

Supporting Information

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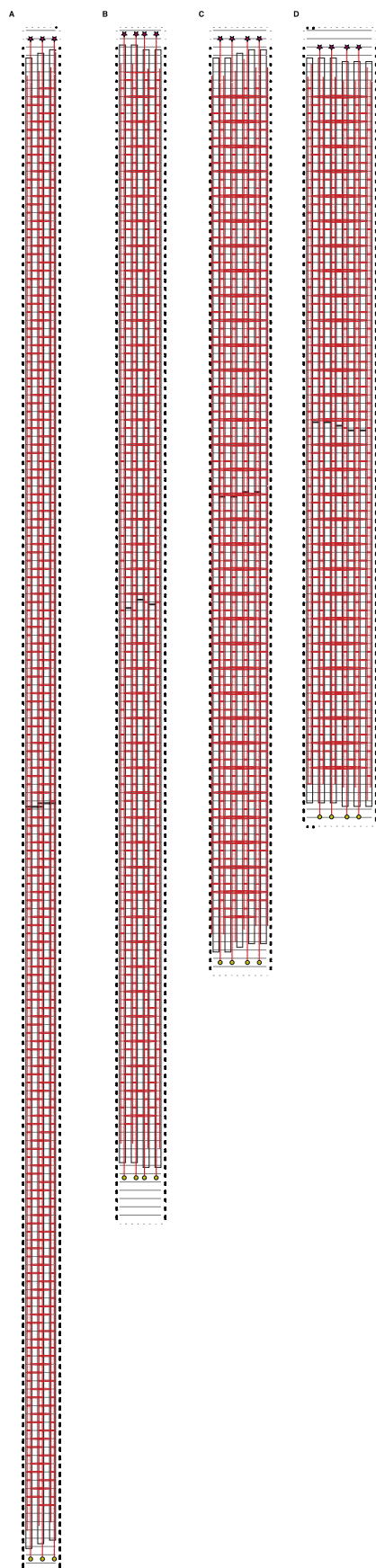
69451 Weinheim, Germany

**Rigid DNA Beams for High-Resolution Single-Molecule Mechanics\*\***

*Emanuel Pfitzner, Christian Wachauf, Fabian Kilchherr, Benjamin Pelz, William M. Shih, Matthias Rief, and Hendrik Dietz\**

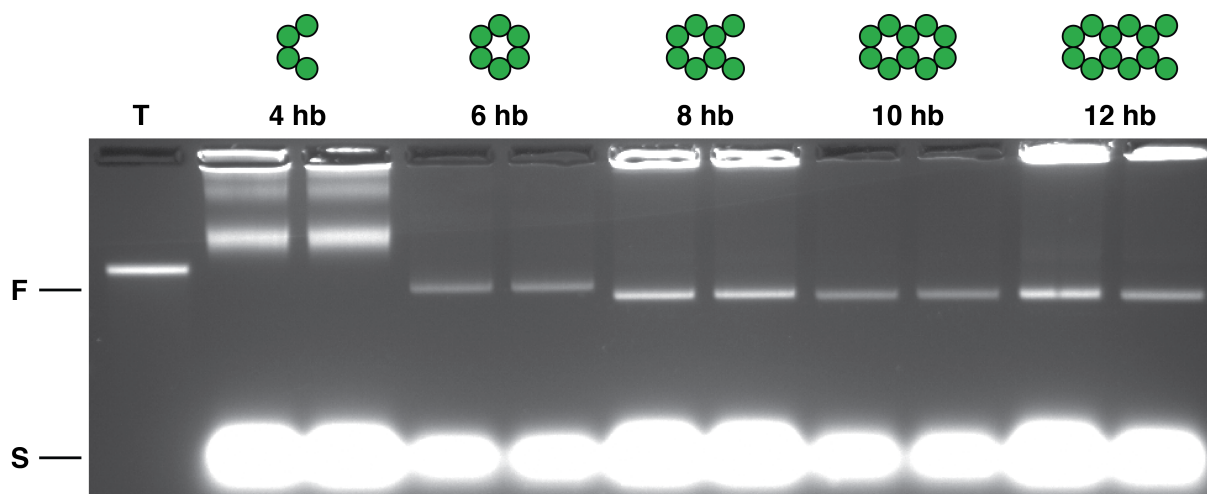
anie\_201302727\_sm\_miscellaneous\_information.pdf

Figure S1



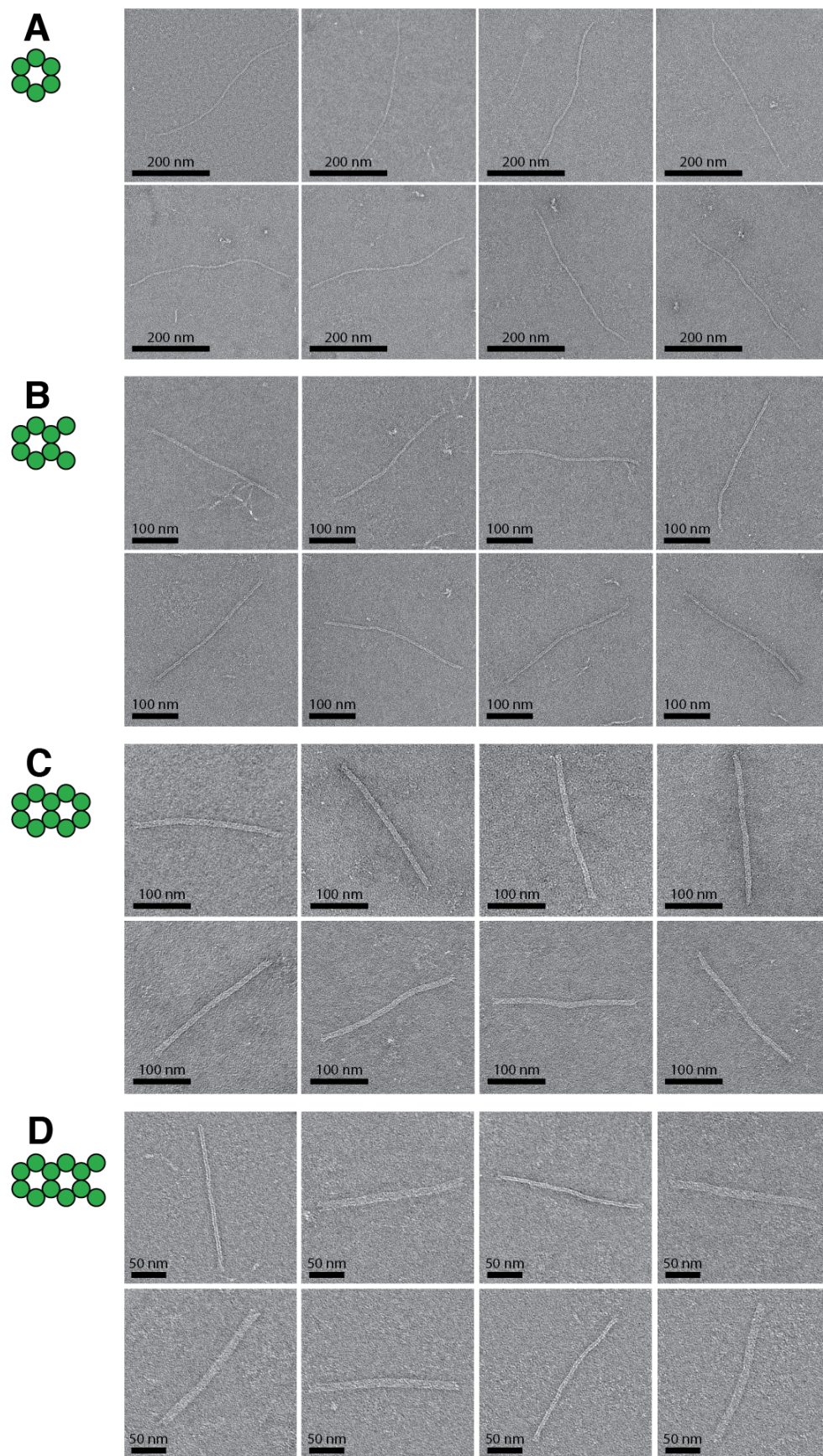
**Fig. S1:** caDNA design diagrams. Scaffold strand path is depicted with black lines, and staple strand paths are shown as red lines.

Figure S2



**Fig. S2:** Gel-electrophoretic bundle folding quality assessment. Photograph of a 2% ethidium-bromide stained 2% agarose gel on which a reaction products were electrophoresed. The gel also includes a four helix bundle not described in the main text. The low electrophoretic mobility of the folded species as compared to the other bundle types is due to the high flexibility (random-coil-like) of this bundle type. The four helix bundle was thus not considered for usage as a linker system. "T" marks the lane on which the template strand was electrophoresed. "F" marks the folded species, "S" excess staple strands.

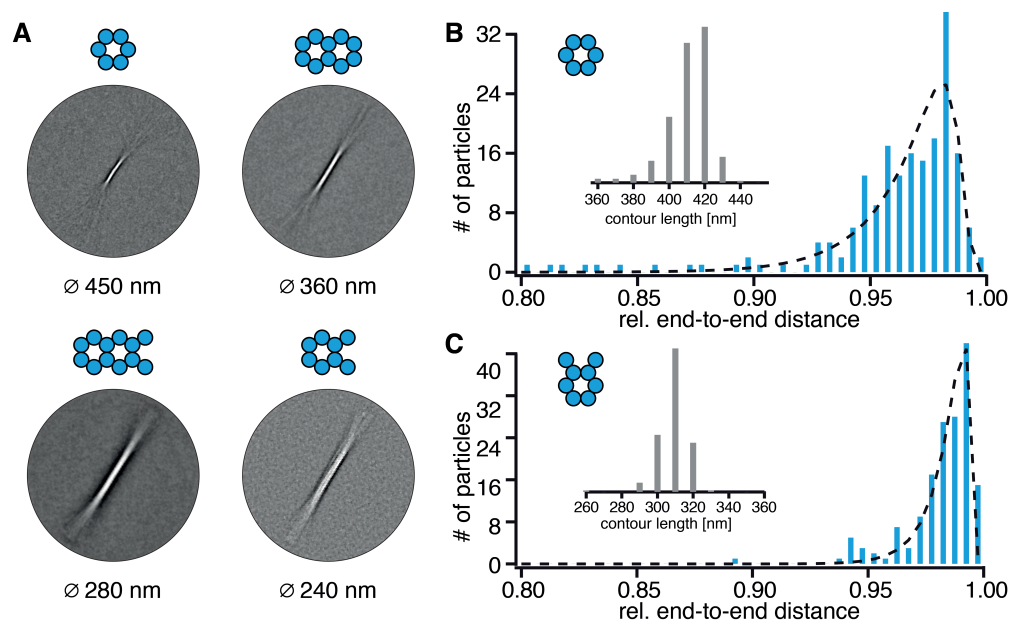
Figure S3



**Fig. S3:** Exemplary negative-stain transmission electron micrographs of purified DNA helix bundle particles.

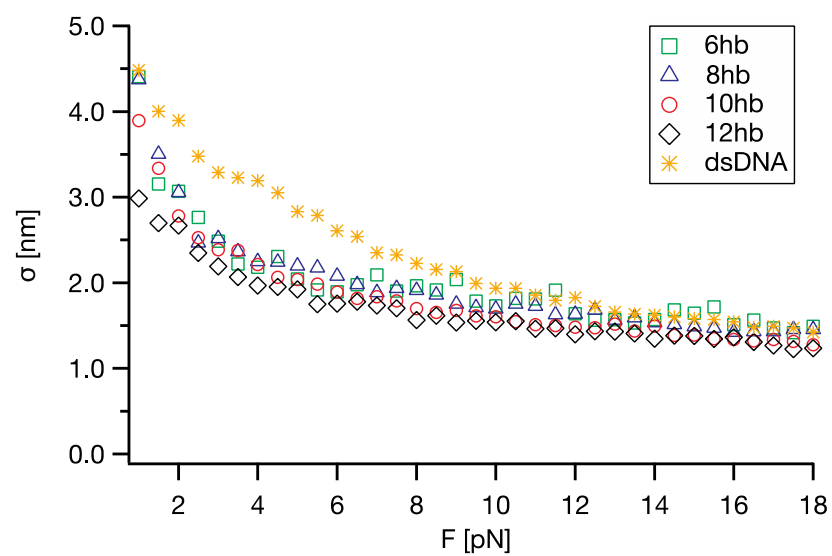


Figure S4



**Fig. S4:** (A) Average TEM-micrographs obtained by rotational alignment of single particle micrographs of each bundle type against a randomly chosen reference micrograph. (B,C) Blue Bars: Histograms of the relative end-to-end distance observed for individual six (B) and eight (C) helix bundle particles. The absolute contour lengths (inset, grey bars) were determined by manual backbone tracing, where the two endpoints of a backbone trace were used to determine the absolute end-to-end distance. Dashed lines: fit by a semi-flexible beam model. <sup>[10]</sup>

Figure S5

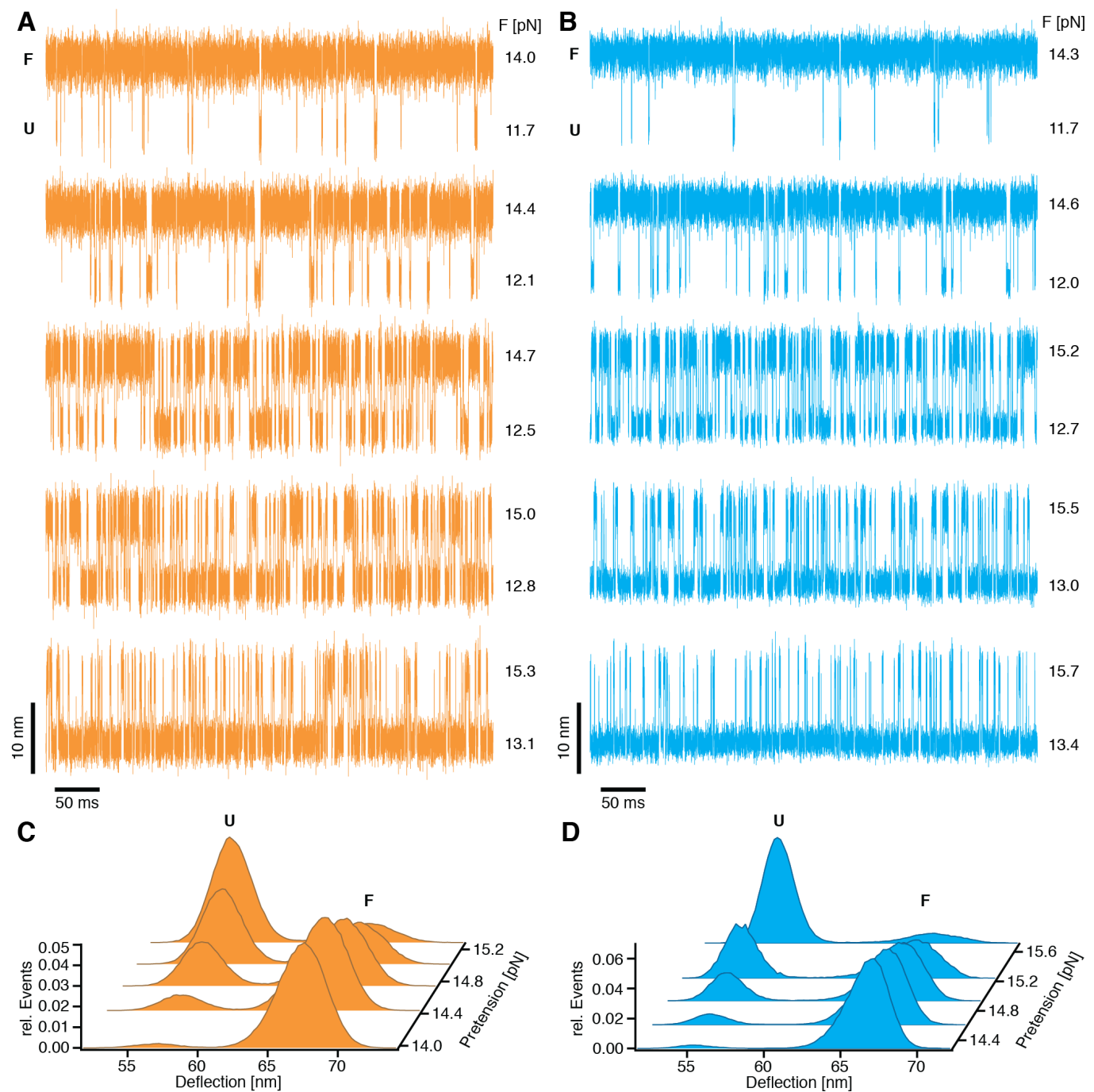


**Fig. S5:** Force-dependent noise-amplitudes for stiff DNA helix bundle linkers vs. conventional dsDNA linkers.

Diagram illustrating the structure of a circular plasmid map and its corresponding linear sequence. The plasmid map shows a circular DNA molecule with a 5' and 3' end, and a 3' end. The linear sequence is shown below the map, with the 5' end at the top and the 3' end at the bottom. The sequence is color-coded: blue for the 5' end, red for the 3' end, and green for the rest of the sequence. The sequence is flanked by two blue circles, each containing a 'T' and a 'G'.

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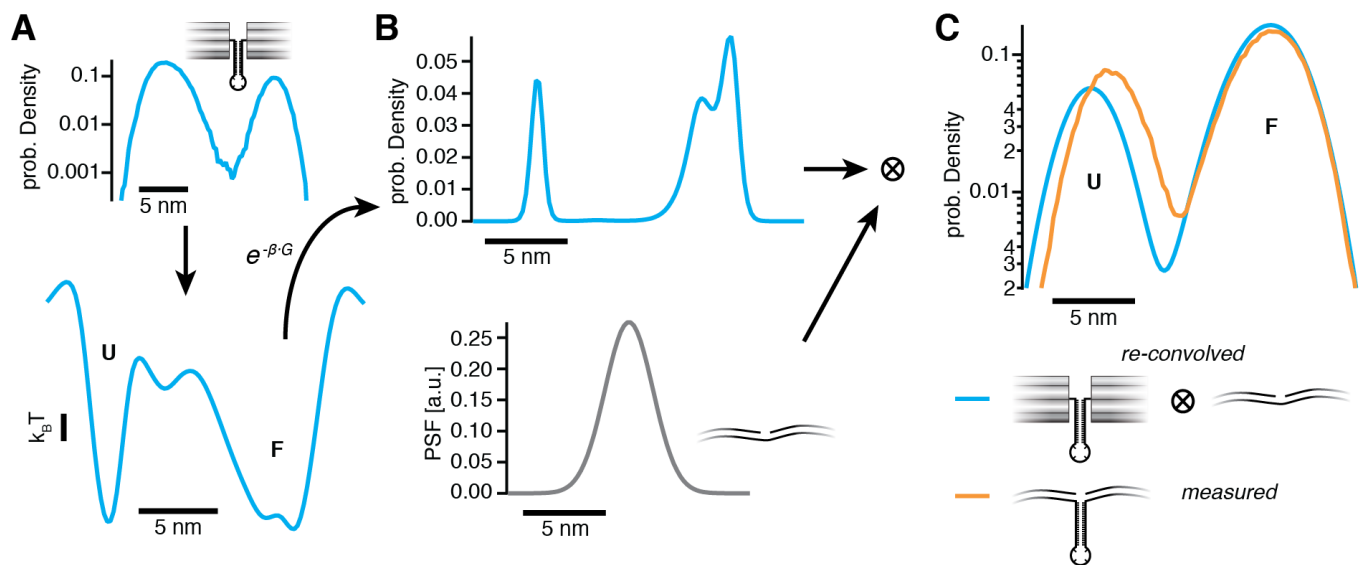
Figure S7



**Fig. S7:** Additional constant-distance data as in Fig. 3 obtained for different loads. Orange traces: data collected with conventional linker system. Cyan traces: data collected with stiff ten helix bundles.



Figure S8



**Fig. S8:** Schematic explanation of the workflow for deconvolution of a deflection histogram to extract the energy landscape for the 20-bp DNA hairpin from figure 3, followed by reconvolution using the point-spread-function of the conventional dsDNA handle system.

## Sequence Information

### 6 helix bundle

Start	End	Sequence	Length
0[370]	4[357]	CCGCGTTATTAAAGTGTATCAAAAGGGTGTAAAGAGGCCAACTT	42
0[412]	4[399]	TAATGAGTAAAAGCTCATATTTTAAAGAGGCTTCGGAACGA	42
0[454]	4[441]	AACATACGTGGATGGAAGCCTTTATTCAGGATTAATTTGT	42
0[496]	4[483]	CTGTTTCAAGTGTGTGTACCAAAAACATTTTCGGCGAA	42
0[538]	4[525]	GATCCCCCGCAAGCAAAATTAAGCAAGCTGTAGTACATA	42
0[580]	4[567]	GGCTAAGAGGTAGGCAATTAAGTCTCAAGCTTGGACACTAC	42
0[622]	4[609]	TCCTCGAAAGCAACGCCAGCTTGAAGAGCAACCTTTATG	42
0[706]	4[693]	AATTAAAGCTTATTAATTCATTAAGTATAGCAATGCGGGA	42
0[748]	4[735]	ACTTTTGTAGGGGCATTTCAACCGATTCACCGTATATTCG	42
0[790]	4[777]	ATGCTGATGTAATTAGAAAAATCATATGTCAGACTTCGCCG	42
0[832]	4[819]	TTTTAACTATAAAGCGCAAGAACACCCCTTATTTGCTTTC	42
0[874]	4[861]	CAATAGTCGCAATCGTAGAAAAATACATGAACCAAGAGGCTC	42
0[916]	4[903]	TGAAAACATTAACATGCGCATGATTAAGGCCCTCAGATTCGGA	42
0[958]	4[945]	TCGTGAACATCTAGGAACCCGAGGGAACCAACAGACAGATT	42
1[385]	3[398]	TTTTATACGGTCACTCGGCTTTTAATCTTGCATTCGACGTAC	42
1[427]	3[440]	TACGCGATCCGGGAAACATCCAACGGTAAAGCAACAGGAGA	42
1[469]	3[482]	ATAACATAACACGGCTTTTGATGAAGAGGATATGATCATTCG	42
1[511]	3[524]	TGTCGGCATCTTTTGTGGAATAAATTAAGGCCCTTATGA	42
1[553]	3[566]	ACAGGAACCAACCTAATTAAGTACGGTTAAATCATTTACGC	42
1[595]	3[608]	ATCGCTTTTCATTTGAATTCCTCAATTCGGTGGCAGCCCTGC	42
1[637]	3[650]	GAACGCTCTCCAGAGCACTTACCATGGAATTTGTATGAT	42
1[721]	3[734]	AATATTAGGCTTGACATCTGATAGCGAGAGAGGCGCAACG	42
1[763]	3[776]	AGGCCCTACAAACAGCTTGCTTTTACGGTGTACTTAAACAC	42
1[805]	3[818]	GCAGAAAGATTCTTTTTCGGTATAGCCGGGAATAACAGTAAT	42
1[847]	3[860]	CTGCGAAAGCTTTAATCAAAATCAGCGACATAAATCTGTCT	42
1[889]	3[902]	TCATAAGTTTTCCTTCAGAGGCGGCCAATCTCTTCAGAACG	42
1[931]	3[944]	ATCAATAGTGAAGAACCCCTCAGAGCGCCGCAATACAGAAAT	42
2[384]	0[371]	AATGTGTTCAAATAATCATAAAGGAACCAATCTGCTCACTG	42
2[426]	0[413]	AATTTTCGGAGGCCCTGCTCATGTTTAAAGAGATGGGTGCC	42
2[468]	0[455]	ATACCTTATGCTCAGATTGTATCATCTCTTTTGTGCACAC	42
2[510]	0[497]	ATAAAGCAGCTTAAGAGCCCGACGATTTCAACATGTCATAG	42
2[552]	0[539]	AAGGCAAAATATGCAAAACGAAAGAGGCCGCCAGCGACGGAG	42
2[594]	0[581]	TACTAATTAACAGTAAAGGGTAAATACATTTTGTAAACAG	42
2[636]	0[623]	TATTTTTCGACATTTGCTTTTGGAGGATACAGATTAACATAT	42
2[720]	0[707]	AAATATTCACATGACGAGGAGCTTAAGGTACGTGGTGAACCTA	42
2[762]	0[749]	AAGCAAGAATACACATCTCGCCACAGCATATGCGCGGAGAAA	42
2[804]	0[791]	TGTTCACTCGCATAAACAGCTTGATACGGAACGATATGATA	42
2[846]	0[833]	ACATATATTTTCAATATTTGATTCGGTTTTCAGTATTAATCT	42
2[888]	0[875]	AGTATGTCGCTCACTTGTGAAAATCTCCAGCAAAAAGAGGT	42
2[930]	0[917]	AATACCCGAGCCTACTGAAGAAGCAACATATACAGAACTCT	42
3[357]	5[370]	CAAGGGCTCCAGTGAACGGTACGCCAGGAACTGAAAGGCCG	42
3[399]	5[412]	CAAGCTGTGTGAGCTAGAGGGCCACCGAGTATAGCCAAAGCA	42
3[441]	5[454]	AGCCAGGGAGCCGGCTGTGAGCAATACGCTTGATAGAGAT	42
3[483]	5[496]	CACGACTCTGTGTGCTGAGTAGGAAGAACATACATACGCGAT	42
3[525]	5[538]	CAATGTTCGGGTACCCAGGAACAATATACAAAAGACTCAACA	42
3[567]	5[580]	TCGCGCTGCTACGTATGGAATAACCTCATGAATGAGTTTCA	42
3[609]	5[622]	CATCTGACTCTGAAATTTATCATTTGGAGGACTTAGTAGAT	42
3[651]	5[706]	ATCATATTTGTTTGTGAAGCGCTAAGAACCCGTTTGGCGGGA	42
3[735]	5[748]	CTCAACAAATATGCTATTAGCTTTTAAACCGGATATACATG	42
3[777]	5[790]	GCACACTGCAAAATATGCTATTAATAATACGCTAGTATGATGC	42
3[819]	5[832]	AAGAACCTTCGCGAGAGGTGAGGGCGATATCAGCTTAGGTT	42
3[861]	5[874]	CAGCAGAGAATTTAAGCCTTGAGAGGCCAGCAAAAAGAGCAAC	42
3[903]	5[916]	CGCTGTATACGCGATTGCTGAACCTCAATAATGGAACACCGCC	42
3[945]	5[958]	AATATCATCTCGTCTGAGTTGGCAAAATCAAACTTTCAAAACCC	42
4[398]	2[385]	GGCGCAGAATCAGTACTCAACATTAATTTGCGTTTTCCTCGAT	42
4[440]	2[427]	GTGCAAAAATTAACAAAGCATAAAGTGTAGCTTTCTGGATAAA	42
4[482]	2[469]	ACAAGGCTCACTTGCAAATTTGTTATCCGCTGTGAATCCCTGTA	42
4[524]	2[511]	AAACACATTAATATCGAGGCTCGAATCTGTGTCAACTCAGAGC	42
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4[734]	2[721]	CTGCGCTTGAATGATTTAGTTAATTTGTATAAAGGAAGGT	42
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4[62]	3[62]	CGGAACATTTGGGGAAGAGAGCTGGGATTCACACATTAATG	42
4[104]	2[91]	AGGACGTAGGGAGCAGGTTGAGTGTGCGTGGGAATAATTT	42
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4[314]	2[301]	CTGGCTGAATCAGACCGGGGAGGCGGGCTGCTGCTAGCTG	42
4[356]	2[343]	TGAAGATAGACAGGGGAACCTGTGTCGGGGAGCCGGAGA	42
4[398]	2[373]	AATTTTCGGTTATCTCAATATATGTTGAGAGAAACAGCGGAA	42
4[1028]	2[1015]	AACGATCTAGAGCCATTTCAATTTGAATTTGAACGGGTAGCTAT	42
4[1070]	2[1057]	CCACGCTAGACTTATGAACAAACATCAACGCTAGTTAAG	42
4[1112]	2[1099]	CCAATGTCGGAACGAGGCAATTTATCAATAGCAAGTAATG	42
4[1154]	2[1141]	AGCCACCTTTGCGGATTCGCTGATTGACAGACCGTAGACGG	42
4[1196]	2[1183]	CGCCACCCATATTCATACAGTAAACAGTATTTTGAACAGGCTT	42
4[1238]	2[1225]	AGCCCGGCTGATTTAAAGAAATTTGGTGTGATCAAAAATAGA	42
4[1259]	5[1267]	CGAGAGGGTTGATACCTCAAGAGAGGATTAGGATT	36
5[38]	2[49]	CATCAACTAATGCAGATACATAACATACGCTCAATTT	39
5[77]	1[90]	GAGGCAATTCGCAACCGTTCGATCTCTTCCAGTAATCGGA	42
5[119]	1[132]	AGGCAATAGAAGTATGGGATAGGTACTTATAAGGAAGC	42
5[161]	1[174]	GAAGTTTCCCGGTTTGGCATTCGCCAGTTCGAAATAAAGGAG	42
5[203]	1[216]	ATAGCGTGATGAACGGAAGATGCGACTTGAGAGTTCTGCGG	42
5[245]	1[258]	TCGCCCTTATCAGGGCCGGAACACAGGCGAGACGCGAGGGC	42
5[287]	1[300]	CATAAATAATGCCGAACTGTGGGAAGGGTTTGGCGCTTCC	42
5[329]	1[342]	GAAGCAAAATCACCAGCCAGTGGCGGAAGCCAGTGTATTA	42
5[959]	1[972]	ACCAAGAGACCAAAATTTACGAGCATTTGTGAATAAAGGAAT	42
5[1001]	1[1014]	AGACGATAAGCCCTCTTATCATTTCCAACCTTTTAAACACT	42
5[1043]	1[1056]	AAAGCCAATAAGAGCATGCGAGAAACAGCAAGAAAGGATTTA	42
5[1085]	1[1098]	GTAAGCGATATCAGCATTCGCCGCCCAATTTCAATATTA	42
5[1127]	1[1140]	GTAATAAAAGTAAATCCGGATTTCTATTGTAATGATTAACA	42
5[1169]	1[1182]	CGTATAAAGAAATAGCACTTGCGGGAGGCCCTTTAAGCGGAA	42
5[1211]	1[1224]	ATTGTATTGTTCTATTTTGGCCCAAGATTTTTCATCA	42

#### biotin-modified

GTAGAAAGATTTCATCAGTTAGGGGTGGTGGTGTGT - Bio  
TTACCAATAGGACGCCATCAAAAATTAATGTGTGGTGT - Bio  
GGCGAAAACGCTATACGGTTTGGGT - Bio

#### digoxigenin-modified

Dig - TTTTGTGAGTACCAGGCGGATTAAGTCGGT  
Dig - TGTGGTTTGTGATTTTCGCGATTCAAAATAAACAGC  
Dig - TTTGTTTGGTGGGCTTACCATATCAAAATATTTTGC

## 8 helix bundle

Start	End	Sequence	Length
0	51551	ASCGTAAAAATGCAACCTCATATCAAAATATCTTTAAATGGCCGTG	49
0	51476	CAATTCAGTAATAATTCGGCTGATGCTTTGAAATACCAATGAAGAAC	49
0	51503	TGTGAAAGCGGTGATTAATCTTTCATACCGTTGCCATCTT	49
0	51916	GACGTGTGTGTGGGAATAGTAACAGAGTAGATTAATCTTGAAGGGAAC	49
0	11944	ATTAAGTGTGGTAACGCGAGCGGGGCTCTTCGCTATTACGC	42
2	71621	ATATCAAGTTTTTTCCTTTTAAAGAACTGGCAT	35
2	71482	TCAACTACATACGATATCTTCGGTCATAGCCCCC	35
2	51496	TCATATGTACCCTAGTTTAGACTGGATAAGGGGGTCTTGATA	42
2	71916	TTCTACTAAGGCGGTAGGTGTCAAGAGA	28
2	71937	GGGGCGCGAGCTGAAAGGGTGCATTAGATACATT	35
4	21321	ATACCGGAGGAAGGTATCAATATCT	24
4	21917	CTTCATCAAGAGTAGTTTGACCGTCAAA	28
5	01321	GAATCAATAATTACTAGTAAGACTACCCAGGGCGGAAA	35
5	01491	TAGTATCACCAAGGGGTGCGAACGTGGACTCCA	42
5	51475	AATCAGATAAGGGAATAGTAAAGGTACAAAATCGAAGCA	42
5	01490	CCCATAGGGTGAATAGAAAAGTAGCGTGTCTTCG	35
5	01910	CGAATGCCCGGAAATCGGGTGTTCCTCCAGTAC	35
6	51940	AGTATAGACCACTTTGAAGAGG	24
7	41421	CGCAATAATACCGGAATACCTTAAGAAAAGAAAAGTTGAA	42
7	21561	GATTAAGCTATCTTATATGCGCTCTTCGACCTAAATAGGAGACTCAA	49
7	01469	TATATAGACAAGGCTTATCCGTTGCTGCTGTGTATCCGCTCA	42
7	21497	TCATAATCTTTGATTTGCGCGAGAAGTTTTCACAGGCTCCAATGCTCAA	49
7	71940	AGGATTAGGATTAGCGGGGTTTTG	24
9	51971	CAGTTTGGCGAACCTGAAAAAGCGCTCAATAGATTTTTCATAAAGCC	49
10	51139	ATAAATCAAGCCGGAACACCGACTTTTACAAAACATGCTGATGATTTAA	49
10	51181	AAATCTGGAGCGGACCAACGAGCTTATTAATTTTAACTGAGCCAG	49
10	51223	AGTTGCACGGGTAAACGTGATCGGGAACAAAAGAAATAGTGAGTAATTC	49
10	51265	ACGGCGAGGCGTACACAGACATCTCTGATTATCAAAAACATAATGCAG	49
10	51307	TGCGTATTTCTCGAGAGATAGATTTGTGGATTATGTAATAACAGAA	49
10	51349	AGTCGATATAAGGGGACAGTGTAGCATATCAAAAACAGTACCAACAT	49
10	01394	TGCGTCAAGTCTTTACGCTGATAGATTTTACATCTGGGTAT	49
10	51423	CTTGGGTTCAACATATGCAATACAGTACTTTTCAAAAGACGGTTT	49
10	51517	CTGGTCTCCCAAAACATAGTATCTCGGAAGTGTGCAAGACAATG	49
10	51559	TAAAGACAACGTTGAGTCTGATGCTTTTAAACAGTCGTTACGGTCTGT	49
10	51601	TGAGTAAATCAGCTATTTTAAATCAGGCTTTTAAAGAAATATCGTCA	49
10	51643	CATAAATATTCGGGTTCAACCGGATGTCATCAAAATAGGAATAGCAACG	49
10	51685	TGGGCAACATGTAGTAGGAATAATCGGCTTTTGAACAACGAGGAAG	49
10	51727	ATACCGGAAACGCGGAAATGCAAGCAAACTCAACGAGCTTGGCAAGCG	49
10	51769	CCGCTTCATGGGCGTTCACGGGCTCTTTTGAATATACCTTAAAAACAC	49
10	51811	GCTGAATACGACGAGCAATATTAATTCGTGAATATGGGCTTAAACAA	49
10	51853	AAAGTGTTCGGCGATCAATAAATCAACTTAAGATATAAGGCTTGTGCA	49
10	51895	GTGCCAATCGCCATCAATAAACAGTGTATCCCAATCATACAGGCGCA	49
10	71104	GCGCAAACTAAAGGAGGACATAGCAAAAGTAGAA	35
10	71146	CAGTATTCGACGCTTACGAGGAAAGAAACGCAAG	35
10	71188	CGACGAGCGGTAGTGTGACCAATCAATAGAAAT	35
10	71230	ATGCGGCCACACACATCAATAAGGCGACATCA	35
10	71272	TACGTGCAATATGGGTTTAAGACGGAATATTC	35
10	71314	GGCCCACTTAGAATCAGCCTCTTGGCGCATTTGG	35
10	71356	CAGATTCATTTAGCGAGGCTACCATTAACCAITA	35
10	71398	CATTTGTATAAATTTGACCCCATCGATAGCAGCA	35
10	71440	CGCCAGCCGCAATTCGGGGAGTGCCCTTTAGCGTCA	35
10	71524	ATCGTAAACACGAGACGCTTGCAAAATCAACGGGAA	35
10	71566	TGCTTGAAATATTTTAAAAAACAGAGCGCCACCC	35
10	71608	GGGTAGCTCATTTTAAAGAACTTCAGAGCGGCCCA	35
10	71650	TATGATATCTGGGCTTTCAACCATGTGACAGGAGGT	35
10	71692	CAAAAAGGGGAGTACGTTAGTATTCGCAACCAAT	35
10	71734	ATATTTTGATTGACCATAGTTTGCAGCTCTGTAAT	35
10	71776	CTTTATCATGACAGTACATCTTTTATGATGATA	35
10	71818	TACCAAAATTCGGGCGCAATAGGTCGTCGTTGA	35
10	71860	ATTAAGCAACCTTCGCGACACCGCTCTGCTATTC	35
10	71902	TAACTCTCAGGCTGAGGTTTATTAAGAGGCTGA	35
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10	01133	CAACGCGCTAATAGGCGGAGCGGCAAAATCCCTT	35
10	01175	TAATAAGAATTAACGGCGCTGGCCCCAGCAGCGGA	35
10	02177	TGTCAGAGAGAGAACCCCGCCCTGGCCCTGAGAG	35
10	02599	AACGCGCGATTTTTTGTGTTTTTTCACAGTGAG	35
10	03011	AAATAATAATAAACAGAGCGGGGAGAGGGCGTT	35
10	03431	CAATAATACGCTAAACAGGAAAACCTGTCTGTC	35
10	03851	TAAACAGCTATTTTCAAGTGAACATTAATTCGCT	35
10	04277	TATTTTCCCGACTTTAACCTGTCAAAAGGTGAAG	35
10	05111	ACAACAAGGTTTATAGATTGCTCGAATTCGTAAT	35
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10	06371	GCTGCAAAACATCTTCGTGTGTCAGCGATTCTCA	35
10	06791	TTTCAATTTCAGACAACCCCAAGCACTCTGCG	35
10	07211	ACCAACCCAGCCCTCTGTAATGGGAGTGACTTATG	35
10	07631	TCATCTTTCGTACACCTGTGCCCCCAAAATAAAC	35
10	08051	TACAACGATAGCAACTCAGGATAAGTGCTTAAGT	35
10	08471	ATCCGCGTCAGAACTGTGCGGTGGATGTTCTTCT	35
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10	21440	ACACCACATTTAGCAACATGTAATATATGTAATTCGACAAGGGCGGT	49
10	21182	TCATATGACGGGAGAGATATGAGAGACTACCTTTTAAAGAGAAAATAC	49
10	21224	ACCGATTCTTTTACACGACGAGCTGAGAGAGTACCACAGGCTCTTA	49
10	21266	ATTAAAGAAAGAACCTGTTTACCTTAGAATCTTGGATGATGGTAAGAA	49
10	21308	GAATTAGAGTTACAAATCCCAATAAACCTTCTTACTTCTACATCTCT	49
10	21350	GCAAGGCTTACCAACGGTGTCTTTTAAATGGAATATTTGACATTTGG	49
10	21392	CGTAAATATAGTTAGTACGGGAAACAAAATTAACGAGTTATACCTA	49
10	21434	GACTGTAGAACCTCATAGTTCATTAACCTGAGACAATGGATATTAC	49
10	21518	CCAGAGCTTGTATCCCATCGATAGCGAGGCTTTCATAAAACGGTA	49
10	21560	TCAGAAGTTGAAAGAGGGAACATCATAAACCTTCAGAAAAGGTCAAT	49
10	21602	CCAGAAGCAAGGACAGCAATACATAACGCCAAACCTTGACCGGAGA	49
10	21644	TGAGGCAATTTGCTTAGGCTTTTCACTAGTTGAGATTAGATTAAACCTCAA	49
10	21686	AAATCTGTCTGCTTTAAACGGGAAACGAACATTAACGATTCGAGAAAGATT	49
10	21728	TTACCGTCCACAGATAAACGATTATACAGTCAAGAGGTGACCCCTCAT	49
10	21770	GGAGTGTCTGAGTTTGACCCCTTAATCATTTGTAAGTGGTTCAGGAGAG	49
10	21812	GTAACAGTTCAGGGGAGTTTAAAGAGTAGTAATTAATGCTTCGGTGT	49
10	21854	CGGAACCGCCACCTGCTGCTCATTCAGTGAACGGTGTCTTAGCA	49
10	21896	GACTCTATACCCGAAATCATACAGAACCGGATATTTCTGCGGTAGCAT	49
10	51761	TATTAAGAGTGGCTTGTGACACTAACCACTAATATTCATTATACA	49
10	51118	AGATAGGCGATTAAACGCTGATTGAGGATTAGAAAAGACAAAAGGCGTTA	49
10	51160	CGGAAATTAAGGAAGAGGGTGTGCTTAAATCCGTTATATAATTTAG	49
10	51202	CTGGTTTTCAGTACCTGCGGAAACAAAATTAACGAGTTATACCTA	49
10	51244	TTTCAACGGGCTTAGCTATTAAGAGGCGGAATTAAGAACATAAA	49
10	51286	GTTTTCGACGAGCTGAAGCGCAATTCATCAATAAATTTTCTCAACAA	49
10	51328	AACGCGCGAGCTAAAAAGGGGAATAATGGAAGGGGTGAGTGCATAAT	49
10	51370	AGTCGGGCGGTACGATTTTTCAGTGAACAGAGAAATACCTTTCTCT	49
10	51412	GCTAACTGCCACCGGATGAATAACGTCAGATGAAGATTAACCACTCAT	49
10	51454	CCGGAAGTGTAGCAGAGAACAGAGAAAACAAATTAACGATTCATGAATCAT	49
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10	51580	ACGTGGATTTCGAGGCTATCACGAGATGACCCATGAGCAACGTTAAAG	49
10	51622	CTGACCTATAGGAATTAATGATTATAGTCAGAGAAATGACAGAGACAG	49
10	51664	TGAATCGAGCCAGCAAACTAGAGGAAGCCGAAAAAAGATTGAGGAGCT	49
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10	51748	TCGCCCTGGATAGGTTTGAAGAGATTAGAGAGTACCTGGCTCAAGAGG	49
10	51790	CAATGTATCTGCGTTTGTGCTTTTGGGATGCTGCAACTTCAGGAT	49
10	51832	CAGGATAGTCTTCAAGAGTTCACACTGATGACAGGAGTATCAT	49
10	51874	AACCGAGGAAACCAAGAATGGAAGTTCATTCGCAAGCTCCATGTT	49
10	71831	CATCAACGTAAGCGCAATAGACTCTTATTACGC	35
2	71125	CAGTCCGAGGCTGATTTAGTATACATAAAGGTGGCA	35

21601	71167	ATAAAACGGAAGAGAGGGTAGGATAAGTTTATT	35
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21644	71251	TTGAATGATGGCCCATAGCAGGAGGAGGGAAGGT	35
21286	71293	TCGACCACTGATATCCAAATGGAATTATCACCG	35
21328	71335	CAGTAATAACAGGAATTTGCCAGCCAGCAAAATCA	35
21370	71377	GAAATGGCGAGAACTCTGTAATGGGAACGCTACCA	35
21412	71419	AAACGCTAGTAAAAATCTAAGCTAGCGACAGAA	35
21454	71461	TAATATCATACTCTTTTAGTGCGCGTTTCTACG	35
21538	71545	CAAGAGAATAATTTCTTTAAACACCCAGGAACCG	35
21580	71587	CTACAAATTAATTTTTCACGCCACCTCTAGAG	35
21622	71629	CTGATAACGCGCATCGAAGATAACACCACGAGCCG	35
21664	71671	AGACAGTTTTCATCTATGGGAGGTGAGACATTGG	35
21706	71713	AGTAATGCTCCGTAAGTTTTCATTAAGCGCAGAA	35
21748	71755	AAAATTTTCACGTTTACGATTTCAGTAAGCGTCA	35
21790	71797	GTAATACAGTTTGACGTAAACAGCTTGCTAATAGTT	35
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21874	71881	GGCAAGGAGGCAACAGCAACCTATTATCTGAAAC	35
51771	01701	AATCTTTGAAATAACTAAATGAAACAGGCTCCAC	35
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51245	01238	CACATGTAATGAAAGCTACAGACAGCTGATTGCC	35
51287	01280	TAGATAAATCCCAACGCTGTGGGCGCCAGGGTG	35
51329	01322	TACGAGCGAGCTTAGGCCGATTTAATGAATCGGCC	35
51371	01364	TATCATATTATCTCTGGAAGTACGCCGCTTTC	35
51413	01406	CGAGAACAGCTTAGAGTCTGTGCTAATGAATGA	35
51455	01448	TACCGCGCAGGGCGTTGATTACACAACTACAGAG	35
51539	01532	TAACCGAAAAGGAGAAATTTGGGAGGATCCCGCGG	35
51581	01574	GCCGCTTAATAATTTTGTAAACAGGGGCTTAAGCT	35
51623	01616	ATCGGAAGCGGAGTAAACAACTTCTTCGCAAGCT	35
51665	01658	AAAGACTTTTCTGAACTATGAGAAATAGGGGCTC	35
51707	01700	ACGTAATCGATAGGGAACACAGTGGGCGCTGC	35
51749	01742	CAAAAGAACGCTGGGTAGTAACTTATTTACG	35
51791	01784	TATACCAATGACGGGGAAGTGTCAACTTCAG	35
51833	01826	CGCTGACCAACACCGAGCTTGTGTAATTCATGG	35
51875	01868	ACTTAGCCCAACCTGCGCCATGCTTCTCGAGAGA	35
71841	21777	AGTATGTGAACAAACACAGTAATAATTTTAAAGT	49
71126	21119	ACATATAGATAACCATGCGCAAAATCAATCGAGT	49
71188	21161	TGTGCACTTGACCATTTTCCCGGTTAGGTGTGTT	49
71210	21203	AAGACAAAAAAGGAGTAAATTTATCAAAATCAT	49
71252	21245	AAATATTCTGTAATTCAGTACGAGGATAGCT	49
71294	21287	TCACCGAATTTTGTGCTGAGCTGCTTAAATTT	49
71336	21329	CCAGTAGCTTCCAATGAGATAAATCAATATTT	49
71378	21371	ATGAACAGCTACACCAAGATAACCAATTTCAAT	49
71420	21413	TCAGTGTGTTTGAAGCAAGAGATAGTAAACAA	49
71462	21455	GCAITTTAAGAACGCCCAATAGGCGAGAGGGAAT	49
71546	21539	CTCCCTGGCTCATATATTCAGACGACGATAAAC	49
71588	21581	CCACCCTGCGAATTTGCGGGACGAGGCAATG	49
71630	21623	CCGCCAGAGTTTACAGAGGTTACCACTTCAAC	49
71672	21665	CGTGATAAATGAAATTTTCAATATATACAGGT	49
71714	21707	TGGAAAGAGGCTAAGCACTAGGAGAAAATTC	49
71756	21749	TACATGGAATACATACACTTGCATTTTAAGAA	49
71798	21791	TTAACGGAGGAACCAAGCGAGAGATGGTTTAA	49
71840	21833	GTTAATGCTCAGAGTATTTGGCCCTGACGAGAA	49
71882	21875	ATGAAGAGTACCCGGGACCGCAACATCAACGT	49

**histin-modified**  
 CCGTGTGATAAATAAGGCGTTAATTTTTTTT-Bio  
 AGATAGCCGAACAAAGTTACCGATTTTTTTT-Bio  
 Bio-TTTTTTTTTTACGGCCACTACGTGAAC  
 Bio-TTTTTTTTTTAAAGGGAATTTGA

**digoxigenin-modified**  
 CAGCTGGCGAAGGGGGTTTTTT-Dig  
 CGCAAAATGGTCAATAACCTGTTTTTTT-Dig  
 Dig-TTTTTTTTTTTCGAGCAAGGGCATAGGCTGGCTGAC  
 Dig-TTTTTTTTTTAAAGTGGCGCTGAGAGGGTTGATA

# 10 helix bundle

Start	End	Sequence	Length
[131]	[838]	CCACTACGTGAACCATCAACCAATGGGATATACCAAGCGCG	43
[1385]	[8399]	AGAAGTGGCGGTTTATGTCAGGACGCACACCAAGCGGCTTTA	42
[955]	[638]	AAAGAACCCCTGACCAATAAATAACCA	32
[969]	[976]	AAGAGTCTTTGGGGTCGAGGTAAACATCTTCCATG	35
[990]	[997]	AGTGTGAGACATAAATCCGAAGAAGGACGAGCGG	35
[1111]	[9118]	GAATAGCAGGGAGCCCCGTAATAGGCACTAAGGGA	35
[1132]	[9139]	AAAACTCTTGACGGGGAAGCAAAATACCTTTGAA	35
[1153]	[9160]	GATGGTGGCTGGCGAGAAAGGGGAGTTCCGGTGA	35
[9174]	[9181]	CAGCAGGGAAGCGGAAAGGAGGACATAAGGCTGG	35
[9195]	[9202]	CGCGCTTAAGCGCGCTGGCAACAAAGCGGATAT	35
[9216]	[9223]	CGCTTGAGCTACCGCTGCGGACGAGATGATATC	35
[9237]	[9244]	CTGATTGCAACCCCGCGCGCTGACCCGTGAAC	35
[9258]	[9265]	CACCAGTGGCTGTAAGGCGTTAAAGCGGTGAATA	35
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[9279]	[9286]	CGCGAGGTGGTTGGCTTGACGCTCGGTAGAGAAAC	35
[9300]	[9307]	AGAGGCGATAACGTGCTTCCACGCGATATAATG	35
[9321]	[9328]	TGAATCGAATCAGAGCGGGAGCAATGACTAATTTC	35
[9342]	[9349]	CTGTGCTGGAGCGCATTAATATGATCTGAAAT	35
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[9489]	[9496]	TGAGGATCACTTGCTGAGTAAAGCGGAGAAATGGAA	35
[9510]	[9517]	CGCGTCACTAACTATGCGCTTAAACAATTTACC	35
[9531]	[9538]	ACTCAATAATAATCCAGAACATTTCTTCATACAT	35
[9552]	[9559]	AICTAAACGCCAGCTATGCAATTCACGAGAGT	35
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[9594]	[9601]	CGTCAGTTCTATTGACGCTCAACAGCCCGAGTAAC	35
[9615]	[9622]	CAAAACCCGAATGGATATTAATCAACGCTGATCTT	35
[9636]	[9643]	ACCTTGGCAGATTCCACCGCTTCGTCAATTCGAA	35
[9657]	[9664]	AAATGAAACAGTAATAAAGGGGCACTGTAACATGAA	35
[9678]	[9685]	GCCACGGCGGCAACAGAGATAGATGACAGAGACTCC	36
[9699]	[9706]	ATTAACATCTGACCTGAAAGCGGCACCATAGGATT	35
[9720]	[9727]	ACAGAGATACGTGGCACAGACTCAGAACAGTACC	35
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[3155]	[434]	GTCTTAAGAGATAAGAGTACGATTTTGTAGAGA	36
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[3148]	[4742]	ACAGTAACCGGATCGATAGGA	21
[324]	[212]	ATTGCTGAGATGCTGCGGCAACAGTGGTGA	21
[563]	[3169]	CGGTAAATCGACGATCTATGA	21
[5525]	[31531]	GCATTTGCAAAATCATATCA	21
[469]	[863]	AAATTAAGAGGAGGAAACACCGCATGATCTTTG	35
[490]	[884]	TATTCACAAATATCTCAAAATCGCGGAAAAGAT	35
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[4132]	[8126]	GGGTGAGGGAAGACAGTCCAATGACCAAGATATG	35
[4153]	[8147]	TGTGTAGAGGATTAATAAGGTAGATGAATCCATT	35
[4174]	[8168]	TTAAATGTTGCTCTTTTGCAGGCGCATAGCATTT	35
[4195]	[8189]	TTTTAGCATTTTGAAGGCTTCATCAATACAGAG	35
[4216]	[8210]	ATTTCAAGCTTAATACGATAAAGAACCGGGAACG	35
[4257]	[8231]	ACTTTTGGCTAGCATAACCAATCAACATGAGA	35
[4258]	[8238]	AAAGATTAATGCAATGATGATCTACGCTTTT	35
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[4300]	[8294]	AGCAATAAACAGTTTTCACCTCGAGTAGACGATA	35
[4321]	[8315]	GGCAAGCGCAAGCATTTGAGATGGTTTAAACACC	35
[4342]	[8336]	TCCAATAACCATTAACAGGATATCATTTGGATGTT	35
[4363]	[8357]	CTAATAGATGGTCAAACTAACCGATTTTCTTTA	35
[4384]	[8378]	GAAAGAGGTATATTAAATCTTTACCATCATGCTT	35
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[4426]	[8420]	AGGGAAAGTGAGCGCAGAGGCAACGAGGAAAAAA	35
[4447]	[8441]	CAGAGAGTAACCCACACCTCGACGATTTTCTTCA	35
[4468]	[8462]	AAATGAAAGCCCAAAACCGCCCAACCAAAAGGAA	35
[4489]	[8483]	CGATTTAAACATGCGGAACCAAGCCAGTGAGAT	35
[4510]	[8504]	TATCCCAATCTTACTCTTTCTCTCTGACTTCAA	35
[4531]	[8525]	AAATTAAGAAAGGCGCCCTTTAAGCGGTGATAGG	35
[4552]	[8546]	AAAGCTTAACAAAGCTGCGCATGATAAGCTGTT	35
[4573]	[8567]	AACGCTTACGAGGACGTCAGATAAAGAAAGTTT	35
[4594]	[8588]	AATTTAGGAATACAGAGATCGTGCTTTTATAGT	35
[4615]	[8609]	TGCTATTATGATACAGCAGCTTAAAGTAGCAT	35
[4636]	[8630]	AAGCCTTGCAGTATACCAATTCGCTATCCAGTAC	35
[4657]	[8651]	CCCGACTAAAATACCAATAGCTTCTGACACCGTAAC	35
[4678]	[8672]	GCGAGGCAACATAAATCACAGAGGCTAGCCCAA	35
[4699]	[8693]	CTTATCCAGACACCTTTGGGAGAGGATCCTTCAT	35
[4720]	[8714]	AAGCAAAATTTGTCACCCGCTTTTGTCTCCGCCAC	35
[4741]	[8735]	ATCATTAATCTATATATTCATTAAAGTGCTCAGAAC	35
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[798]	[3111]	TAACTGCAACATCCCGGTTGATTAATGAAGCCAGCTGGT	42
[7119]	[3132]	CAGACCAAGAGCGGCTCAAAACAGGAGCTGGCGGCTTT	42
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[7161]	[3174]	CCTTTAACTATGCTTAAAGCTTAATTAACGGCACGAACT	42
[7182]	[3195]	AAGAGGTAAACCTCAATTCGCAATTAATGCAAGGGGTAA	42
[7203]	[3216]	GCTTAGACGCAAGGAATCAGCTCATTTGCTGGCGTAAGTA	42
[7224]	[3237]	TATAATGCGGGGAGAAATAGGAAGCCCATGGCTCTCTCGAT	42
[7245]	[3258]	GTTTAAATAGACCAATTCGGGCTTGGCTTGGGAACCCGGGT	42
[7266]	[3279]	TACGGTGAATCGGTTAGCGAGCTTTCATGCCAATTCGTAATC	42
[7287]	[3300]	TCCATAAAGCCTCAATGTGAGCGAGTAAACAGTTCTCTGT	42
[7308]	[3321]	AATCTGAAATAGCGTTCGGATTTCCGGGGACCCCGCTCAC	42
[7329]	[3342]	TAGTTTGAATCAATAAACCGGCGGATTTGAATCGCATACGAGC	42
[7350]	[3363]	TTGCAATAGTAGCGGATAGGTCACGTCAGAGTAGTAAAGC	42
[7371]	[4385]	GTTTAGCTGGCATGATGGGCGCATCTGAACCGACGAGGCT	42
[7392]	[2392]	GGCACCTTGAACAAAGCCACAGAACTTTTGGGACACCGGA	42
[7413]	[3426]	GGGTAACTGATATATGATCAATATGCGATTAATCGGAGAT	42
[7434]	[3447]	AGAGCAAAATACAAAATCTTACCAAGGTTTATGATATAT	42
[7455]	[3468]	TGAGTTAAATAGCACAGCGCTCAACGATATCTTCTTATCA	42
[7476]	[3489]	AGCAAGATTGTTTAAATGAGATCGCAAAATATTTGGAT	42
[7497]	[3510]	AATAGCTATCAAAACAAAGCCCAATAGAGAACGCATGGAG	42
[7518]	[5524]	CTTTTAAACAGCCAGGAGAG	21
[7539]	[3553]	ATAGCGAATTTGCGTAAATAAGAGAAATAACTATAAAACAG	42
[7560]	[3573]	AAGGAAAACGAGCGCGCAACAAAGGTAAATCCGGCTGATAGAT	42
[7581]	[3594]	TAATAACTCTGGAATCTGCCAGACGCTGAGAGATCAGATG	42
[7602]	[3615]	AACCTGGCTTGCACCAACAACTGTTCAGCAATTTATGACCTT	42
[7623]	[3636]	TTATTACAAATCAAGAACGCGGCTGTTTCGCTGAGACAAATA	42
[7644]	[3657]	AACGTAGTGGCGGAATAGATAAGTCTGTAGCGATTTGCTTTG	42
[7665]	[3678]	AAGGTGGGTTTGAAGAAATAATCCCACTTTAGAAATCGC	42
[7686]	[3699]	AACGCAAGGTATCTTACGAGCATGATGCTTCGCTTCATTT	42
[7707]	[3720]	AGGTTTATCAGATATCAATAATCGGCTGGAAATAACAAAGAG	42
[7728]	[3741]	ATAGAAAAGCGGCTTATCATTCACAGCATAAATATCAAGA	42
[7749]	[3762]	GAAGGTAAATGGAATATTA	23
[944]	[256]	TAAATGTGTCGAATCCGGAGAAATGGAGTGAACCTTAAGT	40
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[998]	[491]	CAGACGGTTTATGAGCACTGGGTGGATGTTCTTATATGTAATATGA	49
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[9140]	[4133]	AGAGGACTTAGACTCTGACGCTGTAAACAGCGGCAAGATTTGTTCAAA	49
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[9182]	[4175]	CTGACCTTTTGCACCTGGTTCGATTAAGTGGGTTTGTATAATATAT	49
[9203]	[4196]	CTTGACAAAACCAACAGGGCTAAAGGGGGATGTGCTTTTGTATAAAAA	49
[9224]	[4217]	ATTACCTCGTTTACTTGTATCGCTATACGCCAATTAACAGCGCTTT	49
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[9287]	[4280]	ACCAGAAAATGCAAGTAGCTGTGCAAGCGCCATTCACCAATAGAGCAT	49
[9308]	[4301]	GGCTTGATTAGGAATGTATTCGCTTCTGGTGGCGGAACACCAAAATTA	49
[9329]	[4322]	AACTTTAGAAAGATCACAACTCCAGCAGCTTTTCGTATTTAAGAGCAA	49
[9350]	[4343]	CTTTAGGGAACAAATAAAGTTGCGGCTCAGGAAGCGTAAATTAACA	49
[9371]	[4364]	GGCTCATAGTTAAGCTAAATTTGAGGGGACGACGGTGTAAATCTGA	49
[9413]	[4406]	AGCATTCACACCGGCAAGAAAAGGCTTTAAATAAAGCGTGTGACGGGA	49
[9434]	[4427]	CAGGTCAAGAACCGGATTTTCAATACCGACCTGGTTTATACATAAAAC	49
[9455]	[4448]	ATATTCATCCCTATGGCAATGACTTAAATTTAATAAAGCGGCTTTA	49
[9476]	[4469]	CTCATTAAGAGGCACTGATTTGTTTATTTAATTCAGGGCTTACGTCAA	49
[9497]	[4490]	AGCGAGATAACTCACTGAATAGAGAAAACCTTTTCAATTTAAGAAGAA	49
[9518]	[4511]	GTTCAGATTAGCGACCTACCACATCGCAAGACAATAATATATATAT	49
[9539]	[4532]	GGCTTTATTTTCGTGCAAGTATGTAATGCTGATCGAGGCAAGCTTAC	49
[9560]	[4553]	GTACTGGCTGTAGCGAAATTTGATGTTGGGTATATAAAGATCTTCCC	49
[9581]	[4574]	GGGGTCAAAGTTTGTTTTAAAGCTTACCTTTTAACCAAGTAAATCTTACC	49
[9602]	[4595]	AGTGCCCGTAATCAAGTAACACAAATATCAGGTCACATAACACGCTAC	49
[9623]	[4616]	TGCCCCGAAACAGGGAAGAAAGAGTCAATAGGTGAATACAGAGATTAG	49
[9644]	[4637]	CTTATTAAGGGCGGCTCATGAGCTAGATTAAGAAATCAACAGGTTTGT	49
[9665]	[4658]	AGTATTAAGTAGCAAGTTACAAATCTTGAAGAACCACTATCTCAACAT	49
[9687]	[4679]	TCAAGAAATAGAGGAATATATAATTAATTAATTTCTCAATTAAGAAC	48
[9707]	[4700]	AGCGGGGACGACTTGAGCACTTGCTTCTGTAAGAAAACCAATAGAGG	49
[9728]	[4721]	AGCGGATAAAGGTACAACAAATATATGTAAGTCTTTTCCAATAGC	49
[862]	[956]	ACCCCAAGTTTCTCAAT	21
[883]	[776]	ACACTAABCCGTAATTCGATTGTGGAACCTGACAGCAGTTACGCCGA	49
[8104]	[797]	AAACGAAACCTTAACCGATAGAGGTTGATCTGTAATAATCCCGGCTT	49
[8125]	[7118]	CACACGTTAGAGCTTATAAATCAAAAGGCGACGATGCTCAAGCGAA	49
[8146]	[7139]	AACGGGTGCGGGAAGTTTCCGAAATCGCGGAATCGGGGATAGCACTCCA	49
[8167]	[7160]	TTTATGAAAGGGGAACGAAATCTTTAATGATGAGAGTA	49
[8188]	[7181]	GCTTTGACGGGCGACGCTGGTTTGCCCTGACCTCAAGAAGATTTTGTAT	49
[8209]	[7202]	AGGGTAGGTGTAGCGAGAGTTGCGCAAGAGTAAATAATGCGGGCGATG	49
[8230]	[7223]	CGCAAGTAATACCAACCTTCCACGCTCGGTGGTGGCAGAGCTGTGAA	49
[8251]	[7244]	GCGGGATTTAATGCGAGAGCGGCAACAGAAAGAGCACTATCTCAACAT	49
[8272]	[9266]	AGGGAGTCGACTAGTGGTT	21
[8293]	[7286]	TATTCGGAGACGCTGTTCGGGTATGGGATGGTCAATACATAGTTTAT	49
[8314]	[7307]	ATCGGCTGCTTAGGCAAGCGCGGGGGTGAATAATACACAGATTCCTC	49
[8335]	[7328]	CGCGGACTTAAACGACGCTGCTTAAATTCCTATCTAGGTAGAT	49
[8356]	[7349]	AACAGCTGGGATTTTCCAGTCGGGAACCGGAAGCAATTTATGATACAT	49
[8377]	[7370]	GCTTTCGACCGAGCTGGGCTGCTGCTGGGGTTGAGCAACGATAACCT	49
[8398]	[1384]	ATTGATTTTATACATATATCATACATTAATGGAATCTG	42
[8419]	[7412]	AGGCTCCCGAGTATTAATTTAAAGTAACCACTCCAGAGCAGTCAGA	49
[8440]	[7433]	CGTTGAATCAGCAAACTCTTTGCGCGTATCATCTCCACCTTAATATC	49
[8461]	[7454]	TTGCGAAGCAATACAAACATTCGACCAACAGATAGAGGCGCGCAAGAT	49
[8482]	[7475]	AGAAAGGATAACATTTAGAGAAATTAGATATTAATCCCAACGGTAATAG	49
[8503]	[7496]	CAGTTTCGAAGAAGCATAGATACATTTATATCAATCAACCAATAGC	49
[8524]	[7517]	ATTTTGCTTGTGGACTAATAGATAGTAGAGGTAGATTGTCGACGAGGCC	49
[8545]	[7538]	TAAATGAATATTACATATCTTTAGGAGCAATTTATGTCATAGTAAGCAG	49
[8566]	[7559]	TGTCGTCACAGGAAATTTGAGGAAGGTAAATAAAGCGTTTATTACAG	49
[8587]	[7580]	TAGCGTAATACCTTAGGCAATCAACGATTTTCAGGCTTTAGAACCGCA	49
[8608]	[7601]	TCCACAGATCGTCTTCAATCAATATGATATACGTAGTACGACCAAGAG	49
[8629]	[7622]	AAACTACACATTTGGTGAACCTCAAAATTTTACATCTCGATGACAGCACTC	49
[8650]	[7643]	ACTGATACAGGACAAATCTAAAGCATTCGGATTCGAACGCTGTAGCA	49
[8671]	[7664]	TAGCAACACATCTTGAGAGGACAGCAAGCACTACCACTACATACATA	49
[8692]	[7685]	TTTCAGGGAACCTCCCGCTGCAACAGCTCAGAGGCGAGCAATAACATA	49
[8713]	[7706]	CTCAGAGTAAGAAGTAGGCGGTCAGTCAATTAAGGCAACGGAAAT	49
[8734]	[7727]	CGCCACCAATATTTCAGGACAGCAAGATAAATGATGAGAATTATACAATCA	49
[8763]	[9748]	AGGAGGTTAGTACCGCCACCGGCTCGAG	29

biotin-modified  
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ATTGTCAACCTTATGACAAATGTCGCCGCACTC-Biotin  
Biotin-ATTCTCATACCTTAATATATAGTCAGAGCAAAAGC  
Biotin-ATATCTAATCTCTGAGATTGTGATATCGGCTCGA

digoxigenin-modified  
Dig-TATTACCCACCTTACCTTAAACATCGCCATAAAAA  
Dig-TACCAAAATCTGATTGGAATACCTTTTAAATGGA  
CCAGCGCAAGAGCAAAAAGGGGCGATTACTCTTCTA  
AGGGTTGATATAAGTATAGCCCGGAATAGGACTCTCT-Dig



## 12 helix bundle

Start	End	Sequence	Length
0[69]	5[69]	CGTCAAAGTGGCGTTATTACCTGAAAGGATGTTAGAAT	42
0[321]	5[321]	GGGTATTAGCTAAACGGGTACCGATTGGTATCTAAATATATG	42
0[342]	11[356]	ACCGCGCAACAAGCGGTATTAAGTAAATCAACAGAGCGCCG	42
0[594]	5[594]	CTGGTGGCCATCTAAACGGGTCTAAATCATCAAGCGCAAGC	42
0[622]	5[615]	GTGTAATGAGTAAAGTGGCGCGTGGGAACCAATATACTAATGACCAT	49
0[639]	1[650]	CTCTGACCTCTTGAAATGGTGGTGACGATTT	31
1[131]	11[62]	CCATCACCACAAATCAAGTTTGTGGATTATTCACGCAAAATCACCA	46
2[76]	11[83]	AATGGATAAAGACCGGAAGGTCAATACCATTAGCA	35
2[328]	11[325]	GACAGGACAGAGAGAGCGGCAACAGAGCGCGCACCA	42
2[349]	9[349]	GGGAGTGTCTGGGAATCAACCGTCTTAAGGACAGAAACAAC	42
2[601]	11[608]	TGGGAAGCTAAGCAATAAGTATTCTTGAACAATG	35
2[629]	11[623]	AGCGAGTAAACAACCGCAATAAGTGGCGTGAAGG	35
3[42]	8[39]	CAGAGATAGAACCCTATCTGAATTTACCA	31
4[629]	9[629]	GGTGGCATCAATTCCGGAAGCAAGAGCTGTCACTCCAGAG	42
5[42]	6[39]	TTGCACGTAAACACGACGGTGTGATAAATA	31
5[70]	2[77]	CGAATAGAAATTTATGGCTTATACAAAGTTGCTAATACGTGCTGTA	49
5[322]	9[328]	TGAGTGAATAACATAGATTGAGGAAGGTCAATATCATACGAGCAATGAGC	49
5[343]	2[350]	AAGCCCGGATGTACAGAGCAGGCGATCAAAAGAGCTTCG	49
5[595]	2[602]	AGTAGATTAAAGAGCACTAACGGAACAACAGGGAAGTTTATTTATCTCCG	49
5[616]	7[629]	AGATACAAAAGGACAGGTATTAAGAAAGAT	28
6[83]	9[76]	CTGACCTTTTCAGATCAATAATATCTCGTAAGATTTTGCAATCCCAA	49
6[328]	6[329]	TTCTGTACCATCTCAAGAGCAGGCGATCAAAAGAGCTTCG	42
6[356]	5[342]	CGAAAGCAAGAGCTATCAACCACTAATGATATAGTAAGAGG	49
6[608]	9[601]	GGCATAGTTAGTTAGTAGTAGGATTAACAGCTCATTTCCATAAAAGGCC	42
6[636]	5[639]	AGATACAAACGCGCTTCGCAAAATGGTCAAT	31
7[49]	2[49]	AAAGCGCTGTTAGACAAATTTTCTGACATGGGAGATTAC	42
7[63]	4[56]	TATCATAATGGTTTGAATACCAAGATAATGGATTA	35
8[636]	3[639]	AGGCTTTGAGGACTCTCAAAAATAATTCGC	31
9[49]	0[42]	AAATAAACAGCCATGACGGAAGGTGGAGGGCGGAAAAACCGTCTACAG	49
9[77]	0[70]	TCCAAATGAGGGAGTAAATCAAGCGGTGATCTCAA	35
9[329]	2[329]	AGTGGGATTTGCTATGAACGGGTGTATTACAGTGGATTITA	42
9[350]	0[343]	TTTCAACAGAGCTTTGGGTAAATGAATCGGCCA	35
9[602]	0[595]	GCTTTTGGGTGTAGCTCTGTGACAGGCGTTAAGCTA	35
11[39]	10[28]	GAATTAGAGCATTAATAGGTGAATTAATCAGC	31
11[63]	8[63]	GTAGCAACAAATATTATTTTCCACGT	28
11[84]	6[84]	AGGCGGACCGGATTAAGAAACAGGATTATCTTACCTCATCTT	42
11[336]	0[322]	GAAACCAAGTAATTTCTGTAAATCCGATTAGGAGAGGCGGTTT	42
11[357]	6[357]	CCAGATCTTTCCAGTTTCAACCACTCTTCAATCGTCAGAA	42
11[609]	6[609]	AAAGATTGAGGAGCGGGATCTTCAATATTATTAATTACGA	42
11[623]	0[623]	CTGACACTCTCAAGTACACGCGGATATAGCGGCTGTGTTG	42
0[90]	5[90]	ATTAAGGAACCTCTGCTCAATGACAGAGCAATCTGCTTAAC	42
0[132]	5[132]	GATAGGGCGCGCGTTCGCAAGCACTGAAAGGAGCGGGAGA	42
0[174]	5[174]	CGAAATCAGCGGGCGGCTTAACCACTTTGAGTAAGTTAC	42
0[216]	5[216]	TGGTTTTCGCTAACTAGTAATAACCACTCGTATCTCGAGC	42
0[258]	5[258]	TCACCGCGCGGTACGTCATCATGAAATGAGGAAATTAAT	42
0[300]	5[300]	TTTTCTCTCTGTTAAGTTTAACTCCCACTAACTAGGAAAT	42
0[384]	5[384]	CTCGGGAACGAGCTGCGGGGCCATTTTAAAGGTTGGAGCTTC	42
0[426]	5[426]	CATCTCGAAGCACTTCGCAAGGCTCTTTTAAACAGGATT	42
0[468]	5[468]	CGGAAGCGCAGATTCGCGCAACATGATTATTTCTCATTTT	42
0[510]	5[510]	GTGAAATGCAATGAGGACGAAAGGCGCAACAAAGCGGTAG	42
0[552]	5[552]	ACCAGGCGCTCGCGGATGGGCTTGAATAAGCAAGCTCTGGA	42
2[97]	11[104]	TTTTCGAAAGGAAACATCTCAAAACGTCACCAATG	35
2[139]	11[146]	CCAGCAAGAGTGGGGAATAATAGGCAACAGATCA	35
2[181]	11[188]	AAACTAGCTTAGCGGCAAAAGCGCTTTCATCGGCA	35
2[223]	11[220]	CTTTGATACCACACCTGAGATTAAGTCACTTTTCA	35
2[265]	11[272]	AGAGTCTTATGGTTCTGGGATCAACGGGAACCGCT	35
2[307]	11[314]	TCTTGAGAGATAACGGAACCCACTCTCAGAGCA	35
2[391]	11[398]	GATCGGTTGTAAGAAAGTTCAGGCGACGAGTGTGCT	35
2[433]	11[440]	AGGCGCAGGTGGGACCAAACTTAAAGCCAGAAATGG	35
2[475]	11[482]	CGCAGCTCTTAAGTAAATAGGAATGAAGCGTCAATC	35
2[517]	11[524]	AGGGGACTCCGCCACCCCTAGGATTAAGATTTTA	35
2[559]	11[566]	TGGTGTACTGGAGTTTGTAGTACCGTATAAAGCTT	35
5[91]	2[98]	GTCAGATGTTAAATAGTATAAAGCAAGCTGATTTGAATACCTACA	49
5[133]	2[140]	AACAATAGCAAGCGCCATATTAACAAAGCGGAGAAACATATTACCG	49
5[175]	2[182]	AAAACTGTTGGTATTTCTTGAGGCGTGAAGCAGATATAAAGAACTC	49
5[217]	2[224]	AAAGGAATCATAGGTAAAGTAAATCTGTCGTTTACAGTGCAATACTT	49
5[259]	2[266]	TACATTAAGATTAAAGCTAATGAGCAAGCGGTATGATACAGAGTAAA	49
5[301]	2[308]	CAGTACAAATATCTGACAGAGAAATCAACCAATCTGGTGGCAGAGA	49
5[385]	2[392]	AAAGGCAATATCAGCGGATATCAATACAGGCGCATCAACAGAGGCG	49
5[427]	2[434]	AGAGAGTATGCTTTTCAGTAAATAGGTTGTGCAACAGAGCAGGCA	49
5[469]	2[476]	TGCGGATATCTGCAATAAATGGGCTTAACAAAGATCATACACTCA	49
5[511]	2[518]	CTCAACACAGAGGGGTGTGAATTACCTTAAACACAGGAAGACAGTTTG	49
5[553]	2[560]	AGTTTCAAAAACCCAGTACGAGCGTTCGCAAGGCAATTTGTGTACGT	49
6[104]	9[97]	TATTTTGAATAATATATATGATAGATGCAATATTTTAAATCGATTTT	49
6[146]	9[139]	TCCAAATCAGGATTAAGAAACCAACAGTACGCTCGGTTTGAAGAGA	49
6[188]	9[181]	CTTAGGTCGACAGATTAATTTTAAAGAGCAAGAAATCAGAAATTAAC	49
6[230]	9[223]	ATCAAAAGATGATGAACCAATTCGCAAGCGTGCATATTTTCGCTAATA	49
6[272]	9[265]	ATAGCTTAAACAATTAATAGATAATACATATCTAATAAACCAGAAATA	49
6[314]	9[307]	GCTATTATAATCAAAATCTTTAGGAGAAATCAATCAATAATACGGAAG	49
6[398]	9[391]	AAATCAAACAGAGAGGTAAGATTCATAGAGAGAGAGCGTCAACACTA	49
6[440]	9[433]	CCCTCAAACTTTAAGAACCTCTATATAGGAGCAATCCGGATCTCCA	49
6[482]	9[475]	GCCTCAGGCTTAGTGGCGGAGAGAGCTCATGTCATACAGCGGTTTAT	49
6[524]	9[517]	GTTCCTGCTGTTTATAAATCGGTTGACCAAAACCTCATCTGTATACCG	49
6[566]	9[559]	CGACGATTTCCATAGAAATAGCAAAATGTTTAAATCAACACCGCATAA	49
9[98]	0[91]	TGTTTAAAGGGCGGCGCCCGGAACAAAGAGTCCACT	35
9[140]	0[133]	ATAACATAATCAATCGAGAAAAAGATAGCCGA	35
9[182]	0[175]	TGACACAAAGAAACCGTGGCTTTGATGGTGGTTC	35
9[224]	0[217]	TCAGAGATAGCAAAACCGCGCGCAAGCGGTGCACGC	35
9[266]	0[259]	AGAGCAAAAAGAGCTTTGACAGCTGATTTGCCCT	35
9[308]	0[301]	CCCTTTTACCAAGAGAGCGGGGGCGCCAGGCGTG	35
9[392]	0[385]	AAGGAATCTCATACGAGCGCTTGCCTGCTTTCCA	35
9[434]	0[427]	AAAAAAACACGAGTTGCTTTCGCTTAATGAGTGA	35
9[476]	0[469]	CAGCTTGAAGCCGCTCTTACACACATACGAGC	35
9[518]	0[511]	ATAGTTGAACCGCGCAAAATATAGCTGTTTCTGTT	35
9[560]	0[553]	CCGATATCAGGAGGGACTCTAGAGGATCCCGGGT	35
11[105]	6[105]	AAACATAAGGCAACGTCGAAGAGGTTTGCCTCACTCAAAATA	42
11[147]	6[147]	AGTTTGTGTCACAAAACATTAAGAGCGCAACATGCAAA	42
11[189]	6[189]	TTTTCGACATACTGACCCCAATATAAGAGACTCTCCG	42
11[231]	6[231]	TAATCAAAAGTATGTAATCAACGAAGCAGACGTGAATTT	42
11[273]	6[273]	CCCTCAGATAACCGGAACCAACGAAGCGCGCTGATAGCG	42
11[315]	6[315]	CCACCTCAAAAGTTTAAAGAAATGTAGATAATATCAATCGTC	42
11[399]	6[399]	TGATATTAGACAGTGGCAATCGGAACGCCAAATCTGACCAT	42
11[441]	6[441]	AAAGCGCGTCTTCTGGGCTCCATAAATTTGTCCTGTGAATCC	42
11[483]	6[483]	ATGGCTGGGATAGCTTTCGAAGCGCGAGAGATGGCTGGATA	42

11[525]	6[525]	ACGGGGTCTCCAGCGCGGACATACACTTGGGATTAAGAA	42
11[567]	6[567]	AATGCCCGGTACTATTTCGGTGCACATAGGAAGATACCCAGA	42
0[111]	5[111]	AGTTTGGATTAGATGGAATGGCTATTATCTCTGACGATAC	42
0[153]	5[153]	TAAATCAGGAAGGGGAACAATCGCATTTGCGGAACCGCTGA	42
0[195]	5[195]	AATCTTGAAGTGTAGAGTAGACAGAGGTGAAGCTTGGCAAT	42
0[237]	5[237]	GTTCAGGCTTAATTTGAGTCAAGCTGATTTTACAAACAAA	42
0[279]	5[279]	CGGCAAGCAGACAGCGCCAGCTTCTGAGCGCTCTCAATTG	42
0[363]	5[363]	GCTGCAATACCGCAGGAGTGAATTAATGAGTCAATCAATA	42
0[405]	5[405]	GCCTCAGTGGCCCTTGTGAGGCTATAATGTTGTCGGAAGC	42
0[447]	5[447]	CTGGGGTCTAAGTGGGGAACCAATGATATTTTATGCTC	42
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0[531]	5[531]	ATGGTCACCCCGCTCATCTGCTGTATATAAAGCAATAATGC	42
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2[118]	11[125]	ACGCTCAGCTTACAGCGCCAGCATAGCAGCAGCG	35
2[160]	11[167]	ATATCCAAAGAAAGGTTTATCTTTAGGCTGACAG	35
2[202]	11[209]	CTTGCCCGCGGTGAGGTGACATATAGCCCCCTTA	35
2[244]	11[251]	TTAACCGCGCGCGGATACGCAATACCGCCAGCA	35
2[286]	11[293]	TCAGTGAGTATAACATAACGAGGCGCGCACCTCA	35
2[370]	11[377]	TATTACGGGGTTTTTTGTGCTGACAGAGGTTGA	35
2[412]	11[419]	TGCGCAAAAGCTTATTCACCAACAAACAAATAAA	35
2[454]	11[461]	CTGGTGGCTTGTGACACTGAGAGCTTGAAATTTA	35
2[496]	11[503]	CCTCAGGATTTGCAATTTCTTGAATGATACAGGA	35
2[538]	11[545]	AACCGTGCTATACCGCCACAGGCTGCTGATTA	35
2[580]	11[587]	CCGTAATGACAGTGTGATTAACCTGCTTATTCGG	35
5[112]	2[119]	AGTACCTAATCTTATAGTAGGGCTTAATCTCCCGAAATGCGCAGGAAAA	49
5[154]	2[161]	TTGCTTAACTGTGATGATTAATTAGGAGGCTTAATCCGAAACCGGTGTA	49
5[196]	2[203]	ATTCATTTTTTAAATAAAGTACCGAGATATCATTCAGTATAACATCA	49
5[238]	2[245]	CATCAAGTCAATAGACAGCAATAAACCAACTCATGACGAAACGCCAA	49
5[280]	2[287]	AATTACCTTGAAATTTATCAACAAATAGCTTTCTTAATCATTTATAA	49
5[364]	2[371]	TCGGTTTATTAATAAGAGTAACTTGAAGGGAACAGGATAGTCTTCGC	49
5[406]	2[413]	AAACTCCAGGAGAAAGATCAACAAAGCTCCAGTTTTCGCTGTTCAGGC	49
5[448]	2[455]	CTTTTGATTTTCAATACGAGGAACACCGAGTATCAATGCTAATCGTTCG	49
5[490]	2[497]	TTGCTGAGTTAGATTAAATTTCAACTACGAGATAACAGAGTATCGG	49
5[532]	2[539]	AACATAAACTTTTGTAAAGACTGCTCAAGAGGATTAAGAATCATG	49
5[574]	2[581]	TGATTCCTCGGTAAATCTAGCTTAATGTAATAAATTTTGGATGTA	49
6[125]	9[118]	CGGAGAAATACATGGAATATCATATGAGTGTCTTCTGCGGAATAAA	49
6[167]	9[160]	ATATGTAAGTAAACCAATATCATTTTAAATACCGGTATGGGAAGC	49
6[209]	9[202]	GACTACCTCAATATAAATTCGTTTGGCCGAGCGCGTACCGGAAAGTCA	49
6[251]	9[244]	AGAGAGAAACAAATTAAGAGTATTAGAGGACCGAGAAACCAAGTA	49
6[293]	9[286]	AGAACTCTTTTAAACATAAGATAGAAGCTCATATCTTTGAAATA	49
6[377]	9[370]	CCCTGACTTAATTCAGAAAGCGCGGAGCGCGGAGCAATCGCGGAGT	49
6[419]	9[412]	TCAGAAAAACAGGTTGCAATGCGCGAGTACGCTCAATATGCAATAAT	49
6[461]	9[454]	TCATAAATAAGAGGAACCAAGGATTAAGAAAGCTGCGCTGAAAGGAG	49
6[503]	9[496]	GTAAAAATATATAATATGATGACCTGTAGGTTGATATACAGGTTGAT	49
6[545]	9[538]	CGAGAGGGTACGGTTAAAGGCTCAGAGCAGCAAAATCAAAAGAAATGAG	49
6[587]	9[580]	TCATAACCAATCTTAAATCATATAGGCGCCAGTAACTGTAATCGTGTAG	49
9[119]	0[112]	ATAGCAGGTTTACCGGGGAATTTGAGTTGTGTTCC	35
9[161]	0[154]	GCATTAGGGAATAACGAAAGGGGCAAAATCCCTTA	35
9[203]	0[196]	GAGGGAATACATAACGCTGCGCGCTACGAGCGGAA	35
9[245]	0[238]	ATTGAGTACTCTTTACAGGCGCTGGCGTGGAGAGA	35
9[287]	0[280]	GCAATAGCGAATATAGCTTTTTCACCGGTGAGA	35
9[371]	0[364]	GAGAACTAATAAGTCCAGCAACCTGCTGGCCA	35
9[413]	0[406]	TTTTCACTGTAGGCTCAGGAACTAATTTGCGTT	35
9[455]	0[448]	CCTTTAATACCGTAATCTAGTATAAAGGTGTAAGC	35
9[497]	0[490]	TTCTTAACACCTTACCTTATTTGTTATCCGCTCA	35
9[539]	0[532]	ACAACTACCTAGATATTTACTCGAATTCGTATTC	35
9[581]	0[574]	GCTTGCAGAAATAGGCGGCCCTTGTGTTACCTCGAT	35
11[126]	6[126]	TAATCAGTCATATGCTTTTACAGCAAGCAGAGATAACAAAGAC	42
11[168]	6[168]	TGTAGGCAACACACAGCGGAGTATAGAAAGGCAATATAACT	42
11[210]	6[210]	TTAGCGTAAATCATATTTGAGGATCTGAGCAAAAGGCTGTGAGA	42
11[252]	6[252]	GAGCCAGGATTAAGTAAAGCCAGTACCGCATGTCGAGCGTG	42
11[294]	6[294]	GAACCGCGAGGAAACTTCTTCGGGTGTATAGTCTTCCTT	42
11[378]	6[378]	GGCAGGTAAAGTAAAGAAAGAAATCATACAAAGACGCTTTA	42
11[420]	6[420]	TCCCTACACAGCGGTGAAAGCACTGCTGCTCATTAACAGCT	42
11[462]	6[462]	CGGTGACCACTGTTGTGAAGGATTTAACAGATGGAATCG	42
11[504]	6[504]	GCTGATCGAGCCACAGCTTTGACCCCTAATCATGATGATA	42
11[546]	6[546]	ACAGTCCCGCCACTGCGCAATTAACAGATATCAAAATAG	42
11[588]	6[588]	AACTATTAGCCCGGGGATTTAAACGGAAACGAAACACTA	42

### biotin-modified

GTCAGAGGGCGAAAAACCTCTCCCTCC-Biotin  
ATTGTCACCTTATGACAAATGTCCCGCACTC-Biotin  
Biotin-ATTTCATCATCTAATATATATGTCAGAGACAAGC  
Biotin-ATATCTATCTCTTGAGATTGTGATCTGCTCTGA

### digoxigenin-modified

Dig-TATTACCCACCTTACCTTAAACATGCGCATATAAA  
Dig-TACCAATCTCTTTTGAATTAATCTTTTAAAGC  
CCAGGCGCAAAAGCAAAAGGCGACATCTCTCTTA-Dig  
AGGGTTGATATAAGTATAGCCCGGAATAGGACTCCCT-Dig