

## Description of Supplementary Files

File Name: Supplementary Information

Description: Supplementary Figures and Supplementary Notes

File Name: Supplementary Data 1

Description: **GenERA CRISPR/Cas9 sgRNA libraries corresponding to data in Fig. 1-5.** Detailed representation all sgRNA spacer sequences together with corresponding protospacer genomic coordinates, forward/reverse oligonucleotides used for cloning in pAc-Cas9-sgRNA vector, and PAM identity/position.

File Name: Supplementary Data 2

Description: **NGS library primers and amplification conditions used for generation of GenERA amplicons.** Targeted PCR conditions including annealing temperature, minimal non-saturating cycle number and amplicon-specific primer sequences are displayed for each amplicon in the study

File Name: Supplementary Data 3

Description: **Top expressed S2 miRNAs.** Seed sequence identity of most abundant S2 miRNA families is shown together with their ranked expression levels as reported by Ruby et al.

File Name: Supplementary Data 4

Description: **Assembled miR-184 target network.** TargetScanFly 6.2 prediction of conserved 8mer and 7mer-m8 sites and miRanda prediction of conserved 8mer and 7mer-m8 sites with good miRSVR scores were included. \* = putative lincRNA.484 MRE was excluded from further analysis due to genome misannotation. miRNA-MRE base pairing interaction are shown for miR-184 and most abundant S2 miRNAs.

File Name: Supplementary Data 5

Description: **Expression levels of putative miR-184 targets in S2R+ cells.** FPKM data from two previously published studies (Berkely Drosophila Genome Project (SRR070279) <http://sra.dnanexus.com/runs/SRR070279/experiments>; Berkely Drosophila Genome Project (SRR124149) <http://sra.dnanexus.com/runs/SRR124149/studies>) and an independent profiling performed in this study were used to compile an average FPKM value for each miR-184 target gene (genes with FPKM < 0.015 were not included in the analysis).

File Name: Supplementary Data 6

Description: **Putative RNA regulatory elements (RRE) and their occurrence within selected miR-184 targets.** The count, sequence and position of each RRE encoded within a 200 bp window centred on the miR-184 MRE seed sequence is shown.

File Name: Supplementary Data 7

Description: **GenERA CRISPR/Cas9 sgRNA libraries used for miR-184 MRE network analysis.** Detailed representation all sgRNA spacer sequences together with corresponding protospacer genomic coordinates, forward/reverse oligonucleotides used for cloning in pAc-Cas9-sgRNA vector, and PAM identity/position.

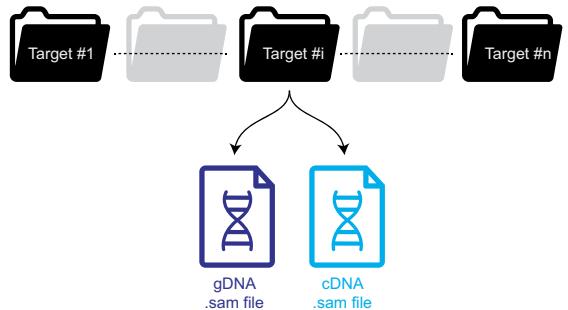
File Name: Supplementary Data 8

Description: Synthetic reference genome sequences underlying the miR-184 MRE network.

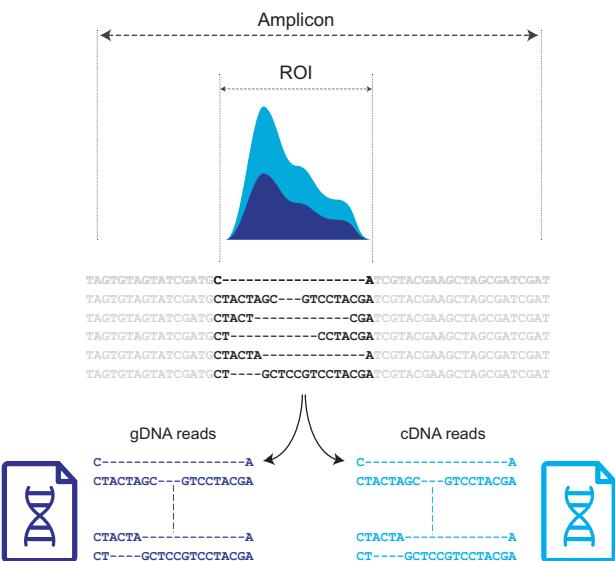
File Name: Supplementary Data 9

Description: **MRE-score distribution across miR-184 targets.** Weighted normalized MREscore (MRE-score), corresponding standard deviations, UDP counts and MRE-score group partitions are shown. miR-184 MREs reported by (Kertesz et al.,)31 and (Hong et al.,)32 were used for validation purposes.

File Name: Peer Review File



**1** Reads from gDNA, cDNA sam files are reconstructed from CIGAR strings. A deletion profile is created by counting for each nucleotide along the reference sequence the number of gDNA, cDNA reads with deletions at that position. A region of interest (ROI) is determined which contains the majority of NHEJ deletions.



**2** gDNA and cDNA reads are cropped to the region of interest (ROI).

**3**

### Read filtering

Reads only partially covering the ROI and those containing insertions are excluded from the analysis. Additionally highly mutated reads are not considered for the downstream analysis due to reduced reliability. In order to increase read counts sharing the same Unique Deletion Pattern (UDP) we reverted back mismatch of lowly mutated reads to the corresponding WT sequence.

CGTCGATCGTCGGCATCCCTCCAGCTAGCGATCGAT ✓ M.....  
 \*\*\*\*\* TCGTCCATCG ----- CGGATCGAT X discarded (partial ROI coverage)  
 CGTCGATCTGGT-----TCCAGAACCGATCGAT X discarded (high mismatch prevalence)  
 CGTCGATCGTC-----ACCGAAGGAT ✓ M.....-----N.....

Unique Deletion Pattern (UDP)  
 '\*' = cliped reads, 'M' = match, '-' = deletion

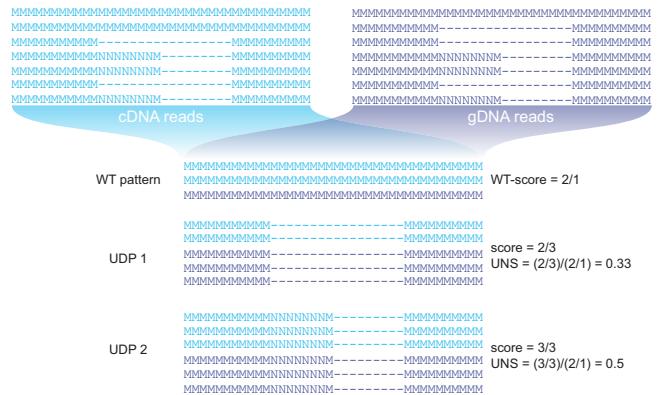
**4**

### Reads sharing the same UDP are paired across libraries

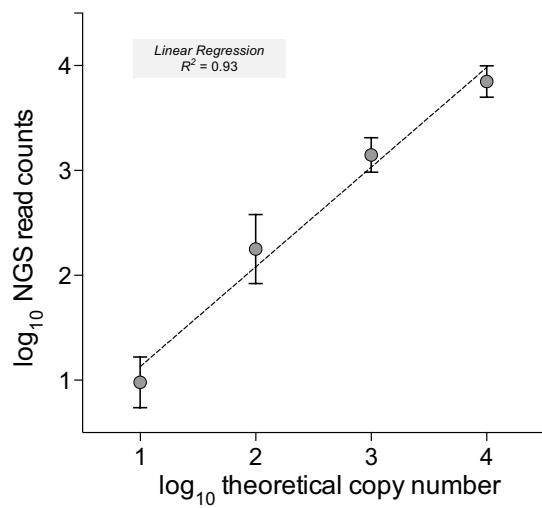
Post filtering reads are converted to a binary string encoding their deletion pattern (see above). Reads sharing the same UDP are compiled within library and then paired between cDNA and gDNA libraries. A UDP normalised score (UNS) is then computed using the formula below, which reflects the phenotypic impact of a given mutation (transcript abundance).

$$UNS = (cDNA_{UDP} / gDNA_{UDP}) / (cDNA_{WT} / gDNA_{WT})$$

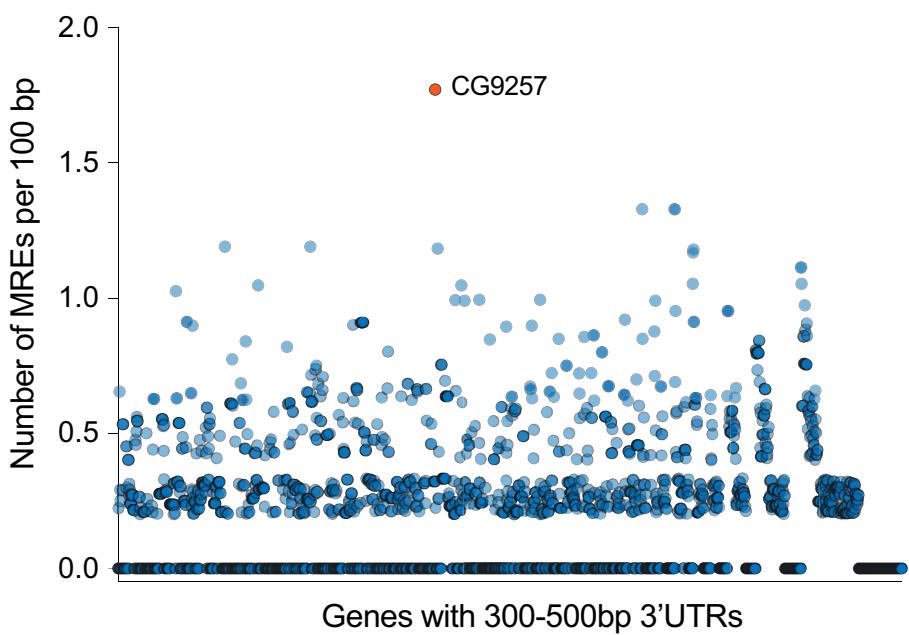
cDNA and gDNA refer to read counts in the corresponding libraries.



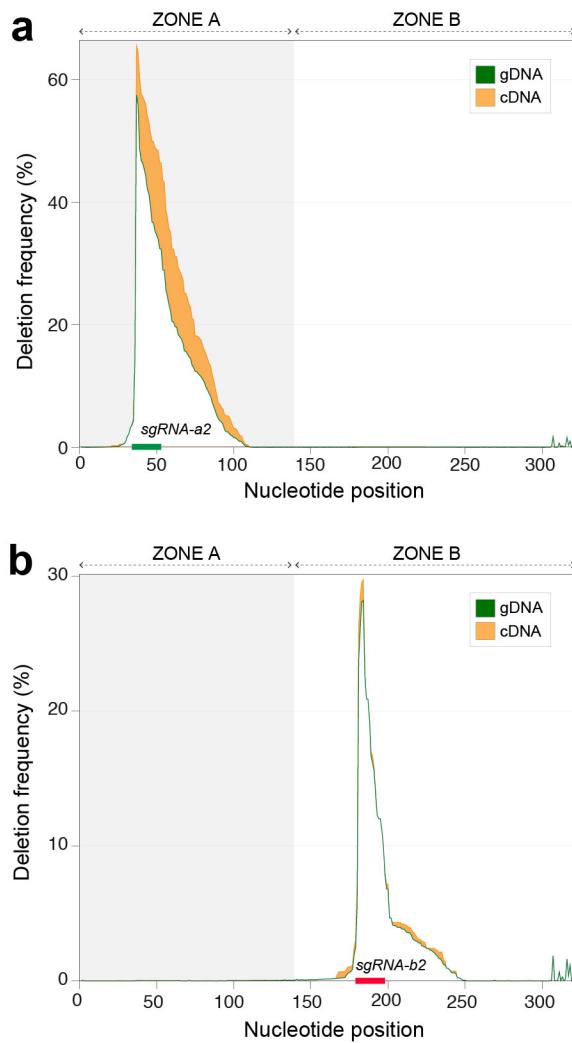
**Supplementary Figure 1. GenERA computational pipeline .** Stepwise overview of the computational pipeline used to derive the UDP normalised scores (UNS) from gDNA and cDNA library sam files containing mapped sequencing reads.



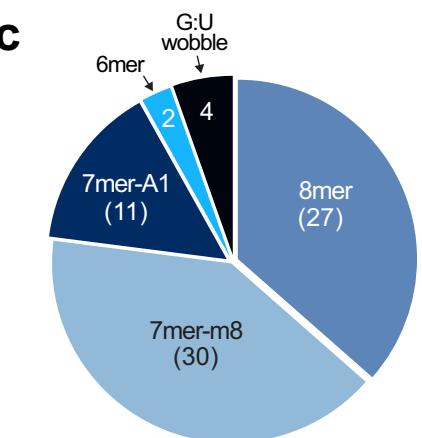
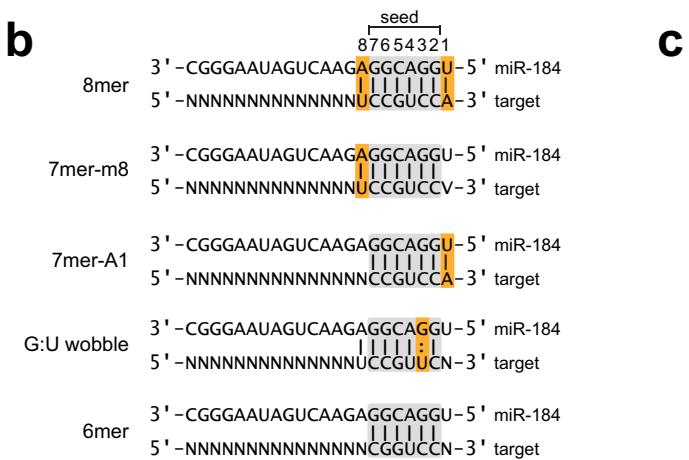
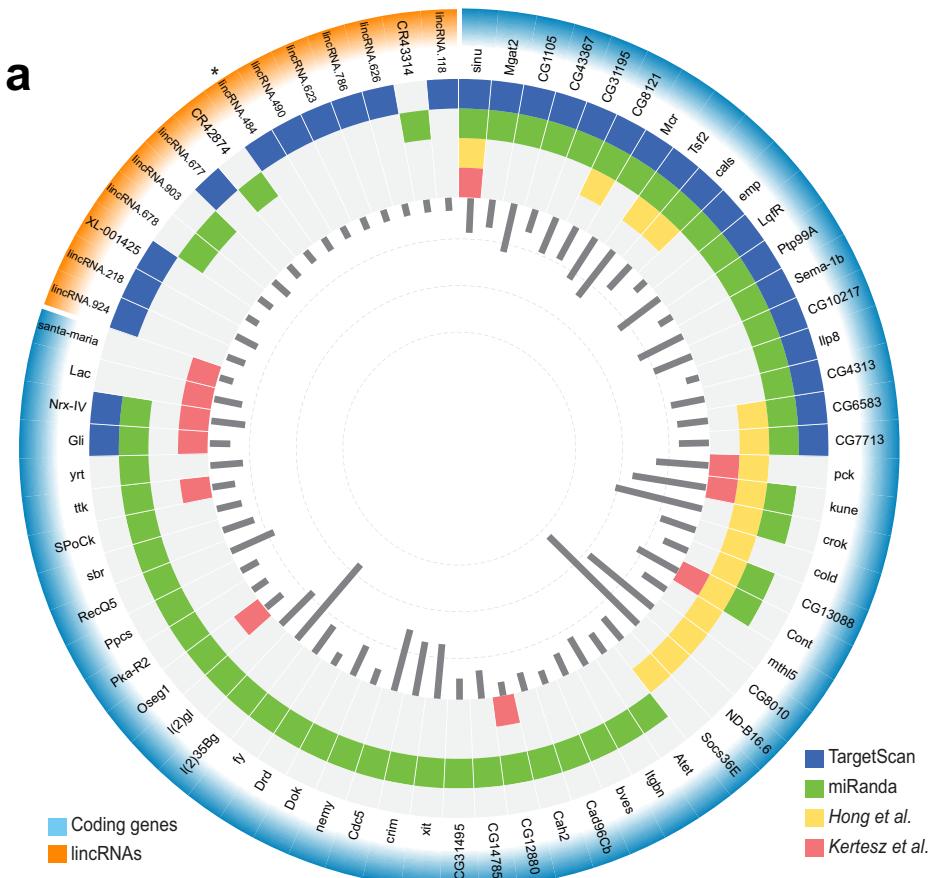
**Supplementary Figure 2. Reliability of deep sequencing datasets.** Estimation of NGS copy number resolution and dynamic range. Each dot represents average barcoded RP49 read counts following serial dilutions in gDNA and cDNA libraries. For each dilution, 3 barcoded RP49 triplicates were generated and sequenced. (Error bars = SEM).



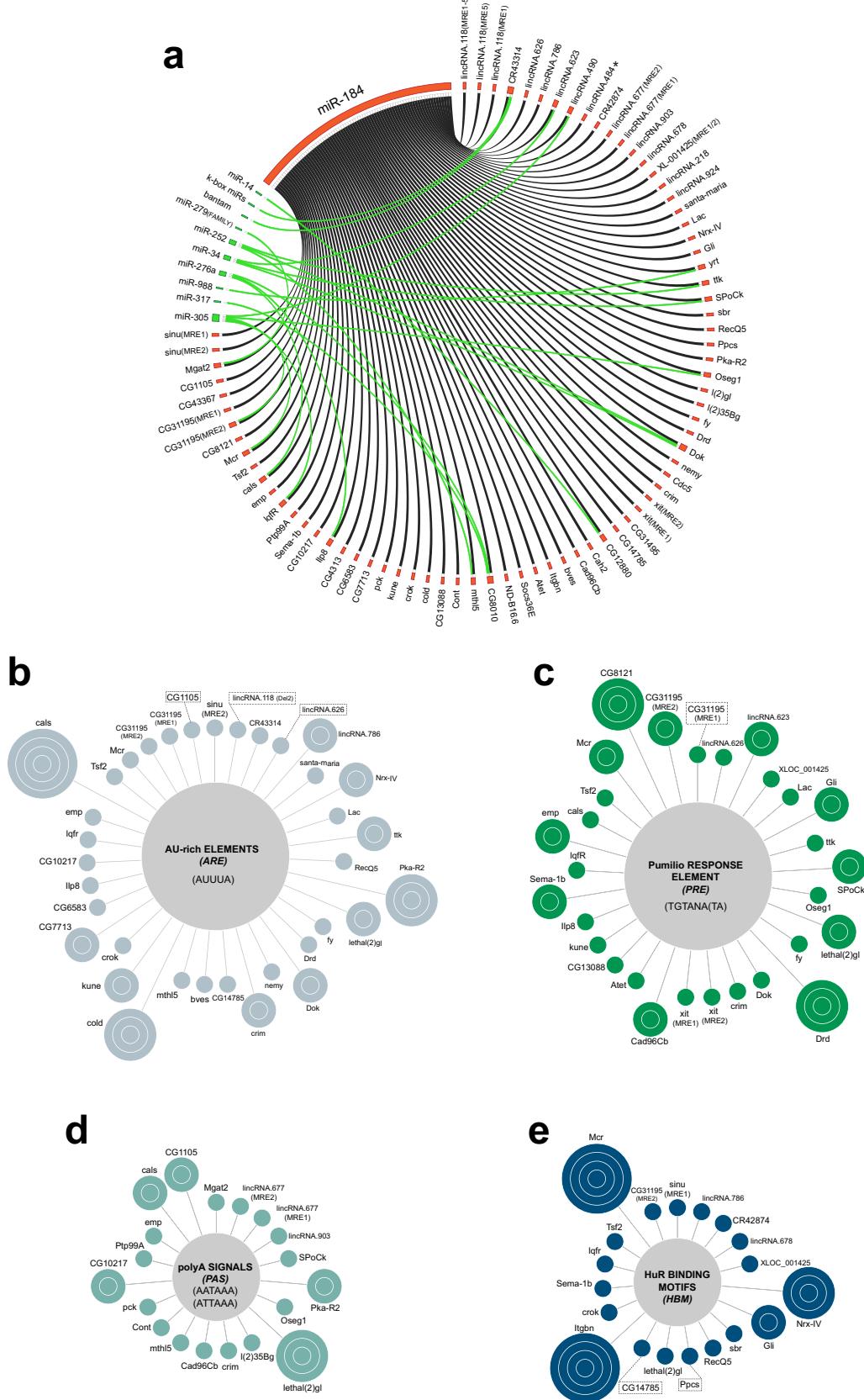
**Supplementary Figure 3. Selection of a candidate 3'UTR for GenERA-based unbiased survey of regulatory activity.** An MRE enrichment score, defined as the number of MREs per 100bp, was calculated for all S2R+ expressed genes containing 3'UTRs between 300 – 500 bp. The MRE predictions included the top 10 most abundantly expressed miRNAs in S2R+ cells (see Supplementary Data 3). The 3'UTR encoded in CG9257 (orange dot) displayed the highest MRE enrichment score among the 2311 genes subjected to this analysis.



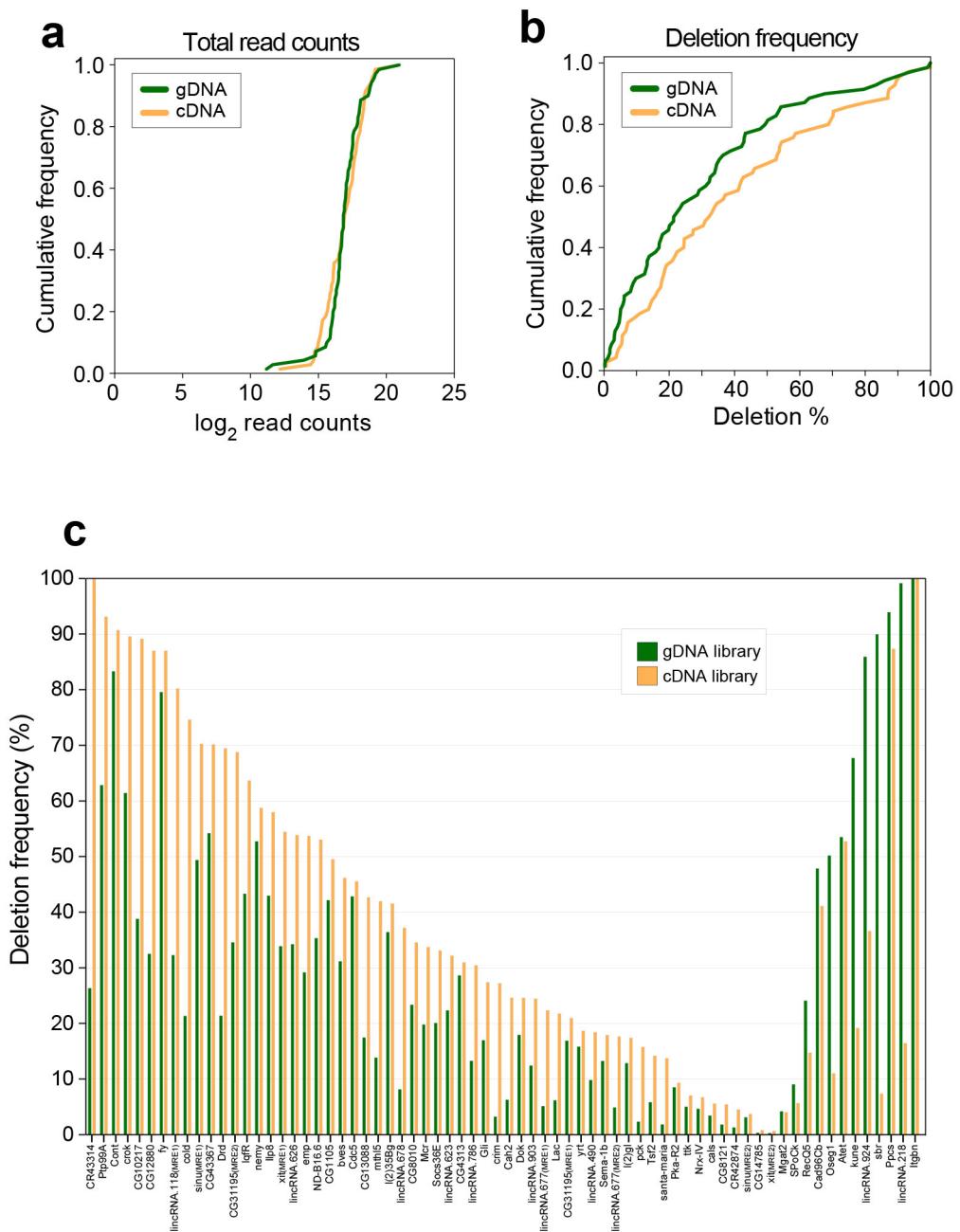
**Supplementary Figure 4. Validation of differential regulatory potential associated with CG9257 3'UTR zone A and zone B .** (a, b) Second set of individual sgRNAs separately targeting zone A (a) and zone B (b) of CG9257 3'UTR (relevant to Fig. 3 d, e). The cDNA and gDNA deletion frequencies across the 3'UTR are shown in orange and green, respectively. The precise positions of the sgRNA protospacers targeting zone A (sgRNA-a2, green) and zone B (sgRNA-b2, red) are displayed on the x axis.



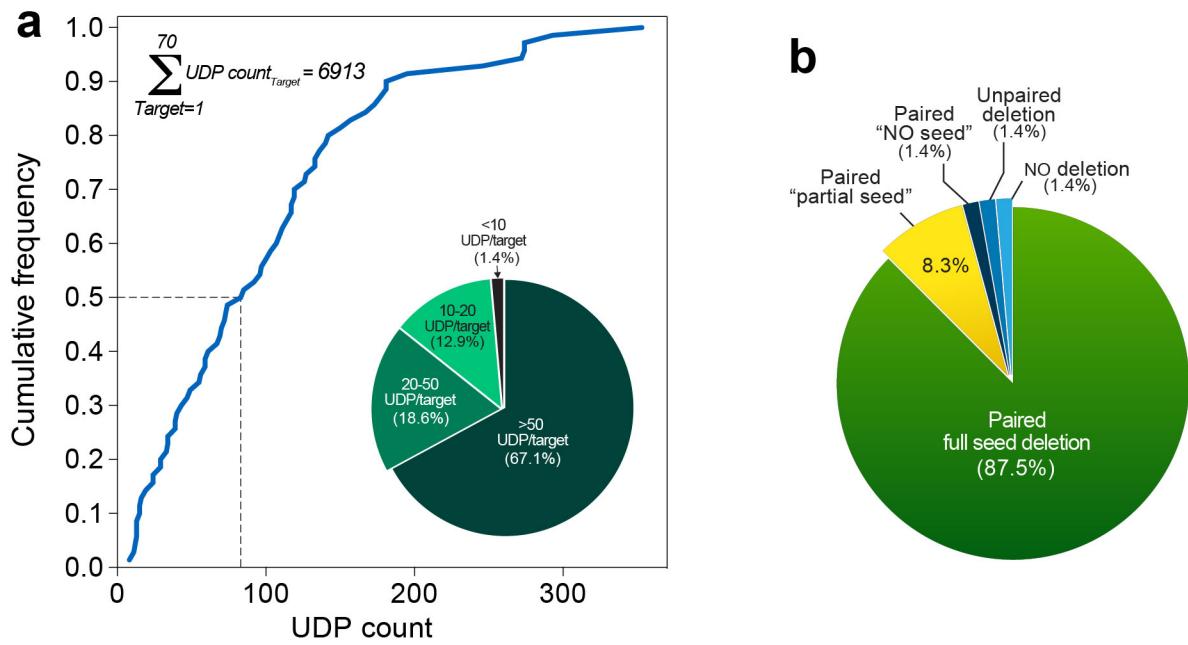
**Supplementary Figure 5. Predicted miR-184 target network.** (a) Graphic representation of the putative canonical miR-184 MRE network outlining the identity of coding and non-coding targets, their source of prediction, and relative expression levels (histogram) in S2R+ cells (\* = putative lincRNA.484 MRE was excluded from further analysis due to genome mis-annotation). (b) Classes of representative canonical seed types found in the predicted miR-184 MRE network. Unique base pairing features defining each class are highlighted in yellow. (c) Corresponding incidence of single MREs within the miR-184 network.



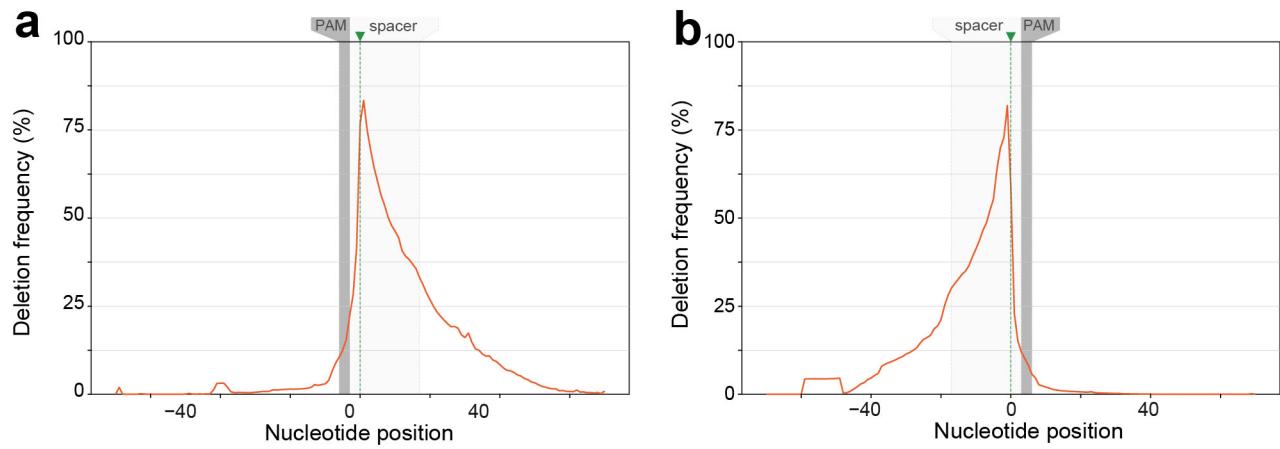
**Supplementary Figure 6. Coincidental RNA regulatory elements across the predicted miR-184 MRE network.** (a) Prevalence of predicted interactions between most abundant S2 miRNAs (green lines; *TargetScanFly* 6.2) and putative miR-184 targets (black lines). (b - e) *In silico* identification of AU-rich elements (b), Pumillio response elements (c), polyA signals (d), and HuR binding motifs (e) encoded within a 200 bp window centred on the miR-184 MRE seed sequence. Circles reflect the number of predicted RNA regulatory elements (RRE) in each target.



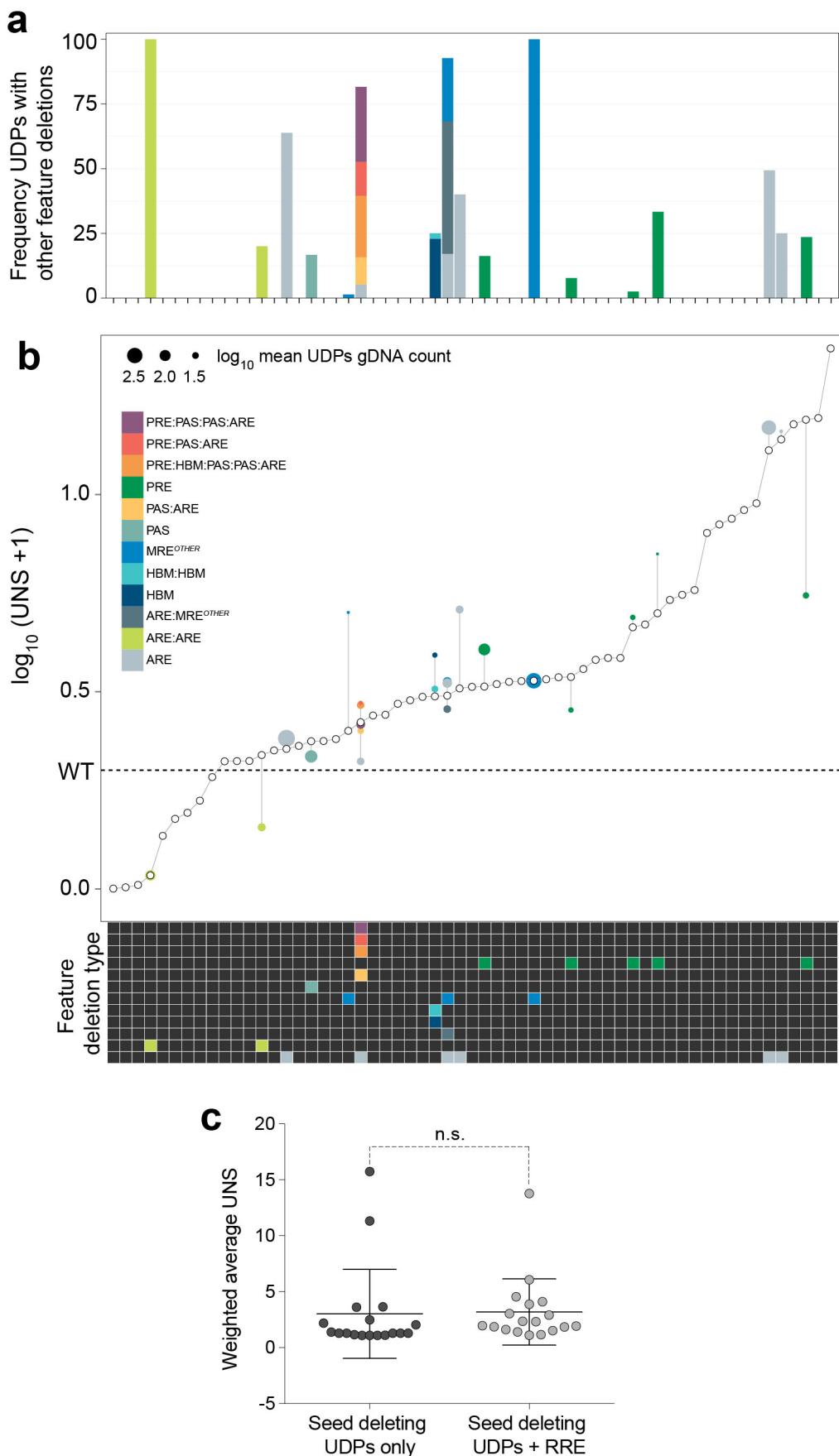
**Supplementary Figure 7. Effects of CRISPR-induced genomic deletions on transcript levels.**  
**(a)** Cumulative read count distribution across all amplicons in pooled gDNA and cDNA NGS libraries. Most gDNA and cDNA amplicons were sequenced at high depth, ranging between 16,384 (~214) to 262,144 (~218) copies per amplicon. **(b)** Comparative analysis of deletion frequencies in cDNA and gDNA libraries at the level of the entire network (Wilcoxon rank sum test,  $P = 0.037$ ). **(c)** Comparative analysis of deletion frequencies in cDNA and gDNA libraries for every individual target underlying the predicted miR-184 MRE network.



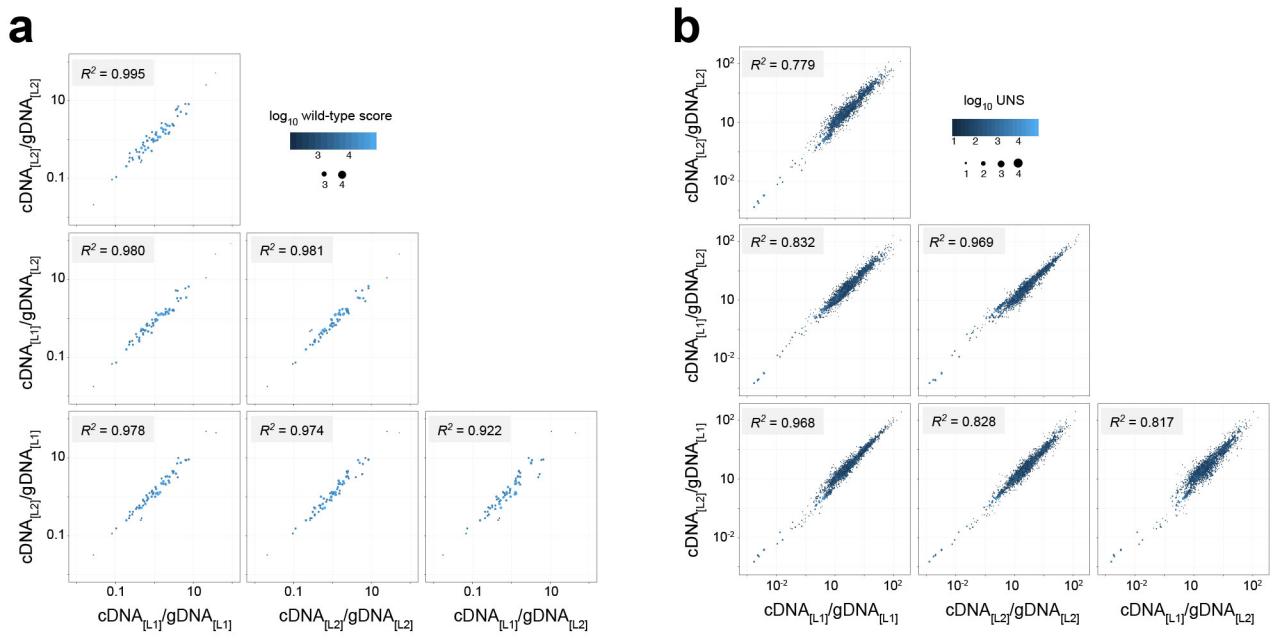
**Supplementary Figure 8. UDP repertoire complexity.** (a) Cumulative frequency of UDP counts across the miR-184 target network. A partition of the data into UDP count intervals is shown in the inset pie chart. (b) Classification of deletion patterns relative to seed coverage. Paired full seed deletions = amplicons carrying the same full seed deletion type in both gDNA and cDNA libraries; Paired “partial seed” = amplicons carrying the same partial seed deletion type in both gDNA and cDNA libraries; Paired “NO seed” = amplicons carrying the same deletion type outside the seed in both gDNA and cDNA libraries; Unpaired deletion = deletion types found only in gDNA or cDNA amplicons; NO deletion = ineffective sgRNA.



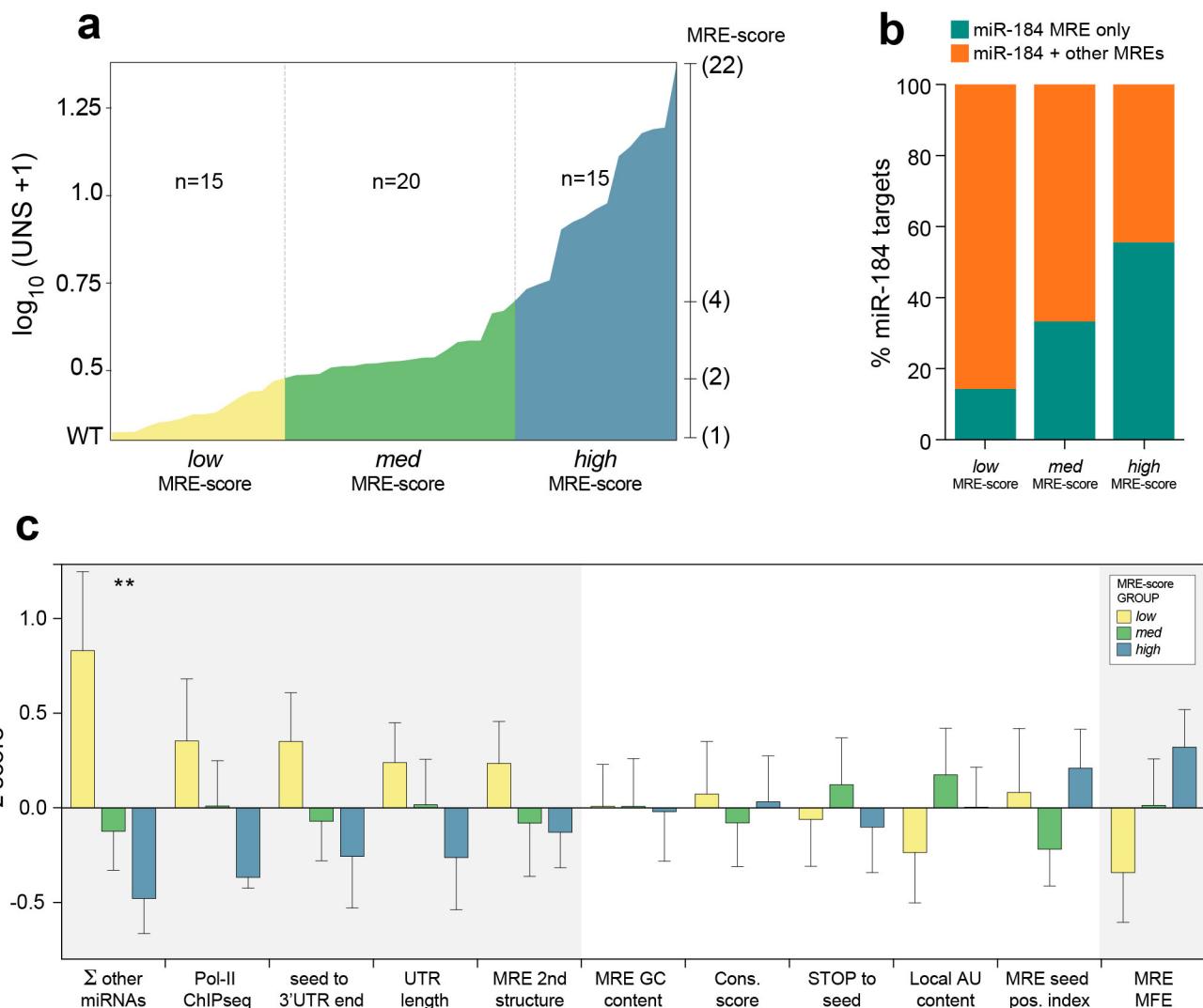
**Supplementary Figure 9. Summary of gDNA deletion frequencies across the sgRNA library used for GenERA-based MRE network analysis.** (a, b) All deletion coordinates were centered by the predicted Cas9 cut site (green line). A biased asymmetrical deletion pattern (away from the PAM) was consistently observed independent of the protospacer being oriented on the sense (a) or anti-sense (b) strand relative to the direction of transcription. A total of 55 (a) and 22 (b) genomic loci were considered accounting for 5357 and 1538 unique deletion patterns respectively.



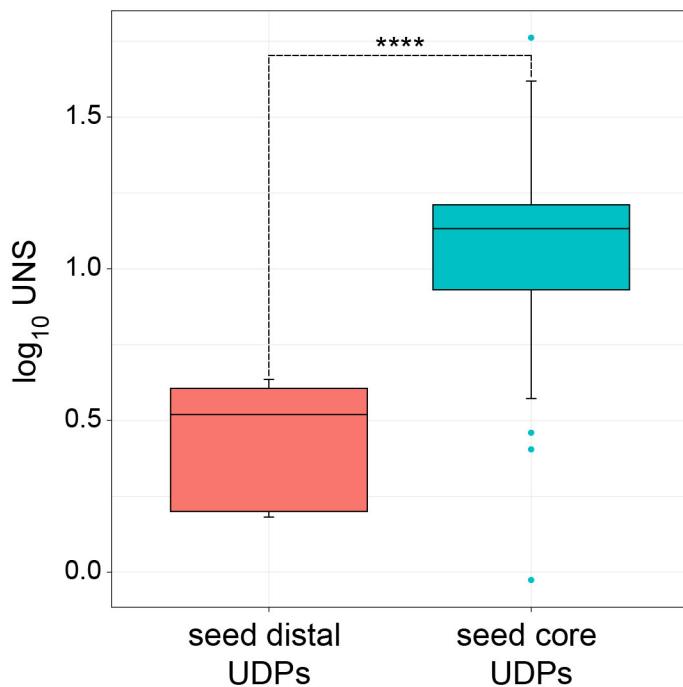
**Supplementary Figure 10. Evaluation of coincidental RNA regulatory elements (RRE) deletions on MRE-score calculation.** (a) Identity and frequency of all coincidental RRE deletion occurrence in UDPs considered for MRE-score calculations. (b) Comparative analysis of weighted average UNS values for seed deleting UDPs sharing RRE alteration events (colored dots) relative to corresponding MRE-score values (white dots). (c) Analysis of UNS values from seed deleting UDPs with or without coincidental RREs (paired *t*-test,  $P = 0.82$ ). ARE = AU-rich elements; PAS = polyA signals; PRE = Pumilio response elements; HBM = HuR binding motifs.



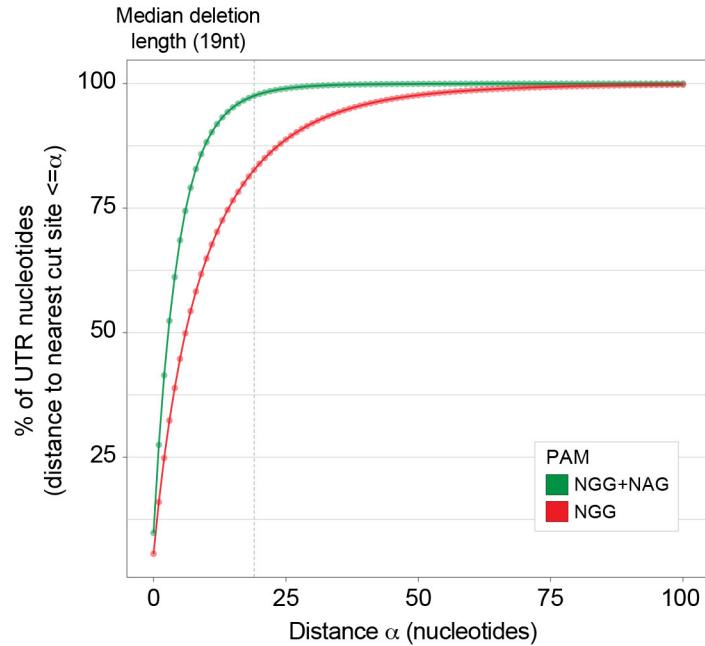
**Supplementary Figure 11. Reliability of MRE-score analysis.** (a-b) Pearson's correlation analysis of wild type scores (a) and UNS (b) between all gDNA and cDNA library replicate permutations.



**Supplementary Figure 12. Correlation of MRE-scores with contextual features.** (a) Partition of miR-184 targets into MRE-score groups: *low* ( $1 < \text{MRE-score} < 2$ ), *medium* ( $2 - 4$ ), and *high* ( $4 < \text{MRE-score}$ ). (b) Percentage of miR-184 MREs co-targeted by top 10 endogenous miRNAs across three MRE-score groups. (c) Comparison of average z-score values for 11 contextual features across MRE-score groups. Features with positive or negative trend across the three MRE-score groups are highlighted by gray shaded areas ( $n = 15$  [*low*], 20 [*medium*], 15 [*high*]), Error bar = SEM, \*\*  $P = 0.0065$ , one way ANOVA). Cons. score = conservation score; MRE MFE = MRE minimum free energy.



**Supplementary Figure 13. Role of *crok* MRE distal seed nucleotides in miRNA-mediated repression.** UNS of all seed deleting *crok* UDPs were partitioned into those deleting core seed nucleotides (nt. 2-7, teal) and those affecting only extended seed distal positions (nt. 1 or 8, red orange). Core seed deletions (teal) displayed significantly higher UNS values than those only affecting distal seed positions (red orange) (Bonferroni rectified \*\*\*P < 0.0001).



**Supplementary Figure 14. Analysis of Cas9 cut sites frequency across *Drosophila* 3'-UTRs.** Graph displays for a given distance  $\alpha$  (x-axis, in nucleotide) the fraction of nucleotides in the *Drosophila* 3'UTR repertoire at a distance equal or shorter than  $\alpha$  from the corresponding nearest CRISPR-Cas9 cut site (y-axis). CRISPR-Cas9 cut sites considered to derive this graph were either NGG PAMs only (red) or NGG + NAG PAMs together (green). Dashed line = median genomic deletion size in the GenERA screen (19nt).

## Supplementary Note 1: CG9257 3'UTR saturation editing

**GCCTTATACCACTGCTGT**TCTGCCTCTGCAAGCTGTGA**TCCCCA**CCGTTGCTCAC  
TCAACG**CTTTCG**CCTGC GT**TGG**TTT**CCT**GCTTGATTTGTGATTGATAAAAG  
**ACTT**AATTATATA**ACTT**ATCGTTAGTCGTACGCACT**CCTTAGAG**T**CGG**TT  
**CCCT**TATT**CCCAGG**ATGCAAAGCTCTAACG**CCC**TGTACTTT**CCA**TATTGATAAG  
CAATT**CTT**TTGTGTT**CTT**CTCGTTGA**CCACTGCCG**T**CCCAGG**CTCGTGT**ACTT**AATC  
GTT**GAGTC**ACTT GATTCAATT**CCCGG**ATCGTTG**CCG**CTGACT**AGG**ACAATGTG  
CGCAAAAGGT**TAG**ATCGGTTTT**GAGACAG**TTTG**CTTGGG**ACTGTTGGTTTG  
ATGACATTGAATAAA**ATTAAC**CTT**GTTTGT**TATT**G**aacaaaaagctttgattattt  
tcatgatgg**CTGGATTGTGCGACGACT**

**NN:** ORF exon

**NN:** 3'UTR

**NN:** PAM sequences

**NN:** polyA signal

**NN/NN:** Fwd / Rev gDNA NGS library primers

**NN/NN:** Fwd / Rev cDNA NGS library primers

To maximize the recovery of NHEJ-induced deletion events, the gDNA 5' and 3' PCR primers for targeted NGS library generation were designed outside but immediately flanking the UTR sequence. For cDNA amplification, the position of the 5' primer overlapped that in gDNA while the 3' primer was placed immediately adjacent to the predicted polyA signal at the end of the transcript (Fig. 2b, Supplementary Data 2).

## Supplementary Note 2: CG9257 multiplex MRE editing

**GGGCCTTATACCACTGCTGTTCTGCCTCTGCAAGCTGTGA**TCCCCATCCGTTGCTC  
ACTCAACGCTTCGCCTGCGTTGGTTT**CCT**GCTTGATTGATTGATTTGATAAA  
AGA**CTT**AATTATATA**CTT**ATCGTTAGTTCTGTACGTTACGCACTCCTTAGAGTTCGG  
TTCCCTTATTCCCAGGATGCAAAGCTCTAACGCCCTGTACTTCCATATTGATA  
AGCAATT**CTT**TTGTTCTCGTTGAC**CCA**CTGCCGT**CCCAGG**CTCGTGTACTTAA  
TCGTTGAGTCACTTGATTCAATTCCCAGGATCGTTGACCGCTTGACTAGGACAATG  
**TGCGCAAAAGGTTAGATCGG**TTTTGAGACAGTTTGCTGGAACTGTTGGTTTT  
GAATGACATTGAATAAAATTAACCTTGTGTTATTG

**NN:** ORF exon

NN: 3'UTR

**NN:** PAM sequences

NN: polyA signal

NN: predicted MRE seed seqeunces

**NN:** Fwd gDNA/cDNA NGS library primers

**NN:** Rev gDNA/cDNA NGS library primers

### Supplementary Note 3: DNA sequences of all genes comprising the miR-184 target network.

**NN**: ORF exon

NN: 3'UTR

**NN**: miR-184 MRE

**NN**: Cas9 PAM

**NN**: sgRNA

**T**: G:U wobble

#### miR-184 MREs in coding RNAs

##### 1. sinu

TATTACAGTGGGCCACACTGCAGTCACACCCAAACAGCAAACAAACATTCAGGAAACCAGCGAAATTCAATAGAATTTCCACTCAAAGGATGCAAAACCATCCACTTCGACGAGGATTCCCAGGAGCACCTGCCAACAGC CACACGCAAAACAGTCACACAGTGAACATGAAGAGACGCACACTATCCGAAGTTGTGGCCTGGCGTCTTCGTGTC CTTGCCTCATCGTATTGCGTTGCAACGCCAGTTGGTGGTCAGTGATTACCGCATACGGGCCAGCAGCTGGATC GCCTGGGATTGTGGGTGCACTGCTCCGGTGCCTGCCAGGTGAACGACGATAGCCAGGGAGATTCTCGTGGCTGC CGTTGGGTGATGACCCATTACACGGGTTACGATGAGATCCGTGGATTCCCTGCTGCCCGCCTTCATGATAGCCACTCA GTTCTTCTACACTTGGCCTCATCGGATGTTGGTATCTGCCATCGGAGTACTGGTCTTCATCCTGTGTGCTGGTCCGG ATCAAAAGCACTTCATTACGCTCATTAATCGTGGGATATGTGCTCTCGGTGCTGGAGTAAGTGCAGCCATTGCA GTG ATTGTATTGCTGGATTGGCAACCGGAATGGCTGGATGCCGGAGCATGCGAACAAATTGGTTGGCTGGCCTTATCCT GGCCTGTGTGGTACTGTGTTAACCTGGCTGGCTACTTGTCTCAGTGAAGGCCACGTGCAGCACAAGAAGAGAA TTCAGTTCAAGGAATCTCAGACGCATTGAGCTGGTGCCTGGATAGTCGATGCCGTGGCAGCTGGCACACTTGACCAA GGAAAACACAAGCCAACCAAGAGGAAAACAAAACAAAACAT **TCCGTCC** CCTCAAAGAATTCTCACCTAAACATTACAG TCGAACTTCTCAAGCGAAATATTCAACTAACAGTTCTGGTAGTGCACGTTTAAAGTTGGTTATGTTGTTTTA TATTTTATTCAATGGTTCCCTGTGCATAGGTTAAAGTATTGTCGGTAACATTGTTTATATTACCTTAATCGTT TAAATAACTCCCCACTGGCGGAGATTGCAAAAAATTAAACGAACCAAAGTTATTGGATAAAAATAATTTCGCCAT ATATTCACTGCTCTCTCAACATAACCTAAAGAAATTAAATTAGTCAGAGAACGTTGACTGTACTCGTAGTTTTAAT TTAAAAAGTTATGCAATTGACGCCGCACCAGATTACAAGCTTCAGTTGGTACTTATAATCATCCATCCAATCGA CCAGATCAGTCCAAGAGAACGACTGAATGTTTCCGAATGAAAACCAGAACCAAGCACAGCTAACATCAAACGAAGACG CTCAACTAGATGACTCATGTTCATGAGTATTCTACAGTATTGCAATTTCGCTCAATTACGCCAGACAC AAAGAGCATACCCCTCCGAATTGACGTTATTAGTCAGGTTATTAGTCAGAGAACGTTGACTGTGAACTAGCATCGAAA TATTACACTCGCCTTAATTAGATAGAACATTAGGTTAGTCTGCTGAAACAAATTATTGTTAAACAT ACCTGAACGACTGACCTGCGACTAACATAGAGATCATTGAATTGCAATTACCTCAGCTAAGCTAGTGCCTTAATTAGT GACCCAGAGAAAGGTAAGCGAAACTCCGCCATTAGTTAAGAACAAAACATTCCCTTGAAACTCGAACATTCCGCAC AAGTACCTCTTTAAAGGTAACATTAGGT **CTG** TATTAATCA **TCCGTCC** TTTGCACGTATTGTAACATTCTAAAC CGAGAATTAGCAATTGATGTTAAAGCAATACATACGTTACCGTCCCATATACAATAATATGTGTAACATTGTTATGT ACAAACAGACTTTAGCTAAAGCGAAGAATAAAAGTTTATTGTATAACACAGCTGAATGCAATTGCAATTGCAATTGCA

##### 2. Mgat2

CAGACTACTCAGTTACTCAGCTACTCAGATACTCGTTCGCAGCGGACGGCGAGCTGAGAGATACTCGATCGCAGCTGTTG TTGCCCCAGCCAGGCAGGAAAGTGAAAACGGAACGCCAGATGAACGCCAATAAATCAAGCGAAGTAGCAACAAATGCCGG AAATGTGAAAATGCGCGCCAGTGGTTGAAGCGAGACGGTAGAGCTAGAGAGATGCCGTTGCCGCTGCGTGT GTGTGTGTGCAGCGGGCGCGTAGCAGCTGCTGATGACGAGATGAAGGCCTCACCCAGTGAAAGAGCAGCAGTACTCG AATCCGAGGAAGCGAGGAGCGACGAGATGTCGAAATGAGGGTAATTGTAATTGAACTCGAGTTCCGGCTTAGAAAAGC CAGCCTGCTAAATCCATTAAATGTTGCCATTGAAATAGGTCGATCTGATCCCCTGCCAACACGGCGATGGGT CGCAAGCGAACAACTTCATCGCAGCTGCCAGTCTGTTCTGCTGGCCCTGGGCAATTTCGGCTCTGCTCCAGTACAACAACTT CAATTATCGTAAGTAATTAGGCGCCAGCAGCTCGCATCACCAGGTAATCTTACTTCTGTTGCTGAGTGGACTCGCGGGACA ATGTGCTGGCGACGCGGTGACCAACGACTCGGACGACGCCATCTGGCCATGGTGCCACCTGCACAAGTATCTG ACGCCCCACTCGCAGAACATCACAGCGCTCGGGAGCAGGAGCCCTCAACGGCGCGGGTAAGCCTTCAGCGAGTTAGCCAAG ATAATGAGATAGTTAACTGGATTGAAATCACCTCTAGGCCCTGCTCTGAATGCCAGTAGTCCGGCGCAGCTACTGCC AGCACCACATCGCTCGACGGTGTACCATCCTCGGAATATCACCGAGATCAAGCGCCAGATCGTCAGGTACAACGACATGCA GATGGTGTCAACGAGGACGTCTGGACCGCTGCGAGAACGACTCTGTTGATAATCGTGGTCCAGGTGCACACGGGATCA CCTACCTGCGCCACCTGATCGTCAGGCTGGCGAGGCCGGGACATTGCAAGGTGCTGGTGTCTCGCACGACTAC TACGACGACGACATCAACGACACTGGTGCAGCAGATCGACTTCTGCAAGGTGATGCAAGATCTTCTACCGTACTCCATTCA GACGCACTCTAACGAGTATCCGGCGTGGATCCAACGACTGTCCCGAAACATCAAAAAGGAGCAGTGCCTATTGGCCA TCGCTATATGTATGGCTATTCTATAAGTATCTGCTGTTCCAGAGCTGCTGATCACCAACTGCAACAAATGCGATGTATCCC GATCTCTATGGCCACTATCGGGAGGCGAAGTTACCGCAGACGAGCACCATTGGATCTGAAAGGCCACCGGGTGTCAA CGAAGTGGAGGTGACCCGCTATCACACAGGACTGGTCTGCTGAGGAGGACCAACTACGTTGCCGAGGACTTCCTGT ACCTGCTGGCTATGATGCAAGCACCAAGAACGACTTGTGCCCCAGTGCAACGTGCTGCCCTGGCACCTATCTCAAG ACCTTCAACTACTACACGTACCAAGCAAGGTGGAGGTCACTGCCCTGGTCAGCAGCAAGCACAACATGGGCTTCGCCCT CAATCGCACCACCTGGTCAAACATCCGGAAATGCGCCGCCACTCTGCACCTACGATGACTACAACACTGGGACTGGTCCCT TGCAGCAGCTCGCAGCAGTGTCTGCGCGAAAGCTGCAAGGCCATATTGTAAGGGCCCGAGTCTCCACATCGGA

GAATGCGGGCGTTCATCACAAGAACAAAGAACACTGCGAGTCCAACCAGGTGATCTCGAAGGTGCAGCACGTGCTGCGCATAGC  
 CCGCAACTCGCACCGCTTCCCAGCTCCCTCACGGTGCCTGAGTCAGCTGATGAAGAACAGTCGAAGCTGCGCAAAG  
**GCAACGGCGGCCGGCTGGGGCACATGCGGGATCACAGAGCTGTGCCTCAACATGACCGTGGCACGAGGTAA**GGCGGGGAGCC  
 CACGGAATGGAACCAATCAGTCAGCAGCAAGGAAGCCTAGCTAATAGCGTTATCTAAGTAGGATTATAAGCAGTAGTC  
 TTAGTCAGTAAACGAACATTAGCAAGTACTCTGCATTAGCTAACTAACGAAAATAATCAAAGGCCAGAAGGCAGAA  
 GCCGGAAGCCAGAAGGCATGAGCCGAAACCGGAAGGCCAGAAGGCCAAAGACCTCGCCTCCGGCGCTGCACA  
 ACTCATACTATAAGTAACCTCTAGTCACACATTCAATTGGAATGCCGTGTTGCAACCTCAATCTTGCGCTTC  
 GCCCTCAGTTGCGATCATCACTGGAAGTGT **TTAGGCTCCGTCCAGATCCAAG**ATCGACAGAGTCCAAGCGACGG  
 GCGGAATCCTCAAACGAAACCTAACCTAGGAATAACTAGCGATAGACCATACACATACATATGTACT  
 AACTCGTATCCCCAGATTGCGAAGCGTTCGAAGGCCAACTTAATTGAGTCTAGATCTAGAGGCAGCAAATCG  
 CATAAAACTAGTCATAGCAATGTATTACAGCACATGCGCTATAACATCATAGATCAAACCTTTGATATATCTACGAGCC  
 TACGTTTAAGCCTGTACATAGAGGAATTGCGTTGATACATCACATAAGGTCTTTATGATATTAACGCGGAACTAAT  
 GTAATTGTTATTTTAATGTTAGTCAAAGCGCTTTATGATAGGAATTGATTAATCGAACGAATAATCCTAG  
 TGAATAAACGCCAAGTAAATGCTCTACAGTATAGACTTTAAAAAC

### 3. CG1105

ATCTCTAACTCTGAATTGGCGACCGCAGGGGTTCAAAACAGACCGAAAAGTCGTTAACTTATTGGTGCCTTGAGTC  
 ACAGAAATCTCTAACATGGCCTTAAAGGTTGTAATTGCAATTGGACAACCCGTGGAACACCTACTACGCCGGTCAGACG  
 GTCAACGGCCAGGTGAAGTTACGTTGACTGCCCAAAAAAGTCCGAGGTATTATCATTGGTCTCTGGGAGAACGCAA  
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 TATCCATTACCGTGTGCTGCGCTTAATCTACCCCATCCTTGAGGGCGAATTGGCATGTGCGCTATACCATCAA  
 GGTGACGCTTGATCGTCCGGAAATTGATCAGGACATGAAGATGGCGTTACAGTAATAGGCCCGTTGATCTAAATC  
 TCAATCCTCGAGTCAGGAGCCATTCAAGCTCGAGCTAGAAAAGTCATTGCTGTTTGTGCTCGCTCAGGACCGCTG  
 GCTGTCATCAGAACATCCGCAAACAGGATTGTTGGGAGGTGCTGCCATCACCTGCGAGGTGGACAACACAG  
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**AGATGATCGGCACGGGCGCTCGCACCTCGGTATCCACTTCCAGTTAATAATGCCACAGCACGCCAGCTAGCCAG**  
**TGATTGAATCTCTAACTGACCCACAATGATATTCTAAACTTGTGACTATTCTCTGGCTGTCTGTGTT**  
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### 4. CG43367

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## 5. CG31195

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## 6. CG8121

CCTGGAAATATCTAAGCCCTGATTGCTATGTTTGTGATGCGTTGCGTGGTGAATTAAATATTGTTTGGGCT  
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**TTTGTGACTGTCTATATGGATGGCGTGGTACTCTGTTGGAACTGGTGGATCTGACCAAGCTGGAGGAT**  
**CTAGCCCATTACGCAATGGCACAATGGCTCCCTGCACCGCATCGACAGCAAACCTACTTGCAAGGATCACGGCAACACCAC**  
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## 7. Mcr

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## 8. Ts<sub>f</sub>

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## 9. cals

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## 10. emp

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### 11. **IqfR (Epsin)**

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### 12. **Ptp99A**

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 TTTCTAGGCACTAGACCCACTAGGCTAAAGACTACTTAAACGCTAGAAGGGCTGTAATTCTCATACGACACCG  
 CGGTGGTAACAAATGCTCATCTGAATATAACGACAAGATCATATTATTCGATACAACAGCAGCAACTA  
 TGCATTAACTGTTAAACACACATAAAATAGTAGTGCATGCAACTATAGGTGGATTTCATGCAATTG  
 TAATGTTGTAAGTGAAGAAAGAAAGAAAAGCTTAATCAACTGCAATTGCAATGAGCAACGAAACACCTTGTG  
 TGATCACACAGAGTTCTACCGAAATCTCCAATTAAATGTGCAACTCTTGTGAGCAACAGCTTCCCGACTA  
 AACTACCTTCACACATTAACCTAAAGTAATTGAGAACGATAGAAACAAGGCCACAGATAACACAGATGATTATCT  
 GACGGTGTATGCAATTATAATGTGATGAAAGTAAAGCTAAGGTTAAACAGAAATTGAGATTATGAGTATATG  
 AAACACAAAAAAATCAAATAAAATATAATAAATATAAAACAAACGATTGATGAGAAAGAGCTGA  
 ATGTAGCGAGAACTTAAGTATCTAACGATGCCACGATTAGAGCTAAGCAAAATAGATGAGAACATCAAATACA  
 AATACCCAGAGGCAGACCATAGGTTAAAGAACGAAACGAAGCATTGCCCCACATCTGACAGGTTACCGCAGCTG  
 GAGGCCATTGCAAGCCGGAGGAGTCGCCACAGAGATCGGACAAAATCTCGCATAAACAGAAATTGAGTATATG  
 TATGATATGTATGACACCTAACGCTAACGCAATGAAAGTTGAAATGACTAAAGAACATGGAACACAGACA  
 TTGAAAGCGTCTAAATAAGAACATCAAATTG

### 13. Sema1b

TCTAGCCCACGTTCGGGTCCGCTGGTCAAACACTCATGATAGTAAAAAAAAACAAAAGAAAACAGCCGAAAA  
 CACAAGACACAAACACAAGTAAGCCGAAAGAAAACCTGGCGAGAAGAAAAGTGTGCAAGAACATTCAAGTAGCCGTATATA  
 AGTGGATTTAATTATCTGATTGTTGCTTCCCTTTGCCACCAACAGCTAATTGCAATTGCAATTAAATAAACAA  
 GTTGTCTGCTGGCGTAAATCATTGTCGCTGACACTCGAATACTATAGCCAGATGCTGAAGCAGCAGCAG  
 CAGCAGCAGCAGCAGCAGCAATAACACCAAGTAATCAGCACAACACTATCGACGACAATGGCAGACATCGGACATCAGAATG

GAGGCAGTCAGTATTAAAAGCAGTCGGAGAGGCCAGCCATCCAACCAGCCAGGCCAGCCCCATTGGCACCAAG  
 AGCAAAAGCAGCTGAATCATCAGAACTCGGAGGAGTAGTGGAGTACTAGTTCGCGCATCTCTACTACAAGCCAGCAG  
 AATGAGCTCCAAGAGCAGACCTCATCGCGGGCATGTTAGTTAACCGATACCCATGATACTGCTGATCACCTATCCG  
 GCCTGACGATCGTGGCAGGTTGGATGCCGACGTAAAGCCTGATTTGCAAACAAAACAAGACAAAGTCCCTAGCCCATTT  
 ATAGGCAACTCGACGGACTATTCAAGATTTGGACCACAACGATGAGTTGTTCTAGTTGGTGCCTAGGGAGCTGTGCCC  
 CAATGTGAGCCTAAATGGCCTGAAGGAGATTGCGCGTCTGGAGTGGCACAGCACGGATGCGGACCGGGAGCTGTGCCC  
 TAAAGGGGAAGCACGAGTGGATTGCCATAACTACCTTCGCGTCTATGCACTGCGTCCCATGGCAGGTGCTGCTCTGC  
 GGCACCAATTGGTATAAGCCGCGTGTGTCACTACACGCCGGTGGAAAGTGTCACTCCGAGGAGGCCGCTCTGCTGGTCA  
 CGCCCATGCCATGCGTACGAAGTCAGCCGAGATGTGGAGGCCAGGGCTGTGCCCTACAGCCCCCCCACAATAGCA  
 CGTACGCCCTGCCGACGGCATTGTACAGGCCACCGTGGGACTTTCGGGTGAGATCCTCTACAGGGAG  
 AACCTGCGTACCGAACAAATATGATCTCAAGCAACTCAACCAGCCGACTTTGTTGGCGCATCGAGCGAACGGTTACGT  
 GCTGTTCTCTCCGCGAATCTCAATGGAAGTCATGAACTTGGCAAGGCCGTTATTGCGCGTGCCTAGGGTTGCA  
 AGAACGATAGAGGTGGTCCCTACAGTCACGGCAAGAGTTGGACTTCCTTAAGGCGCGACTCAACTGCTCGTGC  
 GGCAGGTTCCGTTCTATTGACGAAATCCAGGCCATTAGTCCGATTGTGGAGGCGATCCAAGTCAGTCACTGATTATGC  
 TGCTTCACAACCTCTGTGAATGCCATTCTGGCTCGGCGTATGCCATTAACTGAGACGATATACTGGCCGCTTTG  
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 GGACAGTGCCTGAGGACTCTAGAACCTGACCAGCATGCCGTTAATTCTATTAAAGAACCATCCCTGATGGAGGAGGC  
 TGTGCCAGCAGTCACGGCGTGTGACCAAGGTAATCTGCACCACCGACTTACTGCGATCGCAGTCATCCGC  
 AGGTAAGTCCCTAACTGGAGCTTACTACGATGTTATATACAGTGGCACCGATGACGGCAAGGTGACCAAGTTCATCAAT  
 ATACTGAGCACTCACCGAACACTAACGTTGATGCCCTGAAAGACAGTGGTTATATCAGAGATGCAGGCTTCCACTTGG  
 AACGCCAATTAGGGAAATTGGTGAATCTCCACTCGAAAAACTCGTTGGTTGTTAGCGATGGCAGCTGGTTAGCGTGC  
 CCCTGCACCCTGCTCACATTGCGACTGTCCTGGCTGTTAAGTCTCAGGATCCCATTCGCGCTGGACTTACAG  
 ACACACGAATGCAAGAACCTGGTACGAGTCAGCACAAGGTTGGAACAAAGACTTATTGCAAAGCTGAAATAGCACCAA  
 GAAAGCCGCTGCCCTGCTATGTCCTCATATTCCCCGAGATGCAACCCGGTGGAAACAGTAAGCTTGTACCATGGCTC  
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 CCAACTGGTGGGATGACTTGGTTGAAACAAGATCACCTTACGTCATGGATCCCAGCAGTATTATGCCGAAATTGAA  
 CGATCCCCAAAGTCTACGGCCAGTGTAGATGAAATCAAAATATGCTGTAAGGCTAAAGAGCCGCTGATGGAGGCCAAATG  
 GATGAAAGTCACCGCAATCATTCGCAAGCCAATGCAATTAAACATCAAAACTGGCAAAGACATCAACCTGTTAAT  
 GGGCTCCAATGCCATACGACCACACAAAAACACCTAATCTGGATTAGAAAAGGATCGCAGTCATGAGTGCAAAACT  
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 TGTGAGGGCAGCCAATGCAAAATTAGTCGAATTCTATCTAAACATATTGTTGTTATGTTGCTGTTAGTTA  
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 AGAGCTGTTGCGTCTTTTACTTAGCTGTATTGTAATTAAATTGATGTAAAATTGCGTAAGCTAGTTCTTGTC  
 AAAATTAGTTAAATTATTGACAACGTTAGGTGCGGGCTGCCGAGCCTGGAGCTAACCGAAGCATTCCGATTCCCC  
 TGACGTCGACCTTGCTGCCAAATAGGAGTGGCGCTTGGCTACGCAACCCGATTGTTGCGCCCTATGTTACT  
 CTTAATGTATAATCTTAAGTCGAAATATTAGAGCTGTACAAATTAACTCACATATCATGCATACCGAAATCGAACCGA  
 AGCGGACCAGCCCCAATTATTCTCGAAATTCAATCGAATTTCACAACAGTATTGTTAGACTGAATGATTAAATTATC  
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 AAATAAAATGAAACACTGCAACACAGCAAGCCTTAAATAGTATAAAATCGTGTGAAATCTAGGCCTAACGCTAAACTT  
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 AGCTGTTG  
TCCGTCC  
CACTAAAGAAGTGTGCATAAATACAATGTACAATGGCAATTGTTATTGTTAAACGCTGAAATGCA  
 ATTTCATCGTAATTAATATTTTACGTATATGTCAGCCTAATGTCATTAACATAAGAAAGTCTTAAAGT  
 GGAAAATCCCAGATAAAAGCTATATATAAAATTTTAAATCTACAAAAAAACGTAATGTAATGATTCTCATATTG  
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 TCTGATTGCGATGCGATGATAATTGCGTAGGAATGGGCTTCCACTCGGAGGATCAAGAACCCAGAG  
 TGTTAGTTTATTACTGTGTATGCTCTGTAATGTTGTTGAAACATGATATATGACGAAACGAG  
 AAAGTCGTATTTCATAACTC

#### 14. CG10217

CCAGTGTAGTAAGGTCGCCATTGCCCCATCATGTTGTTAGTTCTGAGGACACCAACACGCAATGCATTATTACCTGG  
TGCTATCTGCCCTGGCTTAATGGCTCAGTACACCTGGTCAGAGGACAACGTCATCATTCAA  
AGATCCTGCAGGGATCGTGGTCTCGTGGAAACGGGCCCTGCCACCGCTGACGGTGTACGCCACCTCGATGCCAGC  
CGAGGCTACTGCATCAACTACATGCCACAAAGGGCAGCAGACTCATTGCTCTCAAGGAGCGATCCAAGGACTGCTA  
TCACTGTCAGGACCATCATCCGACCCCTTAATGTTGCGAAAGTTGAGGGGCCCTGCGTGGCATAACCGCTGGAC  
AGGAACCCACCGTGGACTATGTTGCGTGGTGTAAAGGATGACCGAGCAGTTGATAACGCTCTTAACGAGAACCTTG  
CCCACGTCAGTGCCTGGAGGAGTGTGGCATTTACGTAACGAGCAGTCGATCCGATTCAACGCTGGCTATGCGA  
TAAGCCGACGCTGCATTGCTGAAACGGCAGGAACGCGAGTCCTCATCCAGAACGAGTCACGTCACCT  
ATCAGCAGTGCAGGGTATGGAGGGCACATTCAACGGCACCCTGAGGTTGAGCTGCGCTGGCGATTGTTGTTG  
GGCAAGAACCAACTTTGCGTGGCCAACACAAAGGAGTCGCGTAAGGATGAGAAGTACCGCTGCTTCTTAAGAATCGT  
GACGA  
TGATCTTACGTGGCGTTCCATTACGGCGAATGTAACACACTGAAAGACGCCAGAAACGTCACCGGAGCGTCAAAGC  
TGACGCCAGTGAAGGCTGAGTTGGAACCCGGTTGCACGCTACCAACAAACTTACGCGAGAACACTG

AACATTGATGCCGATGTCGATCAGTGAGACCCACATAAACGAGACGTACTATCCGGATAAAAGCTCGATACCGAAAGAC  
 AATCTATGTTGCCGAGAGCAGCGACGGGTAATCGTGTGATGGCTGCCCTACTGTCAGCGGTTGCCAGAAGGATTACG  
 TGTGCTTGACTTATGCCCGCCACCACAACATCATTGCTACCGCAAGGGTTGGCTGTGATCAAGGACGATTTAGT  
 ACAGCTGCTCTGGGTTAGTCCCAATTGGAGGCATGGAATATGATCTGTTCTGCTCGAAATCCTGTCCTGT  
 TCGTTGCCCGTTGGAAAATTCAACTTACGCGAGGAGAGCATCCCTCAGAACGAGAATTCTGGTGGTCA  
 CTCTGAGTCCCGTCCCAGATATCCATTGCAAGAAAACATCTCGGATTATCGTGTGCGATACAGATCAAAGAATTG  
 CGGGTGGATGAAAATTCGCTGCGTGGATCACTTGGGCGTGTGATATCTACAGTGTGATCCGATTACCGCAT  
 GAAGTGCATTGGCTCTGGAAGGAGAACCTCAAGTCCACCTGATTACGACGATTTGGATCCGCTGTCCAAGTATC  
 GCTGCTGGGTCTATCAGCGTCCGACCTCAACCGTGTCCATGTCGAGGCTGTGGCGCATTCTGCAAACACTGGAGCAA  
 GATGTGACCTCGTGGACCCTCGGAGGGCGCTGCCGTAGCCATCGATGCCGTGGAGTACGAAAGGGAGCGTGACGACTG  
 TCCCAGTCAGTCGATGACGGATTGAATCCTTGAAGCCGTCGGATGCCCAACATCATATTGATGGACTTTACC  
 GAGCAGGAGCAGCAGACTAAAATTCAACTGACTGGCTCGTTATCGTGGTCTAGTCCAGGTAGGAGTTATTAGCT  
**TAA**ATAATCAAAGTATTTTAACTATTCCATCCAAGAAATTACAAAATTAACTAACACAAACATAAAAGTCACTTT  
 AGGAGCAATTGCGTGCCTAGAATGCAATTAGAAATTAAAGCAATTAAACTTAGTTGTAAGATCATCACAA  
 TTTTACGTGTAACAG ***CCAT TCCGTCC CAAACGAATCTCAAATGTATTGCAAATAATTAAAGCAGTTAAATAATGCC***  
 TTGAAC

### 15. IIp8

CCGCTCGTGATTATCAGGTAAACGTCGATCGGACGGACGGGTTAACCAATTCAAGCAAGTTAGCTCTGCAGCAAGTTGCCCG  
 CCAAAAGAAAACACGTATACTGAGTGTGCGAGCCAGAAAGGTGATCGACGTGCTACTGATCTATTGATCTATTGAGAAT  
 TGTGATCCTAGAACGCCACTAAAT**ATGAGTTCAAAGTTGCATATGTGCCGCTGGATGTCGTCGGTATCGGAGTCTGTTGC**  
 CTGATGGGACGCTCGTGGGACGCTCTGCTCCCTGGAGCGGATGAAGAAAGTTCGCGATGGAGGCGTGCAGCACCTCTT  
 TCAGGCGACGAGGGTGCCTGGCGACAGAAAGGTCCATCGAGTTCGCGCACCACCATCTGAATCGACTTGGATCCGGCA  
 AAACGCACAACAAGCATCACTACATCAGCCGAAGCAGCTATCCAATGGGTGGCTACCTGAAGGTGACCCGGAGCACTTC  
 AATCGCCTCAGCGAAGTGGACATCTTCCGCGTACAAGCCCCTCAAGCCGACCCACGAAAAGACATCGCTTCAAGAG  
 GGATCACTCGAGCAGGTCTACAACAAACATTCCCTACTGCTGTCAAACCAGTGCAGGAGGAGTCTTCTGCTAAAGAGA  
 GCTCCAGTGTATGCTCATGCCACGCTCCAAGTCGGTGCCTGGTCAGAATCTCTGAA**TCCGTCCATAGTTATA**  
**GTTAGG**CTATTATGTACATTCTACAAGTCGCGCTCACTTTATAGAGTACCTTATTGTCGCCACTCCCAGGGACACAG  
 ATCCATTACAGTATTACACTTAGATGATTGAAATAACAATCGAACACGGCAACTGATCGG

### 16. CG4313

GCGACTGCCTCGGGCCCACTTTGGGAAAC**ATGTCGGTGTATGGATGCCGAGAATTCAAGTTAGTACAAATT**  
 GGCGCAGGAACGAGCAACCGCTTTGCCAACACAAAGTGGCGCGATGTCCCCACCAAAATCCGGACCAACTCCGC  
**ATCCGATCTCGATAGCGCTGCCCTTTGGATCCCTATGCTACTGATCACATGGCGGATCAGGACGCCGTTGCCGGTA**  
 CCCCTTCCGACAGCCTGGACCTGGACGTGGACGCTCAATCTGAATATGAATCTTAATCTCAATGTAGACGACAG  
 GCGCCAGGTGCTTCGAGGGCTATTGGACGAGCTGTCGACCATGCCCTGGGTGGCCTGCATAGTGGTCATATTGTGG  
 GCGTCCCGGAAATTACTGACCATAGTGGCCCTGTCGCGAGGTGTCGAAACCCGCAACTCCACGGCCATATTCTATAATC  
 AATTGTCCTGCGATCTACTCTTGGCTGCTTAATCTGCCATTGGCGCTCCACTTCAAGGAGCGGGCGTGGAC  
 CCATAGTGTATGCTGTGAGGTTGTTCCGATGCTGCGTACGGCTTGTGGCGGTGTCCTGCTCCTCGTGTGCTGA  
 TCACCATCAACCGGTACATCATTATGCCCATCCCAGGCAGTATCCAAGAATTACCAAGCAGTATTGGCTTGTGATG  
 GTAGCCGGACATGGATCACACGTTCCATCATGCATGATCGGTATGGTAGATCCCCAAGGAGTCTCTCATGCCGCTTCATGG  
 TGCCGTGCATCTGCATCGTGTATGCCGAATCTCCTGTCGGTGCAGAAGGCCATTGCGCGGGAAACGCT  
 GGCAAGACCAATGTCAGCGATGTGACTCCAGCTCCGCCGCAGCATCAGATTCAAGCCATGGCCACACCAAAGAAGCC  
 GGAGAAGGTGACCACGTCAGCGGCCAGGCGAACGAAACCGATCGCAGGACGACCTTGTGGAGGAAAACCTAGCGT  
**ATATCGATGACAATGCTCGACGGACAGTCTGCCATCTGTACAGCATACGGAGGAGGGATCAGGACCAGCAGCCACCG**  
 CCTGGTGTGCCAATGTGGTGTCAAGGAGCGGGAGAAGGAAAGGGATCGGGATCAGGAAAAGGTGAGCCTGGGACGAG  
 TCAAACGCAACTAGAGATGGCAAACCCATGAAAGAACCCCTATCACTACATCCCTGCCGACCCACCTCCTCACACGGT  
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 AAGGACGGAGATTGCTTAAGATGATACTGGTAATCTCGTGTGGTTGTAATTGTTATGCCATACGGTGGCCAA  
 AATATGGAAAAGGCCACCGAGGTGCACTGGTCAACATTGCTGGCTATCTACTTACTGACCACTGCACTCAACC  
 CGCTGATCTACGGTGTGAGCTCCGAATACCGAACAGAGCTTACTGGAACCTGCTGCGCTGCCATGGCTCTCCAGATACC  
 CAGAACGCGGAATCAGCCAATGCAAAGCGAACAGCACCTGGAGTCAATCGCAGGTGAAGGCCACGACGTGACTCCG  
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 AGAAGTACCCAAATGTCATGGTGAATGATCTGCCGAAGAAAAAATTCTGATCCCCTGGTATTAGTGCCTCCAAA  
 AAAGATTCTAGTCACATGATCCGGCGGTGTCCTCCACGCACACACGAAAACACAGATCACCCATATATCTATA  
 TCTATATCTATATAATGTAGGAACCTCAGAAGTATATTGATTTATACCAAATGACTTAATATCTGGTTTACATTG  
 ATTCCATGTTAAGAACATCGATGTTGTTAAATTGTTTCTTGTGCTGCTAATTGTAACCAATA  
 CATAATAGATATATCCTATATGGATACGCCCTTCTATGCCCAATAATGCTCGAATAAT

### 17. CG6583

TTTTATAGCGATCTACCATATGGTACACAGTAACCTCGCGTTATTATTTCTGTTTAATTCTATTAAAGTT  
 AACAAAGCGCACAAACAGTTAAGTAAAAAATGAATGCTCTGCCGTTGACGTTCTGCTGCTCACCCGCCATA  
**TGTGGCAGCTGGCCATTGATGCTACCAAGTGTGCTCCGACCAGGATCGCAAGGGACACGACAGCTGTGGCGCTACAA**

GAGATTCAACCGCACCGAGCACATTCCATTGAGTGCAACAGCGACGGAGGCCACATGCCAGGATCCTTCTGCATGAAGG  
 TGGTGCAGCAGGGACCGAGAGGATTCTGGACGGCTGGGCCAGGTGATCCGACGATGTGCATCCGTTCCGAT  
 ACCGGAGTGGTGGCGTCGTAACTGGGAGCTACGAAAACGGAGTGTACTGGGAGGAGTGTACTGCTCCAGCGACAG  
 CTGCAATGGTCCAGCTTAGTCAAATGCACGGATCCCTGCAGCTCTGTTGGACTCTGTTGATTGGCGTGCCTCCA  
**GAATATTAGAAGTTAATAGGAAAACCAGGCTATCCTCACCAAAACGAAACAGCAATAAGTGCCTTTCGCCACTCC**  
 ACATGATATCTAAGCTAGGATTGA **CTTGTGATATCAAAAACACATCCGTCC** ACCTTTTCTGCCTATAATTAGTCGA  
 AAAGTTGAAAACATTTAATCGTATAGGTACATTAGTTATCTACAATATCCAACAAAGAGGCCAGGTAAATTACAGACCA  
 AAACTAGCTAAATATTTAAATGAAAAGTAACGTTAACAAATTGAAATTTCCTATGTCGTTGACTTTATATAGTT  
 AACTTAATGAACATGTTCAAAATGATTACTACATTGATTAAAAATGACCATCCGAAAATGCTTACGTTG  
 ACATATTATTACTAGTATTACGCCATATCGAATAACCACTACCATATCGAAAACCAGTAAATTGTACACACAGA  
 ATATGAAAATATATACTATATTAGAGTATT

### 18. CG7713

ACGCACGCATGCCAACCCATTGCTCAAGCCGTTGTTTTCTAATTAGCGATTATTATTTAAAAAGCAAC**ATGG**  
 TAGTGTGAGCTCCTGCTGGCGCCATAATTGGCGATAATGTCGGACGGAGCTATGCCGTGGCGGTATACG  
 GCTGCCTATCCGCCGTAAATGATCACGCTGATTCGTATGCTAGCAGGCGGGAGTCTGCGCAGCTCTATTCCCCGCT  
 CTTGAGACGGATATACGTTGCGATGCCGTGGCAGGGCTTCTTATCATCTACTCCCTGCTAATCATCTGTCTT  
 CCTATTGGTCTACTACGGTATTAAAGATCAGCACTCGAGGATGGCTGCTACCCCTGGCTGGGCTAATTGGACTAGCCATT  
 CTCTTCAGTTAGCTGGAGCCTTGGCTTATTGGAGGCTACTATATTATCTATTGCTGGCTGGGTCTTCGCA  
**GTACAGATATTCCGGAGATCCAGAACATCCGACATTGAGCTGCTCATGCCATGATGGAAGAACCTCCATACAGAAGCTC**  
 GAATAATCTATTAGAATACTTTGA **CTTGACACGCAACAAAGAACCCGTCC** ATGCGCATTTCACACATTACACGATCAATTACT  
 TATGTGTAATTGAATGCCCTAGTCTTCAAACGACAATTAAACAAATAGATT

### 19. pck

CGAAGATCAACAAGCCTAAAGTAATAAAATTAAATAAAAAGCCCCACCTAGTTGTTGCTGCATTTCGGGTGTGCGTT  
 GTATTGTTACTGACAATCCCCAGAAAAGAAACTCCTGCTCCAATGCGATCCCTCGAGACCCAGGACAGCAGC**ATG**  
**CGCGAACTTAACAAGCAGCAGTCGAGGACACCACGGACTCGGGCGCGAAAAGGGCGACTATCCGGGGCCACCAATGG**  
 CGTGGCTTCGGGCCATAGTTACATTGCCAGCTTTCTGCTGATGATGTCCTTGTCCGCGTACTGGATCGAGT  
 CCTACGAGGAGACGGCCAGCTCAAGAACATGGGCTGTGGCAGTACTGCTTTAAGGACTTCGTTGACCCCAAGTAC  
 GCGTCTTGAAGCAGTTACCGGTTGCCACAACATCTCTCGCACGTGAGTACAGCAAACACTGGCCGGTCTTACCTTGT  
 CCTGATCTCTCCACTCGTACAGGAGTATTACGTTACCGTGAGTATCTGCTGCCGGCTGGCTATGGCGGTGCAGGG  
**CTTCGTGACCATGTCCTCATCGTGTCCCTGTGCTGGCCCTGCTCGCTTACCATCATCCGACTGCCTCTTAAGG**  
 CGTCTGCACTGAGTGGCTGCTCGCCCTCTACATGGGACCGCCATCTCGTCTGTGTTACGTTGCTGTTGCGCC  
 GTATGCATCTCGGGGCTGCGCTACCGCCGACTGGATGATGACCCCAAGTTCAACGTGCTGGCTGGCTACGC  
 CCTGCCGTTGTCACCTCATGTCCTGGTAGCCGCTCTGATCCCTGAGCCGAGGCCGGCAAGCCTACGACGCC  
 GCGGGAGCAGAAGAACCTGGAATGCAAATGGAGATGCAAGGAGCCGGTTACAGCCGCGCGCCACCAACAGCCAG  
 TCGGGAGCCTGAGGGCTACATATAACAGCTGTCGCCACTTCATACATTCCGTTGATGGATCTAGCTTAAG  
 TTTTCAGACGCAATCTGCGCTGAAT**TGCGCAATCGACCCGTCCATAG** TGCCCTTAACAATGACCATATGTATCT  
 ATATACACGCCACTCAGCGAGATCAGCTGATCCGTTGTAATTGATTAAATAGCACGTACAGCTAACAGTAAGAT  
 CAGTGCAGCAAAAGATCCACTAACATTACCATCTACATTATAGGCTAAAAATGTTACAGCT  
 TAAATTAGACCATATCCACCTTATAAGCAAATGGGTGAGAGACATATCAACATTGAAATTAAATGTATTATTG

### 20. kune

CACACTTAGGGCGTGTATTTCAGCCACTTGTAAAGAAATTATAGAAATAACAA**ATGGGTGCGTCAACCG**  
**CATTTACAAGTTGCCCTTGCCTCGCTGTTCCATAATGTTGTTATCGCTTCCACGCCATTGGCTGG**  
 TTACAGACGGCAGATTACAAAACCCGCACTAACCTAGGCCTTGGGAGTTGTTCAACAACTCCAGGACATA  
 CACCGATTTTGACAATTCTTAACGGATGCCATGGGTTTCGAAGAGGAATTACATCATAACAGACTCCCT  
 GCCTGGCTCTACATATCCGTTCAACTTTGCTACGCTATGCTTGTAAATGTCCTGTTATACCAACTCAGTAG  
 CATTCTCGCACTCTCGCGATGATGACAGGTACATGGTGCTCTGCTGGCGATAGGATCATGCCAGGTTGTTGGCTCT  
 GTTTGGGTTATTGCTGTGGTCTTGGGCCAAAGGAGATTCTCGGGATTGGATGCCAGGATGGCAGAACATGA  
 TATGGGCTGGTCTTGCTGGCGTTGCGCTGCTGACTGCGCTGGCGTACTCTATCTAGTAGAAAGCAGC  
 GTCGGGAAAGGTACAAGCGCTAAACGAAATCAGCAATGGGAGATCAGTGAATACGGAGATGAGTATTACAGAACATCAG  
 GGACCCCGCAGTGCCTCCAGCGCTCCATGCCATATCAACCTATTGCACTGAGCCAAGCCGACCAAGAACACC  
 CCAACAAAGTAGTCATCGAATTCCGCTGTACAAGGAGGAATTCAAACAGATATATAAAATCAAAATGAATTGAAAGGAA  
 TTA**CCGTCC** ATTACACAAATTTAAATAATAATTAAACAAATACAAATATACGATGTAAGGTAGAATAGTCATTCC  
 ACTAAATCCACTACAAATGTTTAAATGTATCGAATTGAAAGTAATAATTGTCATGAAATAATTGTCATGAAATAA

### 21. crok

TCGAACGCTGCTCACACGGTCACACTGCTATTACACGCCAAAAAATTGAATCCAATTGAAACGCGAAGAAAAACATA  
 AATATAACG**ATGAAAACACTGGAGAAATATATCCATTGCCATCGTCTGCTGCCTGCTGCAATTGGGCCAGGCAT**  
**CAAATGTTGGACTGTCGCCAGCGATAACGATCCAAATGCCGGGATCCTTGTACAACACAGCACGCTGCCATACCGATT**  
 GCCAACAGGCCGGAGCTGGAGCATCTGAAGGGCGTGCAGCCGACAATGTGTCGCAAGATCCGGCAGAAGGTGACGGT  
**GAATGGCGCTACTTCGAGCTGCCATGGCGAGCCGGCATCGAGGGCGACGAACGCTCTGCCTGATGCGCAC**

TGGCAGCTACAACATCTTATGGAGTTCTGCACGTGCAACAGCAAGGACGGCTGCAACTCGGCGGGCATCCATAGGCTAG  
 GACTGATGGGCGTCCTAAGTGGCACACTGCTCTCGGTGATTGTGGCTCATCTTCTGCGGAATAGGTGGATCCACCTGCT  
 TCAGCTGTAGCCAATGCCACGCCATTGCCACAAGGCTGATCAGATAGTATAAAACTGTATTAGCATGCTTGCTTGTGT  
 GTTCAACGTAGATATAAAACTGAAGAATTCTATAATTGTAACCGAGATATTAGCAAATATTTAAAATTGTAAATT  
 ACCTTAATTAGCAGTAACCTACGCGCGAACAGTCTTGTGGTGAATGAGCGATGTGAATCAAGTGCCTTCTGT  
 AACGGACTCGTATTGTAACCTAAC **CCTGGAATACTATTCCCGTCC**AAAAAATATCTTACGCCATGACATTCCATCTCGT  
 AAACTGAATCTGTAATACTTCTATTTCTACATTATTTGTAATGTTTGCAGAAGATGCCAGTGGTAGTCAC  
 TTACAAGTTTATTGTAATGCGATGTCAAATAGAACATTGAAATAATTAAAAAAAAACGGTTACACAC

## 22. cold

TTGGTGTGAGAACGCGGTTCTTCAATTGTTGCTAATGTTTTGGAGTGGTTGTTGTTAATTGACTTGT  
 TTATTACTTGTGAGCTTATCGGCTCGATTGCGTTAATTACTCCAGATATACGATGAAATCGTGGAGATAGCTGTGGTG  
**CTAGTCGCCGCTGTTACTTGTGTCGCAAGTAAATTCTG**TGGCCGGTCTGGAGTGCTATGTCTGCTCCAACCAGACGGG  
 CAACACGGAGAAGTGCCTGAACACCATCAAGACCTGCGAGGCCATTGAAAACGTCGAGGAGATCCGTTGGGAT  
 CCCAGCGTACTTCTCGGAGGGTGCCTTAAGCAATACTACGTCAGGAGTCATGCCAGAAGGAGCAGTGCAGTCC  
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 TCGGTGCAACTACTTGTCTTGTGCTATTAGTGGCGCTCTCAAGACAAGGCTATGGTGCTGTTAACCTGCTCACCGCACTAT  
 TAGGACTTGGATCTGGCTCATTCCACGCTCTAGCCTCAAGAGCCTTGCTATACCAAGCACATTGCTTTGTATT  
**ATTTAAACGAAGCCGTCCATAAG**CTAAGTAGATTCCACACTCCATTAAAACACCCGTCAAAGTGCCTTGACTCAGTT  
 GGGAAAGTAATGCCATTACTATCCTGATGCCTGCGAAGCGAAGTTCTTTCTAATAACTTACATTGTAACAA  
 AATTGTTCTGGAATTCAAGTTTGTGATTAGTTTACGTTATGGAGCAGTTTACGTCATGACATGACAACAA  
 GAGCTATTGATTATTGCTTATTGCAATTGCTGATGAATACGACATATCCTAGACAAATTAT  
 CTAAGAGAAAGCTAAACATGAACATTAAGCTACATATAAGAATCAAAGAATGCCCTAAAATAACAATCTGTTCATA  
 TCGAATCTCTCCCACAATAAGCAACTTGTGACGTCTTGGCGCTTAATAT

## 23. CG13088

ACAGTATTGTTTGGTAATCTGTTGAACAAAATTGATGTGAGATGGAGCAAATAATGAAGATGTTGGATA  
**GATATACTAAAGTACCTACCAATGAGGGATCAGTGTGCGTGGCGAA**GTCAACGAGAAATATATCGGCATACGTGAAGTA  
 TCATTGGAGTCATCTCAAAACAGTGA
 AATGTCGGGGCTGGTGGCGAACGGTGGAGCGTTGAATTGCAAAGTGCACAGCAGTGA
 GAATACGACTTCCCCACCTCCGATCGCTGACTGCCACATGGACTACAATCTGAGGAGGCGGACGAGGAACCCCTCT  
 GCTAACCGAGCTGTTCCCGCTTGTGACACGGCTGTCACTCAATAGCAGCACAAACAGGCTGTTACTTGCGACTGGAAAG  
 AACTAAGGGAGTTGAATCTCACCTGGTGTGAGTACCTGGACCCAGACCACCTCGAGGAGATCTTGGCAACCTACAGCTG  
 ACCAAGCTGACTATGCTCTTACGATATAACGTAACCTAGGCGAAAAGTGTGGACATAACGCGATGTACACGCT  
 TGAAGAGCTGACATCGATGACCACATTGCTGGGTGATTCTGCCCCGACTGATGAATCTGCCACTTCCGCC  
 TGGCTTTTACACCGAGACTATGAGTATTGCTAGGTTCTGGTGGCCAGGCACAAGCCGCTGAAGGTTCAATCGCTT  
 CTCTCAACGACTTTCTGGAGCAGTGAACGGTGGCGGAAGTATACTGCACATGACCAATCTCGTCCGCTGGCCT  
 GCAGGAGGATGACATCGATGCGCAGCAGCTGACACCCATCTGCCACAAATTGCAAACATTGAGGAGCTACACCTTGG  
 CGATGCGAGAAGTCCGCTCCAGCCACCTGTGGAACTCTAGGTCACTGCACATCTTCTAGGATTCTAACCTATCG  
 TCTACCAAGCTGGTGGACCTCCGAAGCTCCTGCCGACAAAAGTCTGCTAAGTCGAGGATTCCACTGACCTGCATCT  
 CCACAAACACTGGATTGGATCCTAGTAAAGTCTACGGGATTGATAGTGCCTCTTGAGAAATCTCTTGGAGCCGATCC  
 ATTTGAATATTGGAGTCCCGGATTGTGGAAATCGAATTCACTGAGACTCGTAGTAGTCATTGAAAAGACTAAATG  
**ACATTCTACCTCAA****TCCGTCCGG**TATACGAATATATGAGATGGAGATCCAAATGATATATCCTGAGTAAATGTT  
 TACCCCTTGCAAGGGTATAGTCAAATAGCTTAACAAAACCACAAAATCTTATCTGCTACCTACCAAAATCGTATT  
 AAATTCAAGAAAAT

## 24. Cont

GCCCACAGTCGGCTTGCACATCTAATTGCTGGTCACTCAGCTTCTGGACTGTTGTTTATTAAATTAAACTA  
 TAATTAAAAAGCTGTAATTGCTCTGATACATTATTGTCGGATCCGCACAACTCGTGGTTTTTTGTCAGCA  
 GGAACACCTCGCAGTGGTAGCGTTCAACCCTCGTGTGCTGCCCTGGAGTCGAATTAAAGAAACTATCGAATTCCA  
 GAGAAAATGTGCTTACTGCTGCGAGACATCTAAACCAGAACCCACGCTATTAAAGCTAAGGACCAACT**ATGCTAG**  
**CAAAATCGGCCCTCTGGCGAGTATTCTGTGCTCAACTTAGTGGCCAGATCACGCCCAATTTCGAAAACCTGCCG**  
**GATCCGGATCCGAGTCGGGACAGCAGCCTCAGAACTATCAGCCTAGCTACAACAAGGATTACTCTCCGCGATACAACCC**  
**CCTGTATACTGGACAGCAAAGCGCTGACCCCAACCAGTTGACAATACGCTGGTCAGGGACAGAGCCTAATACCTACA**  
**AGGGTACTATGATGGCAGGGCAGGAGTGGGGCTTGGCGCAATGTAGTCGGGCTGAAACACCTCGCGGACTC**  
**GGCCACAGTACGACCCCTTAATCGGAACAGCATAGGTCAGCAGGAGTCAGCTACCGGGACGCCTATACGGATGAAGA**  
**TAACCTTGTGCCGGAGCAGTGGCTCGTTGACAGACAGCTGCTATCGCTTACCGCTGCCAAGCGCAACTGGCAG**  
**AGGCTAAAAGATCTGCAAGGCACACAATGCCACCTGATAAACGTTGATAATGTAGAGAAGCACTCGTTATACTCAAG**  
**AATCTCATCTGCGAGAACGAGACCGATTCTCATCTCCGCCGCCAACGGGCCGTTGAATTGGGTCACGCA**  
**TGACAAACACCCAGTTAGTCCAGATCGAAGACTCCTTTCATGGACAGCAGCAAGTGCCTTGGAAAACGAAGATCTGCACG**  
**ACAATAGATTCTGTGCAAAACGACCTCAACAAACAAATCAATAATCAAATCAGTTTACAACCTCTCCGGGG**  
**ACTGTTAATCAGCGCAATCAAATAACCTCGCGCTTATAGTCCAACCAGCCGTATGGAGACAACGATATGTGCG**  
**CGATCGCGTTGTGACGCCCTCTCAAAGAAAAGGGATCGGTGGATGTCATGCCAGCCTATGAAATTGAACTAAATCTCT**

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 ATACCCGACATGCCACAAATATCAATACGGCTTATCTAAGCACATTGCTACTCATTGACACCTTCTTACAAAGC  
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 GTAAGTGAAGTTAAAAAGAAACTACATTCCATCCGTCTTGCACAGGCCCTAATTGATAACAAATGCAAT **TCCGTC**  
**CATTTTATAGACTTTATC**AAAAATAATTCTAAAATTGATAACAAACAAATAATGAATACCAATTATGTTGCAAGT  
 ATGTTGCTGAACAAAAATAAC

## 25. mth15

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 AAAAAGTGTGCCCTACTTGAGTTCTGGCATGCAAGTCATGATCTGGCTCACAGCCGGCACCTGATGACGCAATTCT  
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 TAGCCTGGCTATTCTCATATGTCTGGTGCAGATGGAAGGCCGCTTACGCCACATCGTAGTTAACGCCCTGCAG  
 ACGCCGCTGGCTATA CATATGCGTGCCTGCCAGCGACATGTGACATTCTGCTGAAAAAGACCTGCTACAATGA  
 GCCACCGTCGGCAAACGACTGGGCGATGAGCTGCATTATGAACGGCAACGATTACTGAGGAG **CCACCGTCCACACTA**  
**ATATAAGT** GTTTCTGCTAGTTAGCAATGTCCAACATTGAGATTGTACTTAATTAACTAATTATGCTGACAGATA  
 GTTATTAAATGAGAAGTCTGACACCT

## 26. CG8010

GCCAGTGCCAGTGCAGCCAGGGCTGCATGCCATCAAGAACATCTGTGGCAGAACAAATAAAAAAGGAAAATACTCGC  
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 GCTACTGCAGTGCACAC **ATGGACCACACACGTACGCCAAGAGCGGATCCAAGAAGTGGTCCGCGGAAGAGAACGCG**  
**AACCTGGTGGTGCAGCGGATCGCGGAGCAGCTCTCCCCTACTCACCGAAGCACGCCAGGAGCCATGGAAAAAATT**  
 CAAGGAAATGCCAAGATCGCGATTTCAGCGAGATTGCGCTACGCAAGCAGTGGTCTCATGGTGCAGCGGTACCGCA  
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## 27. ND-B16.6

ACTACTTGGTATTTACCTGCCCAAGTCAGGCCACACTGCCCTGGTTATGCAAAAACGTATTTTGACAATT  
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**GCTGCA** ATATAGCCCACGGCAGTGCCTCAAAATGCCAAATTAGATGGATTGGCAGGACCAACAGGGGAATCCCAG  
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## 28. Socs36E

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 ACTCTGCCACCTCGAGCTTCA **ATGGGT** CATCACCTAGCAAGTCTCAGCACGCCAATAGCCAGGAAAGAGAAA  
**TCCTACCGCCAGCTCGAGCCCCAGGGAAAGCAGCAGCAGCAGT** TAGAGAGCAGTCAGCCTGGACGAGGAGGAGCTGCA  
**AAGAGATCCTCACAGAGGCCACTGTC** TCGCCACATGCCGGCGGAAGTGCAGTCAGATCATCATAAAATCCAGCTG  
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 CAGTGGGCTTCACCTACTACGAGGACGTGGAGGAGCAGCGAGGTGGACGAGGACGACGAGGGGGTCAACATAGC  
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## 29. Atet

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 TCTTGTGAACCTTGACACCATTCCGGCTACCTGCAGTGGGTGACCTACGTCACTATGCGCTATGGCTTCGAAGGT  
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 CAGACAGACAGACATACACACACGCATCGCATCGCATAGGTATCTGGAAGACGATTCTAAATATACCGATGGA  
 TATATATACACCGTACAGTAGCTTCTGCATACCACAAAAAAATGGAATTACATAGAAACTATATAGAGTTAAT  
 ACTAGGAGTACCCACGTAGAGCCCCACTTCAACACACACTCAATTGAGGTACATATAATTCTGTGACCATGCA  
 GAGTCTGAACCTAAGTAGGTTGAAAACATAACAAATTGAGACAAGATCCTTAAAGGAAACCGAATCGGTGAGAGT  
 TTAGGACACGACTTCTC **CCAACGTTCTCA** **TCCCGTCC** TAAGTTGACTCTC AAAACACACTATAATCTTCTTGG  
 CAATCGATTCGAGCCCAGAAAGGCAGCAGCGTGCATCAAACCGAAGTTAGGGCAGCGTGTAAAGCGGAAACGGAAAC  
 AGAAATCATTCCCTGTGATAGCAAGCGGTGCAGTCCGGCTGGAGAATCCACATACCGGACGGGAAAGGTAGAAC  
 AAATCGGAGAAGGGAACCTGCCAACGGTATATAAGCAATTACTATTTCAGATGGGATGCAATCGGATTGTT  
 AGCTGCGCGTATTGTTGATTATTACATTAACTTTAGTACATGTTAGTTAATTTACGTCTACATTGTCATGCCA  
 AAATAAATGCAATTTCATTAAACAGCACGCTCTCCTTCGGCAGCCACACCAAAGTAAATTATTAAATGATAGC  
 CTTGGATGTAACACATGGATTGATACACAAACCAAGTTAACTACAGAAATCGGTGGCAACATATTCAAGCTGCTT  
 CAAAGGACAGTACAGCATTGTAATTATTAAAGCAATCTGTTGAAAACCCCCGGGAAATATCGTTAAGTATG  
 GTGTGTTGATATTCGGTAGTTGTTGAGCCATGTTACACTGTACATTAACGTTAAAGAATGGCATTACTGTAC  
 ACCATATTAAATTACTGTCAAAATCTGGTAGAGTATTAAATAAGAACCCAAACATCATCGAT

### 30. Itgbn

ATATTAGTTACGCTGGACTTCTAAGGCCAACGGTCGCTAGCTCCCTAGCAAAAAAAATATATGTGATCGTTGGTAGTG  
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 GATCTCTCATTGGATGCAATTGATGATCAGTGCAGGCACCGCAGACAGCTGCGAAAGATGTTATCCGCTCATC  
 TGGAATGCGCTTGGTGCACGGACAAGGAATACCAAGGTGGATATCGTTGTCTCTCGCCGGCAACTACTTAAC  
 TGCAGTAAACGGATATATACGAGAACCAAGCCGGTCTAGATGTCCTCAGGATAAGCCCTAAAGGACTACGAAAC  
 TGATCAGGCTGCCAAGTCACACCCAGCAGGGCATACCTTAAGTGGTCAAAGGAAACACGCAAAGAATGAGCTGAGC  
 ATAGAACTGCACGCAACAATCCACTCGATTGAGTATTGATGGATCTAACCTGGACAATGAGGGATGACAAGAAC  
 CTGGAGGAGTTGGAGGCCAACTCAGTCAGACTTAAACAGGAAACTACAGATTGGGATTGGTTCTTCGC  
 GGACAAAGCCAACCTACCCATGATTGTCACAGCATAGGGAAACCCCTGTGCTGCGAGCGGGCACGTTGAACCAA  
 CCTATGGATATAGACATCAACTTCGCTAACCGATGATATACCAGCGTTACCTCAGCGTGGCAACAGCAAAC  
 GGAAACTTGGACAATCTGAAGGTGGCTTGGATGCTTAATGCAGGTTATAGTGTGACCAAGGAAATCGGCTGGAAGG  
 ACAAGCTGAAAAGTGGTATCTTAGTGCACCGATGGCTTATGCACCTAGCGGGGACGGTCTTAGCAGGTATTATCC  
 AGCGCAATGATAAGCAGTGTCACTGAATAAACGAGGTGAGTACACGGGTTATTGAACACTACGACTACCCCTG  
 GAAATCTATCGGAGTTGCTCGCAAGATAATGTAATCTTGTGTCACCGAGGAGGTTGAAGTAGCTATTGG  
 ACTCACTGCCATAATGAAGGAAATTAGTACGTGGACATCTAAGTGTGATTCTCTCTAATATATTGAAACTAA  
 AGAGCTATGAGAGTTGATTAACGCACTCAGTTGCCGATAACAGCCGGATTATTGACATGGCTTACTATACGG  
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 GGACGTCACTCTCAAGAAATATCCGATAATCAGTTATACACACAAAATCGGGTGGAGGAAACCTCCTTGAGCG  
 AGTCACTGGATTGGATGTTGAGCTCAGCGGCCGTGCTCTGTGAGGAAACACAGATCCGAAACAGGAGGACGTT  
 CTGTGCGATTATAAGGGCTATTGTACTGCGGAATGTGCGATGAGGGTGGACTGGGACCTATTGCAACTGT  
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 ACCTCCG  
 AATTGGTTGCTCCAATCACGGGACTGTGACTGCGAACCTGCTTGTGATCTGGATATACGGGCTTTTGCGAG  
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 AACAAACTTGT  
 TTGTTCTACGAGCCCTGTGTCACATGTCTGAGCAGCAAAACAGGAAATGGCGTCTGCGAAA  
 ACTGACTGAAATATG  
 CAGCAGTTGGATAGACAAGAACCTATCCATACAACCTGTCCAGGAGCTGGATCTGAGCAGGATCA  
 ATGCTGGTAC  
 GACTGGTGAACAAACATGGCATCCAATGCGACAGCTCTCGTCTAGGTCACTGACCAACTCAA  
 ACTCTTGT  
 CACCGAGTTGATTGCGAGCCCCGGATTACGTGGCTCTGGTGGATATATCGGCAATTACGCTGCT  
 CATTGGACTTT  
 GATCATTTCATAATTCTGTGGTACATTGAGCTAACGGATGCCAGGGAAATATGCCAAGTTGAGGAAGAT  
 CAGAAGAATA  
 GTGTCGACAGGAGAATCCAATTACAGGGATCCAGTGGTAGATGAGTGGCCAGGGCTAAGTGT  
 TAAGTATG  
 GAGAATCCTTCGCAAGTTAACATGAAAATGTTTTTTAAGATATTAGATTGACTTTGACT  
 TAAATTTCTTCA  
 AAATTGTTGATACCGTTACTCGTAGAGTGAAATTACACTAGATTGCGTGAAAAGATGTAACCG  
 GTAGAAGTAA  
 CGTGGCTTCTGATATTAGTTAGTTATTCTGATCACAATACATAGCGAATTGATCTGGCTTGTCTG  
**CCG**  
**TCCGTCCGTTGTTTTTTTTTTCTAATTGTTCCATTCTGCGTATTCCAAAAAAATCTCTCCTCGCATT**  
 TACTTGATGAGTAATGGATAGCTGATATTGCGTTACTCGACTATAGCATTCTCTTGT  
 ATTAAAGCTATTGTA  
 ACTCA  
 AGAATTACGAGTTCACGAAGACATTAATACAAATATAAATTAGGTGTTATTAAAAGTATT  
 AGCCCCGTT  
 CAGAAAT  
 AACATTCCAATTCCAAGACTGGTCACAACAATAGCACTATTAGATCAATAATT  
 TATAATTAGAACATAAAAT

TTGTTTTTATACATAGAAATTATGAATTAGTCAAAAACTCGTGTATATCTACGCTAAATGAGCTAAAATTATTAC  
AGCTAAAAAACGAGTAAACGTATAACAAAGTAGCTTCAGTTCTGATTGGACAGCTCTTCAGCGATTGTATTCA  
ATAAGTACAAAATAATGGCTTAAGACATCGTTACATTAATCCGCTTGGGGAGTATAGATGACTACTGGCAAATCGGAA  
ATGCAAAACATTACCTTAACCTAAGAGGCATTACAAAATACAATTGCGATTAATTAAGGC

### 31. bves

ATCGGTAATAGGGTGACAACCTGACCGCTGGCGGGTTTATTGTTCCCTTAACCACCTCTTGGCTGCTATTG  
CGGTTATTCGCAGTTAATTCTCGATTATGATTAGTACTATTAGTGAATCAGCTAAAGCACGGCTGGTATTCA  
GGCAATTCTAAACCGCTTCTGGCTTATATCCCAGCTGGTACTCTCGTGGGAGCGAGCACGAGTAGTGAGC  
AGAGAGCGAAAAACACATTCAACCACACAGTGGGGTACGATTATTATTTAGAAAGTTACCAAAAGTTGTCTAA  
ATACGTTAGACCAAAGATGTCAACTACGCCAAGCAACCCAGCTGCCAAGCGATGCAAAATGTGACAGGCGGCAGCAG  
CAGTAGCCTCATCAAAGGTTAGCTACCAATAGCCCCAGCTAAATATTAAACAATACGAAATAAGAGTCTAAGGCCAT  
TAACAAACGGCTGACTAAGCTGAAATGCAAAGTAATTCAAATTCAACTAGCTAGGATATGAACCGCAAGCAAAAAA  
TACATTCAAGACACCAACCGCAAAGCACAAACATTGAAAAGGGCAGCAGCAATAACAAAACACCACCAACAAA  
GGTTGAGGAAAACCGCGTAAGCAACAACGAATTAGCGAAGGAGAGAAAGGTGGAAGATAGAAACAAAAGGCAG  
ATTTTGGCATTGCTAGCGTGGACAACAACCAGTACATCAACCAATCAATTGTAATTAGGCAAAATACGTG  
TGCAAAAATAGCTAACGATCAGGTGAGAGGAGAGTGACAGACTGAAACGAAGGCAAGTTCGCTGGATCAACGGA**ATG**  
**CCCAGCACGGCGGGCAGTCAGCTGGCTGGTATGGCGCCCTGATCAACAGTGCAGCAGCAGTAGTGTCT**  
GGGCATTGGCATGGGAGCAGCTGCTACGGGAGCAGGAGCACCAGGCAGTTCGGGATCTGGAGCAGATGCCTCCGCG  
CCGGCACTTAATGCCAGAGCACGGCGGGAACAGGCGCCAGCAGTGGAACGATCACCTGGACAACAATGGAACC  
CTGCGATCGATCAATCCCGCGATTGGTCATCGAACAGTGCCTCGGCCACATCACCTGACTTCAAGCTCGGCTGGC  
CTTTCTCTCCAGCCTTTGGCTTACACGGCCCTACGGGCCCTGTTGATGCGTCATGCTGCTCATTGGCTGCC  
TGATGATGGCATGCATGGCTACTGGTGGCTTGCACCGAGCAGTGGGATCTGGGACTGGGACTTTGTCAAC  
TTCATCTATCTCGTGTGGTGTATGCCGGCTGAGGCCTGTGCGATTGAGCAGGAAATCGAACGGTCTACCTGGCACT  
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AGGAGGTCTACGCCAGGAGAACGGTACCAAGGTGACAGCCTGCTGGTGTGAGCAGGAAACTGGTGGTGCAG  
CATCAGCGCCCTGCACATTGTTCCCCATCAGTCTAGACTGCCAGAACGGTCTCGACCGATGACTA  
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GTAGCGTTCTAACATGAACCTAAACAAATGATCTTAAGAACAAACATACCAATTGATGACCTTTAAC  
TGATTATTGATGCACAGAACATATGATGCTGGTATTCTTAAAGTCATAATTACTGACTAAACCGCTTAAT  
TTCCTGTTCTTAAATATTAAACAGAAATGGCTATTAAGGATGCTGTACGATTCTGTATCTGTA  
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TTGATTCTCCACTTCGACCTTGTGTCTCCCTCCCTTGTGCTATCACACTCTGCTCCACTATCTATCTGTTG  
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ACAAACAACAACAACAACACTACTGCTACTGCAACAAACAACTAGAACAAACCGTACAAGAAGCAACAACC  
AGGCCAACAAACATCGAGCAAAGTGCATGAAATTATGCAACCGCCGCTGCTCAACTGCCAGAACAGAAATC  
AGAGCA**CCCA****TCCGTCC****CTTCACCTTGAC**CTGCACAGAAACTACCAAGGAAACACCCACACATGCA  
ATCAGCCATCAGTCATCAGTCAGTCATCAATTGCGCAGCCACACCAACAGAACATCGATCACGATCATCAGAA  
AAGAGCGAAAAGAGCTGCGAAGATATGGAAGCTATCCATCTAAATCCGATCCTAAAGTCCATATAAATCTCA  
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CGGGAAAGTGTAAACACCGTGTACCTTAAAGGAAACTGGAGAAGTTACTAAAGAAGTTACAGGCGTGGAAA  
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ACACTGACTAAATCTAGGAAACTGGGAGACTGCTAAATGATTGATGTTGACTTCACTGACAACACTCTATTAGCAGTGT  
AGAGAAAACGTAACATCGTAGACCGTTATTAAAAAAACAGAAAAAAACTATTCAATTAAACCGTAGTAATA  
TACTGTATTCCATGAAGATAAGAAACCAATTATTCACAAAGAAAACACAAAACAGTAGGAAACTTACATTG  
CAACACTGTTGAATTCAATATGCTCCCTGAACTCCCTGCAAAACTATCGTTACTGCACTGAAACCCAGTTGAAA  
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TCAGTTGAGGGGTTTCAATTGAAACACGATCAGAACGCCAACAGTTAAATTGAAATTATGATGTTGAGT  
TGAGTTGAATTGAAATTAGTAATATTGATATATCACGCACTACAAACATTATGTTAGGAAAGTTGAGT  
ATCTCTACAATATAACGATTATTACACTCGTGTAAATAAGGAAACCTTGTAGACGCAACTATTCCAAGAACCTA  
TTATAAGTATAATGACTGACAGGTGATAGGTAAATTGAAACTGGTTGATTACAGCTCTCAGCTCAATTACGATTCTAGATCTCAAGATC  
GATCAGTCTAGTGACAGGTGATAGGTAAATTGAAACTGGTTGATTACAGCTCTCAGCTCAATTACGATTCTAGATCTCAAGATC  
TTCTTATTCTACCGCTTGGCATTACTGGAAATTACTGTGCTGGCACATCAATCAAGTCTCTAAATCTCCATTAAAC  
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AAGCGAAAGCT  
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### 32. Cad96Cb

CGAGGAGCTCGTCAGCGTGC GGAGTACTA ACTGAAAGCGTT CGGCCAACATTGTTCTTGTCCGGT GCAATCGCGGT  
CAAACAAATGAAATAACTAACATTAGTGCACCTGGAAAGATAAGCAGAAAGGATTAAGGCTAAACAAAGGAAAAGTTG  
CACATAAATTCAATTCTGTTATGGCACATCACAGATAGACGAAAGGTTATTGCTTAAATAAAAAACACGTACCGGTTAAA  
CAAGGGCATTACAATGGAGTATGGAAAGTGTCAAAGAACATCAAGTGAAGTGACTGAGCTAAATCTTATTCATGATT  
TGTGTGGAAGTGTATGGCTCTGAAC TGAGGAAGCTGAGGAACATCAACTAACCTAATTGCTAACATAATATCGTTG  
TACAAAATGCGATAATCACCAATGGATATACTATGCGATCTGCTCTCTGCTGGTTAATAAGCTACTCTGGCGC  
CATTCTGGACTCACGTTTACCTGGAGGGCGAGGTTCCCGCGGAAAGTCTTCTGCCACCAGGATTGGCGTGGGCT  
CCATAATCGGCAAGCTGCCGATCAATGGGATGCCAACAGGCGACATCAATCTTCTTGTGAGGGAAAAGAAT  
GCACCAAGTTGAGATTCATCCCGCACCAAGGAGCTAGCCCTATCGGTGGAACGGACAAGGAGGGTGTGAGAGGACCATC  
GTCCATTACGTGAATGTGATCTGCATACGACGGCATTCAACGGATCCGAGCTCGTCATTCCCCTAACGTGCGAGTAA  
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ACCGTACTCGGAGTACGTGCA GTTCTCAATCCTCTGGAAAGGACGGCTGGTCTAAAGAAAGCCCTGACTACGAACAGC  
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ATCACGGATGCCGATGATCAGAACCCAAAGTCCAGCGAGAGTCTACAGCCTGAGATACCAGCGGATGGTAGGCCAGG  
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TCCAGTCGCAAGGATGCCAAGTACTCCGCATCCGACAATGGAGCTATTAGCCTGCTGACACCACCGATGGATATGCG  
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GTTGACGCGACCTGGAAAGTCACAGTGA TTGAGCACTCTGGCTTCTCGTCCAGAAGCGATTGATGCGCATTGCGGAGG  
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GTAAATT CGCAGTTCTTAGCTAGGATCGCTGGAGAACCTGTCTAGCCAAGCCACTCGACTACGAGAAGATGCCAA  
GCACCGAGTTCCAGGTCTAGCCACCGATGGCATGCCAACAGCACCGCGGAACCTACACTGGAGGTGATTGACGTTAAATG  
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GAGGGCGTGTCTATCGGCAAGGTTGAGGCTGCTGATGGGATCGTAATGACAACACTGGAACCTCTCGCTGCGTGGTCAGCA  
CGCTGGACTGTTGAAATTGATTCCACTGGAAATATCTACATGCGTCCGGAGCAGCTACAGAGTCTCAACGAGTCCACGG  
TCCATCTGATAGCCATGCCACGGACACGGCGTTCCACCCAGGAGCAGTCTCCGGTCACTGTCACCAGGAGG  
CTGACGCTGGCCCGTCCGGATGGAACAGCAACATGTTGGCATGTCGGCATGATAATGGCCTCTTGTGATGATCAT  
AGTGGCGTTAAGCTGCTACATAGTCGATGAGCAAGGAGCAGTGCATCCAGGGATTGGACTCGGTGGAATC  
GGGTGCACAGCCAGGCGCACAGCAGCGTGTCTGCCAATCTGTTACCCACGAAAAGCTAGCGGGGAATGCCAACCGC  
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GAACAATGGCTTGATCACGTGGCCAGTGGAGCCAGTAGTAGCATGCCCTCGTAGTTCCGCCCTGCTGGAGCGGG  
AACGAGAGAGGGAGCAGGGACCGGGAGCAGCGCCAGGGAGAGCTATGCGGCTACAGTGCAGCAGCATAGTTCGCGCCCTCC  
GCCAATGGCCAGCTCTCGAGGAGGATGAGATCGAGCACGACTCACTGCAGACGGAGAACACAATCGGAAAGTGGCAGC  
CACGGGAGTGGCAAGCGTGGGGTTACACCATCAGTACCAACACCGCCAACCGGGCTCCAATCTCTATCCGCCACCG  
ATGCCATGGGTTCTGAGAACAACTTGACGGTCACTCTAGAGCGCAGGAGGGAGGGAGCTACTCCGGAGAGCTT  
GTAAATATCGCTGCTAAGTCAGTTAGGTTAGGTTAGGCCACACGTGTATGTACATTGAGTAGACCAAAATATAG  
CATTGTAACAGACCCAGCACCAGATCTAGGCTAAGTCCAAGCGATTCCATTGTAAGTTGCTTC **CCA GATTGTAAG**  
**AGGA TCCGTCCA**ATAAAAGTTCAACTATTC

### 33. Cah2

AATTAGTGA TTGTACAAAAAAAAGGAATACAAAAGAGTCTATTATAACACGGACATACATAAGTCTTCTTGGTC  
TTGGCCCAAGATTTGGCTACGAGGGCAGACATGGACCCGAGCACTGGAGCGAGGATTACGCCGCTGCA GTGGAAAGCA  
CCAGAGTCCCATAACATCGACCAGGT CAGCGCCGTGGAGAAGAAGTCTTCAAGCTGGAGTTTCAATTCAAGTGG  
TGCCCGACAATCTCAA**ATGACGAACAACGGTACACGGTCTGGTGAAGATGAGCTACAACGAGGACGAGATCCAAGT**  
**GTGCGTGGTGGCTTGGCGAGAAA**ACTCCTCTGGCTATCAGTCTGAAACAGTCCACTTCACTGGGCGAGAAGCA  
TACGATTGGCAGTGAGGATTGATCAACAATCGTGCCTATCCCGCCAGTTGCACTGTTGATGGTATTGCGGAATCTAGAGTATC  
CGGACTTGCAGCGCCCTGGACAAGGATCACGGCATGCCGTGATGCCCTTCTTCAAGGTGGGGAGAACAACTCAACT  
GGTGGCTATGAGGGCTTCAACCAACTGCTCTCCAGATTGACCGAACAGGCAAGAGCGTTAATATGACCAATCCACTTCC  
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AGGTCACCTGGATTGATTCACAACGCCATTGACATCACGGAGAACGAGCTGATGCCTCCGTCTGCA GAGCCAAC  
GATGATCATCTGAGAACAAATTCCGACCCATCCAGCGCTGAACGATCGCACCTGTACAAGAACACTACATAGAAATTCC  
GATTACAACATGGGCTCGATTCCACTCGTCACTGCGAAATGCGCTGGCAAGTGGAGGGACAAGCCGAGCAGTGC  
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TCGAAGGTTAAATCGTAAATGCTAGCTGGCCTTAGGGCTCAAATTAGCGCTGTAATTACATTTCGCTTGT  
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TATCCCC**ATATTCCCTTCCGTGAGGAGG**CCCGCAGTGCAGGGAGACCCCTGGATGCTCTCGTATTGACATTGT  
GATTAAATGCTGGCACAAGCAGAGCTTTATTGTCAGTTAGATTAAAGTTGAAACATAAAACCGTAATAA  
ATTATTATTGTTAACATCT

### 34. CG12880

CATTCACTTGGCGAGATTTATCGTGTGGGACGTATCGCGGTGAATTGCGGAACCTGAAATCCAAATCCGAGTGGCAAGATACCGAATTTCGAACACTGAAAACCGTGCTCGTAGTTGACCGAATTTCAGTGCACCTTCAGCATTGGCGCAAGCAGCAAAGACTTAGAAACTCTAACCGATTGCGTTACTCGAAATCGTTGCTGCCACATGCTAAAGTGTCAAGTCGGCAAGGATGTGCGCTGGCAAGGCTGAGGCTGGACAGGCAGTCAGGAGCGGATCGAGTCAGGCTCTGACGGCTATCAAGCAGGCCAAGCAACTGGCCAGAAGCGCCCATGGACTAACAGAAGCTGCTCAGCCAGATGGGCTCATGAAGGTGAAGGCGCCAAACAGCACAACAGGAAGCAGTGGCGTTGAGTCACGACGCCACTCCTCGCGCATGATTACACATGTTCATCATCAAATTGCCAAATATGCACTACTATGCTGGACCCGGTGCAGAAGAATTCCATCGCAGAGCAGCAGGACAAGTCTGGCCACAAAGGTAACGCACCCAGCACGATGGCATCCAGCACACCAGTCGAGCGCATCAGCAGCAGTTGCAACATCACCAGTTGCCAAGCGTCGGCGTTGAAAGCTAACGGAAAAGGCTCGTTCCCTCAGCTCGAATGGCCGGCGGGCGCATCTACCAACTGGAACCTGCCCCTGAAGGAGGCGTGGCCAAGAAGGCCCACTTCGCCCCACGCTCGGCGGGATCGCAATAAGACTGCTGGACATCCAGGGTGTGCCACGTCGGCCAGCCCTGGGAGAACGAGACCGCCAGAAGGGCTTCAGTGCCGGCGGGAAAGGCCAAGTATCGCAAATCCCTGAAGGCCAAGGCTACGTCACCCACATATTATGCAACCCGCCAGGGCGTCAAGCAGCAGGCTTCCACAAGTACTTCGCGGGCAATGGCAAGGCCAAGGGATTCTATGTCATCAAGGAGCACCAGAAGAAGCCGAGTCTACCAGAACATCATCTCGTAATCCAACGGCATACCCCGGTATACCCAGGGAATACCCCTGAAAGACTGCGGTGCCGGCGTTC TATTTCAATATTCTAACGAAGGGAGCTCTC ***CCCTCT TCCGTCCAGTTCCACT***TGGAATACCAATCTGGCAATCCGATCACATCTGGCCAAATGTTAACAGCATCTGGCCAATGTTAACAAATTGCGATGGGTGCGTTCTATTTCATATAACGGCAATGAATACAATAACAGTGGCTGGTCCAGGGGGTGATCTCATATCCATCCATGTGTCAGCCATATCAGCACAAAGAACAAA GTGTCAAAGTGAATGACTTCATTGAGATTCAATGTAAGAACAGAGTTAAAGTTCAAGAAAGGGGTTTGATTGA GCACAAACTGAGTTGAAACTATCGATTATATTAAATTGAAATTAAATCTAAAGTTAATCAGCTTAGCAGCAGA GTGTGAAATATTGAGTGGCACTTCTCATACACATATCGTTGATCGGAAATCCAAATCCTTCATGAGGAC ACCCCATCCGAAGGACCAGTCCAAACTGCATTAGCTAAGCAACTGCGGCAAGCAATAATTATGGCATAATCAAGCTAGTTAAGTGCATGAACGACTTGTGATAAATAAAATTGACTGATTTAAGTGAATTCCCTAAAGA

### 35. CG14785

TAGCATACACAGTCTCTAACAGCTTCTCCGAAACCCACTCCAACTAAATTAGTTTATTGCGGTCAATCTGTGCAAGTTGCAACCCCTCCCTATCCAACC**ATGAAGCGGCTGTCGGACGTCCGGTCTCCGATGAGCAGAAGGCCAGATGG** GCAACCAGATTATGAGCAGGGCGCAATAGAGGTATTATGCGGTGGCCACCGTCTCGCGCTACTCCGAACGCCGGTC CTTCCGAGCTAAAGGATCAACAGGATCAGCTGAAGGACAAAAGCAGGACCGCAGGCCAGGCCAGAGAAACT CGAGAGCGGCGCTCAGGGCGACGCCGGTCCGCTCCTCGACGTCGCCAACGCTGTAAAGCTCTGCCAATCGAGGGAG ATCCAGCCAAGCTGAAGGGCGTGAACAGGCCAGCAATTGCGACGCTGCTCAATCACCTCGCGACGGCGCAGGGC GATGGCGAGGCCAAGCAGATCGTATGGAGCTGCTCGGCAGCGGGGTGCAGCCGACTCGCCATCGAGCGGTTCCGA ACAGCTGTTGCACTGCTGCGGGAGCAGACTCTGATGGAGGCCGATATTGCGCAATGAGCGGTATCGCTCTTTA AGGGCGTGTGAGACTACGCCAAGATATAACGCGCGCAGCTGTTGCCCGCGTGTGCGGAGGAGGTGCTGAGATCCCGCAGCTGCGCCA CGCGATGCGGGTGCAGCGAGCTGTTGCCCGCGTGTGCGGAGGAGGTGCTGAGATCCCGCAGCTGCGCCA GCTGAGGTGCTCCCTGCGAAGATGGAGTTCTCAACGAGGACACGCCCTCAAGGCCACCTGGAAGTCCGTTATCACC CGCAGGCCGTATCTGCGAACTATGCTCACCCGATCTCGCGCTTCTGCACTGCGCTCACCATTCTC CTGTACTTCACCGAGCAGGAGCTCACGACATTTGCGAAACTGTTACCTCAGCGTCAACTCGGAGATCGAGATATTCC TTCCCGTGTGCTCTGGCTGGAGCACAACACTGGCTTGAGCGCGGAGCTGCGCCAGCGCATCTAGCCGAGGTGCAATTGCG CCGTGTGCAACGCTTGATTGCCGAGGCCCTAGAAGACGCCATCACCTCAAGCTAAGGCCGCTTCAGTGGCGCCGCTT ACACCGACAACGAAAGCATCAGGAGGAGAGGCTTGGTCCCGAGCTGGATAGTGGACTTGAGTCAGGCCACCAACATA AGAGCCACTGCGAGCAGTCGTTACCCCACCTACGACCGTTCAAAGAAACCTGGCCAGGATTATCTGTCGCCGCC AACTGCTGGCGCACGTTCCGCTGCCAAGAGGTGTACCGCAGGGACTCCGCTGCTGCTGCCACACTTCAACTA GACTCAGCCCAGCTCCAGTATTCTGTACCTTCTGCTACTCTGGCAGTCGTGATT**CTGTCAGT TCCGTCCATTGTCC** TTTGCCGTGTCCATGTTTTAATGCCATCCACACCACATCCATAATCCTTACAACACCCCTGACCAATTACCGACCTGATTAGAAAATAAAAGCATTGTTGCCATCC

### 36. CG31495

GCAGTTATCGAACATCGATTAGTGGATAACATAATACAAAACATTGTAAGAACAGCTGATACAATTGACCA TTTCATATCCAATATAATTCTCCATAAAAGAACGCAACCCGGCGCAAG**ATGGTCAACATCACTTACCCAAACTGAGGACCGTGTGAACCTCTTAAATCGCGCATCACACAGGCTGGGTGACATTACGGTGGCCGGGATGTTGCGCACGCCAGA** TTGCAATGAAAACAAAAATTTCACGGAGGAGCAGCTGTCGCAACTGGTCCGAGAACGCCCTCCATGAAGCCATTGTCGCG ACTCACGTACCCAGATGCCGAAGGGCCTTCAAGTCACTCAGGTCGATCTCGGCCGCGTAAAGATCATTGCCACG ATTTGGCGGCCAGGAGGAGCTCCAGCTGCTCATGCCATATGAATATCTAGATCGTACCGCTTCTGATCGAATGATC TCCTGTCCACGGACGCCAAGATGGTCTGCGCATCTGGCGAAGGTAAGGCCAAGTCAGGCTTAACACTACGGTGGCGCT CAAATGGCTTAAAACAGACTGCCCTTTATAAAACATCTCGTCGGCGAATTGCTGGACTTAGCGATTCCGAAAA ATGCCAGAGAATACGGGAAGTCTTGGAGGACGCCATTGTTCCAGGGCAGCTGCGTCATATTGACGACTTGTAGCGTG TCATTGGTTACGGCGCCCTGGGTAAGCGGTACTCCAAGGAGTTCTGCAAGAAGCTGACGGTTCTCTGAAGAAACAACCG CCCAACAGCCACGAGCTCATCATATTGCAACAGCAATCGTCTCGATGTCCTGGAGGAACCTGGCTTGCTATCGTTTT CACTTCAGTTACAATGCGAAATGTTCCACGCCCTAAAGAACCTTATGGTCAAGTGGAGCAGCTTCAAACGATTGAGC CCGAGGAGCTACGCCAAATCGAGATGCCATGGATGGCCGAACTGCTCCATTGGCATCAAACGGTTGCTTGACTTGATT CGTGGGTGAATTCCCTAAATCCCGACCGTCAGCTGCCAAATTGCTGAAGAAATTGGCGAAGCAATGGGATGGTAGG

**GAACCATTCTAGTCCACTTGTACCTCAATACATTAA** GACGTTAGGTTAGGAAATGCAATATTGTATCTACAATTCAGGAAAATAATTCCAAGTGATCCAAATAATTCAAGTAATGCTTATATTAAAAGAACCGACGTAAGCATAAAAAAACCACATCTA  
AAAAAATAATATTGCATTATTTGGATGAACCTGCTTCATTGCAATGCAATGTTCTAAGACTAAATGAACAGCCC  
GATAATGTGATTGCAAAGATACTAGATTATTATATTATATATACACCAGGCCAGACCTCTCAGTGGCCT  
TCGCCTGCTGTA**CGGAGAGCAT** **TCCGTCCAGG** AGTTGGTGGCGTGGGCCAACATCTCCTGGCTGCCAAAG  
GCGGGCTTGATGTCGTTGCCAATATATATACATAATTATATATATAAGTAAAGATAAGAT  
TTATATTGTTAGTGTCTTGGTATTGGCATACAGTTTATTACCATATTATATATAAGTAAAGATAAGAT  
ATTGAATTGCAA

### 37. xit

ATCGGACAGCACTTACCGCTGATAAGGAATGCATCGAATCTAAACTACAATCCAATCCCTGAGAGAACGAAGATGAAG  
**GACCTCTCTGGCACCTGGTGGCATAGCCACCGCCTAAAGATCCTGCTGATTCCAGCCTACCATTCCACGGATTTCGA**  
**AGTCACATCGCACTGGCTGCCATCACCCACAGTTGCCGCTGAATCAGTGGTATGTGGACGCCACCAGCGAGTGGACAC**  
**TGGACTATCCGCCGTTCTGCCTACTTCGAGTGGCTCTCGCAGGTGGCAAGTACGTGGATCCACGAATGCTGGTG**  
**GTGGATAACCTGAATTACCGAGTCCAAGGCCACCGTTACTTCAGGGCTCTGGTATTGGATCTAGTGTATGT**  
**GCTCGGCGTGCAGCTGGCTACTGGGATTGGCAGGGATACGCAGCAGTCTTGGCCCTCATGTTGCTGC**  
**TCCTCAACGTGGGATTGATCTCGTGGACCACATTCACTTCAGTACAATGGCTTGCTGTTGGCATACTGCTCCTGAGC**  
**ATCGGTTCCCTCATCCGGCAGCGGTTCCCTGAGCGCCTTGCCTCGCCGTCGTCAACTTAAGCACATCTTCT**  
**GTACATGGGCCGCTTTGGGCTACCTGCTGCCATTGCTGGAGCAAGCGAGTGTGGTCTGCGGTGGTG**  
**CCGTTGTCAAGCTGCTGGCGTTGACTTACTCCGTTGCCGTCGTCTTGGACCCCTCTGGCAGCAATTGCCCAAGTC**  
**CTCTCCCGTCTGTTCTTCAAGCGTGGCTAACGCACGCCACTGGCACCCAACCTCTGGCGCTGTACAATGCCGC**  
**CGACAAGTTGGCGCCGGAGTTTGAAGGTACAGGATGGCGGTGCCCTGACGACGTCAGGATTGGTCAGGAGGTCAAGGC**  
**ACTCGGTGCTGCCAGCCATCACACGCCCTGTGACTTTGCCCTGACCGCTCTTCATGCTGCCATTGGTGAAGCTC**  
**TTTCCGAGTCCAAGAAACAAAGTCCGCTGGCTTCCTGCGAGCGGTGGTTGTGCGGCTGTTCCCTCGTCTCGG**  
**CTGGCATGTGACAGAGAAGGCCATCTTAATGGTGCCTCTCCGCTCTGCTCACGCTGGTAAACCGGGAGGATGCCA**  
**GATACGCCATGTCCTGGCATTGCCGCTATTCTCTCTTCCGCTGCTCTCGATGCCATCTCCCGCTTCCGAGGCTTCCACAC**  
**TATTCCCTGTACATGTCGATGGCCATGATGTACGCCAGCTGTATTCCGCTGTACGGAGCATCTGCTCTTCCGTCACCTGGACC**  
**GCTGGAGTGGCTATACATGCTGGGCTCATGGCTATTCCGCTGTACGGAGCATCTGCTCTTCCGTCACCTGGACC**  
**AGGCCCTGCCCTTCTGCCCTACTGCTGACCTCCGTGTACTCGGCGTGGAGTGTCTACTTCTTGGGCCACTAT**  
**CTGTACGCTCTGGCATTAGCTGGGCAAGGTGCCATCGCATGTCACATCCGCGGCCGTGTTAAGAGAAAACGAAA**  
**AACCAAATAA** ACGGAGCGCTATAAAGGGCTTCAGCAGAAATCCGCAATTGGTACGTCTTCGATTAGTGGTAGTGCT  
GGTGGTGCAGCCGAGCAGTCTCAAGGCCCTGGCGAACAGCCAGTTGGCGCCTGGACGGGATTCTCTGGTCCGAT  
CGGCACAATGGACAACAAATGGCTTTGGCCTAAACAGGTACGGAGGCCAATAGTC**GGCCGGCCGTCAGC**  
**TCCGAGTGGT** GACATTCTGGAAAAAACCGTGAAACACGCTGGAGCTGAATGAATAAGTGTATATCAATTGCCAGCCT  
GGCAGGAATACT **TCAGTTGACGAGCT** **TCCGTCCAG** AGCAACAGATCGACATGCAGATGCCATGGAACTCTCCTATGG  
CCGCGGCTACGATCCAATGGGAGGAACCACCGCTCTCCCTCCGTCGCAACAGCACTGGTCCAGCAATGGTTCT  
CCGACGACTCCGCTTCCGCTGGGGATCCTATATTGAACCTGGCAGCGGAGCCATGCGTCGCGTGGAGGACATA  
CGCACCGAGGACTTCATCCAGTCGGCCTGCGCAGTCAGTCAACTATTGAGCTGCGAGAGGCAGACTAGTGAGGATAGACTG  
GAGTGGATGCCAGGCCGCTGGTACACTCACCTCAGTCAGACACACACCAGCCAAGATGGACTTGCAGGTGCAGCCGG  
GTCATCCAATGTTGTTACGGACAGGGCTGGCCTCTGCGATCCACAGTTGCGCTGCGACTGAGCTCAAATGC  
CAGCAGTCAGGTGGGACATATGTTGCGCTGGTCCAACAGAGCAGGCCGCCACCATGTCGCTCAGCCGG  
GCCGCCACGCCGTGTCCGCTCTCTGCCCTGCAATGGAATGCCAGTGCGATGGTTCTCCGCTACCGGAA  
GCTATGGATTCTGCCGTTAACGATCCGTACGAGATGTACGCCAAATGGCAGTTGTGGCGTGTACACCAGCAC  
ATGATGGAAAGCTAAACTATTAAGCCACAAATTGGTATTAACTGGCGTTCAAGGAATATATTCAACGCTATTCTT  
GAGTACGCCACGCATTACAAATTAAATCTAGGTACAAAATCACAAATGTATTGCTTAATTCTGAAATAAGTTA  
AAAGTAGCTGAA

### 38. crim

CTCATTGCGGTACACTGCTCTAGGGCGTATCTTATTACGTTGTATTGTTATCTGTTTATTGAA  
AAAGAAACCCCGTTTATAACAAAC**ATGCACTACCA** **TACGAATCTCATTGCA** **AGCCCTGCTGCTGGCAGCT** **CTAATTCA**  
**ACGAGGGCTGCTGCAATTGGTGTACCGCTGCAC** **TCGCCACGCCGGCTGCGGAGAAGTTAA** **CTGGCGGGAAATA**  
**GGATTCCCTGGCGAGCACTGCCGGAACCGGACGACATTGCGTCAAGGTGACGGAGCGACGCCAGGGAAACAAT**  
**CACACGTGACTGCC** **TAAGCGCCTGAGCTCCGCAAAGACATTCCCGCAGAAATACGAAGGATGTC** **CTGCCGCCC**  
**ACGATGAAAAGCTAGCGA** **ACTACGTAACACGATCAAGGAGCACGACGTGCCGTGACTACTACACGGACACCACG**  
**TTCTGTTCTGCCCTCGACCATCGATGCA** **ATTGGGCCAGGCCCTACAAACGTCGCCGTAATCGT** **CTCTCACTCT**  
**AATTCCCTGCCCTGCTCTACGCTAGAATCTCTCTGAGTGCAGTATTAAAGTGCCTTA** **ACTACGCTCTGCAA** **CTACG**  
**AAT** **TCCGTCCAC** **ATTCCAC** **CTTGTACAACCTTTGACATGACTACAGCTTTAGAGTCTGT** **TTACTCAACTA**  
ATTAAATACATTACCATATTATAACAGGAGTGAAATCGTTAAAGGACCTACATTCTGGCGCCGGTA  
ATCATAGGCAGCCACCTCGGGTGCACCGATGCTCGAATAGACTTCC

### 39. Cdc5

TATCGCTGTTATCGAAGACTAACCATCGCTAGCGCTACTGCAAAAAAGAGCGGAGTTCGTGGACTCGAGTAAAAT  
TGTGAAAATCCATTGTATACGCGACAGAAAGCAACATAAGGCCAGA**ATGCCGCAATA** **ATGATCAAGGGCGCGTGTG**

GCGCAACACGGAGGATGAAATCCTCAAAGCTGCGTTATGAAATATGGCAAGAACAGTGGAGTCGAATTGCCCTCC  
 TGCACAGAAAAGTCCGCCAAGCAGTGAAGGCCAGGTGGTACGAGTGGCTGGATCCCAGCATCAAGAACCGAGTGGTCC  
 CGTGAGGAGGACGAGAAACTGCTCACTTGGCCAAGCTGATGCCAGCAGTGGGCACCATAGCCCCGATCATGGACG  
 GACGCCAGCCAGTGCCTGGAGCGTATGACTACTTGTAGACCAAGCCAAAGGAGATGGCAGGACACGATGG  
 ATGATCCTCGCAAGTGTGAAGCCGGAGAGATTGACCCCAATCCAGAGACCAAGGCCGGCGCCGATCCCAAGGACATG  
 GACGAGGACGAGCTGGAAATGCTGTCGAGGCACGCCGCTTGGCAACACGCAAGGCAAGAAAGCCAAGCGCAAGGC  
 TCGCAGGAAAGCAACTGGAGGAAGGCCGCCGCTGGCAACTCTCAAAGCGCGCAACTCGTGCCTGGGAAATTGGCA  
 GCGGAAATCGCAAGCGGATCAAGGAAATAGACTACAACGCCAGATTCAGGAAAGCGACCTGCCACGGATTCTAC  
 GACACTCCGAGGAACACTTGCAGAAAAATGAACCCGACTCAACAAGATGCCAGCAGGACCTAGATGGAGAACTCCG  
 CTCAGAGAAGGAGGAACCGAACGCAAGCGGACAAGCAGAAGCTGAAGCAGCGAAAGAGAATGAAGTGCCACCGAA  
 TGCTGCAAAACATGGAGCCCAGCGGAAGCGCTCCAAACTAGTCCTGCCTACGCCAGATTTCGACATGGAACACTACAG  
 CAGGTAGTCAAGCTGGGACGCCAGTGGAGATGGCAAGGAAATTGCCGGTGGAGAGCGGCATAGAGACGACGGATGCTT  
 GCTAGCCGACTACTCCATTACACCCAGGTAGCAGCCACTCCAGTACTCCGGCTCCGTACACCGATCGCATATGCAGG  
 AGGCCAAACATGATGGCCCTCACCCACACAGAGACGCCACTCAAAGGTGGCCTGAACACACCCTGCACGAGTCAGAC  
 TTCTCCGGCGTGTGCCAAGGCCGGCTCCATGCCACGCCATAACACAGTGGATTGCCACGCCGTTAGAACCCAACGCGA  
 GGGCGGAGCAGCCACTCCGGGATTCCAACGCCCTCCGGAGCCCTAGTCCGTAAGGGGGCAGGAGGGGCTA  
 CGGGTGTGTCATAACACCTGCCTACGTGAGGGACAAGCTAAGCATCAATCCTGAGGAGAGTATGGCGTACCGAGACC  
 CCAGCACATTACAAAAGCAACTAAAGTCTACACTGCGTGCAGGCTCTCCACGCTGCCGCTCACGGAA  
 TGACTATGAGATCGTGGTGCCTGAGCAGGAGGAAAGCGAACGCATTGAGACCAACTCAGAACCCGCCGTCAGGATCAGG  
 CGGATGTCACGCCAGGTTATTGGCTGAACAAAGAGGCGCCGAAAGCGAGAGCTAGAGAACGCTCCAGGTTATTCA  
 CGTAGTCTCCCGCACCAACTGAAGTGAACACCAAGATTCTGCACCGCAGTCCGAGAACAGAACCTTACGGAACAGCA  
 GCAGGCCGAAGAGCTGATTAACACGAAATGATCACCATGCACTGTACGACTCGGTCAAGGATCCCGTGCCTGGTCAAT  
 CGCAGCACAAGTGGAACAGCTACAAAGCTACTTAAAGGCAATCCCTACGAGGACATCTCGCAGCAAGAGTTGGCGAAG  
 GCCAAGCAAATGTCACCGAGGAAATGGAGGTGGTGAAGGAGCGCATGGCGATGGCGACTGCCCTAGACGTTATGC  
 CCAGGTGTGGCAGGAGTGCCTCGGCCAGGTTCTTACCTACCCCTCCAAACATCGCTACACACGCCAACATTGGCCAGCA  
 AGAAGGATGATTAGACTGCGAGAACGGCTGGAGACAAATGCCGACATGGCCAAGGAAGCAAGCAGCTGCGGC  
 AAGATCGAGAAGAACGCTGAAGATTCTCACCGGGCTACCGCGCCAGGTGCTGATTAAGCAGCTTCAGGACAC  
 GTACGGCCAGATTGAGCAGAACTCTGTCTCACTCTGACATTCCGTTTTGGCGAACAGGAGGCCATCGCTGCGC  
 GACGCCCTGGAATCGCTGCAAGAGGACGTGCGTCCAGATGGACCCGAGAAGGAGCTGCAGCAAAGTACGCTAGCTTA  
 GTGGAGGAGGCCGATTGCTTACTCTAACTGAACATATAACTGGCTGCAGGCCACGCCAGCAACTCTGCCGA  
 CCAGGAAGCGTAAACCTCTTGGAAACCCAAAGTCCGTTCTATTAAATTACAGTAATTACATGCCAGCAAACAGCTATC  
 ACCACCTAATATCCGATCCATTGCTAGCGCCGACTCTGTTAGACCCACTCTTGTGTCAGCGACAGA  
 CCAGCGCGTTCAACCAGCGCGATGCACTGGTAATGGAATGCGTGGTGCACCGCCACAGTCACCGT  
 TCTGGTTGCACTCGGCGCTGGCACTCGATGCACTGGTCAAGGTTATGATGTGGTTACGGCAGATGGCACAGTTGTCCAC  
 TGCTACGTCCCATCCCCACATGGCGCGAACCCATTCTCACCA **CAAAGCGCTCC** **TGGCTTG** ACAGCTCCTC  
 CACTACAGGATGGCTTCTCGTCAAGTGTGCACTGCGTGGAGATGGACCCGAGAAGGAGCTGCTGCTCCAGTGGTCAACCT  
 GTGTGAAATTACGAATTGTTTCTTCAAAATACAACACTTATTTGCTATTGCAATTGACTATTAAA  
 TTCCATTGTTTGGCTGACAGTTAATAAAAAAGTTGACCACTACCGATTACAGAAATAACACGGGAATA  
 ATTTAAAATAATGCAG

#### 40. nemy

AGTTGTCTAGCTGATGCCCGCATGATCGTGTACACTTCATTGATAGTAAAATCAGTGAATTAAATTAGGAT  
 TTTCGTGCGGTTGTTGTAACCGATATAGAACGACAATTACGCCGAAACGACCTTAGTGGCCATCCCAGATATATT  
 GTATATACACCATATACACCTATAGAGCCGATCCGTGACGCCAGCAGCATTCTACGTTAGTGGGTGGCGGTGA  
 CAGCATAAGCAAGAAAACCAATAAATTGCCCAAATAGCAACACTTGTCTAA **ATGCGAGATGTCAC** **TGGCCAA**  
**ATGCCCGTGGAGGAACGACAACATCGCGTATGAGGAGGATCCAGATAATGCTGCTGGAAATTGCTCCTGGTGC**  
**GAATA** **CTTGTGATCGTCATCTGTCACGATCCTGCTGGTGGCGTGTGGCTAACCCCTCTTCTGGTGTATGTT**  
**ACCGCG** **ATGGCTCGCCTGGTGGAGCTCCAAGCAGCAGTTCAATCTGCATCCCACATCCATGATTGCTGGCTCGTCACG**  
**CTCG** **TCCGGATTTCCATCTTGATCTATCGTCTGTGCCGCTGGTGAAGCAAATCTATGTAAGCTGATCCACATGTT**  
**CTTCA** **CGCCCGTGGCCATTCCGTGCATGCCCTGGGTTCATATCCGTCTCGCTCCACGCGCTTG**  
**CATAAGGTCAACTTCT** **ATAGCCTCCACTCGTGGCTGGCTCGTACAATGGGATGTTGCTGCTCCAGTTGTGATCGG**  
**CTTCTTCAGCTTCTG** **GTAATGCTGTGCTGTGAGAACACGACTACAGCTGCCATGGGCCATGGT**  
**GCCCATCCACGCGCAGCTGGACTGATCGAGAACAGGAGCGCAAACCGTCAACGAGGCGGGCTGAGCA**  
**GTGAGAACAGCTGGTGGAGCACTACATCACCAAGTGCACATGCCGTCGGCGTACCCGATCTTCTATTGG**  
**CATCATCGTCACCTC** **GCGGTGCGCGTTCCAATGCCGCCCTGGCGAAGGTCTACGTACGGAGCC** **CATTAGACGG** **TCCGTCCGG** **CTCGTCA**  
 GAGTCCTGCCGCCCGAATTGGTCCACGCCAGGGTAAGCGAGTCGCAAGCGAGTGTACCCGAGGATTCCGAAACTCG  
 GTGACTTGGACTGGACTTGGGAAACGGACCGGGCTGTGCTCTTCCGCCAGGCTAGCGCACAGAGACACGCTCTGCAA  
 CTCTATCCGAACCCATTGTAACCTTTATGCCAAATTAAACACATCATAACTAGACTACTCCATATGGCTATAA  
 TCAAAACCTTGCCTCAATCTCAGTAGGTGTAGGATTAGGAATCTCAGCAGATGAGCAGAACGACCAA  
 GCAAATCTAATCAAATTGACGTGTTGCAAGCCGAACTTACTGATGCATCCAGGCACTTGCAGCAGCTAGATTAGCCAG  
 TTTGGTACTATAGTAGAATTGATATACTCGTAAGCCGATGTAACAGTGTATGTGATCGTACCGACGATTG  
 AATGTACGCACTGGACTAAAAGTGTGCAATTGATGTTTTGTAATCGTCAAGTATTGATTTCTATAATGC  
 TAAGCAACCTAAGCTGTAAACCAATTACAAACAAATAACTTTGCCCTGCTGAGGGCTCTCATATAATCGCAA  
 TTGTTATGCACTTTAAATATTAAAAACGGTGAGAACACTCGGAAAATAACAAATCATATTCCCTGAAAGGGAAAT

TAAACAG

#### 41. Dok

TGCAACCCTGCAGCGCATAAATAAAAGTCGACTTGCCTCTGTGCTCGTTGGTATTTTTGTGCTGCGGC  
TGCCTTTCGAAATCGAATCTCCGTATCGCTGTGCGATATGATGTGATTTACTATGAAGTGATCAA  
ACCGCGAAGCTATCGTAAGCACAGAAAAACGGATGAATTGAAGGAAACTCCAGAAATCGCATAG  
GCCAACTCCAGAACAAACAATTCGGAATCGGAGGCGCATTGTGAGCTAGATCGGCAAGGAGAGGG  
AAAGAGCAGGAGACGAAGGAGAGGGAGAGTGGAGGAGTGGTGCACAGA**ATGGATGTTGAAATAC**  
**TGTTGGCTGG**  
**ATATCTTAATGTGCCGACCCAAACTGGATTTCGCTAAATCGATCTCGAAAAGAAGCAGTCAGTC**  
**TACTGTTGC**  
**TATTCAAGGCCAGTCGGCATGGATGCCCTGGAGATGTGCGAAAGCAAGGAGGACCGGAA**  
**CCCCAAAATCTTCACC**  
**CTCGAAAACTGTGTGAAGATCACTCAGGAGCCCTCCCGAACGCCCTCATT**  
**CACATTGTCAAGCGGCAGGCTACCGCCCTGC**  
**AAACCGGTGGCTCTGTGACACCTCAGCAGCTCC**  
**TTGATGGCTGTTCAATTACGTTGATGGCTATGCTCCAAATGAC**  
**GCCCCTGACTGAACCTGAA**  
**ATCGAGGATTGGCGCCACAATGCCATGTGGC**  
**CATATCGTTCAATTAGGAAGTACGGCTATCGCAGTGGCAAGTTC**  
**CCTTGAGGCGGGCAGAAAATGCACCAACGGCGAAGGCGTT**  
**CACCCCTGGATCACACCAATCCACAGGAGGTATTCCGC**  
**TGCATGTCCGCCAAGATGAAGTCGATGAAGAAACTGATTAGCGCGATAGTCTGAGCACCTGGAG**  
**TGCGGTGAGAATCA**  
**GTGGATGGCCAGCTGGATGGAGCCCGGTCGAGGCCACCATCACCGTCAGTAAT**  
**CCCCACGCGTGGCG**  
**AGTTCGAGATCAACTCAACGAAAGCTGCATATCCCTGAGGGGCTTCATTAGCTCTAACGATT**  
**CTTAATAACTCTCG**  
**GCAGGGTGGAGGTGGAGCAAGTGGGCCACAGCAACGGGTGGAGGCGTT**  
**TAGGAAGTACCGGAGGAATCACGCTATCCGT**  
**CCCCCAAATAGCAATAGCAACAGCAGCAGCAAACACATT**  
**CCCAACAAACGCCACGCCAAAGCCGACAACCCATT**  
**GTGATAAAATACAGGAATATTGTGAAATATGAACCTGTGGCCATAACTACGATATCTCGCC**  
**CATCCGCCAATTCCGATGTG**  
**AACAGTTCACTGATACTGAAACCCCTGGAAGGCAATCGATATCCCTGCCGAAACAGCAGCC**  
**GATTACCGCCACTGCGAAATGAGAGCGCCTCGAAAGCACC**  
**GCCCTCCGGAGCGGATTACGAGTCATCGAAATATCACAGAGGCGT**  
**GGCGCACAAGGGCGTGGCGATGTTAGGCACAGTGAACGTATGCC**  
**CATACTCTCGCATACTCGCATACTCGCATACTCGCATACTCGCAGAATTGTACGCCAG**  
**CGATCAGTATCGAAATCCACCTGTGGCAGC**  
**AAAATACACCCAAAGATCATAGACATAGACATCGATATCGGTGAGGGTTGC**  
**ACTAG**  
**CGCCACATCCATTAGTGAATCGAATTACGATCGTTAGACTTCTATCGC**  
**GAACAATAAGACATCGAGTGGCTATAAGA**  
**CCATTGTGAATGTCACACCCGTAATCGCATA**  
**CGATGCAGTCCACCGCCACCCAATGAATACGAATTGATCGCTAGTCCC**  
**GATA**  
**CGGAAAGCTCCGTAAAGCGATGACAGTCATCGCGGTTACGGG**  
**GTCGAAAGGCTAGCACCACAGAACAAAGCGGCA**  
**ATCTCGAGGCGGCAGCCATCGATCATCGCAGCTATAATGGCCTCA**  
**ACTACACAATGGTCAGCAAACCGAAGCGAGTGTAA**  
**AATATTAAAGGATTTCAGATCGATAGCAAGGATCCAGAGATCCTGC**  
**ACTCGA**  
**ACTGGGTGAGATAGTCTAGTTTATTG**  
**TCAGTCCAGAACTGTTCAATTGTAATTG**  
**CTGTTCAATTGAAACTGTTAC**  
**TTCTGCGATACTTTAAGTGTGCGATAACAAATGTAATGTT**  
**CAAAAAAATTAAAAACAAAATTGAAAACAG**  
**TAACGTGCCAGTAAC**  
**TTTACATAATTTCATACATTGAGTATAAGTGA**  
**AAAAGAAGAGACTGCCAGTGCCATTACAG**  
**TTCTACAAAAAGAAATTCCCTGGCTGGAAATCGAGGAAAGTGC**  
**GTGCAATCTCAAATCCAGAATTGTCCT**  
**TCCCTTGCCAAGCCAAAACAAATACCATT**  
**CGAATCGTC**  
**AAATTGTTTTGTTGTAATTGTAATTG**  
**TTAAATTGTAATCTGTGATTGTTAATTG**  
**GAAACTGTTAC**  
**TTCTGCGATACTGCAATTGAGTGTGTT**  
***CCGTTA***  
***TCCGTCC***  
***GTCTTCTCA***  
***AGGTTT***  
***GATGCC***  
***CATATAAATCGGTTAT***  
***TCGAATTGTTATTCC***  
***TATAACGTTGATGAGTCA***  
***TAGCTGCTAGGAAACAAACGGT***  
***GTAATGCCATTAC***  
***CTCAAGCCAGGATTGTTGAAACGACAGTTGAGTATTG***  
***CAAAGGT***  
***CACAAAAAAATTATATAAA***  
***TTAAAACGAACAAACGAAGGCGAACCAAATAAGAAAAATACC***

#### 42. Drd

AGTTGACGTTGAATGTTCGTCTTGGCGGTGCTTAATTAAATTGCTGAACGCTGACACAAACGATCCGAAAATCATC  
GATCATGCCCGAGTTGCAAGACAAATTACTATTCAATTGTTGTCATTGTTCTGGTGTATTCTTTGCAT  
CTTTGCTTTGTGTTGGAGTATTGTTGATTATGTCAGAGACTGCAAGCTTATAAGTCGAAAAAGAAAAGTA  
CTTAAAAGGGTTGGTTAATCCAGGATCCTTGACAGG**ATGTCGCGTATGTCG**  
**CATATGCTAGCACTACTGGTACTA**  
**ATTGCCAGTAGTCAGGCCCTCGAGGGCGTGGTCCAGCCC**  
**ACTTGCAATCTAAAGGACAATTGCAATTGGCGATAATCT**  
**TCTGGCCGGAAACAAATTCCGATCCGATGTCGCGATGTCG**  
**CCGATAAGGTCAACTATAGTCTAAAGTTATTGCGGACGATGCGACGACAACAATCTGGAAATATCAA**  
**ACCGCCAATTG**  
**AGTCAGTTGCATCGGTAACGCGCTGAAATTACACGATAACCCAGATGGAACAGGCGGAGGATGATGACGACCAAGTGG**  
**GGAGGATGAAAGGGCTATTGTAACACATTGCAAGCGTTCTGCC**  
**ATTACGCAACCCTCGACGCCAGAAATC**  
**CGGATGCCCTGACTTTGTCGCCAGATGCATCAGTTCTGAA**  
**ACGCCCTGGACAACCTCGATCTGATCTGGCTCTGAA**  
**ATGCA**  
**CGACTCCAGTGGCAAATTGA**  
**ACTCGGGATTCTGAA**  
**CTGATGGCAACATCAACCCGGCGATTGACCGCT**  
**GGGCAATTGCAACGCAATTGAGATCAGGATGCGGGCAGGATCAGGATGGT**  
**GACTCCATTGCGATCTGGCTATTGCGACACCAATTGCAAGCGATTGAAAGAGTTCTTCAAGCTGATCCA**  
**ATCACATCGGA**  
**CCTTTAAGAGCGAGTTCAACGACCCGGCCACCGAGT**  
**GCCCCGCTACTCCTGATCAACTGGGCC**  
**CTGTGCGCTC**  
**CGGCTGCTCCGCCCGATGTGGAGTACAGTGTGGCGAGT**  
**ATCTGGCAACCAGACCGCTTCCACGGCATCACCTCA**

ATGTGCGCGTGGAGCCCCAGATGTGCCAGGTGCGGATCAGCGGCCCTGGGATCGGAACACCACGTGGCGGTGCGCTTC  
 TTTCTGCTCGTCTCCGTGGCCGTACTGTCCACCATCTACGATCGGTCCACCAAGTCGCAGGCCAAGCAGAACCTCCCTG  
 GTTCACGGCCTCTCGCTGGACAAGAACCTGCGCTGGCTTCAGCACGAGCACTGCTCCGGAGACATTGAAGCGGTGC  
 ATGGCATCCGCTTCTGAACGCCATCATGCTGATCTTCGACAAGTCCATGCCATGTTCTCAATCCGTACAACAAAT  
 CGCACCGCCATGTCAGAGAGTCTGGGTAGCCGTGGACGGTCATTGGACGAGCTGCCCTCCGTACACGGATCCCTTCCT  
 GCTCTTCAGTGGAAATGTCGACATCCACTCCCTTTGGCGCTCATGAAGCAGCAACCCATCAGGCTGAAGAACGAGT  
 ACATTAGTCGGTGTGAGGATTGTTCCGCCCTGGCCGCTTGATACTCTTCGACGTATGTGCTGCCTCTGTTGGGC  
 AGTGGACCCCAGTGGAACTCTGGTGGGGCCACCATGCGACATCTGCAAGAAGAACTGGTGGCGAACCTGCTTTCAT  
 TCACAACTAACCTCGGATTAGCAGAAATGTCGCTGACGACACGCACCATCTGGCATCGATAACGGAACTCTTGGGTGG  
 CTCCACTCCTCATCTGGACTGTGGCGGTGGCAAGACGTGGCCTTCTGCCCTGCTGCTCTGCACGGTGGGTACG  
 GCGGCTAGGTACTACACCACGATTGCAACCCAGCTATCCAACACTATATCTACTTGGTACCAACATTGCGCCTGTTCCG  
 CACCGCCGACTACATGTAACCTCCGCCCGCCACCGCTCCACGGTCTACATCATGGCATTCTGCTGGCTATGTTG  
 GCAAGTACCAAGAATGCCGGCTGAGCAGCCTGAGCTGGTGGCTGGCCACCGTCTGTTG  
 CTGCTGGGACAGCAGCCATGGGTGACATCAACTACGTGTACAATTCCACCCATGCGACTATATACGCCCTTGC  
 GATGCCCTGGTGTGTTGTTCTCTGGATCGCTTGTCTCCATAATGGATAACAAAGATAAAACTGACGAAGCTGTT  
 GCCTGGCGCGGTTCCAGGTGTCCAGAAGCTATCGTATGCCATTACCTGACCGAGTCCCGTGTCTTAAATGT  
 CGGCCGAAGGCGTCACATTCAACACTACTACAACCTTGTGTTGATAATTCTGATACCAACGAATTCAATATCG  
 TGGCCTCCGTGCCCTGACGGTGTCTCGATGCAACCGTCCAGAATCTAAAGAAGCTGCTGATCAAGCGACCCACGGCG  
 GCAAAGGTGTAAGGATAGCAAAGCCAAGGCCAGAGCTCCAGGATGCGACGACAACGACACTAGCCTCACAGCCCAG  
 CAGCACTGCCCTTACAATCGCATCATCCGCACTCGGATTAGTCTGGATGCAACACTCTTCTAGCTATGTATAACT  
 TGTAATGTCTTAGCATTAGTGGCAATATGTTAGTGTACCCCCAGCGAAATCCCCCCCCCCAAAAAAAAGA  
 ATCGCAT **CCAA** **TCCGTCC** **ATGTGCTCATCA**CATCCCACAGCCTTCCCCGAATTGAATCCAAGACAGCCCCTAGTTA  
 GTGGAACGCTTATACATTGAAATTGTAATTAGTCAAAACGAGCGACAATTAAAGAACCGAACCGAACC  
 GATCGCTAGCTGCTGGGTACACAACCTCTGGCTACACGGACTGTCTGCTGGCGGGAAAACGCCAGCTGCTGCAGT  
 GTCTAAATGAATTAGTGGCACTGTGGCTCGCTTTGCTGCCCTGGTGTGGACACCGTATCGCTGTGGCACTGAG  
 GGCTGGTGGGACTGGGACACTCGGGACCGCGAGCTGCTGCTGTTCTGTTGAATTGTCGAGCACAAATGCAACACGATGT  
 GCCGGTCTATTGCCCCGTGAAAAGTCCCCATATAGCCTACCGCTTGTCACTATTCTGTGGCGTCGAACAGTGTCTGT  
 GTGTCCTGTGTCGGCGCTGAGAATGCTCGTCCGAGAGTGTGATGCCATGAGTGCCTACCGGAATGAAGGCAAC  
 CTTTGGCAGCCGCCGAGCGATGCGTGCAGCTGAGCATCTGGAGATCGATGAAATGTGCTGGCAT  
 TCTAATTAAACCGACACACCGAAAAAGTCTTACAGAAATCTAAACCGAGTCCGAGTGTCAAGCGCATTTGTGG  
 GCGATCTACAGCGCTGGACATTCTAAAGAAGTTTCGATCAAACATGGATGCACTGCCACCAATTGAAACCCAGTCG  
 TCCAGCAATAAAACTACGTTAGTGGACCAATATAGCTGCTCGGACTACCACAAGTGTCTATGCCCATATGGATGAGCTGG  
 CAATGTCATATTCTGTTGTGGCAGCGGTGCCCTCGCACGCAATGAGATTCCCTGCGCAACGCATCCATGTAATA  
 TACTCCAGGAGAGTCAGTATGTTGGTGAACGATATTCTTCTAACGCTAACAGACTGATTGTTACTACTTTCA  
 TGAAATAATGTTTTTTAAATGACATTATTTAACGAAACATAGGGTACAACCTCCTTCTAAAGTTAAGCGTAATT  
 TTTGACAATCTATATGTATATGCTACATACATAATGGCTTAATG**CTTACAAT** **TCCGTCC** **TTACAAAT** ACTATT  
 TATATTAAATATAACATATGTCCTT

#### 43. fy

TCGATTAAATCGATGTAAGCATCATACGTAGCAGGTTGGCGAAGAAGAACGACTGCG**ATGTCCATCTATTGTTATGTT**  
 TGACAACAAACGGCGATTGCCAGTGTTCACAGAAAGAAGGAGAATGCGAAATCTGCCCTCTCGACGGTGGCCTCC  
 TTAAACGGCTTCACATGTTCTTAAGTCGCTGGGAAATCCAACCTGGAGGCCACGACGACCGAACAGCAGCCTATCCA  
 GAATCAGGCTCCCGCGGATGCCGAGGAGTGGACTTACATATGGCGCACCGGAAATCCATACCCCTGATCGTGTGGC  
 GCGGTGTTGGTGGCAACTGCTCAGACGCTGGCTGATCTGTCTTGAGCGATAGGCATGTTATTACCGGGCC  
 GCAGAAATGGCACATGCAACGCTCTCGATAGACTGAAGAAGGACCCAAGAAGTATGTCCGATTGTTGACGCTATTCT  
 GGAAGCCGCTGTGCGGGTACACAACCTCTGGCTACACGGACTGTCTGCTGGCGGGAAAACGCCAGCTGCTGCAGT  
 GTCTAAATGAATTAGTGGCACTGTGGCTCGCTTTGCTGCCCTGGTGTGGACACCGTATCGCTGTGGCACTGAG  
 GGCTGGTGGGACTGGGACACTCGGGACCGCGAGCTGCTGCTGTTCTGTTGAATTGTCGAGCACAAATGCAACACGATGT  
 GCCGGTCTATTGCCCCGTGAAAAGTCCCCATATAGCCTACCGCTTGTCACTATTCTGTGGCGTCGAACAGTGTCTGT  
 GTGTCCTGTGTCGGCGCTGAGAATGCTCGTCCGAGAGTGTGATGCCATGAGTGCCTACCGGAATGAAGGCAAC  
 CTTTGGCAGCCGCCGAGCGATGCGTGCAGCTGAGCATCTGGAGATCGATGAAATGTGCTGGCAT  
 TCTAATTAAACCGACACACCGAAAAAGTCTTACAGAAATCTAAACCGAGTCCGAGTGTCAAGCGCATTTGTGG  
 GCGATCTACAGCGCTGGACATTCTAAAGAAGTTTCGATCAAACATGGATGCACTGCCACCAATTGAAACCCAGTCG  
 TCCAGCAATAAAACTACGTTAGTGGACCAATATAGCTGCTCGGACTACCACAAGTGTCTATGCCCATATGGATGAGCTGG  
 CAATGTCATATTCTGTTGTGGCAGCGGTGCCCTCGCACGCAATGAGATTCCCTGCGCAACGCATCCATGTAATA  
 TACTCCAGGAGAGTCAGTATGTTGGTGAACGATATTCTTCTAACGCTAACAGACTGATTGTTACTACTTTCA  
 TGAAATAATGTTTTTTAAATGACATTATTTAACGAAACATAGGGTACAACCTCCTTCTAAAGTTAAGCGTAATT  
 TTTGACAATCTATATGTATATGCTACATACATAATGGCTTAATG**CTTACAAT** **TCCGTCC** **TTACAAAT** ACTATT  
 TATATTAAATATAACATATGTCCTT

#### 44. I(2)35Bg

ATTGCCATATCTAACTTGTGAAATAAAAATCGGTTCCGTTGGTGTGAACTTTGAGATATTCTAGAAACACATCCA  
 CAATGGAGAACTTAAAGGGTCTACAAAAATCCCTATACATATGGACGGACAGCGCCGACTGGACAAGCGCGTGGAGCAA  
 CTAAAGCGGCCACTGGCGCGATGTGGCTTGGAAAATGTGCAACGGCTTCTCCTATGCCAACTCCAGTT  
 CGACCTGATTGTGATCGAGTGCAGCCACAGTACGTAACACTGCTGACATGCTGAAACCGAGTGGCAAAC  
 TGCATTGGTGTCTACATCGGTCGGCGCTAGTTGCTGAGGAGATCAAGCTATCCGATTCAAAACTGTCGCGAG  
 GACTCTCAGATGCTTGACCGCGAGAAGCCTGGCTACAGAGACAGGGATCCTCAGCCAGGGTGTCACTCGCAAAGAAC  
 CGCCAGTGCAGTCATGTTGGAAAGATAAGCGGTGACGATGAGGAACACTGATCGATGAGGAGGAGCTGGACGAGGAAG  
 ACAAACAGAAGCCGGATCCGGCTGGTCTGCGTGTCTGAGCACACCGGGCAAGCGCAAGCGTGC  
 GGGCTGGCCGAGGAATTGGAGACTGAAAAACAGAGCCAGAAGGCCACCGAGAATGCCAAGTCAAGCTGTGGAAATTGTTA  
 TCTCGGCAGCCTCCGCTGCTCCACTTGGCCCTATTGGCATGCCCTTAAGCCGGTGAGAAGGTGCAACTGG  
 CCGACAACCTCTGAAATCCGACATTAACTTAGGCTCACCATGGTATACCAATCAACATTCAAATAGTTATT  
 AGCTGACTTAACAAACAATAATGTCATAATGTTAAACAAACAGCAGCGTCAAAACACAGCATGTA  
 AGTGTGACGTTACTAACAAATAATGTCATAATGTTAAACAAACAGCAGCGTCAAAACACAGCATGTA  
 ATATCCAAGTGTGATCACATTGCGATGGTGTACCCCTGATGAGATCGCGGTTCTGGCCCATCAAACATCG  
 CCGAGGGATCGAGTTCTGTACAAATATGTCGGCATCTGTCGGACAGCACCAAATAGTC  
 GCTTCATTGTCGATTTC

ATGTCATTTGAAAGACCAGAGTGTGCGAATCGGAATCGATTAGCCCTGGGTGTCCGGAGTGTATGGACGCTTGATGTT  
 CCAACAAGCAATGGCGTCTTGTTCACCTCGTATAGAACAGTGCAGGCCGGTCTCGGGGTCAAAGACCTCGCGGTGGACT  
 GGCCATTGATGCCACGGTACCAACGTATTTAAAATCGTAATACGAATGCCAGCGGTACGTGTATTGTTGCAGC  
 ACGCCGGTTCGAAACCTGAACTCCTGGTACTGCCAGGGCATGGAAATAATGTCCTGGAGTGTACTGGATTTCATTGG  
 GCCCACGGCAGCC **CAAAAACGCCA** **TCCGTCCAC** **TGG** AGTTCACGCCACATTGAAATCACCATGCAGAGGATCGA  
 AGTGGAAAGAAGTTGTGCTTGACCGGTAGGATTGTCGTTCTCAGGGAGTACACAATCACACCGTAGGCACCCAAATCC  
 GGAATGTAAGCAAAGGCATCTTGGCACTCCGAGCGGTCCGATCCACAACCTAAATATTGATTAATTGATCAGTTGGA  
 ATTTAAATAATACGCCTATGCATACTCACAATATTGAAAGAACGAGTCCTCTGGTCTGATCAGCGGAATGGTAA  
 ACGGCGCAGTAAAGTATCTGTTCAAGTCGAAAACCAGGATTGAATTGGCGTAATCTGCTGGGACTGCCAGGATGT  
 CGGCCAAGCCAGTATCGAGCACCCACAGGCATCGATACGTCGACTTGAATTCCAAGGGAAATGACAGTGGAGTTG  
 TCCTTCAGCTGGATGCCGGCATCTTGAGCCTTGTCCCGTCCAGACGTCCACCGGATGGTGTCTTTGGCCTGTGGCTG  
 CACATCGATGGCAGCTTGTGGCCTCCAACTGGGATACGGATCGAGCTTGTGCGACTTCTCCGTGAATTAAATGTCGA  
 TGTAGTTGAGTGTGGCAGCAACACCAGCCTCCATCTAAGGGAAAATTGTTAGTTACTAATTACATTAAACCTGAC  
 ACTGGCAAGTAACAAATTAAACATTAAAAAGTACCATCACCTGCGCAATTATTATGACTGATTGATATACTGTAAT  
 ATAAGCTATAGTTAATTATATCTAGCACTATCAGTAAAACATTGACGGCAACTATAGCTAAATAATTATGCTGT  
 ATTATCAACAAATTATACTTTTACGTATCTT

#### 45. lethal(2)gl

GCACACTAGACCCACCGATCGCAGCTTGTGTGTTAAAATGACGCGCATATCGTTTGGCTAATAATATTGTCAG  
 TTTTATAAAACAGATAACCGTTAACGCTCTGACCGGTGTTATTATTCTGTGTGTTCTGGTGTACACGTTATTCAT  
 TTATTCTCATTAAAGCAACGATGTGAAGAACAGTGCAGGGCGTTCTCAACACATAAGAACATACACAAACTATGCGTGT  
 GGTGAGCAAATAAAATAGTCAAAAAATTACTCGGCCACCGCTTAACCGTACAAACGGAAATACATAAAAGTTAGAATA  
 AATTGCCACAATTAAATATACTTATTATCATCAAAACCAACCCACAAGAGCATAACCAATCATGTTAAAGTTA  
 TCAGAGGAAAAGGGCAGCAGCCCAGTGCTGACAGACAACGCTACATCGTAAGTTAGAATAAAATACGCCACAATTAGA  
 TACATATTATCATCGAAACCAACCCACCAAGAGCATAACCAATT**ATGTTAAAGTTATCAGAGGAAAAGGGCAGCA**  
**GCCCAGTGTGACAGACACCGCCTACAGAAGGACCTTTGCTTATCGTAAGACGGCACAGCATGGCTTCCTCATAAGC**  
**CTTCGGCTTGTGTATGATCCAGTTGAAACTTATGCCAATAGGGACGCAAACAGGGCTTAAAGTTTCGGTCAA**  
**CCCGGAGTTGAATTGTACGGTCAGCATACTTGTAAACAAATTCAAGCATCGGAGCTTAATGTACAATTACTGAAATGGGT**  
**GTATGAACTGGTCGACACTTGTGACGGCAGCGAATCAATTAAATTCTATGGAGCCAGTTGGAGCAACGTTGCTGC**  
**CAATCAAAACACTACCGTTGACGGAAACTTAAAGTTCATCGCTGTGCTTCTCAGTAAGGATCTGCTATGG**  
**ATTGGAACAGAAGGTGAAACATCTACACTGGATTACATACATTACATTAAAGGAGCCTGTAATTACCATGACGT**  
**TGTGCTAGAGCAGGTGCCACCAGCCTACAAGCTAAATCCTGGTGCAATTGAGTCATCCGCAACTTCAAACCTCCCTA**  
**GCAAACCTCTAGTGCATACAATCGGGCCTTGTGTTTGTGGGATTTGAAAGCGCATCTGTCAGCGAGCATAACATA**  
**GCCCCCTGGACATGGACAGAGCGTTGGTCTTACAGTGAACCTCGAAGGATCTGAATTACCTGGTACCGCTGATGGTTC**  
**ATACGCCACTTGGAGCATAGATAACCCAGAACCGCCGTCAATGTTAATTATGTCCTTATGGACCTGATCCATGCAA**  
**AAACGATAAAATCGACTGTACAAAGGCAAGCGAAGATCCAACGATGTAATTGTTTTCGGCGCATGCCACGGTCAGCAT**  
**ATGGTGTACAAATTGTGTCCGTTACGCCAGCGATGGACACAAAGTGTGTCTGACTTTACGTCTAAAGTGTATTGACTT**  
**TTTGACCTTGAAGGAAATAGAGATGTCGCTGAAGTCTTGTGTTACTACTGAAAGAGGAACCTCGCCTACGATC**  
**TTACTGACCTAATATTGTGCTATCAAGGCCATATCTTCACTCGTCCATGCACTCGTGTCAATTGCAATTACCTT**  
**GCTCTGAAAGTCGTACAGCGGTATGAAAGTATTGAAAGAGCTGGAGATGACAAGACATTGACTATAGCAATTAG**  
**CTGGCCTATCACTGGCGTACTCTCCGGATAACTTAGAAGAACATGTTAGAAGAGGACGCGACTAAGCTTATGAGATT**  
**TGTTAACTGGTCACGAAGATGGTTCTGTTAAATTGGACTGCACTGGAGTGTGCTTAAACCAATTATAATTAAA**  
**ACTAGCAGCATTGGAAAGTGAGTCAGACTCCGAGATGACGCAGCTGCAGATATGAGTGCGGAACAGTCATGAGG**  
**AGAACGCCATTGAAACAGGACTTTGATCCTTACAGATGCCCTGTTAGCAGTGAAGAAAATAGCATTCT**  
**GCCCCAAAACGGACAACCTATTGTTGGCAGCGGGCAAATAGTATAGCCACTCATAGACTTACCCGAAAAA**  
**GTGTCCTTAAACATTCATGAAATTGGTCAGCGATCGTGTGATGGATTGTTGGAAGGGTCACGATCAGTAAACGT**  
**GCGATCGAACTTATTAGACGGAGAAGCAATTCTACGACGGAACGTTGTGTAATATCGGGAGTACTGCAAGTTTGC**  
**CGCCAGCCAGCATAACATGCATGGCACTCGAAGCAAGCTGGGCTAGTATCTGGTGGACTGCGCACGGCTTAGTTCTC**  
**TTTGACTTCAAAAACTTTGTTCCAGTATTGATCGCTGACTTTAACCCAAATGATCTTACTGAGCAGGAGAGCAGCT**  
**GTCTCGTCAAAGTCTTAAAGAAATCATTGAGGGAGTCATTAGAAAGCTTGCACGGGTCATGACCCAGGACCAACC**  
**AGAGCAATCAAGTACCAACACGCTGGAGAAGCAAGACCCGTCAGAGGGCAAATAGAGGCTCGTGTGCAAGATGACGGGCTA**  
**GGATCCATGGTGCATGTTACTATTGCAAAACTTATGTTACTAATGTCACATACGCTGCCAATTGTCAGGAGCAGCT**  
**AACAAATGCCAGTACAGTCTGGTTCTCTGCATTTGCCACCGCGCAGACCGCGGCAAATGCCGTCCGTCGGCAA**  
**GTGGCAATGCACCACACATGCCCGCGAATTCTGCGCAGCTGCTAAAGAAATACAATTAAACATCGTGTCTCCT**  
**GTGGGGTATTCTATTGATCAGCGGGTAGGCCCTGTCGATCAGCTGAACGCCGGTAAAACGGGAGTCCACCGCA**  
**TCGTGTACTTATTGCTTCCGAGGAACAGTTCAAGGTGTTTCACTTCCGCAACTAAAGCCATTAAACAAATATAAGCTT**  
**CCGCTAACGAAGGTGCTGGATTGCCGATCCATTGTTCTGTTAGTTGTCGATATCCCGGAAACACTGCAGAGT**  
**ATGCACTGGTTGAGCCAACTAAGTCCACGCCTCACATGGCAGTGAGAGGCGGATCCTAATATCAGTGGAAAGCTTGGC**  
**TGTAAGTCGTGGAGATGTATATAACGAAACAGCATTGATATGTTAACGAAATATGGCGATATCATGGTTTATCAGTAC**  
**CTGAATTAAAAAGACAGCTGAATGCCGAGCAGTGCAGCGGGAAAGACATTAATGGAGTTCGTCACTTGTCTTACAAAC**  
**TCTGGAGAAGCAGTGTATATGATGTCCTCTGAACTGCGTATTGCTTACGCCACGAGTCGTGCAACCCAC**  
**TGGCGTTGTTCCAGTAGAACCAATTAGAAAATGAAAGAGTCGTGTTGGAAGAAAATGATGCAAGAGAATAAAAGGAAACCT**  
**ACGCATGTGATGAAAGTTGAAATACATATGAAATTAAACATCAGGCATTCAATATGCAACAGGCCTGCAGAGGAA**  
**AAACGTTGGAAGAAAATAGTGTGAGTCAGCAAGTTAATGGAGTCACATTCAACCTAATCAAGCTAACGAGACTATCAG**

CAGCTCTATTGGCGATATTACCGTTGACTCGGTGCGCACCATTAAATATGACGACCACACTTGTGTTCTATTAATA  
 CAGAGGAAACCATTGAAATACGACGAAACGAGTACTAGTTCTGTTGTAATTAAATCTATAATTACAAACATTCTCATG  
 AAAAACGAACGGAGACAACAAATAGGAACGCCAAAACAGCCCTGAAGAAAGCAATTAAACATTGACAGAAGCCG  
 TAACCTACTAATTATTTACCTTACAGAAATACGACGCAAACGAGTACTAGTCCTGATGTAATTAAATCTATAATT  
 CAAACATGTCTCACTAAAAATCGAACGGGGACAGCAAATAGGAACGCACAAAAATTAAATTAAACGCGATTAC  
 ATACAAACAGAAATGACAGAATGATAATATAAAATTACATTTTATTGGCTTAAGCGATGTTGTTCCAAAACCA  
 TATAATTATTTATGATTTATGTAATGTTTCATGTATTTCGCGTAGTGACTTATACCTTACCGTATGATGAAAC  
 GAAACATGATATTGAGAAAGGTACAAGGATAAAAGTGACATTAAAGGCCATTGAAATTAAACGCAACCTG  
 GATAACATGCCGATTAATTACTCTGTCGCGTATAAACACAAATA **CCGATAAAATTGGTCCGTC**  
**CC**ATTAAATTTAATAAT  
 TGATATTAAATGGCTTATTGGATAAATAATTCAAATTATAGGAACATTGGGTTTAAGGGATAGAAACAGC  
 GCTGGTGTCTTCATTCCGTCGTATGACCGTTCAAAGATTATAAAAAATTGTCGTGCCACATTTTAAATTCTT  
 TTATTTCGGTAAATATTGTGATACAAAAAATGCCCTACAAATTAGGACAACAACAAATTGCTTATTATGATGA  
 ACAAAATTAGGAGTACAAAAAATAGCCTGATGCAAGGATGGGAGAAAAAGACCTGAAAGTCAAAAGAATTGTTGAT  
 CAAATCCTCAATAAGAAAACACTTAAATTAAAGCGTGTTCATTAAATTGAGAATTTCGTAATTGAAATTG  
 AAACATTTTAGCACGAAATTACATGTATTAAATTCTTGCTAATACGTCAGTAAAAAGGCTACATTGATGATAC  
 TCACAAATGTAATTTCGATTCTACACAGGTGCAATTAAATTATGGTTTATGAGTTTAGAGGGAGAAAATAT  
 GTCAATATGAGAAGTAAGTAAATGAGCCTCAAGTACTTGTAGTACATAATGTAATTAAATTAGAAATCCTACGGCTT  
 ATATTCAATTGAAAGTATTAAATTCCCTGACGATTGTTGAGTAAATTGTTGTTCTAGCTTTTGCAATTCTCCTCC  
 GTTACTCCGTTACTCGTATAGTAAAGGTATAAGTATATAACATAGTATTGAGATTGTTGAGAAGTATGTTACAGGAG  
 AAGGAAGCTTCCGACCATATAAAAGTATATAATTGAGTATTAGGATTATAAAACATGTTATTAA

#### 46. Oseg1

TTGTTGTGGTGGTGCATGGAATCTGTTGAAACACGACACCTGCATTTAGGAATTCTACCAACTATGCAGGATTA  
 GCCAATTGACCTCTGCAACA **ATGAGGGGTGTCTCAAGTGGCTGAAACGATAGAGTTCCAAATAGCAAAGACAATGA**  
**GATCAGTGTCCACAATGTGCTACCGACCCGATGGCAGCCAATTGGTGGCTGCCGGGATCGGGTGTCACTACG**  
**ATCCAAACGATGGCTCCTGCTGAACACCTTGAAGGCGCACAAGGATACGGTCAATTGTTGCTACTCGCGGGATGGC**  
**AAGCGGTTGCCAGCGGAGCCTCGATAAAATGGTCATCGTGTGGTCACCTCAGCTGAGGGACTGCTCAAGTATTGCA**  
**CGGTGACTCCATACAGTCATGAGCTCAATCTGTTCGCACCACCTGGCCTCGTGTGCTCTGTCGATTGCTTT**  
**GGTCCGCTGATCAGAAGGGGTGCAAAGTACAAGATATCCGCGCGTGTCAATGGCTGCTGTGACCAATGATGGACAG**  
**TACCTGATCCTGGGCTGCCAATGGCACCATACTCATTAGGAACAAGTTGGCGAGGAGAAGGGAGAATCGATCGGCC**  
**TGGTGGGCCAACAGCAACGCTTCAGTGTCCAATGTTGTCGGCCAGCGGTCTGGCAGTGTGGACACCACCGTGTGG**  
**TGGACTGGAGCCAAACGCTATCCTTCACACTGAGCGGCCAGATGATTGCAAGGAGCGAACCTGGCTTGATCCC**  
**TTGTCGCTGCACTTCCCAATGGGAGTTCTGCTGGTGGCGCTGCACACTGGATCTGGACAGTTGCCCTACATCCAATGGTCAATCCT**  
**ACACCAATTGGGTGTCAGATGGAACCCATTGGCTGCTTCAACATTGCGTCTAGTACCGTGCATGCCCTACCGCGAGCG**  
**TATGCCCTCCGGAAAACATGTGCGACGTGATCTCAGCACCTGATCTGGCCAAAAGGTGCGCATCAAGTGTGAGA**  
**TTTAGTTAGAAAATAGCCTTATCGGAAATCGACTGGCGTTCAAGTGGCGAACGTGTTCTCTATGAGCTGAGCT**  
**CTGGAGAGGATCAACCCATGCACTATAAAAGTGGCGAGAAGATCCAGCAAAGGACTTCACTGAGCTGCTGGTTGTTGT**  
**GGCCGTCACATTGCTGTGCCAGGAGAACGCTTCAATGTCGGATTTCATGGGTGTCCTGCAGCGCAGTGGATCAT**  
**GGACTCGTTATAAGATAACATCAAAGTCACCGGTGGCCCGTGGGAAGAGAACGGACTAATGGTGGGCTTGAAGAACGCC**  
**AAAGTGTCCGAATCTTCACAACAACTCACTACCAACTGCTCATCACCACGCGTGTCACTGGGTCGCTGCCTGGATATC**  
**AATTGCAAGCGCAGCAAGATCGCAGTTGTTGAGTGGACGATGTGGTCGCTGGTTGTCGGGACGTAATCAACGATAATT**  
**GTACCAAGGATACGGGTGTAACACAGCGTCACTTGAACACCGCACCTGGATTCGATGCTCTGCTATACCCACACACAGGCG**  
**GATTAAGCGTTCGCGTGGGAATCTGCCGCCGAGGGCACCACAGAACATGCATGGCGTGGTGGGGCTTGGGAGCC**  
**ACTGCCCTCTGCCCTGCCGGAAATTATGACAATACGCCCTGGCTCTGGTCCACCATGTCAGTTGCTACG**  
**TGGATTGTTGACGAAGCCTACCAAGGTGCGCTGCCCTGGCGTGACCAACCGCGACTGGGAGGGATTAGCTCAATCCGCTC**  
**TGGAGGCAGTGCACATAAATATGCTCGCGATGCCCTACGTCAAGGTGAGGAATTCTCCGTTGCTCAAACACTCATCGGTGAG**  
**CTACCGAGATCAGCAACAGCGATCGCGGTTCTAAGGAGGTACTTCTCGCAGAGAATTGCCCTTGGCGCAAGTCAA**  
**GGAGGCAGCTCGTCTATTCTCAAATGCGGCCAGTCCTCAAGAGACACTGAAATGTACACAGATCTACGTATGTTGATT**  
**TAGCCCAGGAGTATATTAAAGGACGCCACCGACCGAGAGAACGGACTGGTACGCAAGCGAGCGGAATGGGATATTG**  
**GTAAAGGAGCCACGAGCTGCTGCCAACTATTGTTCAAGCTGGCGAAATCAGAACGGCATTGATATTGTCGAGAGCA**  
**AGGATGGGCCGATGTGCTTACGATACGGACGACTGAGTCTACGGAACCGGATTCCTGGCAATCGGTGGCCAGA**  
**ACCTAAAGACCTCAAAGCTTCTCCCTGGCTGCTGAGATTTCAGAAGCTGGCGATGAAGGCCACGGTGGTGCACACTG**  
**CACATCGAGGTACGCGACTGCCGGAAGCATTGCTTGGCGGAATCGCTGCCGAACTTCTGCCACGGTGCACACTACCA**  
**GCATGGCAATTGGCTGCCGAAACGATCAGTTATAGAGGCGCATCAGGCCCTACATAAAGGCAGGACGCACCAAGGAAG**  
**CCAATCGTTGCTTAAGCAACTGAGGAATACGCCATCGGAGGAGCGCTACTGGATGCGAGCTACTTTATTGGCTG**  
**CTGGCCAAGCAACATTAGATGTATACGCCAAGGAAGAACAAATCTGTTGATCACCACTTACGGAGTACAAAAAA**  
**TCATTACGTGTTGCCAAGGTTATTACGCCCTACAGTGTCAATACATATGAAAGAACATTCAACACACATTCCC**  
**CAGTTAGCCTATTTAACGTAAGCCGTTTATAACTAAACGAAGTGGAGCACAAGGTGTCAGCATGTT**  
**TCGGTTCTCTCACTCTGCCAAGCAAGCTAAGTTTGCAAGGCCACAAGTTATGCCCTTATTAAACAGAGACTGCA**  
**GTCGGTGAAGCCACCAAGCTGGAGTCCAAGAGCAGATTGATCTAAATTCTAAATAGCAAGGCTTGCACAAAGTGGCTCA**  
**ATGATCCGGAGGAGCTTGCCACTATGTTACAAGTGCTCAACTACAGTCACATTGAAATAGCAATTGCTGCCCCACT**  
**TGCCGGCAGGATTATATTCTCTTCTGAAATTCTACCCCTAGTACAATTGTCAGGATATCTC**

TGATGCCGAGCGGGAGCGCTTGCTGGCTCCTGCAAAGCACGCCAGGGATTGGATCCATTAAACGAGGATGTGGCA  
 GTGCATTGCCCTTGAGCTGGATCGAATGCCCTGCGCATAGATCCAATCATGTGCTGATTCTAAACGGAGGGGA  
**AAGAATGTGCGGAATGTACTATCGTAATATATTACCCGATCTCAGGTTACCTATTGTGCTCAATGTCTGCTGCTTT**  
 CTATGCCGAGGACTTGAGCTGCAAGTGTGCAAAAAGTCAGTGTCCCTTGTACACACATCCTCCGAAAAGCTGATGG  
**AAGACTTTGAAATCGACATAAGAAGTGTGCAATATATAATT** **TCCGTCC** **AAACTGTACAAAT** **AGGA** ACTATACGCA  
 AATAAACGTTACGCCAGGTTCCCTCG

#### 47. PkaR2

AACTTTGTAATTTCGCTGACACGGAATCGGAATACAGAGGCTCTGCTTGCATATTTCACCGTCGACTTGT  
 TTACGTTGCCATTAAATTTCAAAAATCTAACGAGTAACAGCGGTTGAATGCCGGTGGAAAAATATAGAACCGT  
 GTGCCGTTGTGCGTGGATCGCTGGAAAATTAAATACGAATCCAGCAATCAAATCTATTACCGAAATACTTCAA  
 AACATCAAACACAGCGAAAGCGCAATAAACACCAACATCAACGATCCAAAATAATGTAGAAAACAAGGAAAGTC  
 ATGAGAGTAACAAATTAAATTGGTTTAGCAAAACAACAAAAGCAAGCTAAGGAACAAGGCAAAAAAGTCAGCGAAG  
 AAAGAGACGGAGGAGCAGGTGAAACAGGTGGACGCAGGAGACCCACCCACCGAAAGAAACTATCACACCAGCAGGTGA  
 CAGCAAAGCAGCAGAAAGTCAG**ATGTCGAGCGATTGAGTCAGGATCCAGGTGCCCAGGAGCTGAAGGAGGTGCTGC**  
**TGCAGTTCTGATCTCTTCTGGTGAACACCGCCGGATGTGATCGATTACGCTGTGGAGTACTTCACCAAACGTCAG**  
**TCGAACGCCGAGCGTCTCACACGGATCAAAGCACCGATGACCAGCTGAGCGTCAACTCACAGGATGCCATGCCA**  
**ACCACCACTGATGCCAGCTCGCTCGAAATCAGTTTCGCCAGGCCTACGATCCGGAGGCGGATGACGACGATGATG**  
**GCGCCACGCCGTGTCGCCAAGACAGATGAGCAGCGCCGGCTGGTGCAGTCGGTGAAGAACGTCCTCCTCCGA**  
**TCCCTCGAGAAGGAGCAGATGAACCAAGTCTGGATGCCATGTTGAGCAGGAAAGGTTCAAGCAGGAGTACTTATCATCCG**  
**CCAGGGTGACGGCATAACTTATGTTATTGAATCTGGCTCTACAAAGTCTATATAATGACAAGCACATTAACA**  
**CCTACAATCACACTGGACTCTCGGTGAAATTAGCGCTCTACAATATGCCAGAGCGGCCACCGTGCAGGCGAAACG**  
**AGTGGTCTACTTGGCCATGGATGCCAGACCTCCGCCATCCTCTTGAAGTCGCCCTTCAGGAAGCGGAAATGTA**  
**CGAGGAGCTGTTGAACACCGTGCCCATGCTGAAGGCCCTGCAGAACTACGAACGCATGAATCTGGCGATGCCCTGGTT**  
**CGAAGAGCTACGACAATGGCGAACGCATCATCAAACAGGGTACGCCCGACGGCATGTACTTCATTGAGGAGGGAAACG**  
**GTGTCGTCGCGATGGACCGAGGACGACGCCAGGTGGAGATCTCCAGCTGGCAAGGGACAGTACTTCGGCAGCTGGC**  
**GCTGGTGACACATCGACCCGGGCAGCATCCGCTACGCCACGGGGCGTCGTAAGCTAGCATTCTGGATGTTATT**  
**GCTTCGAACGCTTAATGGAAAAGTGTAGTGGGGCATTAGCGTAGCATTAGGATATCGTATCTGGAACAGGATCTG**  
**CGCGAATATTCGGCAACTTGGCTTAATTAGCTTGTAGTTGAGTAGTAGTAATAGTGGAAATCCATT**  
**CATGCCCTAATTGATTGATCAAATTGATTTAAGTCTCAAAGATGGATTAAAGTTCGCAAGTGTGCTAAGACATGC**  
**GGTCAATGAAATGGTCCTTAATTAAAAAAATACATTAAATTATAATTAAACTCAAA** **TCCGTCCAGTCCT**  
**AGACACAGAGGCCATTGAGCGCATCATGGCTCCTGACCGATGACTCAAGCGCAATATCGTATGAGCAAATGT**  
**TCACCGACATGCCCGCGTAATCGAATTGAGAAATCAGTTGAGAAGGAACTAGTTGAGCCTTATTGTATAGTTATT**  
**CAAATTGCAATTGTACCATATACCATCATTCAAATCAAACACAGAGCTACAAAATTAAAGAAAGTAAATAAGCCAGT**  
**TTCGTTCCGACACAC**

#### 48. PpcS

AATTGTCCTGTCAAACACTGCAATCAGCTTACCTGCCAACCGGTTTTTTTTGGCTGCACTTGAACCCCG  
 AGGTTTCGTGCGAATAAGATACGGATTGAGTGCCTGTCACATTCCCGCTCGTCCGCCATCCATCGCTGC  
 CTATCGCATTTCGGTCAGAACCGCCGGACCTTGCTGCCGTGCACATCGCATCGCATTCCACAGCTGCCTTTCA  
 GATTTCGTTCGCTCTCTCTCCGGTTTCAGAGCGCTCCGTCCTCGCTCCCGCTCGCGCGCTCCCTTAACG  
 GTTCATTGATAAGCAGGCCACCCGAGCCCTGCCACGGCCCTGCCACGCATATTCCGCCCTCGCATTTCGGAA  
 CTGAAAAAAACCAAACCAAAGCAAAGCAAATCAATTAAAGATAGTACGCCGAATAGTGCAGTGCAGTGT  
 GTTAATTAAATTGGTGCCTTCGACATTCTGTATTATTAAATTGGCAGTGCATAGCTCCGGGAAAGAGGGCAAC  
 TTTCCGTCATTTCGAGCTTCCGTTGTCGCCAATTAGCAGCAGGTGACAAAGAAAAAGCAAAACAGCTAAAGG  
 ATAAGCCAAC**ATGACGCACTGGGAGGACTTCTACAACACCGCACCTGCCGCCGGACTTCGAGGACAATCGTCCCTGC**  
**TCAAGGAGTCTGCAACGGACAACAAGCTCCAGAATCGCATCGTCTCGTACGTCCGGTGGCACACAGTCCCTG**  
**GAGCACAAACGGTGCAGGTTGTGGACAATTAGTGCAGCTGGCTCCGCCCGGAAATACTCCCTCGACCA**  
**CGACTATGCCGTGATTTCATGCATGCCACAAGTCGCTGGAGCCCTCACGCCACTTCACCGCCAGCAGTCTTCG**  
**ACATGCTGGACATTGCGGACAACAGCCAGAGTCGACGATAGCCATCAAACCGGATTGGAGTGTGTTGCTCCGGTG**  
**CTGGCCAAATACAAGATAGCCCGGAAACGCAGATGATCCTGTACGTGAACTTCACCGCGTGGACTACATGTGGCT**  
**GCTCCCGGCCCTGCAGTGCCTGCCCTCGAGGAGCGAGCCGTCTACCTAGCCGAGCCGTATCCGATTCT**  
**ACATACCCGAGGACATGATGCCACCCACAAAATGCAATCAGGAGATGGAGGCCGACGATTGCTGCACTGGTGC**  
**AAAATGCTGGCCCCCTGCCAGTTGTTGGGCCACGCCCTTGTTGTCCTTAAACTGGAAACGGACGAGAGCCT**  
**ATTGATTGTAAGGCACGTGACAGTCTCAACAAATACAAGCACAAGCTGGTATTGCCAATGTGCTGCAGACACGCAAGC**  
**ATCGCGTGGTGTGTCACGCCACAGATTCTACGAGCTGCATTGACTCGCAACAGACGCTGCAAGGACTCGAGATC**  
**GAGGAGCCCATCGTGGCCATGTTGCAAAAGCAGGCCAGTTCATCAGTAACGCTCAACAGCGCCAATGACCGTTGTC**  
**GATGCCCGGAGGCCACAGACGCAACGCACTGGCACCCACCAACTGGCACCGGTGACGAGGACGAGGATGGCG**  
**GCGATCTTGATCGGATCCATTGCGTACCGAGCTCGCATTATTGCCGTGTCATGCCAGGCCAGGCTTGC**  
**CGCAAATATTGAGCAGGTTATAGTCATAGTGTATACTGGCATGTGGATGACAGACATTACATTGATTCTCATCCTGCC**  
**CTTGGACTTGTGAGTCCGCCACAATTAGCTATCTATTATGGTCGCCCTCATATTGAGGTTGGCACCGCTCATTAG**  
**TCCATTATCTGTTATTATTATGGTATTAGTTAATGTATTGGCCATTGTCCTGAGAAATGATGTCACCT**  
**GTATGTATCCGCCATTCCCTGGCTCTTCTATCTGTCCTGCCAGTTATGTCATCGAGTACGCAAG**  
**CGTGGTGCAGACATCAAAGCGATTGATTGTCAGAGCTTATGTCATGCTTACATTTAATGAAGCATT**

CGATCGATCCGGCAACTGGAATTAAATCCCACAAACACGTGTTTGATATCCTAAATCGAAATAGCGACAGAGTGC  
 ATGCAATCGAATTGCGGACGTTGTCCTGGACAACGGTGTGTCAGTGGCAGGCAGCGACAATTGATT  
 TGCTGTTGCATATATGGGTGAAAAGAGTGGCAGCTTGCCTGCTAATGTCCTGCGTTAAATTATGGCGGC  
 CCAAGCCTGGCCTGCTATCGAACGAGGTCAGTGCCACCGTGCCTAAAGGTCAGCTAAAGCACCTCAAATTGAGT  
 TTGAAAGCGGCGTAGCAGAACGCTAATTAACATCCCACACTCACACAGGCACTGAGTTATTCTGGCAGGCTGAGATAG  
 ACTGGCCTGGCATGGTTCCACATCGTAACCTCAGGTCTTGCAGGCACTGAGCTGGCAGCCTTCCACATTATTATTGCAT  
 GTTATGTCGGCCACACGCCGTCGGCATAAGCTCATGCCGCCCTAAACGAACCTCTGGCAAGGGTTGGCGGT  
 GGATTGATCCTCCTTCATGGAGTTGTAAC**CAACTTCCGTCCGCATT**TGTTTAATTTCAGCCACAAC  
 TATGCAGTGCCTGTTGTCGGCCAGTTGTTCTGTTGTCGGCTGGCAAAAAAAATTCACATTCT  
 TTCTGACGTCGGCTTTGTTGAGGGATGGTGGCTGGCTCTGCGTTGTTGTCATCAAATTAAAT  
 TTTATTGCGCTCGCACGCAACACCGGCTGGAACAGATAACCGAACATGGGAAACAATCACACATGTGGCTTAG  
 GATTAAGCCAGGCTCCGCATTCCTCGTTCACTTCTGCTCGGTTTATGGCTCGCTACCTTAGTGTGC  
 ACCCACATATCAAGATACCCCCCCAGTGTACTTCTCCCGTATCGATTGTCATTAAGTATAAACATAAAATT  
 AGTTGCCAACATGAGTTGCTACATAAAATATGTATAACTAGCAAGTGTGCAACAAGATGGCAGCTGCAATGGAT  
 GGTATTGCTTTACTGCAAATGGTTTCTCAATGAGTACTAAGAGAGAATACTAAGCATGCCAACATAA  
 ATAAGCTACTAAATACAAATAAGTTGAAACATACATGTAATTGAGAAGAACCTCTCCATATCCGTAACCTAGG  
 ACTGCATTGATGAAAGGTCAAGTCCCTGATAAATTCTAGTGAATTCAAGATTAAACCTAAGTACATGAAAGAA  
 TTTCAATAAAACAGAAAATCAAAGTCACAACTCTCAAGTCATCTACATATATGAAATCTATTCTATACACAA  
 TACATGTAATTATCCAAGAGGGAATAAAACCAAGTGAACCAATTGTTCAATATCCAATTGGACCAA  
 CCGATAAGAACAAATTAGAAAAAATACAAACAAGCTATAATTGACGAATATATATATATATATGTATATGT  
 GTATATCTCGATGTATACCTTAAATATAAAACGAAACAAAGGAATTCCATTGAATATTACTCTCACAAGCATTG  
 CCCTCTAAGCTAGATACATAACTATTATTGTTGAAACGAAACACTAAATCCAATTGACGCAATGCAATTGGT  
 CACTGCACCTGCCCTTCTACCGGAATCCATGAGCTGTTAAGTGTAAAATGTTACCAAAATGTTACCAAAATAG  
 TACCATATGCCGACTAAGGTCTAGTATCCATACATACATCTACAAAGATACATCAATATATATATTGAAATGT  
 GTATTAAACTGAAATGAAATAGCTTAAATGTATAGACAAACGAGTGTGTT

#### 49. RecQ5

ATCTCTAACACCAGCTGTTGGCGGCAACCAGTTGGCTGAAGCAAAACAATGTTATTAAATAAAACGCAATTAA  
 ATAACAGCCCAAGTATGGCGCATGAAAGCGCTGTGCACGAGGCGCTGAAAAGCATTGGCCACTCGAAATTCAAATCG  
**GATCTACAGGAAAAGCCGTGAAGTGCCTGTTAAAAAAAGCAGGATGTGTATGTCGATGCCACGGCTCGGTA**  
 ATCCCTGTGTTCCAATTGCTGGTTGATGTCAGAGAATCAAATAACCATTGTTTCTCGCTTTGGCTTGATTA  
**AAGATCAGATAGATCATTAAACCAACTGAAAGTGCCTGGATTCTTGACTCCAAGATGAGCACCAAAAGAAAGGAT**  
 CGTGTATAATGGACTAAAGGCCGTGAGAACCAACCTGAAGTTCTCTACATAACCCCGAACAGGCCACCAAGTT  
 TTTCAAGACCTGCTGAAACCCCTCATAAACACAATAAGCTTGCCTACTTGCAGTCGACGAGGCCATTGTTAGCC  
 AATGGGGACATGACTTCGACCGGACTATCTAAAGCTGGGGAGCTGCGTTCAAATATTCCGACGTCTGTTGGCT  
 TTGACTGCCACAGCTCAAGGAAGTCAAGGAGGATATAAGCAGTTGCGCCTCCACCAACCCGTGGCTCAATTAG  
 CACTCTAGCTCAGGAAAACCTCTTCTATGACATTGTTACAAGAATTCTATCGAAGATGACTTCAGCACCTGGCAG  
 ATTTGCCAGACACTGCTGGCAATCGAAGGAGTTAAGGATACACCAAGCCACAAAGAGGCTGCCATCGTATAT  
 TGCGGACTCGAGACCAAGTGGAGCCATGGCGATAGGGTAACTAAACAGGAATCGGAGCAGTACGCTATCACGCTGG  
 TCTGAAAACAGGAGAACGTAAGTCCAGGAAGCGTGGATGCCGTTGATCAGCCATTATATGTCACCAAAATAGTT  
 TTGGAATGGGTGTTGACAAACCAAGTGTGCGCTCGTTATCCACTGGGATGTACCCCAAATGTGGCTGTTATTCAA  
 GAATCTGGTCCGCTGGACCGCGATGGCTTGCAAGTCATATTGAGATTAACTACGGACGGGAAGACGTCCGGAGTACG  
 TTTCTTCTCAAAACGATGCCACAGGCCAGGGCTGTTGACAAGAGCTGCTTACGGAGAGAGCTATCAAACAAAT  
 TTGAGAAGATCACGAACTCGCAGCGAACACCTGCAAGGCATAAAACTCTTCGGATTCTTGAGATCCAACCTCA  
 GATTGAGTGGCAATCGATGTCAGCGTCCCAAGAAAGCGAAAAGCAGTGGAGATTTCATCGATTATGTAT  
 GGATGATGCCCTTAAGTCCGACATTGCTACAGGATTGCGCAGATGTATGAGGGTGAAGTTAAAACCTCGTTATCT  
 TTTAGCTATTCTGATCTAAAGCCATATTAAACCGTTACTCGTAGAGTAAAGGCTATACTAGATTGATGAA  
 AGTATGTAACAGGTAGAAGGAAGCGTTATGTATGGATATTCTTACAGGATTACTATCCGAGTCGATCTGGACAT**CCG**  
**TTTGTCCGTCCATATGAACGTCAAGGTCTAAGGCTAGCGGATCTCTGGAAATGATTGTTATTAAATTATGTC**  
 AAAAGTATCTTATAGTCGATTAGGTCGAGTATAGCCTCTCTTGTGTTTATATCCTACTAGGCGGCCGACCAGGTATA  
 AACCGTGTGCCAGGAGTACGCAGCGGTGAATCTGGCTCAGATGACGATTGGCTCAGAGTCACTCATCAATGCCAA  
 GAGAGCCAAAAGGAATCACAGGATTTCATTAAGCAGCAGTTAATTCGAAAGCAAATTAGTCAGCTAGGCAATTG  
 AACAGGAAACAAATTGCTCAAATATCTCGCTTGAATGGCAGAACGCACTGAAAAGAAAATCGCAGGTCTCAGGCCACT  
 CATCGGGAAAAAATTGACAGCTTAATCGACGCACTCAAAGCCAATGTAGACAAGTGCACAGGATGAACCAGGGCAGCA  
 GCCAAAAAGTGTATTAACATGACTATGAGGCAATGTCAGTAAATATGGAATATGACGTGTTGCCAGAATAAGG  
 TGGCCAATATGACCGCCATGCCCTAGTTAAAGAGATATCCACAAATCAAGAAACTCACAGAACAGACTAAATTATTG  
 CTGGACTATATTCCGAAAGCTTAAAGGTGCTGGACAGGTGGAAGCGTGGCTACTTGAGCG  
 GAAACTCAAAGAAACTAGAAGAACAAAGGCCACAAAGCTTAAGGAAGTTCAAAGGCCAAAAGACAGAAAAGTT  
 AACAGGAAATAGCAAGCAGACTCAATTCCCTTTTAAAAAGAAATTAAAGAAGAACCAATAGACTGCCAATT  
 CAGGAAGAACGTGAAATATTAAAGTGAACAGATGGATTAGAAGAAAGCACACCTAAAGATATTGCTTCAATCATGT  
 TAAGGAGGAATCCGTGACCCAGAAAGTTCTCTGATCGTGGAGCTTGAACCTCCAGGCCACAGCATAATTGGAGAAA  
 GGAAATTAAATTGATGTCATGGTGAAGCTATGACACTCACCGCAGGAAAAGAGTTGCGATGAAACTCAACCC  
 AAAGAGAAGGAGTCAATTAAACAAAGTCCAGTATGCAAGCATTTTGGTTCTTAAAGCAGTCAGAGGAGTCTAA  
 CCAATCTTGAATGGTTAAAAGTCTGACAAATGCTGACAAATTCAAAACTCGAGGGGTACCTGAAAAAT

TACCTATTAAGGATGGTAAAAAGAGAGTTTAGAAAACAGGGCCGATTCTATTGGATTTACTTCAGCCAGAGAAATGTTA  
 GAGAGGAGCAAACAAAGGAAGGGGACGATAAGTCAAGCAACAAACTCGAAAACGTTAGGAAAAGAAATTAGAAAGTAGGGAAACCA  
 GCTGGAGAAAACAAACTAACGAAAGATGTTGGACAAAAAGACGTTAGGAAAAGAAATTAGAAAGTAGGGAAACCA  
 AAAGATCGCTACAAACAGAATTACAACATAAGCAAGAGGAAGGACCGTAAAGACAAAGCATCCCACAGGAAACGTCA  
 AAGGAATCGTTATCGAAGAAGGAAGGATCCGACTCCAGAGTTGCAAAGGATACAAACAGAAACAGCAGAAA  
 CCATAAAAACAGATGTCAGCAAAATGTTGTCAGTGCTCAATCCGTATTATAACGAAAATAGCCACCAAAGAGC  
 TTTCAAGCTTGGCAAATTGCTACTCAGCGATTTGATGGAAGCTTAGGAGATGGCAGTGCTGGAAAGTGCTAT  
 ATAAAGGAAACTTTCATGGCCTGCAGATGATAAACACAGATCAGGACATTGAAAATATTCTATAAAATCCAAGTGACT  
 AAAATCTCTAATGATTATCACGACATATTGTTGCCTCGTAAAGTTGCATCATTTTTTCGTTAGTTAC  
 TATTTTGATATAAAGCGATTAATTCAATAACATTAAATAAGACGTTTT

### 50. sbr

GATAGACCTGTCAGAAATTAAAGAAAACAATATCAACATCAAGCGAATATACTTCGCTCGTACATATTCTAGTTAGTT  
 TCGTAACTTTTCGCTTTGGCAATTAAATCAAATTTCCTCGAAGAATAACTTGTAAAAGGTGCAAACATCCGTCC  
 AGTTTCTGCCAACCATCGCCACTGGCTATCGAAAGAAGATCGCTAGTGAATTAAATTTCACAAAATTGCGTTGACTTT  
 GGCGTGGAAATTGCAAACACAGCAGCTACTTGTGCTATATTAGCAAATAGGTTAGGAAGTGTGAGAAAATT  
 TCCAGAAGTGGCAGCAGTTGTGAATGCAAAACATCGCACATAGTTTTGCCCTCAAGACAGTTCTCAGT  
 AATAACAACAGCACCCACACAAACACACCACACTCAGCTAAAACGTGCTGAAATTGCGATATCTTTGTCATA  
 CGTTTGAGTGAATATTAGTAAGATGCCAACAGCGGGTGGCAGTAGCCAGCGGTACAACAACACGTTGAAATGG  
 CGGCGGACGTTACAACGCTCCCGAGGATTTGATGATTGATGTGGAGGATGCCAGCGACGCAAGGATCGAAACAGC  
 GGCAGCTAGCTTAAGCCCTCCCAATGTCTACATAACAAAAGGACATCAAGCTGCGACCCGAAGATTGCGTCGATGG  
 GACGAGGATGATGACATGAGCAGCATGACCACGGCGTTAAGGATAGACCCACCTCCGACGTCGGGATGCCATTCC  
 GCGCGCAAGTTCGGCAAACACTGATGCCAACAGCTTGGTGGTACCGTACAGTTACAAAACGCCAGATATACGAAA  
 AGGAAACACTCTTGAGTGTCTATTGGCAGCGATGTCGCCACATGTCCTTATTCTCAATATTGGCAGTGGAGCGAAAC  
 TCGTAATCTCTTACGGACGACTACGAGGCAACGATTCAACATCTGGCAAGAATGCCATCTCCAGATGG  
 CTATCGTCTGATGCCACGAGTACGAGCGGTACCGACTAGTGGCCATCGACGATGCCCTCAAGGAGAAGATGAAGGTCA  
 CAATGGCAACCGTTACAATATTCAAACCAAGGCGCTGGATCTTCCGTTTACGAGATCCGGATCTTAAGCAAGTT  
 TTCTGCCACTTTGTCAGAATGTGATGGCGCTGCCATTGACATTATGTGCGACAATATAACCGATTGGAGGCACT  
 TAACCTGAATGACAACCTCATTAGCAGCATGGAGGCGTTAAGGGTGTGGAGAAACGCTTACCGAACACTCAAGATTCTCT  
 ATTGGGGGATAACAAGATACCATTGGCCACCTTGTAGTGTCTCGAACCTGTCCATCTGGAAACTCGTTTAAAG  
 AACAACTCCCTGCCGTTCCGCTACAAGGATTCCCAGCAGTTATCAGCGAAGTACGTCGCAAGTTCCAAACTGGTAA  
 GTTGGACGGAGAGACCCGGAGCCGAAATCACATTGATCTACCGAGCAGGGACGTCTCGAAACGAAGGCATCCT  
 ATCTGTGCGACGTCGCTGGTGGCGAGGTGGTGGCCAGTCCTGGACCATTTGACTCGGGCAATCGG  
 CAAGCTCTGCTAGATGCCAACCATGAGAAAGCGATGCTCTCCATATCAATGCCCTCGGCCAGTCAGGGGGCAGATTGAA  
 CAGTTCTGGAAGTTCAATCGAACCTCCGGCTTAAACGGCGAAGAGAATCGCACCCGAAACTTGAAGTACGGAC  
 GCCTGGCATGTTCCACATTGGATGAATGCCAAAACGCGACGACGACGACGACGACCTCACCCTGACCTGACCATC  
 TACAATACTCAATGATGTTTACCGTGACGGGATTATTCAAAGAGCTGAACGACGAGACCAACAATCCGCTCCAT  
 GGAATTATATGACGTTGCCACTTGCCCCCACCTACGTGGTGGGCCACAGAATAATGGCTTTGTATCCGCAACGAGA  
 CGATCTCATACAAACGCTACGACGAGCAGGTGCGAGAGTTCAAGCGATCGCAGCACCAGCCTGCCGGAGCTATG  
 CCCTCCACTTCAGTGCAGTGACCATGCTCAGGCCGGGCAGCGGGGCTCGCAGGGTCTGTAATGCCGTTGGCGT  
 GGCCACTGGACCGGGTGGCTATACTATCAGGAGATCCGTTGGCGGCCACCGCACCAGTTAACAGCGGAGTGCCTGCCATAT  
 CGACAAACAGCAGTGGCACCTGGCGCCAGGATGAGAGCACTAAATGCAATGATTGAAGCCATGAGCGCCAAAGCCAA  
 ATGAATGTGATCTGGAGTCGAAATGCCCTGGAGGAAACGAATTGGACTTTAACATGCCCTTGTGTTGAGAAACT  
 ATTCAAGGAAAACAAACCGCTGAGGCTTTATGAAGTAAATCGCATAGGAGTTCCGTAGGACAGAGCCGCGTGC  
 ACATCCACATACTGAATGCTTTTTTTGGTTTGTAAATTAAATTAAATTAGAGAAACCTCTATATA  
 ATAATAATAATTAAATTATAAGCTGCAAGTTGTGTCACATTGGCAGTAGCAATTATTATCCGACTGCGGC  
 AATGTGCATCAACGATCACAGTCTCGATAGATTAGTTAGCTCTTTA**AGT****TCCGTCGCCAGATCCGC****TGGT**CTACA  
 TTGAGGCCAGGACGAGTCGCAACGGTAGTCCTTAAAGTTAGGTTAGATTCTAAGCAAACACCTTCAAA  
 CACACACAACACGACAACACTATTAAACGTAATGCAACATGTCGAATCGAAAGAAACCCAAATTAAATTAA  
 ATACGACGATACCGAAACCAAGGCGATGGTCAAAGCATGGATCGCCTAACATTCTATATTCCCGCTTCCAGAT  
 CCTCGACTCGCCTACATTGTATAACAAATACCATAGAATAAAAGAAACATTGACCACTGTAAGGAAATTGTAACG  
 ACTCGGAAACCAAATTACCTTATTCTTAATAGAAAAAAATTTCGATTACAATACGCTTAGCCGTAATTCAA  
 TTGAATTAAAGTGTAAAGATATAAAATTATACATTAAGACAAAAGTGTAGATTCTAAATAATTTCAGAATAG

### 51. SPoCk

GCACCTCCATTGAAAGCCTGACGATGTGCCATCTCCGGCCGTTGACCGGTGAATCGAGGCGAAAATCGATT  
 GTGCTCGTGTGATTATTTGTTGATTCGGCCTCCGCTCTCGTAATCCTCGCGCGCCGCGCCACATGACTCAT  
 GCCAGGCAGGAACA**ATGAAAATCACAATAAGTACGTCAAATTGCCAAGACCTCGATCTGAGTTCCACAGTGTATTG**  
**GACCTGCTGCCGATACGGCAGCCGACATCGACGCTAGAACACTAGACCCGGCTGGGACACGGATGTGGCGCGCTCAGTGT**  
**GAACCGTCCCCGGCTCTGGGCTGGACGAGGACGACGACGAGGTCATTATCGCCCCCGCGGGAGGCTTCGAC**  
**TAGCTCCCACGGCGGGAGGCAGTGGCGGACACAGGCAGCATCATCGCAGCACCAGCAGTGCAGCAGCTAAAGAAC**  
**CAGCTGGAGCAGAACGAGATTCTGCCAAGAGCAAGATCTGCGGACTAGGCAGCGACAGCATCGCAGACGGCCTTAC**  
**CCTCACACCTGAAATGCTGTCCACCTCGGAATCATCGACCCACAGTGCCTCCGAGGTCGCCGACGACTGCAGGTGCG**  
**ACGTGCGTACCGCCTCAAATGGACAGAGGCCAGTACGTGCGCAAGATCATCGGGCATAACGAGCTGCTCGTGGCG**

GAGGATCCGACATGGAAAAAGTACATTGAGCAGTTAGGAACCCGCTAATTCTGCTGCTTCTGGTCTGCTCGGTGTC  
 CGTAATCATGAAGCAGTTGACGACGCCGTGAGCATAACCATTGCCATCTAAATTGTGGTCACGGTGGCCTTATCCAGG  
 AGTACCGCTCCGAGAAGAGTCTGGAGGAGCTTAAGAAGCTGGTCCCCCTGAATGCCACTGCCTGCGGGAAAGGTGCTA  
 GACACATTCCCTGGCACCGCACTGGTCCCAGGACATAGTGCACCTCAACGTGGGCGACCGGGTGCGGCCACGTGCG  
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 AACGGAAAGGCATAGTTGTCAGCACTGGGAGCGCAGTGGTCTCAAATGATGCAGGCAGAGGAGGC  
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 AAAGCTGCCACTGTGAGACTTGGCTGCCAACGTTATCTGCTCTGATAAGACCGAACCTTGACCAAGAACGAGA  
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 CGCTTACATACAAAATGGCACCCCTCTAGGACAACCCACTGAAGGGCCCTTGTGGCGTAGCCATGAAGAACGGAATGT  
 ACGCCACAGCTGAGAACTACGTTGCATCCAGGAGTATCCCTCAGCTCGGAGCAGAAGATGATGGCTGAAAGTCATC  
 CACAAGTACAACAACAAGGAAGAGATTTCTCGCCAAGGGGCTCTAGAGACCCCTGTTACCGCAGTGCACCAAGTA  
 TCAGTTGGTACCCAGACGGTACCAACTACCAAGCAGAAGCAGGAGGAGGAGTCCCTAGCCGAGGGCGTACGAGATCGGCCA  
 AGGGCCTTCGCGTGTGGCCCTGGCAAAGGGCGGTCCATGCAAGATCTGATCTACTGCCACTAGTCGGCATCACTGAC  
 CCACCAAGGCCCTTGTGAGACTTATTGAAATGTTGATGCAGAGCGGAGTGCCTGTTAAATGGTACTGGAGATGC  
 CCAGGAAACGGCCTTGGCATTGCAATCTCATCGGTATCGATAACATCCATCACCAGCCTTCCGGTCAGGAAATGG  
 ATCAAATGAACGAGCACCAACTGGACAAGGTAGCCAACACGTTGAGCGTATTCTACCGGTATGCCACGCCACAAACTG  
 GAGATAGTCAAGTCCTGAGCAGTGGCAACATAGTGGCATGACGGCGACGGGGTAACGACGGAGTGGCCCTGAA  
 GAAGCGGACATCGGCATAGCCATGGAAAGAACGGGACAGATGTGTAAGGAGGGCGTGAATGATCCTGGTCATG  
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 CTTAGCACATCAATAGCTGCCCTGGCCCTGATTGCCACTCTGATGGACATTGCTAACCCGCTAAATGCATGCA  
 GATTTGTGGATCAACATCATAATGGACGGTCCGCCGCACAGTCTGTTGCGACAGTGGCTGGAGCCGTCGACCAAGTGC  
 AACAGAAACACGGAACGAAACAGCCAATGATCACAAAGTCAGTGGTGGTGAACGTTCTGCTTAGTGCAGTATAATT  
 GTACTGGCACCGCTGTTGCTCCAGCGGAAAGCTCTACGCCGTATGACATATTCTGGTCTCCCTAACCTCATCTGTGCTGGTGT  
 CAAATGGTTTCCAGACCGAAGCTCTACGCCGTATGACATATTCTGGTCTCCCTAACCTCATCTGTGCTGGTGT  
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 ATAAAAACAAATTGACGAACTCATCCAAATCAACTTTAAAGGAGAGTTCAGATCAACAAAGATTTGTATGTAAT  
 CAAAATGCTATAATTCAGCATCTTGTGGCGTTGATGTGATGTTGATAATTAAAGTCATGATTAATGGCTTAC  
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 CTCATATACTAAATAAAATCATAGGAAAGCATGGTTCTGTACTATTCTGTGTTGTTAAATTGAA  
 CTTTGATGCAATAGATTACTAACGCAAAGAAATTACAACACTAACAGTCTTCCAGATTGTTAGTCAGTCAAAGTATACATCT  
 CAATAAAAATTAATGAATTGAAAGAAATTACAACACTAACAGTCTTCCAGATTGTTAGTCAGTCAAAGTATACATCT  
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 GTTTTATACCGTTATTAAAAAGTAAAAGGAAATTCTGTATTGTTGAAATGTAACCCCTATCAAAGAATATATATTG  
 TTGCTCA**CTCTGTCGTC**GTATGTCATCCGTATAATGTAATGAGAATCTGGCAGCTATAAGCTAGAGGTTAGGCATG  
 TAATTTCAGACCACCAAGCGGGCCATAACTGCCACGGCTCATTTCTAGAAATGATTATATTGTTAAGTTA  
 TAAGTCTTGTCAATTCCAAATGATATATCCAAGTTTTTTAATTGATATTGTTTATAAATTGATAAA  
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 CGTCTTCACTTTTATTGAAATGAGCTGTATTCAACATTTCACAAATTCTGTTAGTCAGTAACTGGTACCGTCAAC  
 CACCTGAGATCGCATTAAAGGACCAATGTTAAATACTCATTGATCACACACATTAAGATCAGCTTACGCTCAAC  
 AAAAACGCACTCATATCATGGAAAAAAAGGAAATTGAAAGAACATTTAAAGGAGGTTACTAACAGGAAACATGGAA  
 AAGTAATTGATGAGAGCTGTTCAACAGATCAAGCAATTGAGGTTCTAAATGATATCAGCGCATAAAAGCTACTAAA  
 GTACATAAGCAGAAAACCAGAGACACACAGGCCACACAGTGGCTCATCCACAATGGTAACAATAATTGAGA  
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 TAGCCAAAGCAGGTATACATTGAAATTAAACTGAACTCTACTCAATGTTATGTAAGCAAGTCCGAAATCGACT  
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 TGTCTAATGAGCTGGCTAGTTAAGAGCTGGAAAAATGTTGCTCAAATTGACCTTACTAAATGTTAGTTG  
 GTGCAAAGAAATGTCACAATTGATGACAATGGTACGAGTTCTAAAATCCATTGTTAAACGCTTACCTATAAAATT  
 ACTATAACTATCTGTGCTATTAAACTAACGTTAAAGTATAAGATAAAACCAAGAGTAATTCAAACATATTGAT  
 ACTTCGAGAATAAGAACCATGAGAAATTGGCATAAAAGTAAATTGAAACCATCATGTTTGTGTTACTAAATCTAC  
 CTACAAGTGTATATCATTGAAATATTGGGACAAGATCTTAGGCATAAAATGAAATTGCAATTGGCAACC

ATACATTAAATATTAATCTCGGAATATCGTG

### 52. ttk

AGTTCTGCGCTGTGATCAAAGCTTCGGTAAGGCTTCGGCTCGGCTTGTGATCCACCTTAGCGGCTCTGTGGTGC  
TCGTTCTGCCTCTCTCTCTCGGTTGGGCTCAGCTCAGGGACAACAAATCGGTAACCAACAAAGAGAAGAACAGAAACAA  
GTCGACATCCCTGCCGTTCAGTCAGGGAGGTCGCGCGGCTTCCTCAGCTGATTCTTGCCTTGCCT  
TTCGAACCCGTTCCGGTCCGTTGCCATTACGTGTAACCAACAACAAAGAGAAGAACAGAAACAA  
CAACAAGTACTTCGAGTGTGCAAGCCTCTCCGCCAAGAAACCGCAAAACGGATTACCAAGAATATTGAAAATT  
GCACGCCCTGGAATATATGTTTAGCAGCTGTCAGGCCCAAGGGAGTAAGAATCCGAACCCAGGACAACGAAAGGACCTT  
TGATGCCCACTTGCATGAGGACATAGATCGCATCGCACACACTCGGACATCGCTAGTGCAGCAACCACAGCCTCCA  
ATGAAGATGGCATCTCAACGCTCTGCCTGCGTGGAAACAACCACAGAGCAACCTCTGTCGCTCTCGACCAAGCTGTT  
GCACCGAGAAACCTTCACAGATGTGACGCTGGCGTCAAGGGCAACACCTGAAGGCACACAAGATGGTGTATCCGCCT  
GCAGTCCTACTTTAATACCCCTTTGTAAGTCATCCGAAAGCATCCGATTGTCATACTTAAGGATGTGCCCTACTCG  
GACATGAAGTCGTTGCTAGACTTATGTACAGGGCGAGGTCTCAGTGGACCAAGGAGCAGTCAGTCATTCTGCGGT  
GGCCGAGAGCCTGCGCATCAAGGGCTCACCGAGGTCAACGACAGACAAGCCCTGCCGGCAGCAGCTGCAGGAGCGG  
GTGCGACGGGCTCTGAGAGCACAGCCACTACACCCAGCTGCAGCGCATCCAACCGTATTGGTGCCCAAGCGGAATCGC  
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CTCGCCCTGATGCCAAGCAGAGGGCAGACCCCGAAGCTATCTGGCAGCTCGAATGGCACGGGAACGACTACG  
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CCACTCCAGAGCTCTCCAGCGGCTTCGGCTCTTCGACGACAATCTCAGCAACGGCTCCAGGGGGCAGCAGCACT  
GGTCCTCGGAAACGATTCCCTGCTGGAGATCAGCGATGAAAGGGAGTCCGCCCCGGTTCACCTGCCTACTATTTGGG  
CCTGAAGATACCGGCCATTAACACTACGACGCGTGCAGCAAGGATCCCCCAGACGCCATCGAAATCCAAGCGAAGA  
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CAACAGCAGCAGCAGGAGTCTCCCTGAGTGCCTGGCAGTTATGGATGACGACCTGGAGCTGCTGCTCTTAATGATCA  
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GACACACGAGAACTTACGCCGCCAGGACATTGGCGCCTCAGCCCCCTGAAATACTTGTACAGTGCAGCGGCTGCAG  
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GCCTCTTGACAGCGGGACAGTCGTTATCTGCCGAGTGAAGATGGAGCATGCGAAGACCAGGATGCCAAC  
GGAGGGGAAGTGCAGCGATGGTGGCTATGACGCCCTCAATCCAGCAGCAGCGCCATGTTGAGCCTGCAACAAAG  
ACGTAATCATCAAGGATGAGATAACAGATCTCCGTCGCCCTGCCACCCGCCGCTTCTGCGCTGTGGCGAGGGC  
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GCACAACACGTTACCTAACGAGCTGAGACTGAGATTTAGTTAATGAAACCGTAAGCAATATTCTAACAC  
CGCAACACACCAACTCAAACGATTGTTAATGTAATGAAACCGTAAGCAATATTCTAACAC  
**CTTGTCCGTCCAG**  
**AACGCC**TTTTAAGCAACACTAGTATTAAATTACCTAGGACATAAGTTTGCCATTGAGCAACGCAATTGAGCAGT  
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GCAAGCAACCTTAACTGAGCTGAACTGAGATTTGTTAAACCTTTATATAGACCTTTGTTTTTATTTTG  
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CAGCGTACAACAAAACGGACTTCCAGTGAAGAGACAACGATATTATACGGAATCCAATCTGGTATTATGATTATGA  
TTTTACAGTTACTTAACGCAACAAACGACTTAAGACTTGTGCGCTCAATGACTAGAAATTATAGTTGCGATT  
GATAACTATTGAAAAACCAACGAGTTTAATAAGAAAATTACTGATT

### 53. yrt

GTCGTGACCACATCTGGTACACTGCTCGATAAGATCAACAAACTACTGCGGCATTCTACATTCTACTCCTAAAAAA  
TTCCGAAAGTGCACCTAAATAGCCAAAATAACAAACATAATAGTGACGGGACGATGCGACGCCAACAAATA  
AAAGTGAAAACAAGTGATTGAATAACCCGGAAGTCGCTCGCGAACGCGGAACTAACGTGTTGGCTGGCTTT  
CTCTCCTCTTTCGCCAACCAACAAACAGGCTAACAGTGATTTTTGCGGGGGAGGGGGTGCAGGCCAAC  
CGTAGTAAAGAAACGCCGAAAGGAGCGCCACTTGGCAGCAACCTTAGCTCAGCTCCCGAACCAAAGTGAAGAA  
AAAAACAAAAAAACTAAAGGAAACTTAAGTAAACGGGGCGTCTATCGCTTGCAAGTGTGTTGGTAGTCAGTGTACG  
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GCCGAGGTGAGCACACAGGTGAATTATCGGTGCAATTGGCTGCCCTGGCAACAAATTCCATAAAACACACATAGAAAAC  
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 AAGCATACAGCAACGATGTTGAAGCATTAAATGTATAAAATATTCAATAAAAGTCATTTATGCATCATGTATACAGT  
 ACAACAT

#### 54. Gli

AGCTTGTGTTGGTTAGTGAATTGTTACCTTTCGGGGCGTGGCAAGAAAGTTATTGAGTGTAGAAC  
 GCGTAAAGTGAACACTAAATATCATC**ATGATGCACAAATTGAAATATCGCATAAAATTAAATGGCTTTAGCCCTTC**  
**TTGTGCTGATCGGCACTTGTGTTATTCA**GACAAGGGACAAACAGAGATCCAGATTTCATTCTCGGCCAGGGCTTGAC  
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 TGGCTATGGAGCCAACATCCAGGAGAAATCCACCGGGACAATATCCACAGGAATGCCAATGAAGATCGCTTCGAT  
 TTGACCGAACGATCCGAATGCGAGAACCCAGTTCCGGAGTGTGGCCGGATGGCGAGAGGATTGTCAGGGCAAGCAG  
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## 55. Nrx-IV

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## 56. Lac

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 GTCATCAAGGATTGCGCTTCGCTCGGTAACGATCCAACTCGTCCACCTACAAGCTGAGATCAAGGACATTCAAGG  
 GACGGACGCGGCACCTACACCTGCCAGGTGGAGTTCGACCCAGTGTACCCGTGCCGCTCTGTTGGACAGGCTC  
 TGCAGTACGACATGGACTGGAGTGCCACATTGAGGCCATCCGCCACGGCATTGTGTTGACAGAACATCCAG  
 CTGGCCAACACCAGCACTACAGCATTCGCACTGCCACCGCCGAGGAGTACCGACTCGACGCTCCGTGATCAC  
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 TCGAGACGATCATTGGCTGCCCCACGGGCTGTGGACAGGCCGAGGAGTGTG  
**TCGCTTGTGGCATCTGGCGCTGCTTCTGCCAGATAAGCCAATGGGCCACGGCCGCTGACTCCAAGCGCTT**  
 CAGGTCCAATCCATCAACCAGCCCTGAATGCAAAACATATATAAAATAGATATATGAAAACAAACATCGTGTAA  
 GCGCAATAGATGCTCAAACGATTGCGACACACCAACCATCCAGTTCTGAGCTTCTGAGCTTCTGCTTCCGT  
 TTCCGTATAGTCCAGCCAATTGGTACGTTAATTATGTTAAGTATGATAAGTGAAGGGCAACGATAAA  
 CTAGACTATAATTATTAGCACCTAGAGCCGAACTCAAACGAAATTAGACTCCAGGCCAGCGTAAAGCATGCA  
**A CCTTGATC CGTCC****GTTTGCTT**CTGATCTGGCACTGTATATGTTAAATTAACTAAGCAAGATACTCACA  
 AGCATTGCGATTGCTCGACAATTGAGAGGATATTATTTATTAAATAAAAGTATTTCGTTACGCAAAC  
 CTATCAGCTCGACACACTATTCCCTAAATAATCTCAACCCCCATAATCGAGGCCAAATCCATATTGAA  
 ACAAAAGTTGTTGACATTAGCTCATAGCACAATGAGAAATTCTTGAATAACAAGACATTTCACCAAAATT  
 GTGTGGGTGTGAGTAAATCAATTGACTTTGTAACAAAAAAAGACAAACGAAAAACTAAAATTAAACAAAAAA  
 AAATGAATAA

## 57. santa-maria

ACAGTCGCATCGAAGGCCGCCGGCGAAGCGGTACCATTTACGGCATTTCCAGGGACTCGTGTCCACTTTCCGGCCTC  
 TGTGCAAAGAAATCATCCACGTGCTCAAGCGTCTGTCACATCCTATCGAATTTCATTCGCAAGGAAAATCAA

AACGTAAAAAAAAACCGAGGAGCCTGTGATTGTTCTATTTAGCAAATAAAAGTCACCACATTAGGTATCCACACA  
 CCATCGCCCAGCGTGAGCCATGCCAACACAAAACAGCGCAATGTGGGCCAAAAAAGTAATCGCAAATTAAATTATCGGCA  
 TTTCCGGCTTGCCTGGACTATTGGCATATTGTGCGGCATGTTCTGGGTGACCTGTCGATTGGATCATGCACAAG  
 GAAATGGCTCTGGCCCCGACACACGGTGTCTATGAAATTGGAAGAGTCCGCAATTGATCTCAGTCAGTCAGGCACATCTACCT  
 GTACAACTGGACGAATCCGGAGGATTCGGCAACCTTCACAAAGCCCATTCTGGAGCAGGTGGGTCTTATCGTTTA  
 TCGAGCGACCCGATAAGTGGACATTCACTGGCATCCAGAAAATGCATCCGTCACTTATCGCAGGCGCAGCTTGTCTAC  
 TTTGATGCGGCGGGCAGTAATGGCAGCCTGGACGATGAGATTACACGCTTAATGCAGTGGCTCTGTCAGGCCAC  
 AGCCAAATATTGGCCACCAGTTAAGCGCTCCCTGTCATGTGGACTAAAGATGTACGGTGGCAGAGATGTCTGCCAAA  
 AATCGATCGATGAGCTGCTCTCACCGTTACAATGACGCAATGATCGACGTGCCATGCCATGCCATTGGTGT  
 GAGGTGAAGGTTCCATTGATAAGTCGGTTGTTCTATACGCCAATGGAAGTGGCAGCTTACCGGAGTTTAATGT  
 CTTCACTGGCGCCGATCAACTGGCAAACACTGGTCAGATGCACTCGGAAATTACCAAGGAGAACACTGGATTCTTGATT  
 CCTATTGCGGTATGACCAATGGATCCGCCGGGGAGTTCAGCCACAGCATTGAAAGCCGGCAGATAGTGGTGGCTATT  
 ACGCCCGATATGTGCGGAACCTATTCCGCTGGATTATGTGAAACTGTGGATATTGAAAGGACTGGAAGGTTATAAATTCTC  
 AGGGGACCAAGATCTGTAGATAATGGCACCCAGTATCTGAGAACCTTGTGTTGTGGCCAATGTGTTCCGTCGG  
 GTGTGATGAACATCAGTCCCTGCGTTGGATCTCTGTTTATGCTCTACACTTTCAACGCCGATCCTAC  
 TACCCAGATCAGGTAGAGGGCTTGAGCTCTAACCGAGAACGATCACGAGTTTATGGTTGTGCAGCCAAGTACTGGTAT  
 TCCCCTGGAGGTGGCTGCCGATTTCAAGTGAATATGCTCGTGGAGCCAATCCAAGGCATTAGCTTATACACTGGAATAC  
 CAAGGATATTTCGACTCGTATGGTCGAACAGAAAGTAAGAACATCACTCCAGATATGGCAGATCAACTGAAGGTGCTG  
 CCCATTGTCATGCTCTCCGGACATATATTGCCCGAATTGCTGATCGTAGGCATAACTCTACTCTGCTGGACTCCGGT  
 TCAGATTCTATTGGCTTCTGCGAAATCGTAGATACGATCTGAGGACCAAAACCAAGAACATGGCAGTATAAGAGTC  
 GCTCTCAATTCTCCAGTGTGAGGAACCTGAAAGTCCAAGGCCCTCCTGGTGTGCGAAAAGAGTGTCAAAGGATCTCG  
 GACAGTTCTCCGCTCTCGAAAAAGGCCGTAAGCCAACCAATTAAATCCCAGACGGGTGAAAGTGTGGCACTGCA  
 CACAGCCATCAGCGACAATAAGCAGGATTGAACAGAACGCCACATTCAAG **CCATTAGATAGTTAAGCCGTCC** TAGTGC  
 TTAGTCGTAGCTCATAGTTAGTGAACACTCTAAACACCCAACCTGAAATTGCAAGCATCCCCTGCCTATGCCAAAGA  
 TTAGATAGCCTTCGAAGTCGAAACGTCTGGTGACTTCGCTTAGCTAACGAATACTCTTTTACATTCTTAGTTAG  
 ATATGTTTGATTAAATGATTCTGACCCATGGCTTGTCAAATAAAATCTCAACTAC

### miR-184 MREs in long non-coding RNAs

#### 58. lincRNA.924

CGCCATCATGCGTTGAAATAGTGAGGAGCTTGCCTTGCGTTGAGGCCTACCTTCAAGCCAGTGTGCGTTATTAAAA  
 CACAGCGTCAATTGGATCGCTTAATTAGCCCCGTTGCTCTCGTCCGGACCGCAATCAATTGTTACGTGGTC  
 ACTACATTCAAGCAGAAACTGCAAGTGCAGAACAGAAACTGGAGTCCCTGACCGATCGTTCAACGTGACTATCTCAATTAG  
 AGGTGATTGGAATGGCGACTCTCCGATCTGCTCCCTCGTGAATTGGAAATC **CCGTGTTGTGGACCGTCC**AAGA  
 ACCCTACAAGATTGAAACACATCCAAAAGAAATTGCAACATAACACCTGCTATGCTAACCAAGAGTAATGACAAA  
 CGCCAGAAATCGGCTTACGCAATGTATGATGAATGGGGACGTACCTATAAGATTCTATCTCAAACAAATATAAAATAA  
 AAACATTAGACATGTACCTACATTATAAAATAAAATTGTTCCGATGCCCTACAATAGTTTC

#### 59. lincRNA.218

GTTTAATTAAATCGCTTATGTTAAAGCCAAGTATTAAAAATATAAAAAAAATTAAACTAAATTAAAAA  
 TCAAGCACCGGATTTGCTTGAGTGCTGACCTTGTGCGATCTACCGCTGAGTGTGGCCGGGCTTGATTGCATGG  
 CCTCGTCTCGTCTGGTTGGCTTAGTCTCATCTCCATGCCGTCATGGCATCGTCCATCTGGCCGAACAT  
 TTTGTTGGCTACTCAGTTCTGATTAACAATCAATTGCGTGCGCCATCTACTGCTGCTCGATGAATGAATC  
 GACTTCGGCCTCGCATCGTCACTAAATTGAGCACAGCACCAACCGCAGCCATAAGGTTATCGAGGGAGGAAACCATC  
 GGGCCACTTAGTCTGCCAGGCTCCAAAGTGCATCAATC **AACGCCGAAAGTCGTCCAAAGGATAAGCAAATGACG**  
 GGATGCAAGTGAGTCCTCAAAGGGACGACAGGTGGCGAGGTGAAGGGCAAGACCCACCTCAACCCAAATG  
 CCGCTTATTGAGGCCGGGATGGCGAAGCCCAAAGTGATTAAATGGGACCCATACACCAAAACTGAA  
 ACACCGAATCACCAGCTGAAACCATGAATCCGGAGTCACAGGGCGTCAAAGCTCTGTTGATGTAGCTCTCGGG  
 ATGTTGTCATGGGAGCTGGCATACTGGGTGAGCTGCCGGCATTAACACGTTGCTAAAAAGAAAACAA  
 AGTGCAGGCCACATGGTAACTTTCCATGCGATTGGAGAACACAGTTAATTGAAAAGAGTAGCAAAGACAGAAAAAGG  
 GATATGTTAAATTACGGCTACGATAACCCAGATTAG

#### 60. XLOC\_001425

TTATATTTATTTAATTTCATTTTACTTCTTTCCAGTGCCTGCCCTTCGTTGGATCGTTTCACTT  
 GCCCAAACCGCAAGCCGGAAACTTCGACCTGTGCGAATGACCAAAAGCAGAGGCAAGGAGCCTGCAAGTGTAGACC  
 ACAACTCCTCGTCGAGGACGATGTGAAAAGCCGATGTCTAACCTTTTTGCTTAATTGCGGTGCCAGAT**CATAGCCATCCGTCCGTGTTGG**  
**TAGCCATCCGTCCGTGTTGG**TGAAACAAAGTAAAACGACAGGAGTCCTGCAATCCGGATCCTGTGAATTGAGGCCGA  
 GTCAATGGTTGATATCGCTTGCATAATTGTAACGTAACAGTCGCGCATGTTCAATTAGGGACACTTTG  
 CGACCTTAACCTCTATCCATGGACTGAACACTACGACGAGCTACATGTTCTGTCCATATGCCAAATAATGCCGCA  
 CATTAGTGCCTCGGTGAC

#### 61. lincRNA.678

GCCGTACATGATAGTACAATACCATAGATCGCATATCTACTTACAAGAAACTAATAAGACTTTGATTATTATCTCT  
 TACAACCTTCCTACTATAGCTAAAGGTTAACCTTAAAGTAAGGATGGTGTGCAATACGCATATAAAATAGGGCAGCT  
 GGCCAGTGTACAAGGAGCCCTCTCGAGGGTCCGCTGATGAGGGTCCGGAATGAGCTGTCGGCTGGAAGGCCAGGCATC  
 AGGCTCGTCGGTTGGGTTCTGTTAAATGCAAAGTAGTGTGACACATGCAAAGTTTCAGGTTCCAATGGTCGGGGAG  
 GTGGCAAACAATGTGACCGCATTAGTCACAGTCACACTCGGGCACACAGACGTTACATTAAATTGAGAGATTCCGAG  
 CCGAGCCGCGTGCCTCACGTGAAGTGAGTGAGGTTGGAAGTGAAACGGTTAGTGCAGAAGAGGCAACTCG  
 GGGAGGACTCCTCGTAAGTGGATGTCGGTTGCTGGTTGTCAGTCAGCAGGGCATAATTGAGTTGTTTC  
 AAATGCTTTTACTAAGCACAACCATAACGCAAATACACACTGGAAAACGATTACGAAAGCTGGACATTGCAAGTT  
 GGGCTATGGCATTACATGTACTGCGTATTAGTCATATTAGTCGCTCCTTAATGAGTTCATCAGAGTCGCTTAA  
 GTTACCAAGTAATAATCTATACTATAAACACAGTAACCTGCCTATCAAATGTTCAACATACTATGAGATTACA  
 TACATGTACAATATGAAATTGATCAATTGAGACAGGTTGACACACTGCTTGTACTCTCCGCTGGTTCTCCCAG  
 TGTATTCTGCTGCTCTGATGTTCTCTGCTGCTGGCATCTCTGGTCTTGTGGTCGTTGCGTTGTCAGTCAGTGGTGGC  
 TCTGCTCTGGCACTTTGACATCAGCAACGTTGCCGTTGTCGTTGTCAGTCAGTGGTGGCAGTCAGTGGTGGCAGTC  
 CAACGGAACAAACACAAATACACACACTCGAACATCCGCTCACTACGGGTGATGGTGGAGCGAGACGGGGTAGGGAAA  
 CACGCATGTCGCCGATAAGTGTGTTGTCGTGCCATATAAAATTACTTCGACATGCAACTGCAACAGAAACAC  
 GACTGGGCCTCGCTCTCAACTCACATCGACATCTGCATTCTGGCAGGAAGAGCCGCTTGCATCCGATGGATG  
 CGCCTGACCGAATGACAGATACGCTGTCATTGAGTTGTTCCACACTCAGAAAAAGATTGTTGTTA**ATCTGTCC**  
**TCA**  
**TTATAAAACAGG**TTCCCCACTTAGTGCTCGATTGACGATTGATATCACACACATTTCCTAGAAAATTTTA  
 ATTCAATTATAATGTTATCGATTAAAGGAAAATATTCCACCGCGTCATAGTTGACCACTTCTGGGAATTTCAGTT  
 TTATGAATTGTTGCTCCTAT

## 62. lncRNA.903

CACGTACACAAATCGCGAGATCGAGCATTGCATATAGTATAGTGTAGTATAGTATCGCATTGACTTGGTAGACGGG  
 CAGGAATAAAATTAAATTCTCTCGGCTGGGAAAAAGCCAACCGTGGAAATGCAAGCCCTAATCGAAGG  
 CGAAGTGGGTTCCCCCGGATTAGCGATCTCTCCGCTTTCACCGTTCCCCAGCGAGGGCAGAAATGAAGCGGCCGTG  
 TGCCACACTTTCCACCGAAAGTCGAAATGCGTTCCAGTTATTTCAAAATGTTCAATTCAATAATTAAATTGTC  
 CTCAACGCACGACGACGTCAGGAACCTATAAGGCCCTGCTCCCTACGATCCCCAACCCACAAACCGATCCGTT  
 CGATCCCATCCCATCCGATCCGAGATGAACTCAACCGATCTGAACTGAGCGTTGCCAATTAGTCGAGATGTTGAC  
 AAATGCCAGACAAATGCAGCAGTGCCGGCGCATATAAAATTCAAGCTCATCGCGTCCGATGGGCCATTGTCAG  
 ATCATGATCTAATGGGTTGATCTGTTAAACACCGGAAACAAAAGGGTTACATCTGCTCAAAGACCTCTTTT  
 AGTCATTACATTATAATCTAAATAGAAACGTAACAAATTGAAAAAATTGCTGAACAGTGCAGATTGTATCTGCG  
 ATCGGAATGCGCTCGATTATTGTTGACACGGGTTCAAGGAGCGTGCCAGTTATATCCATTGCTAGATGCA  
 TTCGTTTGACATCTAAAGCAACATAACCGCGAATCGGGCAGGGAAAACACCTTTGTCGGACACCTATAAAATGT  
 CCGATCTTACGATCGTTAGCCCCACCACGTCGAATGGCAGACCGGGTTATATGGTCGGCGACTCCACGAATTCA  
 CGCCGAAATTATGAGGTCGGCCTTTGAGGCCACAACATGTTGCAAACATAAAATATGAAAATTCACTAAAAACG  
 TAAACAACACGACTCGT**CGTTC**  
**TCCGT**  
**TCTCTGGGCT**CCCCATTTCGACATAAAATCATCCACTGAAAATTGAA  
 AACTAAAAACAGAATGGCAGCAAAAAAAAAAGCAAATCCGAGTGTCCAAGGCGCAAACGCTTACGATCGATCGT  
 TAAATAATTGCAAATAATGTAATGAGGCATGGTCAGAACTGAAACGCCGAACCCGCCCATCCACGACTGCA  
 CATCAATGGCGATTGGCGAACATCCGCGGACATTGGCGAGGTGAGCCAACCAAATGGATCTCGAACGACGCTG  
 TGGAAATGTGAATTATAACTGAATTGCAATTGCTATCATGCAAGTGTGGCATTGCTAAATCACAGCTCA  
 ATGGTTCTTGAAGCTCCGACTGTTAATTGCACTTCGCTTAATTGTCATTGCTACTTGCCTCTGACTTGGC  
 CCAAAAGGGCAAGAGTTGGGAAAGGCCAGCGCCAGAGTGCAGACTTCATCTGTGACTTATGTTTCGCTGGTCT  
 GCAGCTGGCTATTGATCTGGCAGGCCATTGCTCAGCGCGTTTGCAGCTTCTTCAGTATCCCTGCATTGTT  
 CGATTTAGCGGGCAATTAGTCAGGCCAGTAATGATTGCCAGCGAATTATCGATCCGATCCGGTCCGCTGC  
 CAAACCTCCCAGCACATGGTCGAGTGTATCTGGCAGATACTACGGTAATGCGGATGTACACGTCTG

## 63. lncRNA.677

GGCGGTACGATAACACATACATCAGCAGTATCTGGCTGGTGAAAACGGCGGCCACACGGACAGATCTGTT  
 CATCTCGCTGCTGATAACCAAACAGCCGCTATACGAGGAGACAGGAACGTGGG**CTGCGTTCTG**  
**TCCGTCC**  
**GGCTGTCAG**TCCTGCTTTGGAAGGCCCATCTGATGTGCGAAAATAATTCCC  
 CCAATGAAATTGGACAGTCGACGGGACACACCCAGAAGAAAACACTGGCGAAAAGAAATCTGAAAACAATTGAAAG  
 TGTTGAACTTATATTGACTTGAATTGCTGCTACTAAGCTATGAGATGGATTCTAGAAGACGGTGCAGCAGGGTAA  
 GTGGCTTCTTCTGAGCTCCGACTGTTAATTGCACTTCGCTTAATTGTCATTGCTACTTGCCTCTGACTTGGC  
 CCAAAAGGGCAAGAGTTGGGAAAGGCCAGCGCCAGAGTGCAGACTTCATCTGTGACTTATGTTTCGCTGGTCT  
 GCAGCTGGCTATTGATCTGGCAGGCCATTGCTCAGCGCGTTTGCAGCTTCTTCAGTATCCCTGCATTGTT  
 CGATTTAGCGGGCAATTAGTCAGGCCAGTAATGATTGCCAGCGAATTATCGATCCGATCCGGTCCGCTGC  
 CAAACCTCCCAGCACATGGTCGAGTGTATCTGGCAGATACTACGGTAATGCGGATGTACACGTCTG

## 64. CR42874

TGCATTGCAATTGATGGGAAATGCAAGATGACTATGCGATACTCTTGAATCTGGAACTTGACCCGATGGCTT  
 ATCTTCGCTCTGTTAGTCACATGATATCCAGCTCTGGACACTTCTCACCGACATGGCGTTGATAATGAGCCA  
 GCTGCTCAATTGGAATCGAAAACAATGCTCGTAGTGTGTTGGTTCGATTACGGTCTACAGTTCGCA  
 TTGCGATTGCTATTGCTATTGCTCAGCGCGTTTGCAGCTTCTTCAGTATCCCTGCATTGTT  
 CGATTTAGCGGGCAATTAGTCAGGCCAGTAATGATTGCCAGCGAATTATCGATCCGATCCGGTCCGCTGC  
 CAAACCTCCCAGCACATGGTCGAGTGTATCTGGCAGATACTACGGTAATGCGGATGTACACGTCTG

AAGAAATCGCTTATTACATTCACTTGCGCCACGAATCGCTTGAAATGCAATCCGCCAGTGATTGGTGTGGATT  
GGTTCCCTTCGGAGTGGTCACTTTGTGGCTTAAAGACAGTTGATCCGACTGTGTCATCTGAATGGTCGCATA  
ACTGGGGATTCAACATGGTGGAGCTATTAAAGAAGGTATAACAATAGAGCTACTAAAAAAACAACAACTCTCTA  
TAGATCCCCTTAAGCTTGATAATAGCACCAATCTTATAAAAAAAACTAGTCAGTTGATCAGACTTTATATGGCCAC  
ATAATCCACAATCCTAGGAGTGCAGTTAGCTTCACCTCCGGTTGGGCTCTGCGAGTCAGCTGCGAGGAAGGAG  
TCCGCTGGCTGCGATGTTCACTTGATTTACATATGAATCCGCATAAAGCACGGCAGTCCGAAACTGCAAAGGAATGTGG  
GCCAAGCCCCAGTCCATCCTCTTAACCGCCGACGCCGC

#### 65. lincRNA.484 (splicing site mis-annotated, excluded in the analysis)

CTAAAATGACTTTCAATCATTTCTAAATATCATTGATCTAAATCGTACGTATCTGGATCATCTGGGACATTG  
AGGATCCTACCTTAATCCCTGCTCGATTCCATTACCCGCTCCGGTCAATTAGTCATTCCGGCAGGAGATATGCAATT  
**CCG**gtattcgtaactcgactagcttgcataaaactctcaaccatattccg **cttattcaccag** **TCCGGTCAAG** TGCT  
CTGACCCGATCCGATCCGATCGGATCCTGAGGACAACGACGACGACGACCTCGACGAGTGTGACTCAGGCAATTAG  
CGGAGGAAGTGAAGTCGAGGAGTTGGACAGCAGACGGATTGTGTCATTAGCGCACTCGCGAGCTGCCACTGTGTC  
CG

#### 66. lincRNA.490

CAGCATTAAAGGATTGCCGTTTCGATTCAAATTATCTGAACAAAGCAATGTCCTAAGTGGCAAGGGTAAGGCACGTAAA  
TAAAAACGAGGAATAAAATCCGGAAAAAACCCATGCACACATATACGATATGCAAATGAAAGTGAAGTGAGCGCA  
GGATAATGGCGACCAGAGCGGGACGGAAGGCCAAATGAGTGAAGAGAGAGAGAGCGATCTGGAGTAAGGGTAATGAGT  
GTGCATAATGCAGATGCAGATGCA **GGATCCCGTCCAGGATCCCAGG** AGCAGACCAATCCAGCCATCTCAGCACTCAT  
TGAACAAGCGGCTCAATCTGCCATGCCAAGATGCACTGGTAGTAGTACACTGGAA

#### 67. lincRNA.623

CGAATCTAGTACCTGTTAACGCAATGAAAACGTTACTGCTCAAACGTATATGTACATGCTGCTGGCCTAATA  
ATTAACCTGCTGATAACCCCTCATCAAATCA **CACAACTGCCGTCCACACAATT** CCATGGTGTCCATTATCAAATCCCTG  
TTGGATGTCCGGTGCCTGAATCAACACAATTCAAAGCCACCCAGAAGTCGAAACCCGTTTCGGCACCATCGTCATT  
CATCATGCCGCCTGGCACTTGGCAAATTACACAAGCCAGTAAACGGATAGACAGAAGGACGGAGCCAGCCAGCCG  
AAATACAATTAAACCAATCTAAACTCAAATTAAATCACAAAAGTTACAGTTACCTTCGGCAACAGCAGAAGCA  
GCAGCAGTAGCAGCCGACGTAAACTTGGTCAGAATCTGTAAGTTGCACTGAATCTGGTCAAAGATA CGCTACG  
ACTCCCATAACACAGCACCTTGGCAAACACACTCAATTATAGATA CAGACAGACAGTGA GAGACCCACAAACTCCGTGGA  
AAACTTCTCGGGCACTTGAAAATGAAGCGAACCTGGCAGACAGACGGCCGTTCCAAGTTGCAATTGCAATTGCA  
GCTGTGCTGGCAAATGCATTGCCATTCAAACAGTTGATTAAGTTCTGGTTAATGGATTGTTACATAC  
CTAACGAGGCCACTCACAGAAGTGCACACTGCAACTTTGGCAATCACCACAAATGGTAATC

#### 68. lincRNA.786

CTGGGAGGCCTCTCACCTGCCTTCTGACCTGGCTTGGCTTGCCAGGGTCTGGCCCTGGTTTGCCCT  
GGCCGACTATCAGTGGGTAGTGTCACTCATCGAGCCGCACTGCTGAGCATCAAATGGCCGTGGCAAATAAAGCCAAAC  
ACTTGGCCAGGGTGTGGGAAGTCTGCACTGGCACCAGGTCTCTCCGCATCATCTTGGCCGGCTCTAGATTG  
AATGCCCTGAAGTTGGAGATGGGCAGATGGAGGTGCTCACGTACCTTCTGTTGATAAGGATCCGAATCGAGAA  
CGAACAGGGCGAAGGCCGTACGGTCAATATATTGGAATTTTATTATGCATGACAGCTCAAAAAAGAAGAACAAACA  
ATGGACACGACGCCAACACAAGGAGCGCTGGGTCTCCATCATTTCCCTCATTTCTGTTGGCTTGTGATCGCATCGGT  
TCTGTTCTGCATTTGTGATGATTATGTAACATTTTCTGCTGCTGAC **CCGTCCAGCGTTCTCGAAAAGGAAAC**  
ATCGATTATTTGACAAATGATTACCAATGAGTTGACAGTTATTGCGTTTACTTCGCTTATTTCCACTTTTATGGAAA  
CGGTTAATAAGAACACAGGAGGGAACATGTATT

#### 69. lincRNA.626

CTTTACACAATAGTATTGTGTGATTATTAGTCAGAAGATGGCAATGAGCTTTCATGGAATCCATATACGGGAAACA  
AATTAAAGTTGTTGCACAAATCTTAAATTGGGATTCTACCTTTGTAACAGTACACATATCACTCAGTTTT  
TCTTATTGAAGTTCAGATTATTGTAATGATCGGACGGATGGCGAATTAAACACAGCTCATGTTACCATGAAAGTT  
GCCGACTCTAAAGTATATTATTCTGATCGGAATTACCA **CCAGTCGATGTCGTCCCGTCA** GAAAGTTAAGGTGAAGAA  
TGTAGATTAGTACTGTCATCCACAGAACGCCAACCTACCAAAGCGACCACACTGAAAGGGAAATTATTGAAATT  
ATGCAATATTTTATTGTTGTTGGTCAGCTAACAAACAGCAACTGCAAGTTGTCACTCTCAATTCTAAAGAT  
CATTTGATATTTCTTTATTATTACCAATATAAATAGGTAAGTTATATCAAGAGAACTTTAAATTCTCC  
CTGCCCACTATATGTGTGTTATATGTAACCTTACGGCATATGTTTTTTTATTGTTCTTATGTACGAATCCAGT  
TCTAAGGCCGATATTATTAACTTAATAGTCAAGTTAATAAAATTGAAACAATAAGACAAACGAAACATTTCATA  
TGAAAATAGTTAAAAGAATGCACTGAATGTTGAGACATTAGTTGGCACAAACATTGCAATCGTACAGGA  
TGAAAATTTCAGTAAACAAACAGCTTAATCGAGTGAAGCTGAGTAAGATTAGTGAATGAAATGCTTCA  
ACGTTCGCAGAGTATACCTATATACAACTTAAAGCCACATCCAATGACCACAAAGTATCTGGCATTATCAA  
ACTGTACCGAGCTGCTGGAAAATTAGCCAAACTTTCCATCAATCGCAGACGATCCGAGCACAAGCGACCAAGGAG  
ACAAGTCAATTCACTTCCCTAACAAAAACTGATCACCTTAAAAACTCATATAACAAATACATTAGCTTTAAATAT  
TTTGATATCGTTGAATTAGTTATTCTATCATTAATTAAATTGCTCCTTCTAGCTGTGTAATGAGTATC  
TGAAAGCCGAGTAGCTTGTAACTGAAAATAATGATGGTCGCTGCCGATGCTGCGCTGCCGCTGACCGCTGGCT

GCATCCACTGCAGAGACGAAGGATGTGGATAGGCATAGGCACACATGCATAACGCCAACCTGGACACAGGTAGCTCGGCC  
TCCGCAGGTAAGCAAGTCTTGGCCGGAAACACTGCCAGATATGTATGCGTCATACAGATTCCTCTGCTCGTT  
ACTTGTGCAAACAAGCCTGAGGACTGAGGAGGGGGACAGCACACAGACTCAACCGGCTGCCCTGCCCATGCTGG  
GCACCACCGGAAACATTAGCAGCCTGATGATCATCAGGGCCAGGCAGTGGGACTAGGGACTCATGACCAAAGCCAGC  
CGGACTGGTAGGGATTGGCCGAGATTGGTTGGCGCTGCAGAGGAAAGATGCAACATCGGTAGCTCAGGGCTGCA  
GCATCATGGTATAAAAAGGAGACTAGTTATCAATTGGATATATGTTCGCTTTGTTAATACAAATTAAAA  
TAAAATATCTAAAATATTGCTTAAATTTAAAGTTTACGTACAACAAAAGTCGTAATTATTAA  
TAACCTAAACTTGCAAAGTAATTATTAGCGTTGCTATTCT

## 70. CR43314

CAGGCAAGTCGAAAGTGGAACGACGAGCGAACGAGCTAAGAAAGTCAGTTGAAAACAGTGGAAACGCGGTGCGAGTC  
CTTTTTCATTGCACATAGAGAGAGGGCCCTGTATACTCAGCAGCCTACCACAAACAATCAACCAACCAACCGACCAGC  
CCAACCAACATACCTGCGTGTGGCACGTTGTCGCGATCCGATCAGTCAGCAGCGCAGCAGCGCACACACCG  
AACACACAACTCAGCAGTCTGGAAACGAACACCACTGTATATACTATCAAGCGAAGTGTAGAGATAAACTGATAATA  
CAACGGAGCGAGAAAAGAAAAGCACTAACGTGGCGTTAECTCTTTAACTAAAAACACTCACTCACTCAACCAAAAAAA  
AAAAAAAAACGTCAAGTAAAGACAATAACAAAGTAAACAAAACAGAGTCAGCGGCAAAGTGTCAATAAAACTCTGCAAT  
TCATTAACACAACCTCAAACGTTCTATTAGTAAGCACTTGCAAATTAAACGAAACAAAAAAATCTGACAGAGA  
GAGAGAGAGACAGAGAGCGATAAAAGGAGAGGAACATTAGACAGAAATTAGCGTGGCCAACAGTCAGGAGTCAGTCAA  
AAGCGCGTGTACGAGTCAAATGCCGTACACCTCGTCGAGACCCACAAGCCATGTCGAAAAACACATCAACA  
AATGTCGAAACGAGATAAAAAACAAACAACCGAGAAATATGAAAAAAACAGAGACAAAAACCCAGACG  
AGACCGCTGAGACATGCTTGGCTGTAAAAGGGCAG//(11079bp)//GCGAAGTGTAGCAGAGCAATTGTCATTG  
CACCAACCTATAAAAGAATGCACAGGAAGAAATTCTATATTTCAAACTATTATACATAAAAATCCAATGAAAGTAA  
AAAATGTATAATGTAATCAAAGTAAATGTACAGCTAGTTATATCAGGTTCCCACCATCTTAAATTAGTAG  
TTTGGGACAAAAATGGTAAATCAACTGTGGCACCTATTGAGGAAGTTTGTGTCCTGTTGTCGTTACA  
TGAGATGCACGTTGTAAGCGGATAGGCTGTGAGGGTTGTCAGAAATTCACTTAATAATAATTGCAATTATATTCAA  
TGTGAAAGCTTAACGAAATGATTATCTATGAAACATTAAAGAGTATTACATTCAATATGTGATATACAATTAAATT  
CAAAACTCACCAATGCCCTCGCTCGTACCTGATACTTATGCCAATTACCATTCGCTTCAGCAGTCGAGTCGGGCTAGA  
AAATGCCATGGACTCGCTACAATTGGCTCGTCTGTCAATCAATTGGATCGAGTCGGACTAGGCTCTAACTTAATT  
GCCGACAGACTGAAGACTAAAGACTGAATGCCAAGCAGCACAGGACCTCTGGTGCACCTAAAGCTGACGGA  
GGCGACTAATTATCAAATGGCTAGAAAACGAAATACTCACCGATAATTAAATCCAGCTAAAACGAAAATGAG  
TGAAACACCGCGATACTCAAATGCGTTTAAACCGTTGCGAACGCCATTCTAACAGTTAAATTGAGACT  
TTTCTTGGTTTGTGCAATTTCCTTCTTAATTGCCCTATTATCC//(5705)//TTCCCTCTGGCAA  
GTTCCGCTCGAGCACATTATGCCGAAAGAGGGGTCTTTAAATGATGACGCTCCTGTTGGCGGCCGACATATAT  
TTATGAGACACCTTGGCGTTGATAAGTCCAGACGAAGCTCAATTGAAATTGACTTAATTAGTCTGGTATGGCGAA  
CCGGCGTTCTTGATCATCGACTAAGAAGGAATTCTGAGAATTGCAAGTAGCCTACACTGGGAGCAATTATTC  
TAGGCTAAAAACAATATTAAACGCGGAAATAGAATTACAAATATT

## 71. lncRNA.118

GTTTGGTATGTTTATTGCTTACTTATAAGATAGTATAGTTAATGAAACATACTTCTTTAGATTACATTAAAG  
AGAGGGGGTATCATATTACTTGTAGAGTGTATACTTGTGGATTCATATTGTAGTCGTCACATTCCATTGAAACGACT  
TTCGGGCGCTCAATTGCGCATAGTATTATACAAATAACGAAATATTGGTATGCGAGTTAAATTGAGT  
GCTATTATTGTCGCTGCGAAGTCTGCCGTCAGCTCAACCGTCCAAACCGTCCAAACCGTCCAAACTGT  
**CCATCTGTC  
CATCTGCCAATT**ACAGTTCAAGCTGGTAAGTTTGGTCACGGACGTTGGAGGCCAGTTGGCTAGATTCTAGTA  
**TCCGTCCGTCGAGCGAA**CATAACAGCATATTGCGCAGCATGGTCGCTGACTTAATGAACTCAACACGAGACGATCG  
ACGATGGAGCTGAAAGATCATCACAGCTCAGTGGTGGTCATATCGGATATCATGGATGTTGGATGTCGGCTTCAAG  
TGGATCGGATGCGATGTGCGTGCAGCTAATTGATGGGTGCACATCAATTGAAGTTGTCACTTTCAATTGGGCCAAG  
ACGCGGAATTTCAGTCAAGTCGCTGTCAGCAGATTGGAATTCTGCCAATCCGGTTATCTGGCCAGATGTCA  
CCTGGATGATCTGTCGGTCTGGACAAAGTGTGCCATCGATAAGGCCGACAAAGTTAACTGATAAACATGGTCTAC  
ATACATACTACATACTCTTGGGCTTCAAATGGAGAGCACTCGGAAAGGATTATATTCCGGCGATAGTGTGATTG  
AGATAATGCACTGCGAAATCTAACGGACGACAATAATCCCCGCATAAGTACGATTGAGACACCAATTGAGCA  
CTTCCAAGTAAAACGAAACAGATCGCATAACATTCCACGAATAACGTTAAGTATCCCCTGAAATGTATACAT  
ATATCGGCA