAATCATGGTC
 18[55]20[42] CTGACGAGAAACACTTGGGCTTGAGATGTTAGGAATACCACA
 13[154]17[167] GAATTTACCGTTCCAAAGTATTAAGAGGGGAAATACCTACAT
 4[69]0[56] AAAATAATATCCCACTGAGAGCCAGCAGAGAACGCGAGAAAA
 2[125]4[112] GTTTAGTATCATATCTGTCCAGACGACGGTTCAGCTAATGCA
 31[154]35[167] GGGCTCTTCGCTATACCGAGCTCGAATCTCCAACGTCAAAG
 16[153]12[140] CCCCTTATTAGCGTGGA AAAACGCTCATCTGAGACTCCTCAA
 22[111]18[98] TGCCAGAGGGGTAGGAGCTAACAGGATATTCATTACCCAA
 0[181]2[168] AATAGTGAATTTATAACAGTACATAAATGCAAAAGAAGATGA
 2[167]4[154] TGAAACAAACATCAGATTTTCAGGTTTAAGAACCTACCATAT
 5[42]3[55] GCAACAGTGCCACGTCTAATTTACGAGTAATTTAGGCAGAG
 30[139]32[126] ATAGCTGTTTCCTGCGGTTGGTTGGGACGCAACTCGTATTG
 35[168]33[181] GGCGAAAAACCGTCTCCTGTAGCCAGCTGTAATGGGATAGGT
 26[209]28[196] TAAGAGGTCATTTTAAAGCGAACCAGACAGTTCAGAAAACGA
 26[83]28[70] GATAAATTAATGCCAGAATCGATGAACGAACGTTAATATTTT
 22[195]18[182] GACAATGACAACAAGCCACCGAGTAAATTGAAAGAGGACAG
 2[83]4[70] GCCAACGCTCAACAGTAATAAGAGAATAAGTCTGAACAAGA
 0[223]2[210] GAAAACATAGCGATCTGTAAATCGTCGCACAAAATCGCGCAG
 35[42]33[55] TTGAGGATTTAGAAATCATATTCTGATTTTGGCCAGCAGG
 10[111]6[98] CAGCAAAATCACCAGTGGCACAGACAATGCAAGAAACAATGA
 16[195]12[182] GTCAGACTGTAGCGGTCTGAAATGGATTATTTGGAACCTAT
 0[55]2[42] CTTTTCAAATATACCTCTGACCTAAATAGAATCGCCATATT
 26[237]24[224] AGAGAGTACCTTTATTTGCAAAATGGTCAATAACCTGTTTAG

32[167]34[154] GGCAAAGCGCCATTACGGCGGATTGACCTTCATCAACATTAA
34[195]30[182] TTCGCGTCTGGCCTTATCAGGCGATGGCTGCAGGTCGACTCT
33[98]31[111] CTTATAAATCAAAATTTTCTTTTACCAAATGAATCGGCCAA
12[139]14[126] GAGAAGGATTAGGAGAACCCAAGAATGGTAAAGCCTGTACCG
21[56]19[69] GGAATTACGAGGCATCATCAGTTGAGATGTTTAATTTCAACT
19[196]23[209] TGTCGAAATCCGCGCGAACTGACCAACTGAGTCTGTCCATCA
15[56]13[69] GAATTTTCTGTATGGCCCTCATAGTTAGGAACCGCCACCCTC
21[140]19[153] GCCGCTTTTGCGGGAAAACACTCATCTTTATACCAAGCGCGA
9[56]7[69] TTCAACCGATTGAGTGTCACAATCAATAGCAATAATAACGGA
3[182]1[195] TACAGTAACAGTACTTTCAATTACCTGACAATATATGTGAGT
8[167]10[154] ACCAACGCTAACGAGCCCAATAGCAAGCGCACTCATCGAGAA
1[70]5[83] TACCGACCGTGTGACAATCGCAAGACAACAAATGAAAAATCT
25[154]29[167] AGTTTCATTCCATAGTAGCATTAAACATCGCGCTAGGGCGCTG
19[154]23[167] AACAAAGTACAACGGACCAGGCGCATAGCCTGAGAAGTGTTT
27[182]25[195] TTAATTCGAGCTTCTGCGGATGGCTTAGCAATTCTGCGAACG
29[210]27[223] GCGCTTAATGCGCCCAATGCTTTAAACCGGAAGCAAACCTC
12[223]14[210] CCGTATAAACAGTTGGTAATAAGTTTTACACCAGAGCCGCCG
9[182]7[195] AGGCTTATCCGGTAACCCAGCTACAATTAACGATTTTTTGT
28[153]24[140] ATAGTCAGAAGCAAAGCGAAAGGAGCGGCAATAAATCATACA
6[223]8[210] AAGCGCATTAGACGGAATAGCAGCCTCTTAAATCAAGATT
19[70]23[83] TTAATCATTGTGAAGCTGCTCATTAGTACGTGCTTTCCTCG
32[209]34[196] CTCCAGCCAGCTTTTGGGCGCATCGTAAGCCATCAAAAATAA
20[167]22[154] CAAAAGAATACACTATCGTCACCCTCAGTAACCGATATATTC
15[98]13[111] TCAGCGGAGTGAGAGTACCAGTACAAATCATTTTCAGGGAT
32[83]34[70] AGCTGATTGCCCTTGATGGTGGTTCCGAAGAAACCACCAGAA
34[111]30[98] TGAGTAACATTATCTTTGCCCGAACGTTCCGCTCACAATTCC
21[182]19[195] ATCGGAACGAGGGTCTGAAGGCACCAACCCGCTGATAAATTG
9[98]7[111] ATTCATTAAAGGTGCATATAAAAGAAACCTCCTTATTACGCA
15[140]13[153] TAATCAAAATCACCAAATAAATCCTCATAAAGCGCAGTCTCT
15[182]13[195] CCCTCAGAGCCGCCAGGTTGAGGCAGGTATGGCTTTTGATGA
17[168]15[181] TTTGACGCTCAATCCGTTTTTCATCGGCACACCGGAACCGCCT
7[196]11[209] TTAACGTCAAAAATGGAGAATTAAGTGAAGATAAAACAGA
5[210]3[223] AGATTAGAGCCGTCAATTCATCAATATAGAAACAATAACGGA
33[182]31[195] CACGTTGGTGTAGACCGGCACCGCTTCTGAAAGGGGGATGTG
7[70]11[83] ATACCCAAAAGAACTCTTACCGAAGCCCCCTTCTGACCTGA
23[126]21[139] ATTTTAGACAGGAAGCAGGGAAGGCTTTTAGCGAGGTTAAAG
25[112]29[125] AGATTCAAAAGGTTTAGCAAAATTAAGCGAACGTGGCGAGA
20[209]22[196] ACGTAATGCCACTAAGCAACGGCTACAGCCGATAGTTGCGCC
0[139]2[126] TTAACTCCGGCTACTAGAATTAACAAATTACATAAAGCCT
28[195]24[182] GAATGACCATAAATCACGCTGCGCGTAAAAAGGTGGCATCAA
13[112]17[125] AGCAAGCCCAATAGTTAGCGGGGTTTTGAACAATATTACCGC
20[83]22[70] ACAGGTAGAAAGATTAGTAAGAGCAACAAGACTGGATAGCGT
14[167]16[154] TGATATTCAAAACGGAACCGAGCCACTTTTCGGTCATAGC
24[139]26[126] GGCAAGGCAAAGAAGAGAAAGAACTAAAAATATGCGCCGGAG

23[210]21[223] CGCAAATTAACCGTTAAACAGCTTGATAAGGCTTTGAGGACT
15[224]17[237] CCACCCTCAGAGCCAGCACCGTAATCAGCGACCAGTAATAAA
17[210]15[223] ATTCACCAGTCACATAGCGACAGAATCACACCCTCAGAGCCA
27[98]25[111] AGCATGTCAATCATATCAATATGATATTTAATGTGTAGGTAA
11[210]9[223] GGTGAGGCGGTCAGTCAATAATCGGCTGGGCGTTTTAGCGAA
18[97]20[84] ATCAACGTAACAAATTACCTTATGCGATGGAACAACATTATT
26[167]28[154] ATATAATGCTGTAGAAGCCCGAAAGACTTTACCCTGACTATT
28[111]24[98] GATTGTATAAGCAAACGGGGAAGCCGGCAATAAAGCCTCAG
24[55]26[42] GTAATACTTTTGCGCAACGCAAGGATAATTTTTGAGAGATCT
1[196]5[209] GAATAACCTTGCTTAGCTTAGATTAAGACACTAACAACTAAT
11[168]9[181] ATTAATAATACCGATTAAACCAAGTACCAATCAGATATAGA
29[168]27[181] GCAAGTGTAGCGGTCAAAAATCAGGTCTTCAAATATCGCGTT
21[98]19[111] GTTTACCAGACGACTAAAACGAACTAATTTAAGAACTGGCT
20[237]18[224] GAGGAAGTTTCCATACTTAGCCGGAACGAGGCGCAGACGGTC
31[112]35[125] CGCGCGGGGAGAGGTGTGAAATTGTTATATTAATTTTTTGA
16[111]12[98] CACGTTGAAAATCTCTGGTAATATCCAGCTCAGTACCAGGCG
17[84]15[97] AACTATCGGCCTTGCCAAAAAAGGCTAACTTTCAACAGTT
12[55]14[42] TAGGTGTATCACCGTACCGCCACCCTCACGTAACGATCTAAA
18[139]20[126] ATCAAGAGTAATCTGGACGTTTACGCGATTGACCCCGGAAGA
35[126]33[139] ACAAGAGTCCACTACAACCCGAGTGTGAGGGTTGTCGGATT
24[223]26[210] CTATATTTTCATTTGACCATTAGATACAATTGCTCCTTTTGA
28[69]24[56] GTTAAAATTCGCATACTAAATCGGAACCAACATTATGACCCT
2[209]4[196] AGGCGAATTATTCACCTTTACATCGGGAATCCTGATTGTTTG
11[84]9[97] AAGCGTAAGAATACGTAGCACCATTAATGACGGAAATT
33[56]31[69] CGAAAATCCTGTTTCACCGCCTGGCCCTTGCCCGCTTTCCAG
22[153]18[140] GGTGCTGAGGCTTCGGTACGCCAGAATGCTGGCTGACCTTC
30[181]32[168] AGAGGATCCCCGGTTACGCCAGCTGGCGGTGCCGGAACCA
32[125]34[112] GGCGCCAGGGTGGTGAATAGCCCGAGATTTCCAGTAAAAGTT
3[224]5[237] TTCGCCTGATTGCTTATCAGATGATGGCAATAGATAATACAT
8[125]10[112] CATAAAGGTGGCAAAATTATCACCGTCATTGGGAATTAGAGC
4[195]0[182] GATTATACTTCTGAAATATCTTAGGAGCGCTGAGAAGAGTC
6[139]8[126] AAGAATTGAGTTAAGTAGAAATGCCAGTCCTAATTATACATA
18[223]20[210] AATCATAAGGGAACACCTGCTCCATGTTTAAACGGGTAAAAT
19[112]23[125] CATTATACCAGTCATGACAAGAACCGGAGGCCGATTAAAGGG
2[237]0[224] TTGAATACCAAGTTTATTAATTAATTTTCCCTTAGAATCCTT
1[154]5[167] CCTTTTTTAATGGACAAAATCATAGGTCTTGAAAGGAATTGA
12[181]14[168] TATTCTGAAACATGAGTAAGCGTCATACCAGACGATTGGCCT
9[140]7[153] GGAATCATTACCGCGCTCTTCCAGAGTACAAAATAAACAG
6[181]8[168] GTCAGAGGGTAATTCCAATCCAAATAAGTTATCCTGAATCTT
3[56]1[69] GCATTTTCGAGCCAGTAGGGCTTAATTGTTAATGGTTTGA
32[237]30[224] CGACAGTATCGGCCAGGGTTTTCCAGTCACGACGTTGTAAA
23[42]21[55] ACTATGGTTGCTTTTCGTCATAAATATTACATAACGCCAAAA
30[223]32[210] ACGACGGCCAGTGCAGTTGGGTAACGCCTCAGGAAGATCGCA
26[125]28[112] ACAGTCAAATCACCATGTACCCCGTTGCCCAAAACAGGAA

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|----------------|---|
| 29[42]27[55] | AGGTGCCGTAAAGCTAAATTTTGTAAACATTGCCTGAGAGT |
| 10[69]6[56] | AAACGTCACCAATGCAACAGAGATAGAATTTTTAAGAAAAGT |
| 7[112]11[125] | GTATGTTAGCAAACGCCCAATAATAAGAATTTTGAATGGCT |
| 8[209]10[196] | AGTTGCTATTTTGCTTCTAAGAACGCGATCTTTCCTTATCAT |
| 33[224]35[237] | GTTTGAGGGGACGAATCAGCTCATTTTTGTTTTTGGGGTCG |
| 34[69]30[56] | GGAGCGGAATTATCGTATTAGACTTTACTGTAAAGCCTGGGG |
| 5[84]3[97] | AAAGCATCACCTTGATCAACAATAGATATAAAGTACCGACAA |
| 6[55]8[42] | AAGCAGATAGCCGAGAAACCGAGGAAACGAAAATTCATATGG |
| 17[42]15[55] | TTTGATTAGTAATATATCAGCTTGCTTTAGACGTTAGTAAAT |
| 8[83]10[70] | GAATAAGTTTATTTGGAGGGAAGGTAAAATTAGCAAGGCCGG |
| 9[224]11[237] | CCTCCCGACTTGCGCATGTAGAAACCAATATTAACACCGCCT |
| 27[224]29[237] | AACAGGTCAGGATTCATTGAATCCCCCTGCTACAGGGCGCGT |
| 8[237]6[224] | GGAGGTTTTGAAGCTTACAGAGAGAATAACATAAAAAACAGGG |
| 10[153]6[140] | CAAGCAAGCCGTTTCGCGAACTGATAGCGAGAGATAACCCAC |
| 11[126]9[139] | ATTAGTCTTTAATGTTATTTTGAGCCATCCGACTTCATCGTA |
| 33[140]31[153] | CTCCGTGGGAACAACGCCATTCAGGCTGAGGGCGATCGGTGC |
| 21[224]23[237] | AAAGACTTTTTTCATCGAGGTGAATTTCTGTAGCAATACTTC |
| 3[140]1[153] | AAAGAAATTGCGTAAGAAAACAAAATTATTTTCATTGAATTA |
| 3[98]1[111] | AAGGTAAAGTAATTGCGTTATACAAATTAATAAGAATAAACA |
| 14[237]12[224] | GCCACCAGAACCACACGGGGTCAGTGCCTTGAGTAACAGTGC |
| 5[168]3[181] | GGAAGGTTATCTAAATAATGGAAGGGTTACGTCAGATGAATA |
| 14[125]16[112] | TAACACTGAGTTTCATAGAAAGGAACAAAATAATAATTTTTT |
| 13[196]17[209] | TACAGGAGTGTACTAATGCCCCCTGCCTATTACATTGGCAG |
| 10[195]6[182] | TCCAAGAACGGGTAACGAACCACCAGCAACACCTGAACAAA |
| 22[69]18[56] | CCAATACTGCGGAAGACGAGCACGTATAGAATAAGGCTTGCC |

The linear origami:

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| 6[139]8[126] | TATCTTACCGAAGCAAACGCAATTTTTAGAAACGAAGACAC |
| 17[126]15[139] | TAAAAAATACCGAATCACCGGAACTAAAAAGGAACAACCAGA |
| 0[55]2[42] | AAATAAGAATAAACACCGGAATCATAATATATAAAGTACCGA |
| 17[210]15[223] | TGAGAGCCAGCAGCTTAGCGTCAGACTGGAGCCGCCACCAGA |
| 1[196]5[209] | GCTGAGAAGAGTCAAGGTTGGGTTATATATTATACITCTGAA |
| 23[126]21[139] | TTTGACGCTCAATCTGAGGCTCCCTCGTATCATAATGCAGGG |
| 15[182]13[195] | ACCGCCACCCTCAGGGCCTTGATATTCAGTACTGGTAATAAG |
| 4[153]0[140] | TTTTACATCGGGAGTCATATTCCTGATTGAACGCGAGAAAAC |
| 8[167]10[154] | AGCCATATTATTTAGAGGCGTTTTAGCGGCGCCCAATAGCAA |
| 19[196]23[209] | ATAAATTGTGTCGAAAGGGAACCGAACTAGGGACATTCTGGC |
| 15[98]13[111] | TCAACAGTTTCAGCGTACAACTACAACCAGGGATAGCAAGC |
| 32[209]34[196] | TGGGCGCATCGTAACAACCCGTCGGATTAACGTTAATATTTT |
| 33[56]31[69] | GTTTTCTTTTCACAACGCGCGGGGAGAACACAACATACGAG |
| 5[126]3[139] | GAGCGGAATTATCAAAACAATCTGAACAATAAGTCAACGGAT |
| 11[84]9[97] | TATCTGGTCAGTTGGGCCGGAACGTCAAAGGTGAATTATCA |
| 29[210]27[223] | AACATCACTTGCCTATACTGCGGAATCGAATATCGCGTTTTA |
| 0[139]2[126] | TTTTCAAATATATAGTAGGGCGTCGCTTGTAATCTTAATT |

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|----------------|---|
| 14[83]16[70] | ACAGACAGCCCTCATTCTGTATGGGATTAAAAAGGCTCCAAA |
| 5[42]3[55] | GTATTAATCCTTTCATTCCAAGAACGGCAAAAGGTAAAGTA |
| 16[153]12[140] | TTTCATAATCAAAACGAACCACCAGCAGTAGGATTAGCGGGG |
| 19[112]23[125] | CTTATGCGATTTTAAGTAATCTTGACAAGGAAATACCTACAT |
| 25[154]29[167] | AACATGTTTTAAATGCGCGAGCTGAAAAAGAGTCTGTCCATC |
| 4[111]0[98] | CCATCCTAATTTACTTTTGCGGAACAAATCTTCTGACCTAAA |
| 9[140]7[153] | TATTCTAAGAACGCTCCCAATCCAATAGTTTAACGTCAAAA |
| 25[70]29[83] | ATACTTTTGCGGGAGCAAAATTAAGCAAAGGCCGATTAAAGG |
| 30[139]32[126] | CAAGCTTGCATGCCATTGCGTCAGCTTCTCCAGCTGCGCTC |
| 9[182]7[195] | CGGGAGGTTTTGAAAGCCTAATTTGCCATAACATAAAACAG |
| 12[181]14[168] | AGAGGCTGAGACTCGATGATACAGGAGTCAAACAAATAAATC |
| 20[83]22[70] | ACGAACTAACGGAACTAATGCAGATACCAAAGAAGTTTTG |
| 9[98]7[111] | CCGTCACCGACTTGATTTTGTCAATCCATACATAAAGGTG |
| 14[167]16[154] | CTCATTAAGCCAGCGCCTCCCTCAGAGAGCGTTTGCCATCT |
| 3[182]1[195] | GGCGAATTATTCATCTTTTTTAATGGAAGCTTAGATTAAGAC |
| 5[168]3[181] | ATTCATCAATATAACGTCAGATGAATATCAAAATCGCGCAGA |
| 27[224]29[237] | ATTCGAGCTTCAAACCTGGATAGCGTCCAGAGTAGAAGAACTC |
| 7[112]11[125] | GCAACATATAAAAGCCTTTTTAAGAAAAGAAAGGAATTGAGG |
| 27[98]25[111] | CAAAGGCTATCAGGAATGTGTAGGTAAAAACGCAAGGATAAA |
| 33[98]31[111] | TTCACCGCCTGGCCAGTCGGGAAACCTGTGCCTAATGAGTGA |
| 14[125]16[112] | GAGTTTCGTCACCAGGAGTGAGAATAGAGGAATTGCGAATAA |
| 28[111]24[98] | GGTAATCGTAAACACGGTACGCCAGAATAAATCATACAGGC |
| 0[181]2[168] | GCTGATGCAAATCCAAAACATAGCGATAACAGTACATAAATC |
| 24[223]26[210] | CATTAGATACATTTCAAGTTGATTCCCAAAGCAAACCTCCAAC |
| 2[83]4[70] | GAGGCATTTTCGAGACGACAATAACAACAATCAATAATCGG |
| 34[111]30[98] | CAGGCGAAAACTCTGACTCCAACGTCAAAGAGGATCCCCGGG |
| 7[154]11[167] | ATGAAAATAGCAGCGAGCAAGAAACAATCTAACAATAATAG |
| 11[210]9[223] | ATTAGACTTTACAACCGCACTCATCGAGTTAGTTGCTATTTT |
| 23[84]21[97] | CAGCCATTGCAACAGCGAGAGGCTTTTGATAACGCCAAAAGG |
| 9[56]7[69] | GGTAAATATTGACGCCAAAGACAAAAGGAAGACTCCTTATTA |
| 12[139]14[126] | TTTTGCTCAGTACCTGTACCGATTTACCTCTCTGATAACACT |
| 12[55]14[42] | GGTTTAGTACCGCCACCCTCAGAACCGCATCTAAAGTTTTGT |
| 32[237]30[224] | GTAATGGGATAGGTGGGCCTCTTCGCTATTACGCCAGCTGGC |
| 26[237]24[224] | GCGAACCAGACCGGTTCTGCGAACGAGTAGATTTAGTTTGAC |
| 1[70]5[83] | TGTTTAGTATCATAATACCGACCGTGTGAATTTTAAAAAGTTT |
| 28[153]24[140] | TGACCATAAATCAAGGCCACCGAGTAAAGGTGGCATCAATTC |
| 5[210]3[223] | TAATGGAAGGGTTAGTAAACAGAAATACAAAAGAAGATGAT |
| 20[209]22[196] | GGTAAATACGTAAACGAGGGTAGCAACGCTTGATACCGATA |
| 23[168]21[181] | ATTCACCAGTCACAGACAACAACCATCGACCCTCAGCAGCGA |
| 8[237]6[224] | TTTTATCCTGAATCGAACACCCTGAACAAAGTCAGAGGGTAA |
| 3[56]1[69] | ATTCTGTCCAGACGCCAGTAATAAGAGATACTAGAAAAAGCC |
| 22[153]18[140] | ATATATTCGGTCGCGTCTGAAATGGATTGCGCATAGGCTGGC |
| 15[56]13[69] | TTAGTAAATGAATTTAGTTAGCGTAACGCACCCTCAGAACCG |
| 0[223]2[210] | TTAACCTCCGGCTTATAGTGAATTTATCTTACATTTAACAAT |

31[196]35[209] CGCAACTGTTGGGACTGCAAGGCGATTAGTACTATGGTTGCT
0[97]2[84] TTTAATGGTTTGAATGCGTTATACAAATATGTAATTTAGGCA
27[182]25[195] AGATTAAGAGGAAGGAGTACCTTTAATTCGGTGTCTGGAAGT
5[84]3[97] GAGTAACATTATCAGAGCATGTAGAAACCATGTTGAGCTAAT
26[167]28[154] GAGGTCATTTTTGCGTCAGAAGCAAAGCTCAGAAAACGAGAA
17[84]15[97] TTAGTCTTTAATGCTGAAAATCTCCAAATTGCTAAACAACTT
19[154]23[167] AAGCGCGAAACAAAGGTGTACAGACCAGATTTACATTGGCAG
2[209]4[196] TTCATTTGAATTACTTCAATTACCTGAGAAGAAATTGCGTAG
34[69]30[56] ATCCCTTATAAATCGTTGTTCCAGTTTGATAGCTGTTTCCTG
18[139]20[126] TGACCTTCATCAAGAGAACTGTTGACCCCTCATCTGCTCATT
8[125]10[112] CACGGAATAAGTTTAGCCATTTGGGAATTCACCAAGTAGCACC
35[168]33[181] CCGCGCTTAATGCGTAAATTTTTGTTAATTCATCAACATTAA
14[209]16[196] CAGGTCAGACGATTAGCCACCACCCTCATAGCGCGTTTTTCAT
30[223]32[210] GAAAGGGGGATGTGAGGGCGATCGGTGCCACGTTGGTGTAGA
26[83]28[70] AGAAAGGCCGAGAGAGAGGGTAGCTATTATGTACCCCGGTT
33[182]31[195] ATGTGAGCGAGTAACCGTGCATCTGCCACGCCATTACAGGCTG
25[112]29[125] AATTTTTAGAACCCGCGATTAAACATCCAATCCTGAGAAGTGTT
30[181]32[168] AGGGTTTTCCAGTGGCAAAGCGCCATTGTTTGAGGGGACGA
28[69]24[56] GATAATCAGAAAAGGGGAGCTAAACAGGTAAAGCCTCAGAGC
7[196]11[209] GGAAGCGCATTAGACAGAGAGATAACCCGAGGATTTAGAAGT
11[168]9[181] ATTAGAGCCGTCAATAGGAATCATTACCAACCTCCCGACTTG
15[224]17[237] ACCACCACCAGAGCAATCAAGTTTGCCTAAATGAAAAATCTA
29[84]27[97] GATTTTAGACAGGATAGCATGTCAATCATTTTGAGAGATCTA
31[154]35[167] GGTGCCGGAACACACGACGTTGTAACCAACCAACACCCG
8[83]10[70] ATGGTTTACCAGCGGAAATTATTCATTACCAATGAAACCATC
3[98]1[111] GCAGAACGCGCCTGTTAACAACGCCAACTCTTACCAGTATAA
33[140]31[153] TTCGCGTCTGGCCTTCAGGAAGATCGCACCGGCACCGCTTCT
1[112]5[125] AGCCAACGCTCAACTTTAGTTAATTTAGAAACCAACAGAAAG
23[42]21[55] AACTATCGGCCTTGAGTAAATGTTTAGCAGTTGAGATTTAG
16[195]12[182] CGGCATTTTCGGTCATTAAACACCGCCTGACATGAAAGTATTA
21[182]19[195] AAGACAGCATCGGATGCCACTACGAAGGTGTATCATCGCCTG
19[70]23[83] GCTTGAGATGGTTTGTAAACAAAGCTGCTAACAATATTACCGC
34[153]30[140] TAACCAATAGGAACGTCACGCTGCGCGTACGACGGCCAGTG
20[237]18[224] TTTTTCATGAGGAATCCATGTTACTTAGCCGGAACGAGGCGC
2[167]4[154] AATATATGTGAGTGTGAATACCAAGTTAACAGTAACAGTACC
1[154]5[167] CCTTAGAATCCTTGAATCGCAAGACAAAATCAGATGATGGCA
21[140]19[153] AGTTAAAGGCCGCTAATACACTAAAAACCAAGCGATTATACC
12[97]14[84] ATATAAGTATAGCCACCACCCTCATTTTGCCTGTAGCATTCC
28[195]24[182] TGAATCCCCCTCAATTGTAGCAATACTTAACCTGTTTAGCTA
3[140]1[153] TCGCCTGATTGCTTAATAACCTTGCTTCATTAAATTTTC
24[97]26[84] AAGGCAAAGAATTAGAAGCCTTTATTTGATTCAAAAGGGTG
18[55]20[42] GGCTTGCCCTGACGAGAAACACCAGAACGTAGAAAGATTCAT
10[153]6[140] GCAAAATCAGATATATATCTTTAGGAGCAGAAATAGCAATAGC
4[195]0[182] ATTTTCAGGTTTAATCCTGATTGTTTGGAACATATGTAAAT

21[98]19[111] AATTACGAGGCATAGTTGGGAAGAAAAATCATTGTGAATTAC
27[140]25[153] CCCTGACTATTATAGGATGGCTTAGAGCTAATGCTGTAGCTC
10[111]6[98] ATTACCATAGCAAGCAAATCAACAGTTGTAAGCAGATAGCC
14[237]12[224] CGCCGCCAGCATTGAGTGCCCGTATAAACAGTTAATGCCCCC
24[139]26[126] TACTAATAGTAGTATCATATACTGAATATTAATTGTTTTAA
22[69]18[56] CCAGAGGGGGTAATCTGGTAATATCCAGCATTGAGTGAATAA
21[56]19[69] GAATACCACATTACAACATTATTACAGGAGTAGTAAATTGG
11[126]9[139] AAGGTTATCTAAAAGAAGGCTAGCAAAATAGAGCCTATCCGG
26[125]28[112] TGCAATGCCTGAGTTCATTGCCTGAGAGGAGAATCGATGAAC
35[84]33[97] TATTAAAGAACGTGGTTTGATGGTGGTTACAGCTGATTGCCC
22[111]18[98] TAAAAACCAAAATAGGAAAAACGCTCATGAACCGGATATTCA
6[181]8[168] AAGCCCAATAAATACTTTACAGAGAGAAGTTACAAAATAAAC
20[167]22[154] GAAAGAGGCAAAAGTTTGCGGGATCGTCCCCACGCATAACCG
16[111]12[98] TAATTTTTTCACGTGCGAACTGATAGCCCGTCGAGAGGGTTG
18[97]20[84] TTACCCAAATCAACAATTTCACTTTAATCTACGTTAATAAA
10[195]6[182] TTTTATTTTCATCGTAGATAATACATTTACAAGAATTGAGTT
27[56]25[69] ATAAATTAATGCCGCAGTCAAATCACCAATTATGACCCTGTA
6[97]8[84] GAACAAAGTTACCAAAACGTAGAAAAATAATAGAAAATTCAT
15[140]13[153] GCCACCACCGGAACAATGGAAAGCGCAGGTTCCAGTAAGCGT
22[195]18[182] GTTGCGCCGACAATCGACCAGTAATAAAGACCAACTTTGAAA
18[223]20[210] AGACGGTCAATCATAATCCGCGACCTGCGTTTCCATTAAACG
32[125]34[112] ACTGCCCGCTTTCCTGAGAGAGTTGCACTGGTTTGCCCCAG
33[224]35[237] ACGGCGGATTGACCGATTGTATAAGCAATAACGTGCTTTCCT
12[223]14[210] TGCCTATTTGGAAGTGCCTTGAGTAACACAGGAGGTTGAGG
13[154]17[167] CATACATGGCTTTTCTCAAGAGAAGGATAAGATAAAACAGAG
20[125]22[112] ATACCAGTCAGGACGTAAGAGCAACACTTTACCAGACGACGA
16[69]12[56] AGGAGCCTTTAATTTGGCACAGACAATACACCGTACTCAGGA
35[210]33[223] TTGACGAGCACGTAATATTAAATTGTAAGTCCGTGGGAACAA
24[181]26[168] TATTTTCATTTGGGATGCAACTAAAGTAGCTCCTTTTGATAA
24[55]26[42] ATAAAGCTAAATCGGTTGTACCAAAAACCTCAATATGATATTC
32[167]34[154] CGACAGTATCGGCTCCTGTAGCCAGCTATCAGCTCATTTTT
29[42]27[55] GTTAGAATCAGAGCCCCCAAAACAGGAAACCGTCTAGCTG
6[55]8[42] CGGAATACCCAAAAGAACTGGCATGATTGCGACATTCAACCG
8[209]10[196] GAGCGTCTTCCAGGCCTTAAATCAAGAAACAAGCAAGCCGT
23[210]21[223] CAACAGAGATAGAAGAATTTCTTAACAGGCTACAGAGGCTT
32[83]34[70] TTAATGAATCGGCCCAGTGAGACGGGCACCGAAATCGGCAAA
35[126]33[139] TCTATCATGTAGCGCCATCAGTCCACGCAAGCGAAAATAA
21[224]23[237] TGAGGACTAAAGACCTTGCTTTGAGGTCCCTCTGACCTGA
31[70]35[83] CCGGAAGCATAAAGTCGTAATCATGGTCGAACAAGAGTCCAC
29[126]27[139] TTTATAATCAGTGAATAACAGCAACAATCTGGAGGTCTTTA
26[209]28[196] AGGTCAGGATTAGACCCGAAAGACTTCATCATAAATATTCAT
17[42]15[55] AGCGTAAGAATACGGTATCGGTTTATCACGTCTTTCAGACG
13[196]17[209] TTTTAACGGGGTCACCTATTATTCTGAACAACAGTGCCACGC
13[112]17[125] CCAATAGGAACCAAGGCGGATAAGTGCCTAAAACATCGCCA

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|----------------|---|
| 25[196]29[209] | TTCATTCCATATAACGCAAATGGTCAATCTTTGATTAGTAAT |
| 11[42]9[55] | AGCATCACCTTGCTAATCAGTAGCGACAATTGAGGGAGGGAA |
| 4[69]0[56] | CTGTCTTTCCTTATGCCCCGAACGTTATTATAAATAAGGCGTT |
| 13[70]17[83] | CCACCCTCAGAGCCCGGAATAGGTGTATTTTTGAATGGCTA |
| 34[195]30[182] | GTAAAAATTCGCATCCGCTACAGGGCGCAGTTGGGTAACGCC |
| 17[168]15[181] | GTGAGGCGGTCAGTATAGCCCCCTTATTCCGCCACCCTCAGA |
| 6[223]8[210] | TTGAGCGCTAATATCGGGAGAATTAACTTTACCAACGCTAAC |
| 2[125]4[112] | GAGAATCGCCATATTTTATCAACAATAGAGAAAAATAATATC |
| 18[181]20[168] | GAGGACAGATGAACGTACAACGGAGATTCACCAACCTAAAAC |
| 7[70]11[83] | CGCAGTATGTTAGCGAAGGAAACCGAGGAAACCTCAATCAA |
| 30[97]32[84] | TACCGAGCTCGAATTGTAAAGCCTGGGGTCGTGCCAGCTGCA |
| 29[168]27[181] | ACGCAAATTAACCGATGCTTTAAACAGTGGATTGCATCAAAA |
| 31[112]35[125] | GCTAACTCACATTATGCAGGTCGACTCTAGGGCGAAAAACCG |
| 10[69]6[56] | GATAGCAGCACCGTGAACCTCAAATATCAAACGCAATAATAA |
| 9[224]11[237] | GCACCCAGCTACAATATTAACCAAGTAACAATTCGACAAC |