Supplementary Figure S1a

70HRT$_{14}$ Population

```
5'  gggaaaagcgaaucauacacaaga. . . . (70HRT) . . . gggcauaagguauuuaauuccaua 3’
```

SIX

```
70.03  AGATGAATCACAGCCGCTCAACGAGATGCGAAGGAAAACGGGATAACCAAATAGATGGAACGAAGACC
70.04c AGATGAATCACAGCCGCTCAACGAGATCCGAAGGAAAACGGGATAACCAAATAGATGGAACGAAGGCC
70.32  AGATGAATCACAGCCGCTCAACGAGATCCGAAGGAAAACGGGATAACCAAATAGATGGAACGAAGGCC
70.54  AGATGAATCACAGCCGCTCAACGAGATCCGAAGGAAAACGGGATAACCAAATAGATGGAACGAAGGCC
70.76a AGATGAATCACAGCCGCTCAACGAGATCCGAAGGAAAACGGGATAACCAAATAGATGGAACGAAGGCC
70.42  AGATGAATCACAGCCGCTCAACGAGATCCGAAGGAAAACGGGATAACCAAATAGATGGAACGAAGGCC
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FIVE

```
70.50  TGAAAATGCAGACCAAGAAAATCCGAGGTGATAAACGGGAAAACACAGAGAAACA-ATAACGTTGCAAGTC
70.5c  TGAAAATGCAGACCAAGAAAATCCGAGGTGATAAACGGGAAAACACAAAGAAACACATAACGTTGCAAGTC
70.52  TGAAAATGCAGACCAAGAAAATCCGAGGTGATAAACGGGAAAACACAAAGAAAC-AATAACGTTGCAAGTC
70.59  TGAAAATGCAGACCAAGAAAATCCGAGGTGATAAACGGGAAAACACAAAGAAACACATAACGTTGCAAGTC
70.79  TGAAAATGCAGACCAAGAAAATCCGAGGTGATAAACGGGAAAACACGAAGAAAC-TATAACGTTGCAAGTC

70.22  TGAACAAACGCGATGAACATTAGGCTATCCTCAGGCGGAGAGGGACAAAACGCACTTATCCCTAAGGC
70.33  TGAACAAACGCGATGAACATTAGGCTATCCTCAGGCGGAGAGGGACAAAACGCACTTATCCCTAAGGC
70.34a   AACAAACGCGTAGAACATTAGGCTnCCTCAGGCGGAAGAGGGACAAAACGCACTTATCCCTAAGGC
70.65  TGAACAAACGCGATGAACATTAGGCTATCCTCAGGCGGAGAGGGACAAAACGCACTTATCCCTAAGGC
70.76b TGAACAAACGCGATGAACACTAGGCTATCCTCAGGCGGAGAGGGACAAAACGCACTTATCCCTAAGGCAC
```

FOUR

```
70.30  AGCGCGGCACCCAAAATCGAAATCCGAAGGCGAACGGGAGAATGCGACCAAAGATACCCTGTGAATGGC
70.40  AGTGCGGTACCCAAAATCGAAATCCGAAGGTGAACGGGAGAATGCGACCAAAGATACCCTGTGAATGGC
70.45  AGCGCGGCACCCAAAATCGAAATCCGAAG-CGAACGGGAGAATGCGACCAAAGTAACCCTGTGAATGGC
70.72  AGCGCGGCACCCAAAATCGAAATCCGAAGGCGAACGGGAGAATGCGACCAAAGATACCCTGTGAATGGC
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THREE

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70.17  CCCTCCTCGTATG-ACGCTAA-CTGAGAATCCGAAGTCCAACGGGAGAGA---GGACACTTATGACGCGGCGCG
70.27  CCCTCCTTGTATG-ACGCTAA-CTGAGAATCCGAAGTCCAACGGGAGAAA---GGACACTTATGACGTGGCGCG
70.38  CCCTCCTTGTATGGGCGCTAAACTGAGAATCCGAAGTCCAACGGGAAGAAAAAGGACACTTATGACGTGGCGCG

70.62  TTGGAAAAACCGAGATGAATCCGATAAGCAACGGGAGAATGCTTAAAGATGGGATACAACCGATGCTCAC
70.64  TTGGAAAAACCGAAATGAATCCGATAAGCAACGGGAGAATGCTTAAAGATGGGATACA-CCGATGCTCAC
70.80  TTGGAAAAACCGAAATGAATCCGATAAGCAACGGGAGAATGCTTAAAGATGGGATACAGCCGATGCTCAC

70.24  GACGCAACCGAGATCATCCCAAGGAGAAGGAGAAAACCAACGCATGGAGTTGATTAACGTGCCAGCAC
70.35  GACGCAACCGAGATCATCCCAAGGAGAAGGAGAAAACCAACGCATGGAGTTGATTAACGTGCCAGCAC
70.67  GACGCAACCGAGATCATCCCAAGGAGAAGGAGAAAACCAACGCATGGAGTTGATTAACGTGCCAGCAC

70.26  TGCTAAACCAAGTAAGAATCCGTGAAGTCACAGCACGGGATAAAACTGTGTCAAAACGCCATAGCT
70.37  TGCTAAACCAAGTAAGAATCCGTGAAGTCACAGCACGGGATAAAACTGTGTCAAAACGCCATAGCT
70.69  TGCTAAACCAAGTAAGAATCCGTGAAGTCACAGCACGGGATAAAACTGTGTCAAAACGCCATAGCT

70.29  TCAGTGTGTGCACGCAAACACCACACGAGAGATAGAAATCCGGGAGGCTCGCACCTGGGACATACACCGC
70.39  TCAGTGTGTGCACGCAAACACCACACGAGAGATAGAAATCCGGGAGGCTCGCACCTGGGACATACACCGC
70.71  TCAGTGTGTGCACGCAAACACCACACGAGAGATAGAAATCCGGGAGGCTCGCACCTGGGACATACACCGC
```

TWO

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70.08  CGAGACAAGTACCGAAAAAGAGATCTGGCAGTGTCACAACCAGGAAAAAGACACGACGAACACGCCGCAC
70.13  CGAGACAAGTACCGAAAAAGAGATCTGGCAGTGTCACAACCAGGAAAAAGACACGACGAACACGCCGCAC

70.12  TATCCGAGCCAAAACGGGAAAAGATGGGAAAAATGGAATTAGTCCTCTAACCCAACGCGGAAACCAACGT
70.16  TATCCGAGCCAAAACGGGAAAAGATGGGAAAAATGGAATTAGTCCTCTAACCCGACGCGGAAACCAACGT

70.21  CTGTGCTGGCTAGACTATCCCAGAAAGTTGAATCCGAGGCGAAACGGGAAAAGACCGCGACCCACACACC
70.53a CTGTGCTGGCTAGACTATCCCAGAAAGTTGAATCCGAGGCGAAACGGGAAAAGACAGCGACCCCCACACC

70.31  CTAACGCGCCATCCTACGCGATCTGACACAAACAAAACGAGAACTCATCCGAGGCCTAACGGGACAAC
```

70.41 CTAACGCGCCATCCTACGCGATCTGACACAAACAAAACGAGAACTCATCCGAGGCCTAACGGGACAAC

70.36 CCACGCCATACCGAACGAAGATCCGAATCGTCACGGGAAAAGACGAAAAAAAAGGCAACCGAAATGACCGC
70.68 CCACGCCATACCGAACGAAGATCCGAATCGTCACGGGAAAAGACGAAAAAAAAGGCAACCGAAATGACCGC

70.48 TGAACGTACGACGTAGCTAGACCAAGATCCGAAGATGCACGGGAGAAGCATCAATACCACAGTATGGCC
70.70 TGAACGTACGACGTAGCTAGACCAAGATCCGAAGATGCACGGGAGAAGCATCAATACCACAGTATGGCC

70.23 ATGATTCGACTTCGAACGAAGATCCGAGACGCGACGGGAGAACGCGAAGGAAAACACCGAAATGCAACGT
70.34b ATTCGACTTCGAACGAAGATCCGAGACGCGACGGGAGAACGCGAAGGAAAACACCGAAATGCAACGT

ONE

70.01 ACGTTGTGCACGGATGCCCACGGTCGCACGAAACCTTGTGTGGGATAGCGCGAATACTGACGAGTGTGC
70.02 TGAAACAGCATAATGCCCACGGCACAATACCAAAGACAGATCCTGAAGCAGAGCACAGGGACCAACGC
70.04 ATTGCGACTTCCAAAAGAGATCCAGGGAGCAGGCGCACTGGGAGAAAACTGCCTGCAGAAAATGGTAC
70.05 TCCGAGGCAGAACGGGAAAATCTGCGAAGTAACTGTGGAATCCGTGACCTTTGACGTGAAAACCGCGA
70.06 CCTACTGGTAAGCGGCCTTGGCAGATTACCGCCGATCCATGTTACTGATCCGAACGCAACGGGATAATGC
70.07 TGTAAGTATAGCCGTTTTCCAATGTAAAACCGGAAGCTCCGAACCGAGAAAAAGAGCGAACAACATGCGA
70.09 GTGGAGAATCGCCTATAATAACCCAGAAATCGCATCCGAAGGTGAACGGGAGAATCACCGACCTCGATC
70.10 AATATCCGTTACCATTCGGGAAAAAATGGGTTGGTGAAAATACTTGACCCCGAAGGACGACGAATCGGAC
70.11 CCTCGGGTCCCAGCAAATCCGAAGGAAAACGGGATAACCAACAACACAATGAAAACAGCCAGTAAGCGT
70.14 TCCGACGTACGACGGGATAACGTTCAAAGCAAGGGACTTTAAAGTAACCGCTGCCGATGATCAACGCAT
70.15 TCACTTGAAAGACGCAGACCTACGAACCCAGGAGATAAGGGGGAAAACTCTGGAAAACCAACCGGTAGTC
70.18 AACCAAGGTGAAAAATGGGAAAACACATAAGGGAGTGAAATAGACGGTCCTAAACCAACACTGGCGAAC
70.19 ACCGAATAAAACGAGAAAACAATGTGTAACACACCGCTACTACCGAACCAAAGAGGAAAGAAGCGTCCGT
70.20 CTTTTCGCTTGTAAACCCCAAAACGAAATCCGAAAGGCAACGGGAAAATAGCCTAGACCATGTCCGCACG
70.27 CCCTCCTTGTATGACGCTAACTGAGAATCCGAAGTCCAACGGGAGAAAGGACACTTATGACGTGGCGCG
70.28 ACCGGACGGGTGAGAACCGAGACAAACACCCACCAAGAGAAAATGTATCCACTACCCAGACGGCCTCGAT
70.43 ATGAACGCAACCCAAAGCTTAACCGAGGTGTAAACGGGATAACACAAGTTGGAAGGAAGATCAGGGAGCA
70.44 CCCGCACCAAATGAAGATCCGAGACCAACGGGAAAATGGTCACACCAAGGAATAGATGCACGCCTCACAT
70.46 GATAGCTTAGACCACAAAATGGGACGAGCAAGGGGAAAATGCTCAAAGGAACCCGAAACTGAGTGTGCAC
70.47a TGGCGCCATGCCGAAAGCTCCGAGCCGGTAACGGGAAAAAGACGGAGTCAGGCTGCCT
70.47b TCGCCTGCCAAAAAATCCGCGCCGTAACGGnCCAAGCCCGGAAGATAAAGGCTGCCTAATACACAGCGCA
70.49 AGATAAAGTGATGGATTTGCTGAAGTCTCTCAACGATCCGATGACAACGGGAAAATGTCACAAACGACC
70.53b NNGNNNNTNGNATCCNAGGCAGAACGGGACAATCNGNNNNNNAAAGAGGTGTTAANCCTATTCCCTGAT
70.55 TACATTACAGAGGTGAGTATCAAAAATGTTGAAACGTATTCCTCGGAAATCCCAGGCCTAAGGGGACAAC
70.57 ACGCGACCCAGTGAATCCGAAGCTTGACGGGAGAACAAGCAAGACGTGATTAAGCACCACCAATGCCAAT
70.58 ACCAAATCCCGAACTACAAATCCGAACGCTAACGGGACAATTGCGAAATGGAACATACGGGCCTGGTTGAA
70.60 CACCATACCAATGGCATCGCTATGATCGAGCGCTACGCTAGACCAATCTGGAGCTGACCACAGGGAACG
70.61 CCCAGAACTAACCATATCCGACAGCCGAAACGGGAAAATGAGGCAGAGCCGAACCAAAATCTGTGTGCAC
70.63 CGCAAAACCACTCGATCCGAAGGAAAACGGGATAACCAAAGATGGGAAACAACTTACACACCGAATCAC
70.73 GATTCTAACCTGAGATGAATCCGGAAGCAAGAGCCGGGAGAAAACTTGCAAGACTACCAAAATGTCGTTG
70.74 GCGAAAGAACTTAGTGTTGATAGACGATTGACCAACCCAATGACAATCCGAGCCGCCACGGGATAC
70.75 CTCTTACCCAAGACTGATCCGAAGGCAACGGGACAAAAGCCAAAGGAAAAACCTAACTCAGCGAGACTGC
70.77 ATCCGAAGGCAACGGGAAAATCAGTCCGAGAAAAAGGGAGAACACCTTAACTCACTGCTGTACGCAA
70.78 CGCGAACGATAATCCGAACGAGGACGGGACAAACTCGAGCAATGATACCTAATCTCCCTACCATCGCTGAA

Supplementary Figure S1b.

80HRT$_{14}$ Population

5'gggcauaagguauuuaauuccaua. . . . (80HRT) . . . uugauucggaugcuccgguagcucaaccug 3’

THREE

80.08 **AGCCGTTAAGCAGTGTCCGCTACCTTCATCCGTTGGCGATCGGGAGAAATCGCAAGAGGAGGCACCGCGCTCCGTATCAT**
80.72 A-CCGTTAAGCAGTGTCCGCTACTTTCATCCGTTGGCGATCGGGAAAAATCGCAAGACGAGGCACCGCGCTCCGTATCGT
80.90 **AGCCGTTAAGCAGTGTCCGCTACTTTCATCCGTTGGCGATCGGGAAAAATCGCAAGACGAGGCACCGCGCTCCGTATCAT**

TWO

80.02 **TGCAAGCCTGATGATCCGAAGGTGACGGGAGAACACCAAACAGTATGCTAATCCACCATCCGAAACCGACTGCTACTA**
80.105b **TGCAAGCCTGATGATCCGAAGGTGACGGGAGAACACCAAACAGTATGCTAATCCACCATCCGAAACCGACTGCTACCA**

80.46 **GCTCGCCAATCCGTTTTCCAGCAGTCTTCGGGATAAAACACTGCTACTCGAGGCCGTTGCATTCCGGCCCATTACTCCA**
80.100 GCTCGCCAATCCGTTTTCCAGCAGTCTTCGGGATAAAGCACTGCTACTCGAGGTCGTTGCATTCCGGCCCATTATTCCA

80.55 ATGGCTCACCACAAGGGGAACGTTGATGAAATAGAGTTTATCCCTTGGACTCACGCCGGCCGTGCTCCACACAATCCA
80.65 ATGGCTCACCACAAGGGGAACGTTGATGAAATAGAGTTTATCCCTTGGACTCACGCCGGCCGTGCTCCACACAATCCA

80.09 ACCGCATGGAATAAGTTCAACACCCAACTACGATCTGTGCGGCCAACACAGGGAGAAAACCGGCCAAGAGGAGGCTAC
80.87 ATCGCATGGAATAAGTTCAACACCCACCTACGATCTGTGCGGCCAACACAGGGTAAAAACCGGCCAAGAGGAGGCTGA

ONE

80.01 **...TGCGAAGAACACCGTAGTTAAACTAGATTTACAAGGAGAGCGGATCGAGGGAACGGAGAGA**
80.05 CAGAAATCGAATATCGAAGCTCATTGACCCTCAATCCGATGCGAGATGGGATAAACTCGCACCCTTCCGCCCCACGACT
80.07 TCGCACATAAAGTGGATGTATGCCACCCTTCCTGTAATATCCGTTACTGTGCCTCGGGATAAAGGCGCTACGAAGT
80.10 CCNNCTGTCACCTTCCGAGTTTGGTAACGTGTTTTCTGCCAGTAGTTCTTACAAATGAAGTCCGAGGGTAGATCGTA
80.13 CTCCACATTCACTGCTGCACCAAATCCAAAAAAACACATCCTCGAAAGCTCATAATTGGGAATGCATATGTCAAGGTGGGA
80.15 GGCACAGAAATACCATAGCCGAAAATCCCAGGCGAAAGGGGACAAGTCTGCATTCATGTGCTACTGCGATCGAAAAGT
80.16 ACGGTGGTTAGCATATCACTCAAAATCTTCTAAGCAAGTCCTGGCATAAGCCCCTCATACACTCCCTCGGTTTCCC
80.17 **TCGACCCTCACTGTAATCCGTTTTCGTTGGCTTCGGGAAAAAAGCCGGCACATGGCCCGACGTGTCATCAAATTCAAGTC**
80.18 TCTGGCATCGTACAAATGAGTCTGGCCCAGATTAACTGATCCGAAAGCTGCATCGGGACACAAGCACAGCGATAATGAG
80.21 AGCCTAGTACCCTAACGTGATCCGTTCCTGTATAGTCGGGAAAAACTATACAACTAAACTACCGCTATATTCA
80.28 GGGTGCAAGGCACGTGTTAAATCAAGTCGTATGGAAAGAAGGAAAACCTAGGAATGGGACCTCCAGTGACAATGAAGTGC
80.30a **AGCCAACCGCTAATCCGATCGTCCGACGGGAAAATGGGACGGCAATAGTTTACCCnCTGAGCAAGTATTTAAGTAAGTAGTT**
80.30b **TGTCAGCGAA TCTACGACCCACTTTCAGTGGACGATAGGTTACATGTTCCGTTACGTGTTCGGGAAAAAACATGGCTCG**
80.31 TGAACTCATCCAGTAATCCCACTCGCGCGAAGTGGGAGCAATAGAAATTCGCGCCTATCTCAATACGGGATCAGTCTATG
80.32 TCCCTAGCATCCTTGACCAATACCGTTCGGAGGCCAAACGAGAAACACCTCCACGAGCAGGAGCACGGCATGCGTATAC
80.33 GCGTCGTATTGCGGTTGAGAGAAGACTTGACACAAGACATGGATCCGATAGCCAAACACCATGACCCAATCCGTTACAA
80.34 GCCAAAACGATCCGGCCGCGAGACAACCGTGAAAAACCCGTACCTGGCTGTACCGACACTTCGATAGTCACCATTAGACG
80.35 CGTATAGTCCAAACAAGTGATCCGATGTCCACGGGAGTAAGGACCGATGTCGCGCCTGAAGCACCGAACTTGATTGTATG
80.37 CGCACCGCACACCGCACCTCAAATCCGTTCCGTAAATCGGGATAAATTTACGACTTTTACTACCCAACACGATTACCA
80.39 CGGTCCAATTAAATCCGTTTTGACGATCGGGAAAAATCGTCGGCACCGGGGCCTAAGTTTCTTCGCTGGACACTTTCAGT
80.49 TGCCTTATCGAACATCCGTTTGGCTGGCGTCGGGAAAAATGCCACGCGAGTCTGGTGCACTGGTCACACGAGAATCTA
80.51 TCCCAATCTCTGTTATCCGGCGGAGAGGACACCGGGAGAAAAAAAGCTCTCACGACGCGACTGGCACACGGATATAAGGTT
80.52 GCGACCCACGACTGATCCGTTTGCGGGCTCGGGATAAAGCTCGCGGCCATTGCACCGGACTCGCTCATCTCTACAAAG
80.54 CAGTCCATCAAAATCCGTCAAGCCGTAGACGGGAGACAAAATCGGCATGACCCAAGAAAAAGAGAGATGTGAAAGGT
80.57 TGCTTCTTGGGATCCGATGCGCAACGGGATAATGCGCAACTTGACCACGCCTAGGCTAAGATCAAATTTCCCTTCTAT
80.59 ACGCCAACGACACAAGTCTGGCTGGGTCCAAACCAGGACAAGACCCACATAAACGCTTACCCTTCGTGACAACTAG
80.60 GCGTCGGGGGCCTCATCTGAAAATCCCAGCGACACAGGCACGACCCTTCAGGACATGGAAAACCTGTGCTATAAGATGCGA
80.61 CGGAAGCAGGCCGACCGCCAAGTTGTAAGTGCTCTGAGAGTGGCTGGTGCATGAATGGAGTATGGCGTTTGGAGTT
80.62 **GGCCTTATAGACCTCTCTAATCCTGGTAGCATGGACCAGTGAGAAAACTCCATGTTTAACAAAACCTTAACGCGA**
80.63 GCCACACTCCACTCTCGACCGTTTCTTGGGTTCTTCGGGAAAAAAAGCAACCTACTATTGACTATCGACGAAGATCTGTT
80.64 **TTCnAGCnTCnTGATCCGAAnnnCACGnGAGAAnnnCAAACAAnAAnAATGCACTATATTCGAAACCGmTTGTGATGAnnn**
80.66 CGGTGATGCTGACGAACTAAACTGATCCCGTAATGACAACACGGGCGAGCCAAGTTGTCTCCCTAGCTCCTATTCGACTG
80.67 CTGTCTGTCGTACCTCCTTCAGCTTTACCGTTAACTTTCGGGAAAAAAAGTCGAGCTGCATGTTACCAGGACAACATC
80.68 CCGACCAAAACGACCACAGATCCTGTGCCAGTGCCAGGCGAGAGAGTGCTGAGAAAAAGAGTTTGAGCGAAGTATGCG
80.69 CCACGATGATTTTGGAGGTACGCGGGATGGTGGAGAAGGATAACCGTTGGTCATCGGGATAAATGGCAACGCTGC
80.70 **GCCTGAGCCAATATTAGTGCCGCCACTACGTCCCCCCTCTGTCATCGTCAAACCCACCTGAGTTATGTTACCATTCTG**
80.71 **TCATGTTGCGCGCAAACGGTATCACTCAAGTTGCTTCGGCACTTCAACGATGCACTCACTTTGCAAATCCTGGTCAAGAG**
80.73 **GAGCGCAACATGATGTGATGCAATACCGACTCGCCCACCAGATCCTCACCGTACAGGAGGGACAAATTGTACGTCGAGAC**
80.74 **TTGGTCTATACGATGTGTCCTCCCCATCCGTTTTGTTCTCGGGACAAAGAACTAAAATTGGAGCTATTCCACCCAACG**

80.75 **TCCTCACCCAAATTGATCCGACGCGCGACGGGACAATGCGCAACCCTGCTACCACACCGTTCTCGGACTACCCCCCTCAG**
80.76 GTCCATGATCCGTGTCCGAAACACGGGAGACAAATTCGGCTAAGAAGACACGGCCCTGATTATTCCCCTTGCGGATGGA
80.77 CTCACGCTTGACAACCACAAGGGGACAGAGCTAAGTTGTTATCCCTTNGCAACACCGACATTGCCGAGCGATGGTTATCA
80.78 **CGTCCACCCGTGAATACGGTCCAGATAATCCGACGGCAAACGGGAGAACTGCCAAGCGCTGCTAACGGAACGTCCGTA**
80.79 **TGCGGCTAGACTACATCTGCTCATTCAAATGGATTGAAACTTTACGTCACTCCCTTAAACTCCTTTCCCATCGTTTC**
80.80 TGCAGTTGTACAATCCGTTTTACTGTGTTCGGGACAAAACACAGCAACATAGTAGCTATCGCTCAACCGGACATGCAAGA
80.82 **ACCCAAAACCGAATATCCGGGCAGGTCCCACTCGGGATAAGAATAGAGGACAATGAAAAGTGCGAGGCAATAGAAAACCG**
80.83 TCGCTCTGAATGCATTGTTCCCATGCGAAATCCGAGGCCAAACGGGAAAAGATGGCACCAGCGGTTGCGCCTnnn...
80.84 CTCAGCCTGAGAAAACTAATCCTCCCAGTGCCAAGAGGTGCAGATAGATTGGCACTCATTTCCCATCCCAAAGGTNGC
80.85 GCCCGGTCACGAACCGAGGTGTTAAACGGGAAAACACAACGAACATCGGAGTATTCATGAACTGCGGTCCCTTAAAGACA
80.86 **CTGCTCAAACCAGCAAACATCCGTCGAGGATCGGGAAAAACCTCCATATGCAAGAAAGCCGCAACCTGAGATGTACTA**
80.88 **CGTTCCACCTCCTCTGCCTTCTACGTCTATGCCTCCCGAACACACGTTGCCGACCCACCGCTTCAAGTCTAGCCATA**
80.89 **GCCAACTGCATCCGCGAGCGTTACGTGGGACATCATAGCGCAACGAACCTACCGCTTCAGCACACTCATCTGACA**
80.92 GCACTCCACAGGAAAACATGGAGGGCGGGTGGAAACGTGATAAGTGCGCTGTGTGAATTTAAACCTTACGAGGCATCTTA
80.93 CCTCTCCACGACAAATCCTTATCGCATGCATGAGGGAGACCAGACAAGCATGTACAATCACCAAGTTATGATAGTTCGAG
80.96 CGCTTGATCGAACGAATTAGCGCGTAGGCTGCGGCCACACTGCCCACTAATCCCGATCGTAAGGGGATAGTACGGCGAGA
80.97 CCGCGGAGTAAAGTTGTGGTCGTTGCAACGATATTCGGGAGGCTAGTGAGTGAAGTTTGATACCTGTGACACCAAGGC
80.98 **TTGCCAAAGAATGATCCTGCGATTGCTGCCCACGCAGGGACTAAAGAATGGGGCACATCACCGGTAGCCCTGAACTTGC**
80.99 **TCGTGCGGCGGAAACCTATCTACTAGATCTGGATTTATGCTTTTGTACCCGTTATCCCTTTTCCTAGTGGTTGGCTTGT**
80.101 **CCCACGCAACGATACTCGTCGAACAAGGCCGCACACCCTACTTTCTCTCCCGCCCTGTAACA**nnnnnnnnnnnnnnnn...
80.102 **CCAATGGCACAGACATCGTAATCCGTATAGACCTGAGTACGGGACAAAACTATCAGGTCATGAGAACAGAGGACTGCTG**
80.103 **GCAACCGGTTGTCTACACGCGGCGAATAGAGCCCGGTTCAAGGACACCGCCACTGCTGTCGACATTTCCTTAGTGCTAGA**
80.104 **TCGACATGATGTGTTGTAAATTCCCGTTAGTCCGACGATGAATCTGTGCAGTGGAACACAGGGAAAAAATTCCACACCGA**
80.105a **CCAAGACGGTAGAGAATATCAGCGCTTTTCGAAAACATCCCGTGGGCCCGCACGGAGAAAATAACAAGGGACGA**
80.111 **CCTAGGACGAAAGCGATAATCGGGCCTGGAGGATCAAATTAATGCTGGCGCCAGGGGCAACGCATCTATGCACCTTAGG**
80.112 **CGCCTACGTTGAAATCCGTGATTCCCATACACGGGATATGCTATGGGTCGCGTTGCGGTGCCGCAGTATCAATCCTG**

Supplementary Figure S1c.

For $70HRT_{14}$ set:

Reported previously: 54 total reads, 42 independent, 34 sampled once, 5 sampled twice
Lower limit of sequence diversity from resampling statistics: $\mathbf{N_{Chou} = 42 + (34)^2/(2 \bullet 5) = 158}$

31 previously proposed F1Pk in $70HRT_{14}$ with 5 of them sampled twice
11 previously proposed F2Pk in $70HRT_{14}$ with 2 of them sampled twice
4 previously proposed non-pseudoknots in $70HRT_{14}$ with 1 of them sampled twice
(note: there were unrecognized triples in this set.)

29 sequences added subsequently, including 8 totally new species.

For complete LTS data set above: 83 total reads, 50 independent, 34 sampled once, 7 sampled twice.
Lower limit of sequence diversity from resampling statistics: $\mathbf{N_{Chou} = 50 + (34)^2/(2 \bullet 7) = 121}$

For $80HRT_{14}$ set:

Reported in JMB: 45 total reads, 44 independent, 42 sampled once, 2 sampled twice
Lower limit of sequence diversity from resampling statistics: $\mathbf{N_{Chou} = 44 + (42)^2/(2 \bullet 2) = 485}$

19 $80HRT_{14}$ in family 1 with no doublets
25 $80HRT_{14}$ in family 2 with 1 doublet
0 $80HRT_{14}$ in family 3

31 sequences added subsequently, including 28 totally new species.

For complete LTS data set above: 76 total reads, 70 independent, 65 sampled once, 4 sampled twice.

Lower limit of sequence diversity from resampling statistics: $\mathbf{N_{Chou} = 70 + (65)^2/(2 \bullet 4) = 536}$

For Combined sets:

Reported previously: 99 total reads, 86 independent sequences
Current information: 159 total reads, 120 independent sequences

60 additional sequences, 39 of which are totally new.

Lower limit of sequence diversity from resampling statistics:
$\mathbf{Net\ N_{Chou} = (121)_{70HRT14} + (536)_{80HRT14} = 657\ minimum\ total\ species}$