

Supplemental Methods and Data

1. Drosophila culture, stocks, demography

Culture

Flies were reared and aged at 25°C, 40% RH, 12L:12D on standard food media: cornmeal (5.2%), sugar (11.0%), autolyzed yeast (2.5%; SAF brand, Lesaffre Yeast Corp., Milwaukee, WI, USA.), agar (0.79%) (w/v in 100 mL water) with 0.2% Tegosep (methyl4-hydroxybenzoate, Sigma, St Louis, MO, USA).

Oviposition plates. 5 cm dishes with grape juice media and excess live yeast were affixed to the food tube port of demography cages. Plates were changed twice daily, with total eggs summed each day. Grape juice plate media: 25 g Bacto-Agar (BD cat # 214010), 250 ml grape juice (Welsh concentrate), 750 ml water and 15 ml 20% Tegosep.

Lifetable demography

Cages for lifetable demography were made from 1 L clear food service containers with a ventilated lid, a gasket covered aperture to provide access to remove dead flies and a port near the cage bottom that opened to a plastic tube affixed a standard glass media vial with 3 ml of standard Drosophila diet. Each two days, dead flies were removed by aspiration and counted, and fresh food vials were provided.

2. Ends out homologous recombination

Reagents

Genomic DNA extraction: Promega Wizard SV Genomic Purification System. Promega, cat # A2361
PCR: Phusion High-Fidelity PCR Master Mix with GC Buffer. New England Biolabs, cat # M0532
Cloning: Zero Blunt Topo PCR cloning kit, Invitrogen, cat # 45-0245
pW25.2 (Drosophila Genomics Resource Center)
Bacteria Transformation: One Shot Top10 electrocompetent cells (Invitrogen), cat # C404052
Plasmid DNA prep: QiaPrep Spin miniprep kit, Qiagen, cat # 27104
Gel purification: QIAquick Gel Extraction Kit. Qiagen, cat # 28740
Site-direct mutagenesis: GeneArt site direct mutagenesis system, Invitrogen, cat #A13282, using AccuPrime Pfx DNA polymerase, Invitrogen, cat #12344-024
Primer synthesis: Integrated DNA Technologies (IDT, IA, USA)
Sequencing: Genewiz (South Plainfield, NJ, USA)
Restriction Enzymes: New England Biolabs
Embryo Injection: Genetic Services, Inc. (Sudbury, MA)

Drosophila Stocks for homologous recombination

Genotype	Stock
$y^1 w^* ; P\{ry^{+t7.2} = 70FLP\}11 P\{v^{+t1.8} = 70l-SceI\}2B noc^{Sco} / CyO, S^2$	BSC 6934
$y^2 w^{1118} P\{ry^{+t7.2} = ey-FLP.N\}2$	BSC 5580
$y^1 w^{67c23}; sna^{Sco}/CyO, P\{w^{+mC}=Crew\}DH1$	BSC 1092
$w^{Dahomey}$	L. Partridge (UCL, UK)

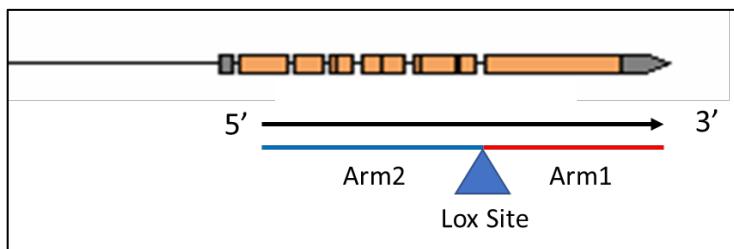
All lines backcrossed to w^{Dah} for six generations.

Targeting arms for homologous recombination

InR gene replacement to produce single nucleotide substitution followed Staber *et al* (Staber CJ, Gell S, Jepson JE, Reenan RA. 2011. Perturbing A-to-I RNA Editing Using Genetics and Homologous

Recombination. Methods Mol Bio 718: 41-73). *InR* genomic DNA from the Dahomey wildtype (*wDah*) was extracted, PCR amplified using *InR* Arm primers (**Methods Table A**) and subcloned into the pTopo vector to generate constructs pTopo-Arm1 (3' end of gene) and pTopo-Arm2 (5'end of the gene). The *w⁺* marker was targeted into the 9th intron of *InR*, dividing this sequence into two segments: Arm1 with 2,686 bp and Arm2 with 3,614 bp. To subclone both arms into the pW25.2 transformation vector, Arm1 was flanked by restriction sites *Ascl* and *BsIWI* and Arm2 by *Notl* and *Acc65I*. Since *wDah* contains an internal site of *Acc65I* in Arm2, this site was abolished by site directed mutagenesis without altering its coded amino acid (from GGTACC to GCTACC, at position 1194 of Arm2). All subsequent cloning was made using Arm2 with the abolished *Acc65I* site. Individual clones containing pTopo-Arm1 and pTopo-Arm2 were completely sequenced using primers listed in Table B. Complete sequence of *InR w^{Dah}* Arm1 and Arm2 in **Methods Fig A, B**.

Placement of homologous recombination targeting arms on *InR*.



***InR* single nucleotide mutations for homologous recombination**

The construct pTopo-Arm2 was used as a template for the site-direct mutagenesis to introduce the *InR^{e19}* mutation, a modification from T-A at position 1890 of Arm2. The construct pTopo-Arm1 was used as a template for the site-direct mutagenesis to introduce the following independent mutations: *InR²⁴⁶*, *InR³⁵³*, *InR⁷⁴* and *InR²¹¹*. *InR²⁴⁶* has a G-A modification at position 291, *InR³⁵³* has a C-T modification at position 537, *InR⁷⁴* has a A-T modification at position 768 and *InR²¹¹* has a G-A modification at position 933 of Arm1. After mutagenesis, constructs were fully sequenced to confirm the single nucleotide mutation. Location of substituted nucleotides are underlined in **Methods Figures A and B**.

Arm 2 = 3614 bp

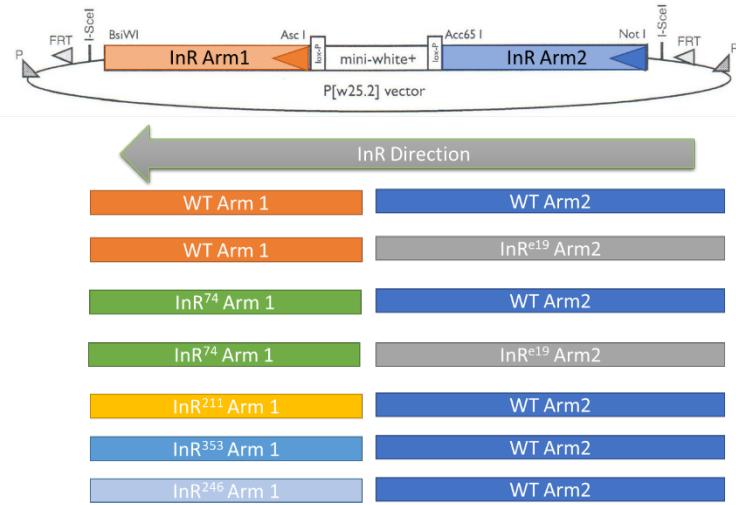
Mutation	Position in Arm 2	Nucleotide change	Aminoacid change
<i>InR[e19]</i>	1890	GTC/GAC	Val/Asp

Arm 1 = 2686 bp

Mutation	Position in Arm1	Nucleotide change	Aminoacid change
<i>InR[246]</i>	291	GTG/ATG	Val/Met
<i>InR[353]</i>	537	CGT/TGT	Arg/Cys
<i>InR[74]</i>	768	ATC/TTC	Ile/Phe
<i>InR[211]</i>	933	GGA/AGA	Gly/Arg

Cloning into transformation vector

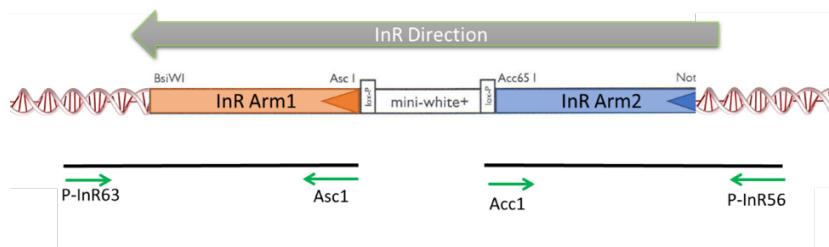
The wild type and the mutated *InR* were sequentially cloned into transformation vector pW25.2. Arm1 wild type or mutated was cloned into the restriction sites *Ascl* and *BsIWI*. Arm2 wild type or mutated was inserted into restriction sites *Acc65I* and *Notl*. Seven different constructs were created, allowing for wild type, single mutants for each one of the five single nucleotide substitutions, as well as one double mutant, *InR^{e19-74}*. Constructs were then transformed into *Drosophila w¹¹¹⁸* embryos by Genetic Services, Inc. (Sudbury, MA).



Flies carrying the *white⁺* eye color marker mapped to the first or second chromosome were used to target homologous recombination by crossing with a line expressing flipase and the restriction enzyme I-Sce1 for excision and linearization of the transgenic construct. Potential lines were selected for *white⁺* insertions on chromosome 3, and confirmed by PCR amplification and sequencing. Accession with successful gene replacement were then crossed to flies expressing *cre* recombinase for removal of the *white⁺* eye color marker.

Confirmation of mutations

Final lines were sequenced to confirm fidelity and homologous recombination of the replacement allele. using primers internal of homologous recombination construct and primers upstream of Arm2 and downstream of Arm1 into the genomic region. PCR fragments were sequenced to completion (primers in **Methods Table B**). The wild type strain *wDah* was also sequence to completion, and the following primers were used for its amplification (Arm1: P-InR63 and P-InR60; Arm2: P-InR56 and InR61).



Methods Fig A: w^{Dah} Arm1 sequence cloned into pTOPO

w^{Dah} Arm1 sequence cloned in pTOPO	
GAATGATCATTAAATGAATTTCATTAATTCAACAGCCTCCGCCGAGCTATGCTAAGGTCTTTCTGGCTACTGGGAATCGCCCTAGC	< 100
GTTCTGATCGTTCCCTGTTGGCTATGCTGTTACCTGCACAAGAGGAAGGTTCCCTAATGACCTCCATATGAACACAGAGGTGAATCCGTTCTAT	< 200
A GCGAGCATGCAATACATCCCAGACGATTGGAGGTGCTGCGAGAACATCATTGAGTGGCTCCACTAGGCCAGGGATCCTTGGCATGGTATGAGG	< 300
GTATCCTGAAGCTCTTCCACCCAATGGCGTGGATCGCGAGTGCTGCCATTAAGACTGCAACGAAAATGCTACGGATCGCGAGCGAACATTCCCTGAG	< 400
CGAGGGCAGCGTCATGAAGGAGTCGATACTATGCTGAAGATTGCTGGTGTGTTCCAGGGTCAAGCGCTCTGGTGTGATGGAGCTAATG	< 500
AAAGAAGGGTGAATCTTAAAGCTCATTGCGTGCCTACGTCGAGATTGCGATGGCATGGCATATTGGCCCAAGAAGTTCGTCATCGTGTGATCTGC	< 600
TGCAGCCTCTACTTATGGAAGAATCTACAGATGCCATTGAGATTGCGATGGCATGGCATATTGGCCCAAGAAGTTCGTCATCGTGTGATCTGC	< 700
AGTCGAAATTGCGATGGTGTGATGATTGACGGTAAATTGGTGAATGACCGTGACATCTATGAGACGGATTACTATCGGAAGGGCACT	< 800
AAAGGGCTGCTGCCAGTTCGCTGGATGCCACCGAGCTGCGAGATGGTGTACTCTAGTGCAGTGATGATTGAGCTTGGAGTGGTCTCTGG	< 900
AAATGGCCACCTTAGCGGTCAGCCATACCAGGGACTTCCAACGAGCAAGTCTCGTGTACGTTATCGATGGCGGTGTTATGGAGAGGCCGAAATTG	< 1000
TCTGTGATTTCTGCATAAACTAATGCAAAAGGTGCTGGCATAGGTCTCGCGAGACCCAGTTTCTGGATATCATTGCTATCTGAACCACAATGC	< 1100
CCCAATTACAATTAAAGGAAGTATCCTCTATCACTCAGAGGCAGGTCTGCAGCATCGGAAAAGGAGCGCAAGGAACGCAATCAGCTAGATGCATTG	< 1200
CGGCAGTCCCCCTGGATCAAGATTCGAGGATCGGGAACAGCAGCAGGAGATGCTACCACACCTTACGAATGGCGATTATCAGCAGAACCTCGTGTGGA	< 1300
TCAACCGCCGAAAGCCCCATGCCATGGTTGATGATCAGGGTCTCACTGGCATTAGCCTGCCATCCGGATTCTGGCAGACTCCTGATGGG	< 1400
CAGACTGTAATGGTACTGCTTCCAGAAATATTCAGCAGCGCAAGGTGATATTCGCGACTTACGTTGCGCTGATGCAGACGCTTGGACGGCACA	< 1500
GGGGATATGAGATCTACGATCCCAGTCCGAATGTGAGAGCTGCCAGCAGCAGAAGTGGCAGTACTGGCGTGGAAAATCAGCGGAGAACACATT	< 1600
GCTGCCAAGAAGGGTCGCCAGCCTACCATCATGAGCAGCTCGATGCCAGATGATGTCATCGTGGCTCTACTGCAACCCCTGACTGCGTCAGCAGCC	< 1700
AGTTCTAATGCCAGTTCGACACAGGACGCCAAGTCTGAAGAAAACAGTGGCGGATTGGCTGCAATAAGGCAAACCTTACGTTACGCCACCTATTTA	< 1800
ACCACAAGCGAACGGGCAGCAATGCCAGCCACAAGAGCAATGCCATGCTCGAGTACCGCAGTAACACCAACTTGACAAGTCACCCAGTGGCTAT	< 1900
GGGCAATCTGGAACATCGAGAGTGGTGGCAGTGGTCCGCTGGTAGTTACTCGGAACACCCGCTCTACTCCATCGCAGCCCTGGAGGGAGC	< 2000
AGCGGTATGCCATTAGCGACAATCTAACTACAGACTACTAGACGAGTCATAAGCAGCGAACAGGCCACCATCTAACGACTAGCAGCCCCAATCCA	< 2100
ACTACGAGATGATGCATCCACCAACCAGTGGTCAGTACCAATCCGAACTATATGCCCATGATGAGACTCCAGTGCAGATGGCGGTGTGACCATTAG	< 2200
CCATAATCGAATTACCGCCCATGAGGCCGCTGTAATGCAAGCCAAAGCCTAACAGTGGAGCGACGCCAGGAGGAGGAGGAGGAT	< 2300
GAGGACGACGACGTGGACGATGAGCATGAGCACATCAAGATGGAGCGCATGCCATTGAGTCGGCCAGGCAAAGAGCGTTGCCAGCAAGACGCAGC	< 2400
CGCCTCGCAGTGCAGCGTCAGCCAAACGAGAAAATCTCTACGAATCCAACTCCGAATGGAGCGACAGGAGCCGAAACCGATCCAACTTGCTTAA	< 2500
AGAGAACTGGCTGCGACCGGGAGTACGCAAGGCCCTCCACCAACCAATGGATTATCGGAAGGGAGGGTAATCGTACGAACTGAGTTCTGTAGAAA	< 2600
ACATGTAATAGATGAGGAGAACGAGCATGGATATAGATAGAGTAGCAAGTCTTGGCGAAGCCAAGCTCTGGAAATTGACT	< 2686

Methods Fig B: w^{Dah} Arm2 sequence cloned into pTOPO. The Acc65I restriction site in region was abolished by site-specific mutagenesis, at position 1194 (marked in red).

w^{Dah} Arm2 sequence cloned in pTOPO	
ATATAGCCGATGGACTGGATAAAACAGCGTTCGGTGTCGGGACCCAATCGCATGGACAAGGAGCGAACATGCGACTGTCAAAAA < 100	
TGTAAAACgtatgttattacccatgtatcacccatgtcaagttaaaggaaacatcccatggcactgacagactataatagctttccattaacacct < 200	
tatagCTTGCACCATGGACATCAGGAACATGGTGTGCACTCAATCAGCTGGAGAACTGCACGGTCATCGAGGGCTTCTGCTGATCGATTGATA < 300	
AACGACGCCAGCCCTCTGAACAGAAGCTTCAAACAGTACAGATTATATCATAATCTACCGTGTACTGGATTGACTCGCTGTCAAAGA < 400	
TCTTCCCAATCTGAGCGTCATTAGGGAAACAAAGCTGTCAGCGGATATGCCTTGGCTGACTCGAACATTGACCTCATGGATTGGACTCACAA < 500	
GCTACGATCCATAACCAGAGCGGTGTGCGATTGAGAAGAACATATAAGCTGTGCTATGATAGGACATCGATTGGCTGAAATTCTGGCGAAACGAA < 600	
ACCCAACGGTGTGCTGACAGAGAACGGCAAGGAGAAGGAGTCAGGCTTCCAAGTGCAGGGGGAGATCAGAATTGAGGAGGGCACGATACACGG < 700	
CTATTGAGGGAGAGCTTAATGCCAGTTGTCAGCTGACAATAATAGGCCCTGTGCTGGAACAGCAAACCTGCGAGCAGCAGtgatgttgccggtgtaaa < 800	
gttataccgttttttcttaacattttcgcgtttttttctccagAATGCCCTGAAAGTCAGAAATAACTGCATCGATGAGCACACCTGCTGCG < 900	
CCAGGATTGTTGGTGGATGCGTGATCGATAAGAATGGAATGAGAGCTGCATCTCTGCAATGTGCTTTCAACAAACATCTGATGGACTCTGT < 1000	
CCGAAAGGCTATTATCAGgtactaatgattcacttttagtataacaacaagttagctaactttcaaacttttgagTTCGACAGCCGCTGCGTAACGGC < 1100	
GAACGAGTGCATCACACTGACAAAGTTGAAACGAAACAGTGTGATTCCGATACACGCAATGTATCACCCACTGTCAAACGGCTTACAG < 1200	
AAGTCAGAGAACAGCGCATGTGCGAACCTGTCCGGCGGAAAGTGTGACAAGGAGTCCTCTCGGTCTATCGACAGTTGGAGCGTGTGGAGT < 1300	
TCCACGGCTGACCATTATAACCGAACCGAGCCCTTACCATCAGCATTAAACGTGAAAGCGCGGtaagtgttttgctgtctaaattaaagtata < 1400	
atctctatattaaaacttctgttcttcagCTACGTCATGGATGAATTAAAATGGCCTGGCTGCCGTCCATAAAATTCACTGTCCTTAATGGTC < 1500	
ATTGACCTACGGATTGAAAGTCTTGAAATTCTTCAACTCCCTAAGTAAATTAGCGGCGATCCGCCATGGACCGCGATAAATGCTTGATGTGCT < 1600	
TGATAATCGCGATCTAGATGAGCTCTGGGACCAACCAAACGGTGTTCATTAGGAAGGGCGCGCTCTCTTCAACCCAAAACATGTGTGTC < 1700	
ACCATTAACCAGTTGCTGCCATGCTGGCTCCAAGCCAAGTTTGTGAAAGTCAGATGTGGCGCAGACTCGAATGGAAACCGCGGATCATGTAAGT < 1800	
AATCTTAAACGAAGATTTAGTAAATTAACTTGTCACTAACTCAAATGAAACATTTCAGGTTGGAACAGCGTTCTCAATGTCACATTCAA < 1900	
TCAGTGGGAGCAACACTCCGCTATGTCAGCAGCAGAAAGTGAATAGGAGAGCCCCAAAGCCGAGCAATGCTACAATTGTTTAAGGATCCG < 2000	
GCGCCTTCATCGGTTCTGTTATCATATGATCGATCCGTACGGGAACTCAACTAAAGCAGTGACGATCCATGCGATGTCGCTGGAAGGTTAGCTC < 2100	
TCCGGAAAAGACCGGGGTATGGTATTAAAGCAATTGCTTCCGATCTACACTAACTACTCCTACTACGTTCCGACCATGGCTATTCCTCGGAATTGACAAC < 2200	
GCGGAGAGCGACGTGAAGAACATTAGGACGAATCCCGACGACCGTCAAAGGTTACGGAGGTGGTAGCAACCGCCATTTCAGATTGAAATTGTGAGTA < 2300	
TGAAAATTGTGCAATAGATTAGTTAGGATTATAATGATTACTAAATGTCAGTAATTGTTCACTTGACTGATCAACATTATTGCTTTCAAGACGT < 2400	
AAACATGGAGCTACCTAGATAACGCTTATGGCGTCAACCGCTATTATAAAAGCCAACCTTAAATCCGCTACTCGAAACAATAACCGGATTAC < 2500	
TGTAATGCAAGCAACACATAGCCAATTGTATCCATAATTAAATTTAAATTTATCTGTCAGCTCTCGTCAGGCGATGGAAAATGACC < 2600	
TGCGACGCCAACGCTTACCAAGAAAATCAGATCCTTGTAGCAGGCGACTGTAAGTGCCTGGAGGGTTCGAAGAAGACTACGAGTCAGGAATACGATGA < 2700	
TCCGAAAGTCAGCGGATGGTTGAGAACCGCTGCAAACATTATTTGTCACGTTGAAACATTCGAAAAGCAAGAATGGATCGCTGCAAAATCA < 2800	
GACGGAGCGGAAGGTGCGCTCTGCTTCAATGCTATTCCAAATGGAGGAGCTACTAACCCCTACGTCAGAAGGAGAGACGTTGCGCTCGAGCGAGAC < 2900	
TCGACGATGTAGAGGGCAGTGTACTCTACGCCATGTGCGCTCCATCACAGACGATACCGCATGCTTTCGAAAGGACGACGAAACATCTAAAGA < 3000	
CGAAGAAGACTGTCTCCAACAAACATTCTATGAGGTGTTGCAAGGAATTGCGACCAAAATCAAACACATTGGTCTTGTGAAACTCGCCCACTTC < 3100	
ACCCGCTACGCTATCTCGTGGTAGCAGAAGAAAATCCCAGCGAAAATTAAGGGACACCCAGTTAAGAAGTCGCTCTGCAGCGATTATGACA < 3200	
CCGTTTCCAACACTAAAGAGAAGAGTAGGTGGACTTGAGAGCGTGTACTTCACTAACTGAGTTGGTTACTTATAGAATTGCGCACATAG < 3300	
TCATGGACCTAAAAGTAGATTAGAACACGCCAACACCCGAGTCCCGACGTTGCGACTGAGGAGAACTGGAGAAATTGTCAACCGAGCTGCGT < 3400	
CTATGAAAGTGGCTACAAGTTGCAAAACCGATCAAGTGGAGAAAGAAGTCGATTCCGGCTGCTGACTTCACCGAGACTGCCGTTATTAATAAG < 3500	
CTCAACGAGGGCTTACAGCTCAGGTGCGAGCCAATTCAATAGCGGAGTACGGCGATTTCACGGAAGTCGAACATATAAAAGTTGAGGTTGAA < 3600	
TAATTTTATCTGC < 3614	

Methods Table A: Primers for cloning InR Arm1 and Arm2, and for site-direct mutagenesis. Sequences 5'-3'.

InR primers – used for cloning and site-direct mutagenesis		
Primer Name	Primer Sequence	Use of primer
P- InR 13	TAATTAATTGGCGGCCGAATGATCATTAAATGAATTTCATTAATTCA	5' HR Region for InR Arm1; forward
P- InR 14	ATATCGTACGAGTCAAATTCAAGAAGCTTGGCTTGGCAAAG	3' HR Region for InR Arm1; reverse
P- InR 15	TAATTAATTGCGGCCATAGCCGATGGACTGGATAAACAGCGCTTCGGT	5' HR Region for InR Arm2; forward
P- InR 16	ATATGGTACCGCAGATAAAAATTATCAACCTACCTCAACTTTATATGTTGACTCCG	3' HR Region for InR Arm2; reverse
P- InR 29	CCACTGTCCAACGGG c TACCAGAAGTCAGAG	Primer for Acc65I modification in Arm2; forward
P- InR 30	CTCTGACTTCTGGTAgCCGTTGGACAGTGG	Primer for Acc65I modification in Arm2; reverse
P- InR 39	GGAACAGCCTTCTCAATG a CACATTACAATCAGTGGGA	InR ^{e19} mutagenesis Arm2, forward
P- InR 40	TCCC ACTGATTGTAATGTG t CATTGAGAACGGCTTCC	InR ^{e19} mutagenesis Arm2, reverse
P- InR 41	ACTTGGAATGACCCGTGACT t CTATGAGACGGATTACTAT	InR ⁷⁴ mutagenesis Arm1, forward
P- InR 42	ATAGTAATCCGTCATAGA a GTCACGGGTATTCAAAGT	InR ⁷⁴ mutagenesis Arm1, reverse
P- InR 43	GGCTCAGCCATACCAG a GACTTCAACGAGCA	InR ²¹¹ mutagenesis Arm1, forward
P- InR 44	TGCTCGTGGAAAGT c tCTGGTATGGCTGAGCC	InR ²¹¹ mutagenesis Arm1, reverse
P- InR 45	TATTTGCGTGC CC AT t GTCGGAGGAGCGGG	InR ³⁵³ mutagenesis Arm1, forward
P- InR 46	CCCGCTCCTCGGGAC a ATGGGCACGCAAATA	InR ³⁵³ mutagenesis Arm1, reverse
P- InR 51	GCCAGGGATC TT GGCAT g TGTATGAGGGTATCCTGAAG	InR ²⁴⁶ mutagenesis Arm1, forward
P- InR 52	CTTCAGGATA CC CTCATAC a CATGCCAAAGGATCCCTGGC	InR ²⁴⁶ mutagenesis Arm1, reverse
P- InR 56	CGGCTGCAACAGAGTGTGAGAGCGGGACGA	Arm2 HR confirmation, forward
P- InR 60	TCGGCCTACTCGAAACAATAACCGGGATTACTGTACTGAA Cgt aagc	w ^{Dah} Arm1, forward
P- InR 61	CTTTCCCGATGCTGCAGACCTGCCTCTGAGTGA TAGAAGGATA CTCCTAAATTGTGA	w ^{Dah} Arm2, Reverse
P- InR 63	CACTATCGTTCTGCTCGTTGTATTGCGTAGAGGCTTAGTGCATTA	Arm1 HR confirmation, reverse
P- Ascl	GTATGCTATACGAAGTTATCTAGACTAGTCTAGGGCG	Arm1 HR confirmation, forward
P- Accl	CATTATACGAAGTTATCTAGACTAGTCTAGGGTAC	Arm2 HR confirmation, reverse
P- SP6	GATTTAGGTGACACTATAG	pTopo sequencing
P- T7	TAATACGACTCACTATAGGG	pTopo sequencing

Methods Table B: Primers for sequencing Arm1 and Arm2. All primers were used for sequencing confirmation of homologous recombination events. Primers used to sequence DNA cloned in pTopo are highlighted in the right column.

InR primers – used for sequencing			
Primer Name	Primer Sequence	Use of primer	pTopo
P- InR 02	GCACAATTACTCTTACTCGCCTGG	Arm2; forward	
P- InR 06	TCAACGAGGGCCTTACAGCTCA	Arm2; forward	
P- InR 08	GTTACAATGGATTGCTAGCCAGTC	Arm1; reverse	
P- InR 17	CACAAGCTACGATCCATAACCAGAGG	Arm2; forward	✓
P- InR 18	TGTATGGACTCTGTCGAAAGGCTA	Arm2; forward	✓
P- InR 19	CAGTCGCCCTAATGGTCATTGAC	Arm2; forward	✓
P- InR 20	ATGATCGATCCGTACGGAACTCAAC	Arm2; forward	✓
P- InR 21	CCGGGATTACTGTACTGAACGTAAGC	Arm2; forward	✓
P- InR 22	GACGAAGAAGACTGTCCCTCCAAC	Arm2; forward	✓
P- InR 23	TCTACAGGCTACCACGAAGATAGCGT	Arm2; reverse	
P- InR 24	CAGGTCACTTCCATGGCCTTGACGA	Arm2; reverse	
P- InR 25	AACCTTCCAGCGATCATCGCATGGAT	Arm2; reverse	✓
P- InR 26	GGACGGCAGCCAGGCCATATTTAAT	Arm2; reverse	✓
P- InR 27	GACACATTCGACAGGAGATGCAGCT	Arm2; reverse	✓
P- InR 28	GAGTGCATCCAGTCACACGGTAGAT	Arm2; reverse	✓
P- InR 31	GGTGGTCATGGAGCTAATGAAGAAGG	Arm1; forward	✓
P- InR 32	GGCGAGACCCAGTTCTGGATATCA	Arm1; forward	✓
P- InR 33	GATCTACGATCCCAGTCGAAATGTG	Arm1; forward	✓
P- InR 34	GACTACTAGACGAGTCATAAGCCAGC	Arm1; forward	✓
P- InR 35	GCCATCTGCACTGGAGTCTCATTCA	Arm1; reverse	
P- InR 36	GACCCACCGATGACATCATCTGGCAT	Arm1; reverse	
P- InR 37	ACTGCCGCGAATGCATCTAGCTGATT	Arm1; reverse	✓
P- InR 38	CACATTACCAAGTCACTCCGATGCGAT	Arm1; reverse	✓
P- InR 64	ACAAAGCATATTTATCCCGTCC	Arm2; reverse	
P- InR 65	CAACACCGAGTCCCCAGTACGG	Arm2; forward	
P- InR 66	CCGTACTGGGGACTCGGTGTTG	Arm2; forward	
P- InR 67	TATGTTGACTTCGTGAAATCGC	Arm2; reverse	
P- InR 68	TTCAACAGCCTCCGCGAGC	Arm1; forward	
P- InR 69	GCAAACACCGAGCAATCTTACG	Arm1; reverse	
P- InR 70	TGGCCGCCAAGAAGTTCGTC	Arm1; forward	
P- InR 71	TGCTGGTACTCGGAGCATTG	Arm1; reverse	
P- InR 72	CACCAACTGACAAGTCACCC	Arm1; forward	
P- InR 73	ATGAATGAGACTCCAGTGCAGATGGC	Arm1; forward	
P- InR 74	GCGTTAGAGGGCTCTAGTCAT	Arm1; reverse	
P- InR 75	TGTGACAGTCGCATTGGGGT	Arm2; reverse	
P- InR 76	CATTGTGCAGCAGGTGCTGCTGA	Arm2; reverse	
P- InR 77	GCAGGACGAAAGATGATTACCTAG	Arm2; forward	

P- InR 78	ACGATATGGCAGCAGCAGCAAC	Arm2; forward	
P- InR 79	TGTTGCTTTGTTGC GG TG GGC	Arm2; reverse	
P- InR 80	GAAGTTT CGCGCT GTTGTCG	Arm2; reverse	

Figure S1. Partial sequence of Drosophila insulin-like receptor (dInR) aligned with human insulin receptor (hIR, HGVS annotation, with signal peptide) and insulin growth factor-1 receptor (hIGF1R). Drosophila sequence translation initiation site of wDah (parent wildtype allele of homologous recombination, GenBank accession MT_563159) and of Fernandez (NCBI Reference Sequence: NM_079712.6). **Red:** Drosophila InR homologous recombination substitutions corresponding to allele. **Blue:** substitution of the Exelisis dominant negative *InR* transgene P{UAS-InR.K1409A} (BDSC). **Gold:** hIGF1R polymorphisms of Suh {Suh, 2008 #5452} enriched in centenarians, and of the *C. elegans* longevity allele *daf-2(e1370)*. Domains: distal portion of L2, proximal portion of FnIII-1, Kinase insert domain (KID), Activation loop (A-Loop) with conserved autophosphorylation tyrosine (Y). The proposed SH2 binding motif of the Drosophila KID is highlighted, with Tyr1477.

dInR	MFNMPRGVTKSRSKRGKIKMENDMAAAATTACTLGHICVLCRQEMLLDTCCCRQAVEAV wDah TIS	Reference TIS	60
dInR	DSPASSEEAYSSSNSSCQASSEISAAEVWFLSHDDIVLCRRPKFDEVETTGKKRDVKCS		120
dInR	GHQCSNECDDGSTKNNRQQRENFNIFSNCHNILRTLQSLLLLMFNCGIFNKRRRRQHQQQ		180
dInR	HHHHYQHHQHHHQQHLQRQQANVSYTAKFLLLQTLAAATTRLSLSPKNYKQQQQLQHNQQ		240
hIR	-----	MAT	3
hIGF1R	-----	MKS	3
dInR	LPRATPQQKQQEKDRHKCFHYKHNSYSPEGISLLLILLANTLAIQAVVLPAPHQQHLLN		300
		:	.
hIR	GGRRGAAAAPLLVAVAALLGAAGHLYP-----GEV-CPGMDIRNNLTRLHEL		50
hIGF1R	GSGGGSPT-S---LGWLLFLSAALSLWP-----TSGEICPGIDIRNDYQLKRL		49
dInR	DIADGLDK----TALSVSGTQSRWRTRSESNPMTMRLSQNVKPCSKMSDIRNMVSHFNQL		353
	*	:	.
	*	:	.
hIR	ENCSVIEGHLQIILMFKTRPEDFRDLSFPKLIMITDYLLLFRVYGLESLKDLFPNLTVIR		110
hIGF1R	ENCTVIEGYLHILLIS--KAEDYRSYRFPKLTIVTEYLLLFRVAGLESGLGDLFPNLTVIR		107
dInR	ENCTVIEGFLLIDLINDASP--LNRSFPKLTEVTDYIIYRVTLHSLSKIFPNLHSVIR		410
	:*.* * * : ^ . **** :*:****:*** ** .** .** .:****:***		
	hIGF1R A67T		
hIR	YLKIRRSYALVSLSSFRKLRLIRGETL-EIGNYSFYALDNQNQLRQLWDWSKHNLTITQGK		452
hIGF1R	YVKIRHSHALVSLFLKNLRLILGEEQ-LEGNYSFYVLNDNQNQLRQLWDWDHRNLTIKAGK		445
dInR	SLMVHLLTYGLSKLFFFQSLTEISGDPPMDADKYALYVLDNRDLDLDELWGPQNQ-TVFIKGG		759
	: :: ::.* * *.::.* * * : .: * : * .****: * : * .: ^ : * *		
	hIGF1R R437H		
hIR	LFPHYNPKLCLSEIHKMEEVSGTKG-RQERNNDIALKTNGDQASCENELLKFSYIRTSFDK		511
hIGF1R	MYFAFPNPKLKVSEIYRMEEVTGKG-RQSKGDIINRNNGERASCESDVLHFTSTTSKRN		504
dInR	VFFHFNPKLCVSTINQLLPMIASKPKFFEKSVDGADSNGNRSGCXTAVLNVTLSQVGANS		819
	::* :****.* * : : . * .: * .: * . * : * .: * :^ : . . :		
	dInR^{e19} V810D		
hIR	FnIII-1		
hIGF1R	LLRWEPY-----WPPDFRDLLGFMFYKEAPYQNVTEFDGQDACGSNSWT		557
dInR	IIITWH-----RY-----RPPDYRDLISFTVYYKEAPFKNVTEYDGQDACGSNSWN		550
	AMLNVTTKVEIGEPQKPSNATIVFKDPRAFIGFVFYHMIDPYGNSTKSS-DDPC-DDRWK		877
	: : * * : . * : . : * .: * .: * .: * : * .: * :^ : . . :		
	hIGF1R N547H		
hIR	GMVYEGNARDIICKGEAETRVAKTVNESASLRERIEFLNEASVMKGFTCHHVVRLLGUVS		1094
hIGF1R	GMVYEGVAKGVVKDEPETRVAIKTVNEAASMRERIEFLNEASVMKEFNCVVRLLGUVS		1070
dInR	GMVYEGILKSFPNGVDRECAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGUVS		1441
	^ : . . : . * :**** * : * *** :*.***** * :***** * :***** * :		
	dInR²⁴⁶ V1384M UAS-dInR-DN K1409A KID		
hIR	KGQPTLVVMEMLAHGDLKSYLRSRPEAENNPP-----GRPPPTLQEMIQMAAE		1142
hIGF1R	QQQPTLVIMELMTRGDLKSYLRSRPEMENNPP-----VLAPPPLSKMIQMAGE		1118
dInR	RGQPALVVMEMLKKGDLKSYLRAH ^R PEERDEAMMTYLNRIGHTGNVQPPTYGRIYQMAIE		1501
	:****:**** :****:****: ^** .: . * : * .: * : .: * *** *		
	dInR³⁵³ R1466C		
hIR	A-Loop		
hIGF1R	IADGMAYLNAKKFVHDLAARNCMVAHDFTVKIGDFGMTRDIYETDYYRKGGKGLLPVRW		1202
dInR	IADGMAYLNAKKFVHDLAARNCMVAEDFTVKIGDFGMTRDIYETDYYRKGGKGLLPVRW		1178
	IADGMAYLAACKFVHDLAARNCMVADDLTVKIGDFGMTRDIYETDYYRKGTGKGLLPVRW		1561
	***** * :***** * :***** * :***** * :***** * :***** * :***** * :		
	dInR⁷⁴ I1543F		
hIR	MAPESLKDGVFVTTSSDMWSFGVVLWEITSLAEQ ^P YQGLSNEQVLKFVMDGGYLDQPDNC		1262
hIGF1R	MSPESLKDGVFTTYSDVWSFGVVLWEIATLAEQ ^P YQGLSNEQVLRFVMEGGLLDKPDNC		1238
dInR	MPPESLRDKGVYSSASDVFSFGVVLWEMATLAAQ ^P YQGLSNEQVLRVIDGGVMERPENCP		1621
	* ****:****: : * :***** * :***** * :***** * :***** * :***** * :		
	daf^{e1370} P1465S dInR²¹ G1598R		

Supplemental Figure S2. Distribution of eclosion time. Six genotypes, emerging males and females from day of egg deposition (0). Sum of two replicate vials. F_1 eggs after 24 hours mating, from crosses of 4 males and 4 females: InR (mutant allele 1)/TM6Sb x InR (mutant allele 2)/TM6Sb.

Emerging adults with marker *Sb* are TM6 heterozygotes with either mutant *InR* allele. Flies without the marker are trans-heterozygote mutant *InR* genotypes.

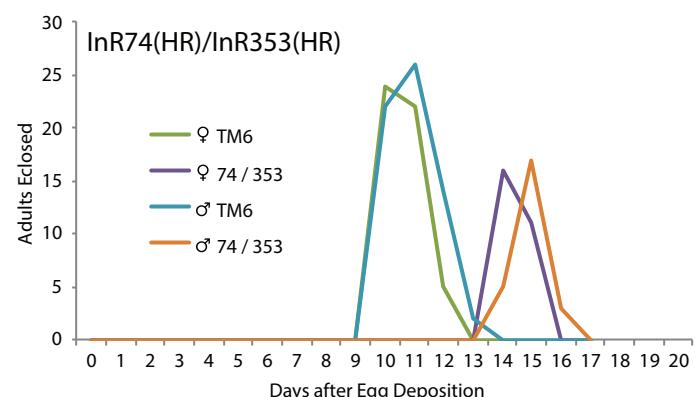
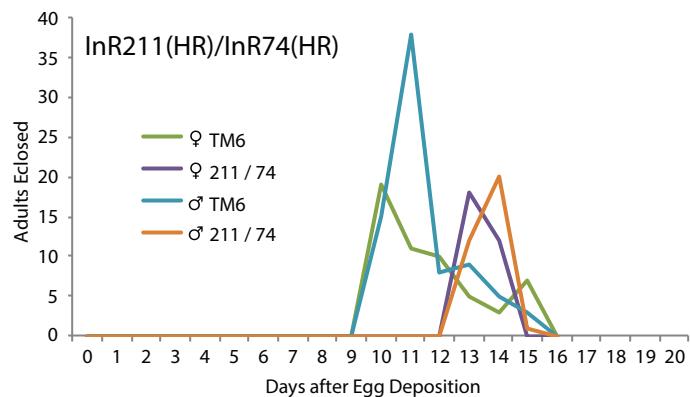
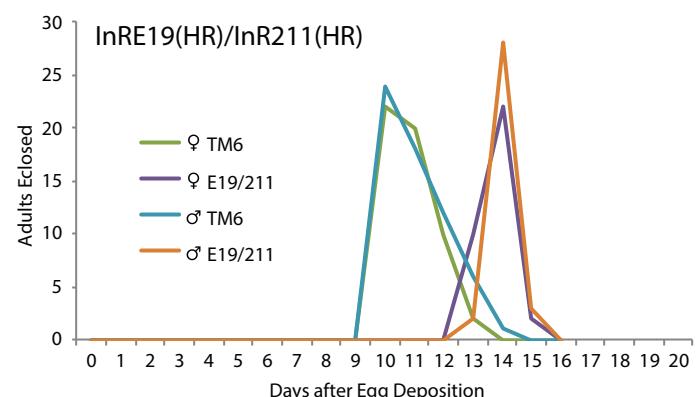
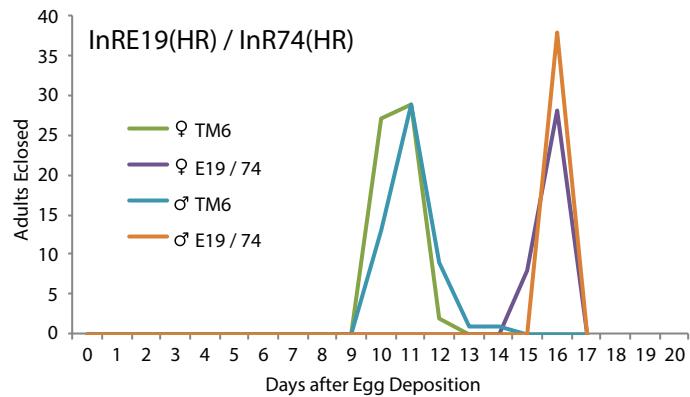
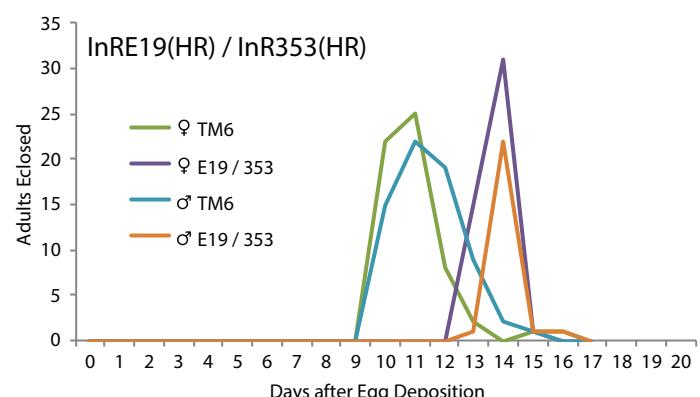
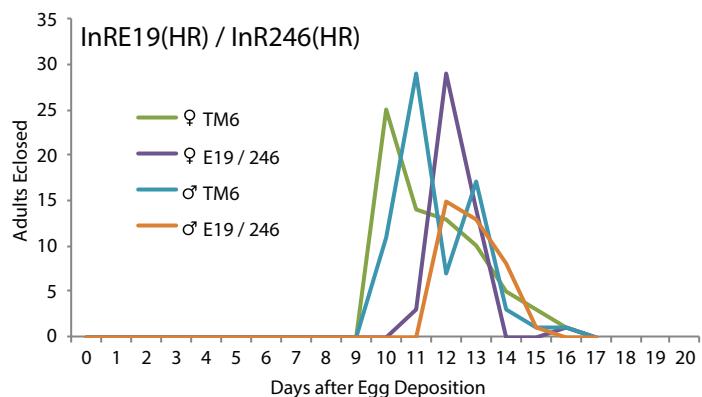


Fig S3. Master DNA alignment HR alleles with wildtypes wDAH and 29B(HR)

DAH	agcgggacgaaagatgattacctagcggggtgagagagacccaaaaagcgaaagcagaa	
29B	agcgggacgaaagatgattacctagcggggtgagagagacccaaaaagcgaaagcagaa	
e19	agcgggacgaaagatgattacctagcggggtgagagagacccaaaaagcgaaagcagaa	
74e19	agcgggacgaaagatgattacctagcggggtgagagagacccaaaaagcgaaagcagaa	
74	agcgggacgaaagatgattacctagcggggtgagagagacccaaaaagcgaaagcagaa	
211	agcgggacgaaagatgattacctagcggggtgagagagacccaaaaagcgaaagcagaa	
246	agcgggacgaaagatgattacctagcggggtgagagagacccaaaaagcgaaagcagaa	
353	agcgggacgaaagatgattacctagcggggtgagagagacccaaaaagcgaaagcagaa	

DAH	ctcgtggaaagctgaaaatttggggaaacaatttcataatacgaatagaaacaccgcaat	
29B	ctcgtggaaagctgaaaatttggggaaacaatttcataatacgaatagaaacaccgcaat	
e19	ctcgtggaaagctgaaaatttggggaaacaatttcataatacgaatagaaacaccgcaat	
74e19	ctcgtggaaagctgaaaatttggggaaacaatttcataatacgaatagaaacaccgcaat	
74	ctcgtggaaagctgaaaatttggggaaacaatttcataatacgaatagaaacaccgcaat	
211	ctcgtggaaagctgaaaatttggggaaacaatttcataatacgaatagaaacaccgcaat	
246	ctcgtggaaagctgaaaatttggggaaacaatttcataatacgaatagaaacaccgcaat	
353	ctcgtggaaagctgaaaatttggggaaacaatttcataatacgaatagaaacaccgcaat	

DAH	ATG TTCAATATGCCACGGGAGTGACAAAAAGTAAATCCAAGCGTGGAAAATTAAGATG	60
29B	ATG TTCAATATGCCACGGGAGTGACAAAAAGTAAATCCAAGCGTGGAAAATTAAGATG	60
e19	ATG TTCAATATGCCACGGGAGTGACAAAAAGTAAATCCAAGCGTGGAAAATTAAGATG	60
74e19	ATG TTCAATATGCCACGGGAGTGACAAAAAGTAAATCCAAGCGTGGAAAATTAAGATG	60
74	ATG TTCAATATGCCACGGGAGTGACAAAAAGTAAATCCAAGCGTGGAAAATTAAGATG	60
211	ATG TTCAATATGCCACGGGAGTGACAAAAAGTAAATCCAAGCGTGGAAAATTAAGATG	60
246	ATG TTCAATATGCCACGGGAGTGACAAAAAGTAAATCCAAGCGTGGAAAATTAAGATG	60
353	ATG TTCAATATGCCACGGGAGTGACAAAAAGTAAATCCAAGCGTGGAAAATTAAGATG	60

DAH	GAAAACGATATGGCAGCAGCAACAACAACAGCCTGCACGCTTGACACATTGTGTT	120
29B	GAAAACGATATGGCAGCAGCAACAACAACAGCCTGCACGCTTGACACATTGTGTT	120
e19	GAAAACGATATGGCAGCAGCAACAACAACAGCCTGCACGCTTGACACATTGTGTT	120
74e19	GAAAACGATATGGCAGCAGCAACAACAACAGCCTGCACGCTTGACACATTGTGTT	120
74	GAAAACGATATGGCAGCAGCAACAACAACAGCCTGCACGCTTGACACATTGTGTT	120
211	GAAAACGATATGGCAGCAGCAACAACAACAGCCTGCACGCTTGACACATTGTGTT	120
246	GAAAACGATATGGCAGCAGCAACAACAACAGCCTGCACGCTTGACACATTGTGTT	120
353	GAAAACGATATGGCAGCAGCAACAACAACAGCCTGCACGCTTGACACATTGTGTT	120

DAH	TTGTGCCGGCAAGAAATGTTGCTGGATACATGTTGCTGCCGGCAAGCAGTAGAACAGTT	180
29B	TTGTGCCGGCAAGAAATGTTGCTGGATACATGTTGCTGCCGGCAAGCAGTAGAACAGTT	180
e19	TTGTGCCGGCAAGAAATGTTGCTGGATACATGTTGCTGCCGGCAAGCAGTAGAACAGTT	180
74e19	TTGTGCCGGCAAGAAATGTTGCTGGATACATGTTGCTGCCGGCAAGCAGTAGAACAGTT	180
74	TTGTGCCGGCAAGAAATGTTGCTGGATACATGTTGCTGCCGGCAAGCAGTAGAACAGTT	180
211	TTGTGCCGGCAAGAAATGTTGCTGGATACATGTTGCTGCCGGCAAGCAGTAGAACAGTT	180
246	TTGTGCCGGCAAGAAATGTTGCTGGATACATGTTGCTGCCGGCAAGCAGTAGAACAGTT	180
353	TTGTGCCGGCAAGAAATGTTGCTGGATACATGTTGCTGCCGGCAAGCAGTAGAACAGTT	180

DAH	GACAGCCCCGCAAGCAGTGAAGAAGCGTATAGCAGTAGAACAGCAGCAGCTGTCAAGCA	240
29B	GACAGCCCCGCAAGCAGTGAAGAAGCGTATAGCAGTAGAACAGCAGCAGCTGTCAAGCA	240
e19	GACAGCCCCGCAAGCAGTGAAGAAGCGTATAGCAGTAGAACAGCAGCAGCTGTCAAGCA	240
74e19	GACAGCCCCGCAAGCAGTGAAGAAGCGTATAGCAGTAGAACAGCAGCAGCTGTCAAGCA	240
74	GACAGCCCCGCAAGCAGTGAAGAAGCGTATAGCAGTAGAACAGCAGCAGCTGTCAAGCA	240
211	GACAGCCCCGCAAGCAGTGAAGAAGCGTATAGCAGTAGAACAGCAGCAGCTGTCAAGCA	240
246	GACAGCCCCGCAAGCAGTGAAGAAGCGTATAGCAGTAGAACAGCAGCAGCTGTCAAGCA	240
353	GACAGCCCCGCAAGCAGTGAAGAAGCGTATAGCAGTAGAACAGCAGCAGCTGTCAAGCA	240

DAH	AGCAGTGAAATCAGTGC GGAGGGAGGTCTGGTTCTCAGTCATGATGATATCGTACTGTGC	300
29B	AGCAGTGAAATCAGTGC GGAGGGAGGTCTGGTTCTCAGTCATGATGATATCGTACTGTGC	300
e19	AGCAGTGAAATCAGTGC GGAGGGAGGTCTGGTTCTCAGTCATGATGATATCGTACTGTGC	300
74e19	AGCAGTGAAATCAGTGC GGAGGGAGGTCTGGTTCTCAGTCATGATGATATCGTACTGTGC	300
74	AGCAGTGAAATCAGTGC GGAGGGAGGTCTGGTTCTCAGTCATGATGATATCGTACTGTGC	300
211	AGCAGTGAAATCAGTGC GGAGGGAGGTCTGGTTCTCAGTCATGATGATATCGTACTGTGC	300
246	AGCAGTGAAATCAGTGC GGAGGGAGGTCTGGTTCTCAGTCATGATGATATCGTACTGTGC	300
353	AGCAGTGAAATCAGTGC GGAGGGAGGTCTGGTTCTCAGTCATGATGATATCGTACTGTGC	300

DAH	CGCAGACAAAATTGACGAAGTGGAGACGACGGGTAAGAGGGACGTTAAATGCAGC	360
29B	CGCAGACAAAATTGACGAAGTGGAGACGACGGGTAAGAGGGACGTTAAATGCAGC	360
e19	CGCAGACAAAATTGACGAAGTGGAGACGACGGGTAAGAGGGACGTTAAATGCAGC	360
74e19	CGCAGACAAAATTGACGAAGTGGAGACGACGGGTAAGAGGGACGTTAAATGCAGC	360
74	CGCAGACAAAATTGACGAAGTGGAGACGACGGGTAAGAGGGACGTTAAATGCAGC	360
211	CGCAGACAAAATTGACGAAGTGGAGACGACGGGTAAGAGGGACGTTAAATGCAGC	360
246	CGCAGACAAAATTGACGAAGTGGAGACGACGGGTAAGAGGGACGTTAAATGCAGC	360
353	CGCAGACAAAATTGACGAAGTGGAGACGACGGGTAAGAGGGACGTTAAATGCAGC	360

DAH	GGGCATCAGTGCAGCAATGAATGCAGCATGGCAGCACGAAAAACAATCGACAACAGCGC	420
29B	GGGCATCAGTGCAGCAATGAATGCAGCATGGCAGCACGAAAAACAATCGACAACAGCGC	420
e19	GGGCATCAGTGCAGCAATGAATGCAGCATGGCAGCACGAAAAACAATCGACAACAGCGC	420
74e19	GGGCATCAGTGCAGCAATGAATGCAGCATGGCAGCACGAAAAACAATCGACAACAGCGC	420
74	GGGCATCAGTGCAGCAATGAATGCAGCATGGCAGCACGAAAAACAATCGACAACAGCGC	420
211	GGGCATCAGTGCAGCAATGAATGCAGCATGGCAGCACGAAAAACAATCGACAACAGCGC	420
246	GGGCATCAGTGCAGCAATGAATGCAGCATGGCAGCACGAAAAACAATCGACAACAGCGC	420
353	GGGCATCAGTGCAGCAATGAATGCAGCATGGCAGCACGAAAAACAATCGACAACAGCGC	420

DAH	GAAAACTTCAATATCTTAGCAACTGTCAACATATTGCGAACATTGCAATCGCTGCTG	480
29B	GAAAACTTCAATATCTTAGCAACTGTCAACATATTGCGAACATTGCAATCGCTGCTG	480
e19	GAAAACTTCAATATCTTAGCAACTGTCAACATATTGCGAACATTGCAATCGCTGCTG	480
74e19	GAAAACTTCAATATCTTAGCAACTGTCAACATATTGCGAACATTGCAATCGCTGCTG	480
74	GAAAACTTCAATATCTTAGCAACTGTCAACATATTGCGAACATTGCAATCGCTGCTG	480
211	GAAAACTTCAATATCTTAGCAACTGTCAACATATTGCGAACATTGCAATCGCTGCTG	480
246	GAAAACTTCAATATCTTAGCAACTGTCAACATATTGCGAACATTGCAATCGCTGCTG	480
353	GAAAACTTCAATATCTTAGCAACTGTCAACATATTGCGAACATTGCAATCGCTGCTG	480

DAH	CTGCTCATGTTCAATTGCGGCATTTCACAAAGCGACGCAGGCCAGCATCAGCAGCAG	540
29B	CTGCTCATGTTCAATTGCGGCATTTCACAAAGCGACGCAGGCCAGCATCAGCAGCAG	540
e19	CTGCTCATGTTCAATTGCGGCATTTCACAAAGCGACGCAGGCCAGCATCAGCAGCAG	540
74e19	CTGCTCATGTTCAATTGCGGCATTTCACAAAGCGACGCAGGCCAGCATCAGCAGCAG	540
74	CTGCTCATGTTCAATTGCGGCATTTCACAAAGCGACGCAGGCCAGCATCAGCAGCAG	540
211	CTGCTCATGTTCAATTGCGGCATTTCACAAAGCGACGCAGGCCAGCATCAGCAGCAG	540
246	CTGCTCATGTTCAATTGCGGCATTTCACAAAGCGACGCAGGCCAGCATCAGCAGCAG	540
353	CTGCTCATGTTCAATTGCGGCATTTCACAAAGCGACGCAGGCCAGCATCAGCAGCAG	540

DAH	CATCATCATCATTATCAGCATCATCAGCATCATCAGCAGCATTTCAGCGCAGCAA	600
29B	CATCATCATCATTATCAGCATCATCAGCATCATCAGCAGCATTTCAGCGCAGCAA	600
e19	CATCATCATCATTATCAGCATCATCAGCATCATCAGCAGCATTTCAGCGCAGCAA	600
74e19	CATCATCATCATTATCAGCATCATCAGCATCATCAGCAGCATTTCAGCGCAGCAA	600
74	CATCATCATCATTATCAGCATCATCAGCATCATCAGCAGCATTTCAGCGCAGCAA	600
211	CATCATCATCATTATCAGCATCATCAGCATCATCAGCAGCATTTCAGCGCAGCAA	600
246	CATCATCATCATTATCAGCATCATCAGCATCATCAGCAGCATTTCAGCGCAGCAA	600
353	CATCATCATCATTATCAGCATCATCAGCATCATCAGCAGCATTTCAGCGCAGCAA	600

DAH	GCCAATGTTAGTTACACAAAATTCTATTGCTGCTACAAACACTGGCAGCAGCAACCACA	660
29B	GCCAATGTTAGTTACACAAAATTCTATTGCTGCTACAAACACTGGCAGCAGCAACCACA	660
e19	GCCAATGTTAGTTACACAAAATTCTATTGCTGCTACAAACACTGGCAGCAGCAACCACA	660
74e19	GCCAATGTTAGTTACACAAAATTCTATTGCTGCTACAAACACTGGCAGCAGCAACCACA	660
74	GCCAATGTTAGTTACACAAAATTCTATTGCTGCTACAAACACTGGCAGCAGCAACCACA	660
211	GCCAATGTTAGTTACACAAAATTCTATTGCTGCTACAAACACTGGCAGCAGCAACCACA	660
246	GCCAATGTTAGTTACACAAAATTCTATTGCTGCTACAAACACTGGCAGCAGCAACCACA	660
353	GCCAATGTTAGTTACACAAAATTCTATTGCTGCTACAAACACTGGCAGCAGCAACCACA	660

DAH	AGACTGAGTTAACGCCCTAAAAACTACAAACAACAACACTACAGCATAACCAACAG	720
29B	AGACTGAGTTAACGCCCTAAAAACTACAAACAACAACACTACAGCATAACCAACAG	720
e19	AGACTGAGTTAACGCCCTAAAAACTACAAACAACAACACTACAGCATAACCAACAG	720
74e19	AGACTGAGTTAACGCCCTAAAAACTACAAACAACAACACTACAGCATAACCAACAG	720
74	AGACTGAGTTAACGCCCTAAAAACTACAAACAACAACACTACAGCATAACCAACAG	720
211	AGACTGAGTTAACGCCCTAAAAACTACAAACAACAACACTACAGCATAACCAACAG	720
246	AGACTGAGTTAACGCCCTAAAAACTACAAACAACAACACTACAGCATAACCAACAG	720
353	AGACTGAGTTAACGCCCTAAAAACTACAAACAACAACACTACAGCATAACCAACAG	720

DAH	CTGCCACGTGCCACACCGCAACAAAGCAACAGAGAAAGATAGGCATAAGTGTTCAC	780
29B	CTGCCACGTGCCACACCGCAACAAAGCAACAGAGAAAGATAGGCATAAGTGTTCAC	780
e19	CTGCCACGTGCCACACCGCAACAAAGCAACAGAGAAAGATAGGCATAAGTGTTCAC	780
74e19	CTGCCACGTGCCACACCGCAACAAAGCAACAGAGAAAGATAGGCATAAGTGTTCAC	780
74	CTGCCACGTGCCACACCGCAACAAAGCAACAGAGAAAGATAGGCATAAGTGTTCAC	780
211	CTGCCACGTGCCACACCGCAACAAAGCAACAGAGAAAGATAGGCATAAGTGTTCAC	780
246	CTGCCACGTGCCACACCGCAACAAAGCAACAGAGAAAGATAGGCATAAGTGTTCAC	780
353	CTGCCACGTGCCACACCGCAACAAAGCAACAGAGAAAGATAGGCATAAGTGTTCAC	780

DAH	TACAAGCACAATTACTCTACTCGCTGGCATTAGCCTTACTCTTTATCCTACTGGCC	840
29B	TACAAGCACAATTACTCTACTCGCTGGCATTAGCCTTACTCTTTATCCTACTGGCC	840
e19	TACAAGCACAATTACTCTACTCGCTGGCATTAGCCTTACTCTTTATCCTACTGGCC	840
74e19	TACAAGCACAATTACTCTACTCGCTGGCATTAGCCTTACTCTTTATCCTACTGGCC	840
74	TACAAGCACAATTACTCTACTCGCTGGCATTAGCCTTACTCTTTATCCTACTGGCC	840
211	TACAAGCACAATTACTCTACTCGCTGGCATTAGCCTTACTCTTTATCCTACTGGCC	840
246	TACAAGCACAATTACTCTACTCGCTGGCATTAGCCTTACTCTTTATCCTACTGGCC	840
353	TACAAGCACAATTACTCTACTCGCTGGCATTAGCCTTACTCTTTATCCTACTGGCC	840

DAH	AACACATTGGCCATCCAAGCGGTGTTGCCAGCACATCAGCAGCACCTGTCACAAT	900
29B	AACACATTGGCCATCCAAGCGGTGTTGCCAGCACATCAGCAGCACCTGTCACAAT	900
e19	AACACATTGGCCATCCAAGCGGTGTTGCCAGCACATCAGCAGCACCTGTCACAAT	900
74e19	AACACATTGGCCATCCAAGCGGTGTTGCCAGCACATCAGCAGCACCTGTCACAAT	900
74	AACACATTGGCCATCCAAGCGGTGTTGCCAGCACATCAGCAGCACCTGTCACAAT	900
211	AACACATTGGCCATCCAAGCGGTGTTGCCAGCACATCAGCAGCACCTGTCACAAT	900
246	AACACATTGGCCATCCAAGCGGTGTTGCCAGCACATCAGCAGCACCTGTCACAAT	900
353	AACACATTGGCCATCCAAGCGGTGTTGCCAGCACATCAGCAGCACCTGTCACAAT	900

52		
DAH	GATATAGCCGATGGACTGGATAAAACAGCGCTTCGGTGTGGGACGCAATCGCAGTGG	960
29B	GATATAGCCGATGGACTGGATAAAACAGCGCTTCGGTGTGGGACGCAATCGCAGTGG	960
e19	GATATAGCCGATGGACTGGATAAAACAGCGCTTCGGTGTGGGACGCAATCGCAGTGG	960
74e19	GATATAGCCGATGGACTGGATAAAACAGCGCTTCGGTGTGGGACGCAATCGCAGTGG	960
74	GATATAGCCGATGGACTGGATAAAACAGCGCTTCGGTGTGGGACGCAATCGCAGTGG	960
211	GATATAGCCGATGGACTGGATAAAACAGCGCTTCGGTGTGGGACGCAATCGCAGTGG	960
246	GATATAGCCGATGGACTGGATAAAACAGCGCTTCGGTGTGGGACGCAATCGCAGTGG	960
353	GATATAGCCGATGGACTGGATAAAACAGCGCTTCGGTGTGGGACGCAATCGCAGTGG	960

DAH	ACAAGGAGCGAATCAAACCAACAATGCGACTGTCACAAAATGTAAAACgtatgtattac	1020
29B	ACAAGGAGCGAATCAAACCAACAATGCGACTGTCACAAAATGTAAAACgtatgtattac	1020
e19	ACAAGGAGCGAATCAAACCAACAATGCGACTGTCACAAAATGTAAAACgtatgtattac	1020
74e19	ACAAGGAGCGAATCAAACCAACAATGCGACTGTCACAAAATGTAAAACgtatgtattac	1020
74	ACAAGGAGCGAATCAAACCAACAATGCGACTGTCACAAAATGTAAAACgtatgtattac	1020
211	ACAAGGAGCGAATCAAACCAACAATGCGACTGTCACAAAATGTAAAACgtatgtattac	1020
246	ACAAGGAGCGAATCAAACCAACAATGCGACTGTCACAAAATGTAAAACgtatgtattac	1020
353	ACAAGGAGCGAATCAAACCAACAATGCGACTGTCACAAAATGTAAAACgtatgtattac	1020

DAH	ctatgatctacccatggataaaaggAACATTCCATTGCACTGAGACTGACAGATACTA	1080
29B	ctatgatctacccatggataaaaggAACATTCCATTGCACTGAGACTGACAGATACTA	1080
e19	ctatgatctacccatggataaaaggAACATTCCATTGCACTGAGACTGACAGATACTA	1080
74e19	ctatgatctacccatggataaaaggAACATTCCATTGCACTGAGACTGACAGATACTA	1080
74	ctatgatctacccatggataaaaggAACATTCCATTGCACTGAGACTGACAGATACTA	1080
211	ctatgatctacccatggataaaaggAACATTCCATTGCACTGAGACTGACAGATACTA	1080
246	ctatgatctacccatggataaaaggAACATTCCATTGCACTGAGACTGACAGATACTA	1080
353	ctatgatctacccatggataaaaggAACATTCCATTGCACTGAGACTGACAGATACTA	1080

DAH	atagctttccatggataacaccttatacgTTGCAAATCCATGGACATCAGAACATGGTGT	1140
29B	atagctttccatggataacaccttatacgTTGCAAATCCATGGACATCAGAACATGGTGT	1140
e19	atagctttccatggataacaccttatacgTTGCAAATCCATGGACATCAGAACATGGTGT	1140
74e19	atagctttccatggataacaccttatacgTTGCAAATCCATGGACATCAGAACATGGTGT	1140
74	atagctttccatggataacaccttatacgTTGCAAATCCATGGACATCAGAACATGGTGT	1140
211	atagctttccatggataacaccttatacgTTGCAAATCCATGGACATCAGAACATGGTGT	1140
246	atagctttccatggataacaccttatacgTTGCAAATCCATGGACATCAGAACATGGTGT	1140
353	atagctttccatggataacaccttatacgTTGCAAATCCATGGACATCAGAACATGGTGT	1140

DAH	GCACCTCAATCAGCTGGAGAACTGCACGGTCATCGAGGGCTCCTGCTGATCGATTGAT	1200
29B	GCACCTCAATCAGCTGGAGAACTGCACGGTCATCGAGGGCTCCTGCTGATCGATTGAT	1200
e19	GCACCTCAATCAGCTGGAGAACTGCACGGTCATCGAGGGCTCCTGCTGATCGATTGAT	1200
74e19	GCACCTCAATCAGCTGGAGAACTGCACGGTCATCGAGGGCTCCTGCTGATCGATTGAT	1200
74	GCACCTCAATCAGCTGGAGAACTGCACGGTCATCGAGGGCTCCTGCTGATCGATTGAT	1200
211	GCACCTCAATCAGCTGGAGAACTGCACGGTCATCGAGGGCTCCTGCTGATCGATTGAT	1200
246	GCACCTCAATCAGCTGGAGAACTGCACGGTCATCGAGGGCTCCTGCTGATCGATTGAT	1200
353	GCACCTCAATCAGCTGGAGAACTGCACGGTCATCGAGGGCTCCTGCTGATCGATTGAT	1200

DAH	AAACGACGCCAGCCCTCTGAACAGAAGCTTCAAAACTGACCGAGGTCACAGATTATAT	1260
29B	AAACGACGCCAGCCCTCTGAACAGAAGCTTCAAAACTGACCGAGGTCACAGATTATAT	1260
e19	AAACGACGCCAGCCCTCTGAACAGAAGCTTCAAAACTGACCGAGGTCACAGATTATAT	1260
74e19	AAACGACGCCAGCCCTCTGAACAGAAGCTTCAAAACTGACCGAGGTCACAGATTATAT	1260
74	AAACGACGCCAGCCCTCTGAACAGAAGCTTCAAAACTGACCGAGGTCACAGATTATAT	1260
211	AAACGACGCCAGCCCTCTGAACAGAAGCTTCAAAACTGACCGAGGTCACAGATTATAT	1260
246	AAACGACGCCAGCCCTCTGAACAGAAGCTTCAAAACTGACCGAGGTCACAGATTATAT	1260
353	AAACGACGCCAGCCCTCTGAACAGAAGCTTCAAAACTGACCGAGGTCACAGATTATAT	1260

DAH	CATAATCTACCGTGTGACTGGATTGCACTCGCTGTCAAAGATCTTCCAATCTGAGCGT	1320
29B	CATAATCTACCGTGTGACTGGATTGCACTCGCTGTCAAAGATCTTCCAATCTGAGCGT	1320
e19	CATAATCTACCGTGTGACTGGATTGCACTCGCTGTCAAAGATCTTCCAATCTGAGCGT	1320
74e19	CATAATCTACCGTGTGACTGGATTGCACTCGCTGTCAAAGATCTTCCAATCTGAGCGT	1320
74	CATAATCTACCGTGTGACTGGATTGCACTCGCTGTCAAAGATCTTCCAATCTGAGCGT	1320
211	CATAATCTACCGTGTGACTGGATTGCACTCGCTGTCAAAGATCTTCCAATCTGAGCGT	1320
246	CATAATCTACCGTGTGACTGGATTGCACTCGCTGTCAAAGATCTTCCAATCTGAGCGT	1320
353	CATAATCTACCGTGTGACTGGATTGCACTCGCTGTCAAAGATCTTCCAATCTGAGCGT	1320

DAH	CATTAGGGAAACAAGCTTTCGACGGATATGCCTTGGTCGTCTACTCGAATTTCGACCT	1380
29B	CATTAGGGAAACAAGCTTTCGACGGATATGCCTTGGTCGTCTACTCGAATTTCGACCT	1380
e19	CATTAGGGAAACAAGCTTTCGACGGATATGCCTTGGTCGTCTACTCGAATTTCGACCT	1380
74e19	CATTAGGGAAACAAGCTTTCGACGGATATGCCTTGGTCGTCTACTCGAATTTCGACCT	1380
74	CATTAGGGAAACAAGCTTTCGACGGATATGCCTTGGTCGTCTACTCGAATTTCGACCT	1380
211	CATTAGGGAAACAAGCTTTCGACGGATATGCCTTGGTCGTCTACTCGAATTTCGACCT	1380
246	CATTAGGGAAACAAGCTTTCGACGGATATGCCTTGGTCGTCTACTCGAATTTCGACCT	1380
353	CATTAGGGAAACAAGCTTTCGACGGATATGCCTTGGTCGTCTACTCGAATTTCGACCT	1380

DAH	CATGGATTGGGACTTCACAAGCTACGATCCATAACCAGAGGC GGTTGCGGATTGAGAA	1440
29B	CATGGATTGGGACTTCACAAGCTACGATCCATAACCAGAGGC GGTTGCGGATTGAGAA	1440
e19	CATGGATTGGGACTTCACAAGCTACGATCCATAACCAGAGGC GGTTGCGGATTGAGAA	1440
74e19	CATGGATTGGGACTTCACAAGCTACGATCCATAACCAGAGGC GGTTGCGGATTGAGAA	1440
74	CATGGATTGGGACTTCACAAGCTACGATCCATAACCAGAGGC GGTTGCGGATTGAGAA	1440
211	CATGGATTGGGACTTCACAAGCTACGATCCATAACCAGAGGC GGTTGCGGATTGAGAA	1440
246	CATGGATTGGGACTTCACAAGCTACGATCCATAACCAGAGGC GGTTGCGGATTGAGAA	1440
353	CATGGATTGGGACTTCACAAGCTACGATCCATAACCAGAGGC GGTTGCGGATTGAGAA	1440

DAH	GAATCATAAGCTGTGCTATGATAGGACCATCGATTGGCTGGAAATTCTGGCGAAAACGA	1500
29B	GAATCATAAGCTGTGCTATGATAGGACCATCGATTGGCTGGAAATTCTGGCGAAAACGA	1500
e19	GAATCATAAGCTGTGCTATGATAGGACCATCGATTGGCTGGAAATTCTGGCGAAAACGA	1500
74e19	GAATCATAAGCTGTGCTATGATAGGACCATCGATTGGCTGGAAATTCTGGCGAAAACGA	1500
74	GAATCATAAGCTGTGCTATGATAGGACCATCGATTGGCTGGAAATTCTGGCGAAAACGA	1500
211	GAATCATAAGCTGTGCTATGATAGGACCATCGATTGGCTGGAAATTCTGGCGAAAACGA	1500
246	GAATCATAAGCTGTGCTATGATAGGACCATCGATTGGCTGGAAATTCTGGCGAAAACGA	1500
353	GAATCATAAGCTGTGCTATGATAGGACCATCGATTGGCTGGAAATTCTGGCGAAAACGA	1500

DAH	AACCCA ACTGGGTTGCTGACAGAGAACGGCAAGGAGAAGGGAGTG CAGGCTTCCAAGTG	1560
29B	AACCCA ACTGGGTTGCTGACAGAGAACGGCAAGGAGAAGGGAGTG CAGGCTTCCAAGTG	1560
e19	AACCCA ACTGGGTTGCTGACAGAGAACGGCAAGGAGAAGGGAGTG CAGGCTTCCAAGTG	1560
74e19	AACCCA ACTGGGTTGCTGACAGAGAACGGCAAGGAGAAGGGAGTG CAGGCTTCCAAGTG	1560
74	AACCCA ACTGGGTTGCTGACAGAGAACGGCAAGGAGAAGGGAGTG CAGGCTTCCAAGTG	1560
211	AACCCA ACTGGGTTGCTGACAGAGAACGGCAAGGAGAAGGGAGTG CAGGCTTCCAAGTG	1560
246	AACCCA ACTGGGTTGCTGACAGAGAACGGCAAGGAGAAGGGAGTG CAGGCTTCCAAGTG	1560
353	AACCCA ACTGGGTTGCTGACAGAGAACGGCAAGGAGAAGGGAGTG CAGGCTTCCAAGTG	1560

DAH	CCC GGGGGAGATCAGAATTGAGGAGGGCACGATACCACGGCTATTGAGGGAGAGCTTAA	1620
29B	CCC GGGGGAGATCAGAATTGAGGAGGGCACGATACCACGGCTATTGAGGGAGAGCTTAA	1620
e19	CCC GGGGGAGATCAGAATTGAGGAGGGCACGATACCACGGCTATTGAGGGAGAGCTTAA	1620
74e19	CCC GGGGGAGATCAGAATTGAGGAGGGCACGATACCACGGCTATTGAGGGAGAGCTTAA	1620
74	CCC GGGGGAGATCAGAATTGAGGAGGGCACGATACCACGGCTATTGAGGGAGAGCTTAA	1620
211	CCC GGGGGAGATCAGAATTGAGGAGGGCACGATACCACGGCTATTGAGGGAGAGCTTAA	1620
246	CCC GGGGGAGATCAGAATTGAGGAGGGCACGATACCACGGCTATTGAGGGAGAGCTTAA	1620
353	CCC GGGGGAGATCAGAATTGAGGAGGGCACGATACCACGGCTATTGAGGGAGAGCTTAA	1620

DAH	TGCCAGTTGTCAGCTGCACAATAATAGGCCTGTGCTGGAACAGCAAACCTGCCAGAC	1680
29B	TGCCAGTTGTCAGCTGCACAATAATAGGCCTGTGCTGGAACAGCAAACCTGCCAGAC	1680
e19	TGCCAGTTGTCAGCTGCACAATAATAGGCCTGTGCTGGAACAGCAAACCTGCCAGAC	1680
74e19	TGCCAGTTGTCAGCTGCACAATAATAGGCCTGTGCTGGAACAGCAAACCTGCCAGAC	1680
74	TGCCAGTTGTCAGCTGCACAATAATAGGCCTGTGCTGGAACAGCAAACCTGCCAGAC	1680
211	TGCCAGTTGTCAGCTGCACAATAATAGGCCTGTGCTGGAACAGCAAACCTGCCAGAC	1680
246	TGCCAGTTGTCAGCTGCACAATAATAGGCCTGTGCTGGAACAGCAAACCTGCCAGAC	1680
353	TGCCAGTTGTCAGCTGCACAATAATAGGCCTGTGCTGGAACAGCAAACCTGCCAGAC	1680

DAH	GAgtagttggccggtgtaaagttataccgtttcttattaacattttcgctttttt	1740
29B	GAgtagttggccggtgtaaagttataccgtttcttattaacattttcgctttttt	1740
e19	GAgtagttggccggtgtaaagttataccgtttcttattaacattttcgctttttt	1740
74e19	GAgtagttggccggtgtaaagttataccgtttcttattaacattttcgctttttt	1740
74	GAgtagttggccggtgtaaagttataccgtttcttattaacattttcgctttttt	1740
211	GAgtagttggccggtgtaaagttataccgtttcttattaacattttcgctttttt	1740
246	GAgtagttggccggtgtaaagttataccgtttcttattaacattttcgctttttt	1740
353	GAgtagttggccggtgtaaagttataccgtttcttattaacattttcgctttttt	1740

DAH	cttcaggAAATGCCCTGAAAAGTGCAGAAATAACTGCATCGATGAGCACACCTGCTGCA	1800
29B	cttcaggAAATGCCCTGAAAAGTGCAGAAATAACTGCATCGATGAGCACACCTGCTGCA	1800
e19	cttcaggAAATGCCCTGAAAAGTGCAGAAATAACTGCATCGATGAGCACACCTGCTGCA	1800
74e19	cttcaggAAATGCCCTGAAAAGTGCAGAAATAACTGCATCGATGAGCACACCTGCTGCA	1800
74	cttcaggAAATGCCCTGAAAAGTGCAGAAATAACTGCATCGATGAGCACACCTGCTGCA	1800
211	cttcaggAAATGCCCTGAAAAGTGCAGAAATAACTGCATCGATGAGCACACCTGCTGCA	1800
246	cttcaggAAATGCCCTGAAAAGTGCAGAAATAACTGCATCGATGAGCACACCTGCTGCA	1800
353	cttcaggAAATGCCCTGAAAAGTGCAGAAATAACTGCATCGATGAGCACACCTGCTGCA	1800

DAH	GCCAGGATTGTTGGGTGGATGCGTGATCGATAAGAATGGAATGAGAGCTGCATCTCCT	1860
29B	GCCAGGATTGTTGGGTGGATGCGTGATCGATAAGAATGGAATGAGAGCTGCATCTCCT	1860
e19	GCCAGGATTGTTGGGTGGATGCGTGATCGATAAGAATGGAATGAGAGCTGCATCTCCT	1860
74e19	GCCAGGATTGTTGGGTGGATGCGTGATCGATAAGAATGGAATGAGAGCTGCATCTCCT	1860
74	GCCAGGATTGTTGGGTGGATGCGTGATCGATAAGAATGGAATGAGAGCTGCATCTCCT	1860
211	GCCAGGATTGTTGGGTGGATGCGTGATCGATAAGAATGGAATGAGAGCTGCATCTCCT	1860
246	GCCAGGATTGTTGGGTGGATGCGTGATCGATAAGAATGGAATGAGAGCTGCATCTCCT	1860
353	GCCAGGATTGTTGGGTGGATGCGTGATCGATAAGAATGGAATGAGAGCTGCATCTCCT	1860

DAH	GTCGAAATGTGTCTTCACAACATCTGTATGGACTCCTGTCGAAAGGCTATTATCAGg	1920
29B	GTCGAAATGTGTCTTCACAACATCTGTATGGACTCCTGTCGAAAGGCTATTATCAGg	1920
e19	GTCGAAATGTGTCTTCACAACATCTGTATGGACTCCTGTCGAAAGGCTATTATCAGg	1920
74e19	GTCGAAATGTGTCTTCACAACATCTGTATGGACTCCTGTCGAAAGGCTATTATCAGg	1920
74	GTCGAAATGTGTCTTCACAACATCTGTATGGACTCCTGTCGAAAGGCTATTATCAGg	1920
211	GTCGAAATGTGTCTTCACAACATCTGTATGGACTCCTGTCGAAAGGCTATTATCAGg	1920
246	GTCGAAATGTGTCTTCACAACATCTGTATGGACTCCTGTCGAAAGGCTATTATCAGg	1920
353	GTCGAAATGTGTCTTCACAACATCTGTATGGACTCCTGTCGAAAGGCTATTATCAGg	1920

DAH	taactaatgattcacttttagtataacaacaagttagctaacttcaaactctttcagTT	1980
29B	taactaatgattcacttttagtataacaacaagttagctaacttcaaactctttcagTT	1980
e19	taactaatgattcacttttagtataacaacaagttagctaacttcaaactctttcagTT	1980
74e19	taactaatgattcacttttagtataacaacaagttagctaacttcaaactctttcagTT	1980
74	taactaatgattcacttttagtataacaacaagttagctaacttcaaactctttcagTT	1980
211	taactaatgattcacttttagtataacaacaagttagctaacttcaaactctttcagTT	1980
246	taactaatgattcacttttagtataacaacaagttagctaacttcaaactctttcagTT	1980
353	taactaatgattcacttttagtataacaacaagttagctaacttcaaactctttcagTT	1980

DAH	CGACAGCCGCTCGTAACGGCGAACGAGTCATCACACTGACAAAGTTGAAACGAACAG	2040
29B	CGACAGCCGCTCGTAACGGCGAACGAGTCATCACACTGACAAAGTTGAAACGAACAG	2040
e19	CGACAGCCGCTCGTAACGGCGAACGAGTCATCACACTGACAAAGTTGAAACGAACAG	2040
74e19	CGACAGCCGCTCGTAACGGCGAACGAGTCATCACACTGACAAAGTTGAAACGAACAG	2040
74	CGACAGCCGCTCGTAACGGCGAACGAGTCATCACACTGACAAAGTTGAAACGAACAG	2040
211	CGACAGCCGCTCGTAACGGCGAACGAGTCATCACACTGACAAAGTTGAAACGAACAG	2040
246	CGACAGCCGCTCGTAACGGCGAACGAGTCATCACACTGACAAAGTTGAAACGAACAG	2040
353	CGACAGCCGCTCGTAACGGCGAACGAGTCATCACACTGACAAAGTTGAAACGAACAG	2040

DAH	TGTGTATTCCGGTATTCCATACAACGGACAATGTATCACCACGTCCAACGGGGTACCA	2100
29B	TGTGTATTCCGGTATTCCATACAACGGACAATGTATCACCACGTCCAACGGGGTACCA	2100
e19	TGTGTATTCCGGTATTCCATACAACGGACAATGTATCACCACGTCCAACGGGGTACCA	2100
74e19	TGTGTATTCCGGTATTCCATACAACGGACAATGTATCACCACGTCCAACGGGGTACCA	2100
74	TGTGTATTCCGGTATTCCATACAACGGACAATGTATCACCACGTCCAACGGGGTACCA	2100
211	TGTGTATTCCGGTATTCCATACAACGGACAATGTATCACCACGTCCAACGGGGTACCA	2100
246	TGTGTATTCCGGTATTCCATACAACGGACAATGTATCACCACGTCCAACGGGGTACCA	2100
353	TGTGTATTCCGGTATTCCATACAACGGACAATGTATCACCACGTCCAACGGGGTACCA	2100

DAH	GAAGTCAGAGAACAGCGATGTGCGAACCTTGTCCGGCGGCAAGTGTGACAAGGAGTG	2160
29B	GAAGTCAGAGAACAGCGATGTGCGAACCTTGTCCGGCGGCAAGTGTGACAAGGAGTG	2160
e19	GAAGTCAGAGAACAGCGATGTGCGAACCTTGTCCGGCGGCAAGTGTGACAAGGAGTG	2160
74e19	GAAGTCAGAGAACAGCGATGTGCGAACCTTGTCCGGCGGCAAGTGTGACAAGGAGTG	2160
74	GAAGTCAGAGAACAGCGATGTGCGAACCTTGTCCGGCGGCAAGTGTGACAAGGAGTG	2160
211	GAAGTCAGAGAACAGCGATGTGCGAACCTTGTCCGGCGGCAAGTGTGACAAGGAGTG	2160
246	GAAGTCAGAGAACAGCGATGTGCGAACCTTGTCCGGCGGCAAGTGTGACAAGGAGTG	2160
353	GAAGTCAGAGAACAGCGATGTGCGAACCTTGTCCGGCGGCAAGTGTGACAAGGAGTG	2160

DAH	CTCCTCCGGTCTTATCGACAGTTGGAGCGTGCCTCGGAGTTCCACGGCTGCACCATTAT	2220
29B	CTCCTCCGGTCTTATCGACAGTTGGAGCGTGCCTCGGAGTTCCACGGCTGCACCATTAT	2220
e19	CTCCTCCGGTCTTATCGACAGTTGGAGCGTGCCTCGGAGTTCCACGGCTGCACCATTAT	2220
74e19	CTCCTCCGGTCTTATCGACAGTTGGAGCGTGCCTCGGAGTTCCACGGCTGCACCATTAT	2220
74	CTCCTCCGGTCTTATCGACAGTTGGAGCGTGCCTCGGAGTTCCACGGCTGCACCATTAT	2220
211	CTCCTCCGGTCTTATCGACAGTTGGAGCGTGCCTCGGAGTTCCACGGCTGCACCATTAT	2220
246	CTCCTCCGGTCTTATCGACAGTTGGAGCGTGCCTCGGAGTTCCACGGCTGCACCATTAT	2220
353	CTCCTCCGGTCTTATCGACAGTTGGAGCGTGCCTCGGAGTTCCACGGCTGCACCATTAT	2220

DAH	AACCGGAACCGAGCCCCTACCATCAGCATTAAACGTGAAAGCGGGCGtaagtgttttt	2280
29B	AACCGGAACCGAGCCCCTACCATCAGCATTAAACGTGAAAGCGGGCGtaagtgttttt	2280
e19	AACCGGAACCGAGCCCCTACCATCAGCATTAAACGTGAAAGCGGGCGtaagtgttttt	2280
74e19	AACCGGAACCGAGCCCCTACCATCAGCATTAAACGTGAAAGCGGGCGtaagtgttttt	2280
74	AACCGGAACCGAGCCCCTACCATCAGCATTAAACGTGAAAGCGGGCGtaagtgttttt	2280
211	AACCGGAACCGAGCCCCTACCATCAGCATTAAACGTGAAAGCGGGCGtaagtgttttt	2280
246	AACCGGAACCGAGCCCCTACCATCAGCATTAAACGTGAAAGCGGGCGtaagtgttttt	2280
353	AACCGGAACCGAGCCCCTACCATCAGCATTAAACGTGAAAGCGGGCGtaagtgttttt	2280

DAH	gctgctcaaattaaagtataatctctatattaaaacttcttgttcttcagCTCACGTC	2340
29B	gctgctcaaattaaagtataatctctatattaaaacttcttgttcttcagCTCACGTC	2340
e19	gctgctcaaattaaagtataatctctatattaaaacttcttgttcttcagCTCACGTC	2340
74e19	gctgctcaaattaaagtataatctctatattaaaacttcttgttcttcagCTCACGTC	2340
74	gctgctcaaattaaagtataatctctatattaaaacttcttgttcttcagCTCACGTC	2340
211	gctgctcaaattaaagtataatctctatattaaaacttcttgttcttcagCTCACGTC	2340
246	gctgctcaaattaaagtataatctctatattaaaacttcttgttcttcagCTCACGTC	2340
353	gctgctcaaattaaagtataatctctatattaaaacttcttgttcttcagCTCACGTC	2340

DAH	ATGGATGAATTAAAATATGCCCTGGCTGCCGCCATAAAATTCACTCGTCCCTAATGGTT	2400
29B	ATGGATGAATTAAAATATGCCCTGGCTGCCGCCATAAAATTCACTCGTCCCTAATGGTT	2400
e19	ATGGATGAATTAAAATATGCCCTGGCTGCCGCCATAAAATTCACTCGTCCCTAATGGTT	2400
74e19	ATGGATGAATTAAAATATGCCCTGGCTGCCGCCATAAAATTCACTCGTCCCTAATGGTT	2400
74	ATGGATGAATTAAAATATGCCCTGGCTGCCGCCATAAAATTCACTCGTCCCTAATGGTT	2400
211	ATGGATGAATTAAAATATGCCCTGGCTGCCGCCATAAAATTCACTCGTCCCTAATGGTT	2400
246	ATGGATGAATTAAAATATGCCCTGGCTGCCGCCATAAAATTCACTCGTCCCTAATGGTT	2400
353	ATGGATGAATTAAAATATGCCCTGGCTGCCGCCATAAAATTCACTCGTCCCTAATGGTT	2400

DAH	CATTGACCTACGGATTGAAGTCCTTGAAATTCTTCATCCCTAAGTGAATTAGCGGC	2460
29B	CATTGACCTACGGATTGAAGTCCTTGAAATTCTTCATCCCTAAGTGAATTAGCGGC	2460
e19	CATTGACCTACGGATTGAAGTCCTTGAAATTCTTCATCCCTAAGTGAATTAGCGGC	2460
74e19	CATTGACCTACGGATTGAAGTCCTTGAAATTCTTCATCCCTAAGTGAATTAGCGGC	2460
74	CATTGACCTACGGATTGAAGTCCTTGAAATTCTTCATCCCTAAGTGAATTAGCGGC	2460
211	CATTGACCTACGGATTGAAGTCCTTGAAATTCTTCATCCCTAAGTGAATTAGCGGC	2460
246	CATTGACCTACGGATTGAAGTCCTTGAAATTCTTCATCCCTAAGTGAATTAGCGGC	2460
353	CATTGACCTACGGATTGAAGTCCTTGAAATTCTTCATCCCTAAGTGAATTAGCGGC	2460

DAH	GATCCGCCGATGGACGCGATAAAATATGCTTGTATGTGCTTGATAATCGCGATCTAGAT	2520
29B	GATCCGCCGATGGACGCGATAAAATATGCTTGTATGTGCTTGATAATCGCGATCTAGAT	2520
e19	GATCCGCCGATGGACGCGATAAAATATGCTTGTATGTGCTTGATAATCGCGATCTAGAT	2520
74e19	GATCCGCCGATGGACGCGATAAAATATGCTTGTATGTGCTTGATAATCGCGATCTAGAT	2520
74	GATCCGCCGATGGACGCGATAAAATATGCTTGTATGTGCTTGATAATCGCGATCTAGAT	2520
211	GATCCGCCGATGGACGCGATAAAATATGCTTGTATGTGCTTGATAATCGCGATCTAGAT	2520
246	GATCCGCCGATGGACGCGATAAAATATGCTTGTATGTGCTTGATAATCGCGATCTAGAT	2520
353	GATCCGCCGATGGACGCGATAAAATATGCTTGTATGTGCTTGATAATCGCGATCTAGAT	2520

DAH	GAGCTCTGGGGACCCAACCAAACGGTGTTCATTAGGAAGGGCGCGTCTTCTTCATTTC	2580
29B	GAGCTCTGGGGACCCAACCAAACGGTGTTCATTAGGAAGGGCGCGTCTTCTTCATTTC	2580
e19	GAGCTCTGGGGACCCAACCAAACGGTGTTCATTAGGAAGGGCGCGTCTTCTTCATTTC	2580
74e19	GAGCTCTGGGGACCCAACCAAACGGTGTTCATTAGGAAGGGCGCGTCTTCTTCATTTC	2580
74	GAGCTCTGGGGACCCAACCAAACGGTGTTCATTAGGAAGGGCGCGTCTTCTTCATTTC	2580
211	GAGCTCTGGGGACCCAACCAAACGGTGTTCATTAGGAAGGGCGCGTCTTCTTCATTTC	2580
246	GAGCTCTGGGGACCCAACCAAACGGTGTTCATTAGGAAGGGCGCGTCTTCTTCATTTC	2580
353	GAGCTCTGGGGACCCAACCAAACGGTGTTCATTAGGAAGGGCGCGTCTTCTTCATTTC	2580

DAH	AACCCAAAACTATGTGTGCCACCATTAACCAGTTGCTGCCCATGCTGGCCTCCAAGCCA	2640
29B	AACCCAAAACTATGTGTGCCACCATTAACCAGTTGCTGCCCATGCTGGCCTCCAAGCCA	2640
e19	AACCCAAAACTATGTGTGCCACCATTAACCAGTTGCTGCCCATGCTGGCCTCCAAGCCA	2640
74e19	AACCCAAAACTATGTGTGCCACCATTAACCAGTTGCTGCCCATGCTGGCCTCCAAGCCA	2640
74	AACCCAAAACTATGTGTGCCACCATTAACCAGTTGCTGCCCATGCTGGCCTCCAAGCCA	2640
211	AACCCAAAACTATGTGTGCCACCATTAACCAGTTGCTGCCCATGCTGGCCTCCAAGCCA	2640
246	AACCCAAAACTATGTGTGCCACCATTAACCAGTTGCTGCCCATGCTGGCCTCCAAGCCA	2640
353	AACCCAAAACTATGTGTGCCACCATTAACCAGTTGCTGCCCATGCTGGCCTCCAAGCCA	2640

DAH	AAGTTTTTGAAGTCAGATGTGGCGCAGACTCGAATGGAAACCGCGGATCATgtaaag	2700
29B	AAGTTTTTGAAGTCAGATGTGGCGCAGACTCGAATGGAAACCGCGGATCATgtaaag	2700
e19	AAGTTTTTGAAGTCAGATGTGGCGCAGACTCGAATGGAAACCGCGGATCATgtaaag	2700
74e19	AAGTTTTTGAAGTCAGATGTGGCGCAGACTCGAATGGAAACCGCGGATCATgtaaag	2700
74	AAGTTTTTGAAGTCAGATGTGGCGCAGACTCGAATGGAAACCGCGGATCATgtaaag	2700
211	AAGTTTTTGAAGTCAGATGTGGCGCAGACTCGAATGGAAACCGCGGATCATgtaaag	2700
246	AAGTTTTTGAAGTCAGATGTGGCGCAGACTCGAATGGAAACCGCGGATCATgtaaag	2700
353	AAGTTTTTGAAGTCAGATGTGGCGCAGACTCGAATGGAAACCGCGGATCATgtaaag	2700

DAH	taatcttaacgaagattttagtaaaattttaacttttcgtactaacctaataatgaaa	2760
29B	taatcttaacgaagattttagtaaaattttaacttttcgtactaacctaataatgaaa	2760
e19	taatcttaacgaagattttagtaaaattttaacttttcgtactaacctaataatgaaa	2760
74e19	taatcttaacgaagattttagtaaaattttaacttttcgtactaacctaataatgaaa	2760
74	taatcttaacgaagattttagtaaaattttaacttttcgtactaacctaataatgaaa	2760
211	taatcttaacgaagattttagtaaaattttaacttttcgtactaacctaataatgaaa	2760
246	taatcttaacgaagattttagtaaaattttaacttttcgtactaacctaataatgaaa	2760
353	taatcttaacgaagattttagtaaaattttaacttttcgtactaacctaataatgaaa	2760

DAH	catttccagGTGGAACAGCCGTTCTCAATGTCACATTACAATCAGTGGGAGCAAACCTCG	2820
29B	catttccagGTGGAACAGCCGTTCTCAATGTCACATTACAATCAGTGGGAGCAAACCTCG	2820
e19	catttccagGTGGAACAGCCGTTCTCAATGTCACATTACAATCAGTGGGAGCAAACCTCG	2820
74e19	catttccagGTGGAACAGCCGTTCTCAATGTCACATTACAATCAGTGGGAGCAAACCTCG	2820
74	catttccagGTGGAACAGCCGTTCTCAATGTCACATTACAATCAGTGGGAGCAAACCTCG	2820
211	catttccagGTGGAACAGCCGTTCTCAATGTCACATTACAATCAGTGGGAGCAAACCTCG	2820
246	catttccagGTGGAACAGCCGTTCTCAATGTCACATTACAATCAGTGGGAGCAAACCTCG	2820
353	catttccagGTGGAACAGCCGTTCTCAATGTCACATTACAATCAGTGGGAGCAAACCTCG	2820

DAH	CTATGCTAACGTCACGACAAAAGTTGAAATAGGAGAGCCCCAAAAGCCGAGCAATGCTA	2880
29B	CTATGCTAACGTCACGACAAAAGTTGAAATAGGAGAGCCCCAAAAGCCGAGCAATGCTA	2880
e19	CTATGCTAACGTCACGACAAAAGTTGAAATAGGAGAGCCCCAAAAGCCGAGCAATGCTA	2880
74e19	CTATGCTAACGTCACGACAAAAGTTGAAATAGGAGAGCCCCAAAAGCCGAGCAATGCTA	2880
74	CTATGCTAACGTCACGACAAAAGTTGAAATAGGAGAGCCCCAAAAGCCGAGCAATGCTA	2880
211	CTATGCTAACGTCACGACAAAAGTTGAAATAGGAGAGCCCCAAAAGCCGAGCAATGCTA	2880
246	CTATGCTAACGTCACGACAAAAGTTGAAATAGGAGAGCCCCAAAAGCCGAGCAATGCTA	2880
353	CTATGCTAACGTCACGACAAAAGTTGAAATAGGAGAGCCCCAAAAGCCGAGCAATGCTA	2880

DAH	CAATTGTTTTAAGGATCCCGCGCCTTCATCGGTTCTGTGTTTATCATATGATCGATC	2940
29B	CAATTGTTTTAAGGATCCCGCGCCTTCATCGGTTCTGTGTTTATCATATGATCGATC	2940
e19	CAATTGTTTTAAGGATCCCGCGCCTTCATCGGTTCTGTGTTTATCATATGATCGATC	2940
74e19	CAATTGTTTTAAGGATCCCGCGCCTTCATCGGTTCTGTGTTTATCATATGATCGATC	2940
74	CAATTGTTTTAAGGATCCCGCGCCTTCATCGGTTCTGTGTTTATCATATGATCGATC	2940
211	CAATTGTTTTAAGGATCCCGCGCCTTCATCGGTTCTGTGTTTATCATATGATCGATC	2940
246	CAATTGTTTTAAGGATCCCGCGCCTTCATCGGTTCTGTGTTTATCATATGATCGATC	2940
353	CAATTGTTTTAAGGATCCCGCGCCTTCATCGGTTCTGTGTTTATCATATGATCGATC	2940

DAH	CGTACGGGAACTCAACTAAAAGCAGTGACGATCCATGCGATGATCGCTGGAAGGTTAGCT	3000
29B	CGTACGGGAACTCAACTAAAAGCAGTGACGATCCATGCGATGATCGCTGGAAGGTTAGCT	3000
e19	CGTACGGGAACTCAACTAAAAGCAGTGACGATCCATGCGATGATCGCTGGAAGGTTAGCT	3000
74e19	CGTACGGGAACTCAACTAAAAGCAGTGACGATCCATGCGATGATCGCTGGAAGGTTAGCT	3000
74	CGTACGGGAACTCAACTAAAAGCAGTGACGATCCATGCGATGATCGCTGGAAGGTTAGCT	3000
211	CGTACGGGAACTCAACTAAAAGCAGTGACGATCCATGCGATGATCGCTGGAAGGTTAGCT	3000
246	CGTACGGGAACTCAACTAAAAGCAGTGACGATCCATGCGATGATCGCTGGAAGGTTAGCT	3000
353	CGTACGGGAACTCAACTAAAAGCAGTGACGATCCATGCGATGATCGCTGGAAGGTTAGCT	3000

DAH	CTCCGGAAAAGAGCGGGGTCATGGTATTAAGCAATTGATTCCGTACACTAACTACTCCT	3060
29B	CTCCGGAAAAGAGCGGGGTCATGGTATTAAGCAATTGATTCCGTACACTAACTACTCCT	3060
e19	CTCCGGAAAAGAGCGGGGTCATGGTATTAAGCAATTGATTCCGTACACTAACTACTCCT	3060
74e19	CTCCGGAAAAGAGCGGGGTCATGGTATTAAGCAATTGATTCCGTACACTAACTACTCCT	3060
74	CTCCGGAAAAGAGCGGGGTCATGGTATTAAGCAATTGATTCCGTACACTAACTACTCCT	3060
211	CTCCGGAAAAGAGCGGGGTCATGGTATTAAGCAATTGATTCCGTACACTAACTACTCCT	3060
246	CTCCGGAAAAGAGCGGGGTCATGGTATTAAGCAATTGATTCCGTACACTAACTACTCCT	3060
353	CTCCGGAAAAGAGCGGGGTCATGGTATTAAGCAATTGATTCCGTACACTAACTACTCCT	3060

DAH	ACTACGTTGGACCATGGCTATATCCTCGGAATTGACAACAGCGGGAGAGCGACGTGAAGA	3120
29B	ACTACGTTGGACCATGGCTATATCCTCGGAATTGACAACAGCGGGAGAGCGACGTGAAGA	3120
e19	ACTACGTTGGACCATGGCTATATCCTCGGAATTGACAACAGCGGGAGAGCGACGTGAAGA	3120
74e19	ACTACGTTGGACCATGGCTATATCCTCGGAATTGACAACAGCGGGAGAGCGACGTGAAGA	3120
74	ACTACGTTGGACCATGGCTATATCCTCGGAATTGACAACAGCGGGAGAGCGACGTGAAGA	3120
211	ACTACGTTGGACCATGGCTATATCCTCGGAATTGACAACAGCGGGAGAGCGACGTGAAGA	3120
246	ACTACGTTGGACCATGGCTATATCCTCGGAATTGACAACAGCGGGAGAGCGACGTGAAGA	3120
353	ACTACGTTGGACCATGGCTATATCCTCGGAATTGACAACAGCGGGAGAGCGACGTGAAGA	3120

DAH	ACTTTAGGACGAATCCCGGACGACCGTCAAAGGTTACGGAGGTGGTAGCAACGCCATT	3180
29B	ACTTTAGGACGAATCCCGGACGACCGTCAAAGGTTACGGAGGTGGTAGCAACGCCATT	3180
e19	ACTTTAGGACGAATCCCGGACGACCGTCAAAGGTTACGGAGGTGGTAGCAACGCCATT	3180
74e19	ACTTTAGGACGAATCCCGGACGACCGTCAAAGGTTACGGAGGTGGTAGCAACGCCATT	3180
74	ACTTTAGGACGAATCCCGGACGACCGTCAAAGGTTACGGAGGTGGTAGCAACGCCATT	3180
211	ACTTTAGGACGAATCCCGGACGACCGTCAAAGGTTACGGAGGTGGTAGCAACGCCATT	3180
246	ACTTTAGGACGAATCCCGGACGACCGTCAAAGGTTACGGAGGTGGTAGCAACGCCATT	3180
353	ACTTTAGGACGAATCCCGGACGACCGTCAAAGGTTACGGAGGTGGTAGCAACGCCATT	3180

DAH	CAGATTGAAAATTgtgagttatgtcaatagatgtttaggattataaatgat	3240
29B	CAGATTGAAAATTgtgagttatgtcaatagatgtttaggattataaatgat	3240
e19	CAGATTGAAAATTgtgagttatgtcaatagatgtttaggattataaatgat	3240
74e19	CAGATTGAAAATTgtgagttatgtcaatagatgtttaggattataaatgat	3240
74	CAGATTGAAAATTgtgagttatgtcaatagatgtttaggattataaatgat	3240
211	CAGATTGAAAATTgtgagttatgtcaatagatgtttaggattataaatgat	3240
246	CAGATTGAAAATTgtgagttatgtcaatagatgtttaggattataaatgat	3240
353	CAGATTGAAAATTgtgagttatgtcaatagatgtttaggattataaatgat	3240

DAH	tactaaatgtcaagtaattgttcacccgtactgtatcaacatatttattgcctttcagAACG	3300
29B	tactaaatgtcaagtaattgttcacccgtactgtatcaacatatttattgcctttcagAACG	3300
e19	tactaaatgtcaagtaattgttcacccgtactgtatcaacatatttattgcctttcagAACG	3300
74e19	tactaaatgtcaagtaattgttcacccgtactgtatcaacatatttattgcctttcagAACG	3300
74	tactaaatgtcaagtaattgttcacccgtactgtatcaacatatttattgcctttcagAACG	3300
211	tactaaatgtcaagtaattgttcacccgtactgtatcaacatatttattgcctttcagAACG	3300
246	tactaaatgtcaagtaattgttcacccgtactgtatcaacatatttattgcctttcagAACG	3300
353	tactaaatgtcaagtaattgttcacccgtactgtatcaacatatttattgcctttcagAACG	3300

DAH	TAACATGGAGCTACCTAGATAAGCCTTATGGCGTGCTAACCGCCTATTTTATAAAAGCCA	3360
29B	TAACATGGAGCTACCTAGATAAGCCTTATGGCGTGCTAACCGCCTATTTTATAAAAGCCA	3360
e19	TAACATGGAGCTACCTAGATAAGCCTTATGGCGTGCTAACCGCCTATTTTATAAAAGCCA	3360
74e19	TAACATGGAGCTACCTAGATAAGCCTTATGGCGTGCTAACCGCCTATTTTATAAAAGCCA	3360
74	TAACATGGAGCTACCTAGATAAGCCTTATGGCGTGCTAACCGCCTATTTTATAAAAGCCA	3360
211	TAACATGGAGCTACCTAGATAAGCCTTATGGCGTGCTAACCGCCTATTTTATAAAAGCCA	3360
246	TAACATGGAGCTACCTAGATAAGCCTTATGGCGTGCTAACCGCCTATTTTATAAAAGCCA	3360
353	TAACATGGAGCTACCTAGATAAGCCTTATGGCGTGCTAACCGCCTATTTTATAAAAGCCA	3360

DAH	AACTTATAAATCGGCCTACTCGAAACAATAACCGGGATTACTGTACTGAACgtaaagcaac	3420
29B	AACTTATAAATCGGCCTACTCGAAACAATAACCGGGATTACTGTACTGAACgtaaagcaac	3420
e19	AACTTATAAATCGGCCTACTCGAAACAATAACCGGGATTACTGTACTGAACgtaaagcaac	3420
74e19	AACTTATAAATCGGCCTACTCGAAACAATAACCGGGATTACTGTACTGAACgtaaagcaac	3420
74	AACTTATAAATCGGCCTACTCGAAACAATAACCGGGATTACTGTACTGAACgtaaagcaac	3420
211	AACTTATAAATCGGCCTACTCGAAACAATAACCGGGATTACTGTACTGAACgtaaagcaac	3420
246	AACTTATAAATCGGCCTACTCGAAACAATAACCGGGATTACTGTACTGAACgtaaagcaac	3420
353	AACTTATAAATCGGCCTACTCGAAACAATAACCGGGATTACTGTACTGAACgtaaagcaac	3420

DAH	aacatagccaaattgtatccataattaattaatthaatattcatctgtgttagCTCTCG	3480
29B	aacatagccaaattgtatccataattaattaatthaatattcatctgtgttagCTCTCG	3480
e19	aacatagccaaattgtatccataattaattaatthaatattcatctgtgttagCTCTCG	3480
74e19	aacatagccaaattgtatccataattaattaatthaatattcatctgtgttagCTCTCG	3480
74	aacatagccaaattgtatccataattaattaatthaatattcatctgtgttagCTCTCG	3480
211	aacatagccaaattgtatccataattaattaatthaatattcatctgtgttagCTCTCG	3480
246	aacatagccaaattgtatccataattaattaatthaatattcatctgtgttagCTCTCG	3480
353	aacatagccaaattgtatccataattaattaatthaatattcatctgtgttagCTCTCG	3480

DAH	TCAAGGCCATGGAAAATGACCTGCCAGCCACAACGCCTACCAAGAAAATATCAGATCCTT	3540
29B	TCAAGGCCATGGAAAATGACCTGCCAGCCACAACGCCTACCAAGAAAATATCAGATCCTT	3540
e19	TCAAGGCCATGGAAAATGACCTGCCAGCCACAACGCCTACCAAGAAAATATCAGATCCTT	3540
74e19	TCAAGGCCATGGAAAATGACCTGCCAGCCACAACGCCTACCAAGAAAATATCAGATCCTT	3540
74	TCAAGGCCATGGAAAATGACCTGCCAGCCACAACGCCTACCAAGAAAATATCAGATCCTT	3540
211	TCAAGGCCATGGAAAATGACCTGCCAGCCACAACGCCTACCAAGAAAATATCAGATCCTT	3540
246	TCAAGGCCATGGAAAATGACCTGCCAGCCACAACGCCTACCAAGAAAATATCAGATCCTT	3540
353	TCAAGGCCATGGAAAATGACCTGCCAGCCACAACGCCTACCAAGAAAATATCAGATCCTT	3540

DAH	TAGCAGGCAGCTGTAAGTGCCTGGAGGGTTCGAAGAAGACTAGCAGTCAGGAATACGATG	3600
29B	TAGCAGGCAGCTGTAAGTGCCTGGAGGGTTCGAAGAAGACTAGCAGTCAGGAATACGATG	3600
e19	TAGCAGGCAGCTGTAAGTGCCTGGAGGGTTCGAAGAAGACTAGCAGTCAGGAATACGATG	3600
74e19	TAGCAGGCAGCTGTAAGTGCCTGGAGGGTTCGAAGAAGACTAGCAGTCAGGAATACGATG	3600
74	TAGCAGGCAGCTGTAAGTGCCTGGAGGGTTCGAAGAAGACTAGCAGTCAGGAATACGATG	3600
211	TAGCAGGCAGCTGTAAGTGCCTGGAGGGTTCGAAGAAGACTAGCAGTCAGGAATACGATG	3600
246	TAGCAGGCAGCTGTAAGTGCCTGGAGGGTTCGAAGAAGACTAGCAGTCAGGAATACGATG	3600
353	TAGCAGGCAGCTGTAAGTGCCTGGAGGGTTCGAAGAAGACTAGCAGTCAGGAATACGATG	3600

DAH	ATCGTAAAGTTCAAGCGGGCATGGAGTTGAGAACCGCTTGCAAAACTTATATTGTTC	3660
29B	ATCGTAAAGTTCAAGCGGGCATGGAGTTGAGAACCGCTTGCAAAACTTATATTGTTC	3660
e19	ATCGTAAAGTTCAAGCGGGCATGGAGTTGAGAACCGCTTGCAAAACTTATATTGTTC	3660
74e19	ATCGTAAAGTTCAAGCGGGCATGGAGTTGAGAACCGCTTGCAAAACTTATATTGTTC	3660
74	ATCGTAAAGTTCAAGCGGGCATGGAGTTGAGAACCGCTTGCAAAACTTATATTGTTC	3660
211	ATCGTAAAGTTCAAGCGGGCATGGAGTTGAGAACCGCTTGCAAAACTTATATTGTTC	3660
246	ATCGTAAAGTTCAAGCGGGCATGGAGTTGAGAACCGCTTGCAAAACTTATATTGTTC	3660
353	ATCGTAAAGTTCAAGCGGGCATGGAGTTGAGAACCGCTTGCAAAACTTATATTGTTC	3660

DAH	CAAACATTGGAAAAGCAAGAACGGATCGTCTGACAAATCAGACGGAGCGGAAGGTGCAG	3720
29B	CAAACATTGGAAAAGCAAGAACGGATCGTCTGACAAATCAGACGGAGCGGAAGGTGCAG	3720
e19	CAAACATTGGAAAAGCAAGAACGGATCGTCTGACAAATCAGACGGAGCGGAAGGTGCAG	3720
74e19	CAAACATTGGAAAAGCAAGAACGGATCGTCTGACAAATCAGACGGAGCGGAAGGTGCAG	3720
74	CAAACATTGGAAAAGCAAGAACGGATCGTCTGACAAATCAGACGGAGCGGAAGGTGCAG	3720
211	CAAACATTGGAAAAGCAAGAACGGATCGTCTGACAAATCAGACGGAGCGGAAGGTGCAG	3720
246	CAAACATTGGAAAAGCAAGAACGGATCGTCTGACAAATCAGACGGAGCGGAAGGTGCAG	3720
353	CAAACATTGGAAAAGCAAGAACGGATCGTCTGACAAATCAGACGGAGCGGAAGGTGCAG	3720

DAH	CTCTCGATTCTAATGCTATTCCAAATGGAGGGAGCTACTAACCTTCACGTAGAAGGGAGAG	3780
29B	CTCTCGATTCTAATGCTATTCCAAATGGAGGGAGCTACTAACCTTCACGTAGAAGGGAGAG	3780
e19	CTCTCGATTCTAATGCTATTCCAAATGGAGGGAGCTACTAACCTTCACGTAGAAGGGAGAG	3780
74e19	CTCTCGATTCTAATGCTATTCCAAATGGAGGGAGCTACTAACCTTCACGTAGAAGGGAGAG	3780
74	CTCTCGATTCTAATGCTATTCCAAATGGAGGGAGCTACTAACCTTCACGTAGAAGGGAGAG	3780
211	CTCTCGATTCTAATGCTATTCCAAATGGAGGGAGCTACTAACCTTCACGTAGAAGGGAGAG	3780
246	CTCTCGATTCTAATGCTATTCCAAATGGAGGGAGCTACTAACCTTCACGTAGAAGGGAGAG	3780
353	CTCTCGATTCTAATGCTATTCCAAATGGAGGGAGCTACTAACCTTCACGTAGAAGGGAGAG	3780

60		
DAH	ACGTTCGCGCTCGAGCCAGAGCTCGACGATGTAGAGGGCAGTGTACTCTACGCCATGTGC	3840
29B	ACGTTCGCGCTCGAGCCAGAGCTCGACGATGTAGAGGGCAGTGTACTCTACGCCATGTGC	3840
e19	ACGTTCGCGCTCGAGCCAGAGCTCGACGATGTAGAGGGCAGTGTACTCTACGCCATGTGC	3840
74e19	ACGTTCGCGCTCGAGCCAGAGCTCGACGATGTAGAGGGCAGTGTACTCTACGCCATGTGC	3840
74	ACGTTCGCGCTCGAGCCAGAGCTCGACGATGTAGAGGGCAGTGTACTCTACGCCATGTGC	3840
211	ACGTTCGCGCTCGAGCCAGAGCTCGACGATGTAGAGGGCAGTGTACTCTACGCCATGTGC	3840
246	ACGTTCGCGCTCGAGCCAGAGCTCGACGATGTAGAGGGCAGTGTACTCTACGCCATGTGC	3840
353	ACGTTCGCGCTCGAGCCAGAGCTCGACGATGTAGAGGGCAGTGTACTCTACGCCATGTGC	3840

DAH	GCTCCATCACAGACGATAACCGATGCATTTTGAAAAGGACGACGAAAATACCTATAAAG	3900
29B	GCTCCATCACAGACGATAACCGATGCATTTTGAAAAGGACGACGAAAATACCTATAAAG	3900
e19	GCTCCATCACAGACGATAACCGATGCATTTTGAAAAGGACGACGAAAATACCTATAAAG	3900
74e19	GCTCCATCACAGACGATAACCGATGCATTTTGAAAAGGACGACGAAAATACCTATAAAG	3900
74	GCTCCATCACAGACGATAACCGATGCATTTTGAAAAGGACGACGAAAATACCTATAAAG	3900
211	GCTCCATCACAGACGATAACCGATGCATTTTGAAAAGGACGACGAAAATACCTATAAAG	3900
246	GCTCCATCACAGACGATAACCGATGCATTTTGAAAAGGACGACGAAAATACCTATAAAG	3900
353	GCTCCATCACAGACGATAACCGATGCATTTTGAAAAGGACGACGAAAATACCTATAAAG	3900

DAH	ACGAAGAACAGACTTGTCTCCAACAAACAATTCTATGAGGTGTTGCCAAGGAATTGCCAC	3960
29B	ACGAAGAACAGACTTGTCTCCAACAAACAATTCTATGAGGTGTTGCCAAGGAATTGCCAC	3960
e19	ACGAAGAACAGACTTGTCTCCAACAAACAATTCTATGAGGTGTTGCCAAGGAATTGCCAC	3960
74e19	ACGAAGAACAGACTTGTCTCCAACAAACAATTCTATGAGGTGTTGCCAAGGAATTGCCAC	3960
74	ACGAAGAACAGACTTGTCTCCAACAAACAATTCTATGAGGTGTTGCCAAGGAATTGCCAC	3960
211	ACGAAGAACAGACTTGTCTCCAACAAACAATTCTATGAGGTGTTGCCAAGGAATTGCCAC	3960
246	ACGAAGAACAGACTTGTCTCCAACAAACAATTCTATGAGGTGTTGCCAAGGAATTGCCAC	3960
353	ACGAAGAACAGACTTGTCTCCAACAAACAATTCTATGAGGTGTTGCCAAGGAATTGCCAC	3960

DAH	CAAATCAAACACATTTGTCTTGAAAAACTGGGCCACTTCACCCGCTACGCTATCTCG	4020
29B	CAAATCAAACACATTTGTCTTGAAAAACTGGGCCACTTCACCCGCTACGCTATCTCG	4020
e19	CAAATCAAACACATTTGTCTTGAAAAACTGGGCCACTTCACCCGCTACGCTATCTCG	4020
74e19	CAAATCAAACACATTTGTCTTGAAAAACTGGGCCACTTCACCCGCTACGCTATCTCG	4020
74	CAAATCAAACACATTTGTCTTGAAAAACTGGGCCACTTCACCCGCTACGCTATCTCG	4020
211	CAAATCAAACACATTTGTCTTGAAAAACTGGGCCACTTCACCCGCTACGCTATCTCG	4020
246	CAAATCAAACACATTTGTCTTGAAAAACTGGGCCACTTCACCCGCTACGCTATCTCG	4020
353	CAAATCAAACACATTTGTCTTGAAAAACTGGGCCACTTCACCCGCTACGCTATCTCG	4020

DAH	TGGTAGCCTGTAGAGAAGAAATCCCCAGCGAAAAATTAAAGGGACACCAGTTTAAGAAGT	4080
29B	TGGTAGCCTGTAGAGAAGAAATCCCCAGCGAAAAATTAAAGGGACACCAGTTTAAGAAGT	4080
e19	TGGTAGCCTGTAGAGAAGAAATCCCCAGCGAAAAATTAAAGGGACACCAGTTTAAGAAGT	4080
74e19	TGGTAGCCTGTAGAGAAGAAATCCCCAGCGAAAAATTAAAGGGACACCAGTTTAAGAAGT	4080
74	TGGTAGCCTGTAGAGAAGAAATCCCCAGCGAAAAATTAAAGGGACACCAGTTTAAGAAGT	4080
211	TGGTAGCCTGTAGAGAAGAAATCCCCAGCGAAAAATTAAAGGGACACCAGTTTAAGAAGT	4080
246	TGGTAGCCTGTAGAGAAGAAATCCCCAGCGAAAAATTAAAGGGACACCAGTTTAAGAAGT	4080
353	TGGTAGCCTGTAGAGAAGAAATCCCCAGCGAAAAATTAAAGGGACACCAGTTTAAGAAGT	4080

DAH	CGCTCTGCAGCGATTATGACACCCTTCAAACTACAAAGAGAAAGAGtaggtggactt	4140
29B	CGCTCTGCAGCGATTATGACACCCTTCAAACTACAAAGAGAAAGAGtaggtggactt	4140
e19	CGCTCTGCAGCGATTATGACACCCTTCAAACTACAAAGAGAAAGAGtaggtggactt	4140
74e19	CGCTCTGCAGCGATTATGACACCCTTCAAACTACAAAGAGAAAGAGtaggtggactt	4140
74	CGCTCTGCAGCGATTATGACACCCTTCAAACTACAAAGAGAAAGAGtaggtggactt	4140
211	CGCTCTGCAGCGATTATGACACCCTTCAAACTACAAAGAGAAAGAGtaggtggactt	4140
246	CGCTCTGCAGCGATTATGACACCCTTCAAACTACAAAGAGAAAGAGtaggtggactt	4140
353	CGCTCTGCAGCGATTATGACACCCTTCAAACTACAAAGAGAAAGAGtaggtggactt	4140

61		
DAH	gagagcgtttacttacattactaacgattggttacttatagaATTGCCGACATA	4200
29B	gagagcgtttacttacattactaacgattggttacttatagaATTGCCGACATA	4200
e19	gagagcgtttacttacattactaacgattggttacttatagaATTGCCGACATA	4200
74e19	gagagcgtttacttacattactaacgattggttacttatagaATTGCCGACATA	4200
74	gagagcgtttacttacattactaacgattggttacttatagaATTGCCGACATA	4200
211	gagagcgtttacttacattactaacgattggttacttatagaATTGCCGACATA	4200
246	gagagcgtttacttacattactaacgattggttacttatagaATTGCCGACATA	4200
353	gagagcgtttacttacattactaacgattggttacttatagaATTGCCGACATA	4200

DAH	GTCATGGACCTAAAAGTAGATTAGAACACGCCAACAACACCAGTACGGTT	4260
29B	GTCATGGACCTAAAAGTAGATTAGAACACGCCAACAACACCAGTACGGTT	4260
e19	GTCATGGACCTAAAAGTAGATTAGAACACGCCAACAACACCAGTACGGTT	4260
74e19	GTCATGGACCTAAAAGTAGATTAGAACACGCCAACAACACCAGTACGGTT	4260
74	GTCATGGACCTAAAAGTAGATTAGAACACGCCAACAACACCAGTACGGTT	4260
211	GTCATGGACCTAAAAGTAGATTAGAACACGCCAACAACACCAGTACGGTT	4260
246	GTCATGGACCTAAAAGTAGATTAGAACACGCCAACAACACCAGTACGGTT	4260
353	GTCATGGACCTAAAAGTAGATTAGAACACGCCAACAACACCAGTACGGTT	4260

DAH	CGCTGGACGCCACCACTAGATCCAACGGAGAAATTGTACCTATGAAGTGGCTACAAG	4320
29B	CGCTGGACGCCACCACTAGATCCAACGGAGAAATTGTACCTATGAAGTGGCTACAAG	4320
e19	CGCTGGACGCCACCACTAGATCCAACGGAGAAATTGTACCTATGAAGTGGCTACAAG	4320
74e19	CGCTGGACGCCACCACTAGATCCAACGGAGAAATTGTACCTATGAAGTGGCTACAAG	4320
74	CGCTGGACGCCACCACTAGATCCAACGGAGAAATTGTACCTATGAAGTGGCTACAAG	4320
211	CGCTGGACGCCACCACTAGATCCAACGGAGAAATTGTACCTATGAAGTGGCTACAAG	4320
246	CGCTGGACGCCACCACTAGATCCAACGGAGAAATTGTACCTATGAAGTGGCTACAAG	4320
353	CGCTGGACGCCACCACTAGATCCAACGGAGAAATTGTACCTATGAAGTGGCTACAAG	4320

DAH	TTGCAAAACCCGATCAAGTGGAGAAAAGAAGTGCATCCGGCTGCTGACTTCAACCAG	4380
29B	TTGCAAAACCCGATCAAGTGGAGAAAAGAAGTGCATCCGGCTGCTGACTTCAACCAG	4380
e19	TTGCAAAACCCGATCAAGTGGAGAAAAGAAGTGCATCCGGCTGCTGACTTCAACCAG	4380
74e19	TTGCAAAACCCGATCAAGTGGAGAAAAGAAGTGCATCCGGCTGCTGACTTCAACCAG	4380
74	TTGCAAAACCCGATCAAGTGGAGAAAAGAAGTGCATCCGGCTGCTGACTTCAACCAG	4380
211	TTGCAAAACCCGATCAAGTGGAGAAAAGAAGTGCATCCGGCTGCTGACTTCAACCAG	4380
246	TTGCAAAACCCGATCAAGTGGAGAAAAGAAGTGCATCCGGCTGCTGACTTCAACCAG	4380
353	TTGCAAAACCCGATCAAGTGGAGAAAAGAAGTGCATCCGGCTGCTGACTTCAACCAG	4380

DAH	ACTGCCGGTTATTAATAAGCTCAACGAGGGCCTTACAGCTTCAGGGTGCAGCCAAT	4440
29B	ACTGCCGGTTATTAATAAGCTCAACGAGGGCCTTACAGCTTCAGGGTGCAGCCAAT	4440
e19	ACTGCCGGTTATTAATAAGCTCAACGAGGGCCTTACAGCTTCAGGGTGCAGCCAAT	4440
74e19	ACTGCCGGTTATTAATAAGCTCAACGAGGGCCTTACAGCTTCAGGGTGCAGCCAAT	4440
74	ACTGCCGGTTATTAATAAGCTCAACGAGGGCCTTACAGCTTCAGGGTGCAGCCAAT	4440
211	ACTGCCGGTTATTAATAAGCTCAACGAGGGCCTTACAGCTTCAGGGTGCAGCCAAT	4440
246	ACTGCCGGTTATTAATAAGCTCAACGAGGGCCTTACAGCTTCAGGGTGCAGCCAAT	4440
353	ACTGCCGGTTATTAATAAGCTCAACGAGGGCCTTACAGCTTCAGGGTGCAGCCAAT	4440

DAH	TCAATAGCGGGATA CGGCATTTACCGGAAGTCGAACATATAAAAGTTGAGgtaggta	4500
29B	TCAATAGCGGGATA CGGCATTTACCGGAAGTCGAACATATAAAAGTTGAGgtaggta	4500
e19	TCAATAGCGGGATA CGGCATTTACCGGAAGTCGAACATATAAAAGTTGAGgtaggta	4500
74e19	TCAATAGCGGGATA CGGCATTTACCGGAAGTCGAACATATAAAAGTTGAGgtaggta	4500
74	TCAATAGCGGGATA CGGCATTTACCGGAAGTCGAACATATAAAAGTTGAGgtaggta	4500
211	TCAATAGCGGGATA CGGCATTTACCGGAAGTCGAACATATAAAAGTTGAGgtaggta	4500
246	TCAATAGCGGGATA CGGCATTTACCGGAAGTCGAACATATAAAAGTTGAGgtaggta	4500
353	TCAATAGCGGGATA CGGCATTTACCGGAAGTCGAACATATAAAAGTTGAGgtaggta	4500

DAH	ataattttatctgc-----	4515
29B	ataattttatctcggtaccctagactagtctagataacttcgtataatgtatgctata	4560
e19	ataattttatctcggtaccctagactagtctagataacttcgtataatgtatgctata	4560
74e19	ataattttatctcggtaccctagactagtctagataacttcgtataatgtatgctata	4560
74	ataattttatctcggtaccctagactagtctagataacttcgtataatgtatgctata	4560
211	ataattttatctcggtaccctagactagtctagataacttcgtataatgtatgctata	4560
246	ataattttatctcggtaccctagactagtctagataacttcgtataatgtatgctata	4560
353	ataattttatctcggtaccctagactagtctagataacttcgtataatgtatgctata	4560

DAH	-----gaatgatcattttaatgaatttttt	4544
29B	cgaagttatctagactagtctagggcgccgaatgatcattttaatgaatttttt	4620
e19	cgaagttatctagactagtctagggcgccgaatgatcattttaatgaatttttt	4620
74e19	cgaagttatctagactagtctagggcgccgaatgatcattttaatgaatttttt	4620
74	cgaagttatctagactagtctagggcgccgaatgatcattttaatgaatttttt	4620
211	cgaagttatctagactagtctagggcgccgaatgatcattttaatgaatttttt	4620
246	cgaagttatctagactagtctagggcgccgaatgatcattttaatgaatttttt	4620
353	cgaagttatctagactagtctagggcgccgaatgatcattttaatgaatttttt	4620

DAH	tttcattaaattcaacagCCTCCGCCAGCTATGCTAAGGTCTTTCTGGCTACTGGGA	4604
29B	tttcattaaattcaacagCCTCCGCCAGCTATGCTAAGGTCTTTCTGGCTACTGGGA	4680
e19	tttcattaaattcaacagCCTCCGCCAGCTATGCTAAGGTCTTTCTGGCTACTGGGA	4680
74e19	tttcattaaattcaacagCCTCCGCCAGCTATGCTAAGGTCTTTCTGGCTACTGGGA	4680
74	tttcattaaattcaacagCCTCCGCCAGCTATGCTAAGGTCTTTCTGGCTACTGGGA	4680
211	tttcattaaattcaacagCCTCCGCCAGCTATGCTAAGGTCTTTCTGGCTACTGGGA	4680
246	tttcattaaattcaacagCCTCCGCCAGCTATGCTAAGGTCTTTCTGGCTACTGGGA	4680
353	tttcattaaattcaacagCCTCCGCCAGCTATGCTAAGGTCTTTCTGGCTACTGGGA	4680

DAH	ATCGGCCTAGCGTTCCTGATCGTTCCCTGTTGGCTATGTCTGTTACCTGCACAAGAGG	4664
29B	ATCGGCCTAGCGTTCCTGATCGTTCCCTGTTGGCTATGTCTGTTACCTGCACAAGAGG	4740
e19	ATCGGCCTAGCGTTCCTGATCGTTCCCTGTTGGCTATGTCTGTTACCTGCACAAGAGG	4740
74e19	ATCGGCCTAGCGTTCCTGATCGTTCCCTGTTGGCTATGTCTGTTACCTGCACAAGAGG	4740
74	ATCGGCCTAGCGTTCCTGATCGTTCCCTGTTGGCTATGTCTGTTACCTGCACAAGAGG	4740
211	ATCGGCCTAGCGTTCCTGATCGTTCCCTGTTGGCTATGTCTGTTACCTGCACAAGAGG	4740
246	ATCGGCCTAGCGTTCCTGATCGTTCCCTGTTGGCTATGTCTGTTACCTGCACAAGAGG	4740
353	ATCGGCCTAGCGTTCCTGATCGTTCCCTGTTGGCTATGTCTGTTACCTGCACAAGAGG	4740

DAH	AAGGTTCCCTTAATGACCTCCATATGAACACAGAGGTGAATCCGTTCTATGCGAGCATG	4724
29B	AAGGTTCCCTTAATGACCTCCATATGAACACAGAGGTGAATCCGTTCTATGCGAGCATG	4800
e19	AAGGTTCCCTTAATGACCTCCATATGAACACAGAGGTGAATCCGTTCTATGCGAGCATG	4800
74e19	AAGGTTCCCTTAATGACCTCCATATGAACACAGAGGTGAATCCGTTCTATGCGAGCATG	4800
74	AAGGTTCCCTTAATGACCTCCATATGAACACAGAGGTGAATCCGTTCTATGCGAGCATG	4800
211	AAGGTTCCCTTAATGACCTCCATATGAACACAGAGGTGAATCCGTTCTATGCGAGCATG	4800
246	AAGGTTCCCTTAATGACCTCCATATGAACACAGAGGTGAATCCGTTCTATGCGAGCATG	4800
353	AAGGTTCCCTTAATGACCTCCATATGAACACAGAGGTGAATCCGTTCTATGCGAGCATG	4800

DAH	CAATACATCCCAGACGATTGGGAGGTGCTGCGAGAGAACATCATTCAAGTGGCTCCACTA	4784
29B	CAATACATCCCAGACGATTGGGAGGTGCTGCGAGAGAACATCATTCAAGTGGCTCCACTA	4860
e19	CAATACATCCCAGACGATTGGGAGGTGCTGCGAGAGAACATCATTCAAGTGGCTCCACTA	4860
74e19	CAATACATCCCAGACGATTGGGAGGTGCTGCGAGAGAACATCATTCAAGTGGCTCCACTA	4860
74	CAATACATCCCAGACGATTGGGAGGTGCTGCGAGAGAACATCATTCAAGTGGCTCCACTA	4860
211	CAATACATCCCAGACGATTGGGAGGTGCTGCGAGAGAACATCATTCAAGTGGCTCCACTA	4860
246	CAATACATCCCAGACGATTGGGAGGTGCTGCGAGAGAACATCATTCAAGTGGCTCCACTA	4860
353	CAATACATCCCAGACGATTGGGAGGTGCTGCGAGAGAACATCATTCAAGTGGCTCCACTA	4860

63		
DAH	GGCCAGGGATCCTTGGCATGGTGTATGAGGGTATCCTGAAGTCCTTCCACCCAATGGC	4844
29B	GGCCAGGGATCCTTGGCATGGTGTATGAGGGTATCCTGAAGTCCTTCCACCCAATGGC	4920
e19	GGCCAGGGATCCTTGGCATGGTGTATGAGGGTATCCTGAAGTCCTTCCACCCAATGGC	4920
74e19	GGCCAGGGATCCTTGGCATGGTGTATGAGGGTATCCTGAAGTCCTTCCACCCAATGGC	4920
74	GGCCAGGGATCCTTGGCATGGTGTATGAGGGTATCCTGAAGTCCTTCCACCCAATGGC	4920
211	GGCCAGGGATCCTTGGCATGGTGTATGAGGGTATCCTGAAGTCCTTCCACCCAATGGC	4920
246	GGCCAGGGATCCTTGGCATGGTGTATGAGGGTATCCTGAAGTCCTTCCACCCAATGGC	4920
353	GGCCAGGGATCCTTGGCATGGTGTATGAGGGTATCCTGAAGTCCTTCCACCCAATGGC	4920
	*****	*****
DAH	GTGGATCGCGAGTGTGCCATTAAGACTGTCAACGAAAATGCTACGGATCGCGAGCGAAC	4904
29B	GTGGATCGCGAGTGTGCCATTAAGACTGTCAACGAAAATGCTACGGATCGCGAGCGAAC	4980
e19	GTGGATCGCGAGTGTGCCATTAAGACTGTCAACGAAAATGCTACGGATCGCGAGCGAAC	4980
74e19	GTGGATCGCGAGTGTGCCATTAAGACTGTCAACGAAAATGCTACGGATCGCGAGCGAAC	4980
74	GTGGATCGCGAGTGTGCCATTAAGACTGTCAACGAAAATGCTACGGATCGCGAGCGAAC	4980
211	GTGGATCGCGAGTGTGCCATTAAGACTGTCAACGAAAATGCTACGGATCGCGAGCGAAC	4980
246	GTGGATCGCGAGTGTGCCATTAAGACTGTCAACGAAAATGCTACGGATCGCGAGCGAAC	4980
353	GTGGATCGCGAGTGTGCCATTAAGACTGTCAACGAAAATGCTACGGATCGCGAGCGAAC	4980
	*****	*****
DAH	AATTCTCTGAGCGAGGCAGCGTCATGAAGGAGTTCGATACTGTATCATGTCGAAGATTG	4964
29B	AATTCTCTGAGCGAGGCAGCGTCATGAAGGAGTTCGATACTGTATCATGTCGAAGATTG	5040
e19	AATTCTCTGAGCGAGGCAGCGTCATGAAGGAGTTCGATACTGTATCATGTCGAAGATTG	5040
74e19	AATTCTCTGAGCGAGGCAGCGTCATGAAGGAGTTCGATACTGTATCATGTCGAAGATTG	5040
74	AATTCTCTGAGCGAGGCAGCGTCATGAAGGAGTTCGATACTGTATCATGTCGAAGATTG	5040
211	AATTCTCTGAGCGAGGCAGCGTCATGAAGGAGTTCGATACTGTATCATGTCGAAGATTG	5040
246	AATTCTCTGAGCGAGGCAGCGTCATGAAGGAGTTCGATACTGTATCATGTCGAAGATTG	5040
353	AATTCTCTGAGCGAGGCAGCGTCATGAAGGAGTTCGATACTGTATCATGTCGAAGATTG	5040
	*****	*****
DAH	CTCGGTGTTGTTCCAGGGGTCAAGCCGGCTCTGGTGGTCATGGAGCTAATGAAGAAGGGT	5024
29B	CTCGGTGTTGTTCCAGGGGTCAAGCCGGCTCTGGTGGTCATGGAGCTAATGAAGAAGGGT	5100
e19	CTCGGTGTTGTTCCAGGGGTCAAGCCGGCTCTGGTGGTCATGGAGCTAATGAAGAAGGGT	5100
74e19	CTCGGTGTTGTTCCAGGGGTCAAGCCGGCTCTGGTGGTCATGGAGCTAATGAAGAAGGGT	5100
74	CTCGGTGTTGTTCCAGGGGTCAAGCCGGCTCTGGTGGTCATGGAGCTAATGAAGAAGGGT	5100
211	CTCGGTGTTGTTCCAGGGGTCAAGCCGGCTCTGGTGGTCATGGAGCTAATGAAGAAGGGT	5100
246	CTCGGTGTTGTTCCAGGGGTCAAGCCGGCTCTGGTGGTCATGGAGCTAATGAAGAAGGGT	5100
353	CTCGGTGTTGTTCCAGGGGTCAAGCCGGCTCTGGTGGTCATGGAGCTAATGAAGAAGGGT	5100
	*****	*****
DAH	GATCTTAAGTCTATTGCGTGCCCCATCGTCCCGAGGAGCGGGATGAGGCCATGATGACG	5084
29B	GATCTTAAGTCTATTGCGTGCCCCATCGTCCCGAGGAGCGGGATGAGGCCATGATGACG	5160
e19	GATCTTAAGTCTATTGCGTGCCCCATCGTCCCGAGGAGCGGGATGAGGCCATGATGACG	5160
74e19	GATCTTAAGTCTATTGCGTGCCCCATCGTCCCGAGGAGCGGGATGAGGCCATGATGACG	5160
74	GATCTTAAGTCTATTGCGTGCCCCATCGTCCCGAGGAGCGGGATGAGGCCATGATGACG	5160
211	GATCTTAAGTCTATTGCGTGCCCCATCGTCCCGAGGAGCGGGATGAGGCCATGATGACG	5160
246	GATCTTAAGTCTATTGCGTGCCCCATCGTCCCGAGGAGCGGGATGAGGCCATGATGACG	5160
353	GATCTTAAGTCTATTGCGTGCCCCATTGTCGGAGTGAAGAATCTAC	5160
	*****	*****
DAH	TATCTTAATCGCATCGGAGTGACTGTAATGTGCAGCCTCTACTTATGGAAGAACTAC	5144
29B	TATCTTAATCGCATCGGAGTGACTGTAATGTGCAGCCTCTACTTATGGAAGAACTAC	5220
e19	TATCTTAATCGCATCGGAGTGACTGTAATGTGCAGCCTCTACTTATGGAAGAACTAC	5220
74e19	TATCTTAATCGCATCGGAGTGACTGTAATGTGCAGCCTCTACTTATGGAAGAACTAC	5220
74	TATCTTAATCGCATCGGAGTGACTGTAATGTGCAGCCTCTACTTATGGAAGAACTAC	5220
211	TATCTTAATCGCATCGGAGTGACTGTAATGTGCAGCCTCTACTTATGGAAGAACTAC	5220
246	TATCTTAATCGCATCGGAGTGACTGTAATGTGCAGCCTCTACTTATGGAAGAACTAC	5220
353	TATCTTAATCGCATCGGAGTGACTGTAATGTGCAGCCTCTACTTATGGAAGAACTAC	5220
	*****	*****

DAH	CAGATGGCCATTGAGATTGCGGATGGCATGGCATATTGGCCGCCAAGAACGTTCTGCAT	5204
29B	CAGATGGCCATTGAGATTGCGGATGGCATGGCATATTGGCCGCCAAGAACGTTCTGCAT	5280
e19	CAGATGGCCATTGAGATTGCGGATGGCATGGCATATTGGCCGCCAAGAACGTTCTGCAT	5280
74e19	CAGATGGCCATTGAGATTGCGGATGGCATGGCATATTGGCCGCCAAGAACGTTCTGCAT	5280
74	CAGATGGCCATTGAGATTGCGGATGGCATGGCATATTGGCCGCCAAGAACGTTCTGCAT	5280
211	CAGATGGCCATTGAGATTGCGGATGGCATGGCATATTGGCCGCCAAGAACGTTCTGCAT	5280
246	CAGATGGCCATTGAGATTGCGGATGGCATGGCATATTGGCCGCCAAGAACGTTCTGCAT	5280
353	CAGATGGCCATTGAGATTGCGGATGGCATGGCATATTGGCCGCCAAGAACGTTCTGCAT	5280

DAH	CGTGATCTTGCAGCTCGAAATTGCATGGTTGCTGATGATTGACGGTGAAAATTGGTGAC	5264
29B	CGTGATCTTGCAGCTCGAAATTGCATGGTTGCTGATGATTGACGGTGAAAATTGGTGAC	5340
e19	CGTGATCTTGCAGCTCGAAATTGCATGGTTGCTGATGATTGACGGTGAAAATTGGTGAC	5340
74e19	CGTGATCTTGCAGCTCGAAATTGCATGGTTGCTGATGATTGACGGTGAAAATTGGTGAC	5340
74	CGTGATCTTGCAGCTCGAAATTGCATGGTTGCTGATGATTGACGGTGAAAATTGGTGAC	5340
211	CGTGATCTTGCAGCTCGAAATTGCATGGTTGCTGATGATTGACGGTGAAAATTGGTGAC	5340
246	CGTGATCTTGCAGCTCGAAATTGCATGGTTGCTGATGATTGACGGTGAAAATTGGTGAC	5340
353	CGTGATCTTGCAGCTCGAAATTGCATGGTTGCTGATGATTGACGGTGAAAATTGGTGAC	5340

DAH	TTTGAATGACCGTGACATCTATGAGACGGATTACTATCGGAAGGGCACTAAAGGGCTG	5324
29B	TTTGAATGACCGTGACATCTATGAGACGGATTACTATCGGAAGGGCACTAAAGGGCTG	5400
e19	TTTGAATGACCGTGACATCTATGAGACGGATTACTATCGGAAGGGCACTAAAGGGCTG	5400
74e19	TTTGAATGACCGTGACTCTATGAGACGGATTACTATCGGAAGGGCACTAAAGGGCTG	5400
74	TTTGAATGACCGTGACTCTATGAGACGGATTACTATCGGAAGGGCACTAAAGGGCTG	5400
211	TTTGAATGACCGTGACATCTATGAGACGGATTACTATCGGAAGGGCACTAAAGGGCTG	5400
246	TTTGAATGACCGTGACATCTATGAGACGGATTACTATCGGAAGGGCACTAAAGGGCTG	5400
353	TTTGAATGACCGTGACATCTATGAGACGGATTACTATCGGAAGGGCACTAAAGGGCTG	5400

DAH	CTGCCAGTTCGCTGGATGCCACCGGAGAGCTTGCAGATGGTGTCTACTCTAGTGCAGT	5384
29B	CTGCCAGTTCGCTGGATGCCACCGGAGAGCTTGCAGATGGTGTCTACTCTAGTGCAGT	5460
e19	CTGCCAGTTCGCTGGATGCCACCGGAGAGCTTGCAGATGGTGTCTACTCTAGTGCAGT	5460
74e19	CTGCCAGTTCGCTGGATGCCACCGGAGAGCTTGCAGATGGTGTCTACTCTAGTGCAGT	5460
74	CTGCCAGTTCGCTGGATGCCACCGGAGAGCTTGCAGATGGTGTCTACTCTAGTGCAGT	5460
211	CTGCCAGTTCGCTGGATGCCACCGGAGAGCTTGCAGATGGTGTCTACTCTAGTGCAGT	5460
246	CTGCCAGTTCGCTGGATGCCACCGGAGAGCTTGCAGATGGTGTCTACTCTAGTGCAGT	5460
353	CTGCCAGTTCGCTGGATGCCACCGGAGAGCTTGCAGATGGTGTCTACTCTAGTGCAGT	5460

DAH	GATGTATTCACTTGGAGTGGTTCTCTGGAAATGGCACCTAGCGGCTCAGCCATAC	5444
29B	GATGTATTCACTTGGAGTGGTTCTCTGGAAATGGCACCTAGCGGCTCAGCCATAC	5520
e19	GATGTATTCACTTGGAGTGGTTCTCTGGAAATGGCACCTAGCGGCTCAGCCATAC	5520
74e19	GATGTATTCACTTGGAGTGGTTCTCTGGAAATGGCACCTAGCGGCTCAGCCATAC	5520
74	GATGTATTCACTTGGAGTGGTTCTCTGGAAATGGCACCTAGCGGCTCAGCCATAC	5520
211	GATGTATTCACTTGGAGTGGTTCTCTGGAAATGGCACCTAGCGGCTCAGCCATAC	5520
246	GATGTATTCACTTGGAGTGGTTCTCTGGAAATGGCACCTAGCGGCTCAGCCATAC	5520
353	GATGTATTCACTTGGAGTGGTTCTCTGGAAATGGCACCTAGCGGCTCAGCCATAC	5520

DAH	CAGGGACTTTCCAACGAGCAAGTCCTCGCTACGTTATCGATGGCGGTGTTATGGAGAGG	5504
29B	CAGGGACTTTCCAACGAGCAAGTCCTCGCTACGTTATCGATGGCGGTGTTATGGAGAGG	5580
e19	CAGGGACTTTCCAACGAGCAAGTCCTCGCTACGTTATCGATGGCGGTGTTATGGAGAGG	5580
74e19	CAGGGACTTTCCAACGAGCAAGTCCTCGCTACGTTATCGATGGCGGTGTTATGGAGAGG	5580
74	CAGGGACTTTCCAACGAGCAAGTCCTCGCTACGTTATCGATGGCGGTGTTATGGAGAGG	5580
211	CAGAGACTTTCCAACGAGCAAGTCCTCGCTACGTTATCGATGGCGGTGTTATGGAGAGG	5580
246	CAGGGACTTTCCAACGAGCAAGTCCTCGCTACGTTATCGATGGCGGTGTTATGGAGAGG	5580
353	CAGGGACTTTCCAACGAGCAAGTCCTCGCTACGTTATCGATGGCGGTGTTATGGAGAGG	5580

DAH	CCGGAAAATTGCTCTGATTTCTGCATAAAACTAATGCAAAGGTGCTGGCATCATAGGTCT	5564
29B	CCGGAAAATTGCTCTGATTTCTGCATAAAACTAATGCAAAGGTGCTGGCATCATAGGTCT	5640
e19	CCGGAAAATTGCTCTGATTTCTGCATAAAACTAATGCAAAGGTGCTGGCATCATAGGTCT	5640
74e19	CCGGAAAATTGCTCTGATTTCTGCATAAAACTAATGCAAAGGTGCTGGCATCATAGGTCT	5640
74	CCGGAAAATTGCTCTGATTTCTGCATAAAACTAATGCAAAGGTGCTGGCATCATAGGTCT	5640
211	CCGGAAAATTGCTCTGATTTCTGCATAAAACTAATGCAAAGGTGCTGGCATCATAGGTCT	5640
246	CCGGAAAATTGCTCTGATTTCTGCATAAAACTAATGCAAAGGTGCTGGCATCATAGGTCT	5640
353	CCGGAAAATTGCTCTGATTTCTGCATAAAACTAATGCAAAGGTGCTGGCATCATAGGTCT	5640

DAH	TCGGCGAGACCCAGTTCTGGATATCATTGCGTATCTCGAACCAATGCCCAATTCA	5624
29B	TCGGCGAGACCCAGTTCTGGATATCATTGCGTATCTCGAACCAATGCCCAATTCA	5700
e19	TCGGCGAGACCCAGTTCTGGATATCATTGCGTATCTCGAACCAATGCCCAATTCA	5700
74e19	TCGGCGAGACCCAGTTCTGGATATCATTGCGTATCTCGAACCAATGCCCAATTCA	5700
74	TCGGCGAGACCCAGTTCTGGATATCATTGCGTATCTCGAACCAATGCCCAATTCA	5700
211	TCGGCGAGACCCAGTTCTGGATATCATTGCGTATCTCGAACCAATGCCCAATTCA	5700
246	TCGGCGAGACCCAGTTCTGGATATCATTGCGTATCTCGAACCAATGCCCAATTCA	5700
353	TCGGCGAGACCCAGTTCTGGATATCATTGCGTATCTCGAACCAATGCCCAATTCA	5700

DAH	CAATTAAAGGAAGTATCCTTCTATCACTCAGAGGCAGGTCTGCAGCATCGGAAAAGGAG	5684
29B	CAATTAAAGGAAGTATCCTTCTATCACTCAGAGGCAGGTCTGCAGCATCGGAAAAGGAG	5760
e19	CAATTAAAGGAAGTATCCTTCTATCACTCAGAGGCAGGTCTGCAGCATCGGAAAAGGAG	5760
74e19	CAATTAAAGGAAGTATCCTTCTATCACTCAGAGGCAGGTCTGCAGCATCGGAAAAGGAG	5760
74	CAATTAAAGGAAGTATCCTTCTATCACTCAGAGGCAGGTCTGCAGCATCGGAAAAGGAG	5760
211	CAATTAAAGGAAGTATCCTTCTATCACTCAGAGGCAGGTCTGCAGCATCGGAAAAGGAG	5760
246	CAATTAAAGGAAGTATCCTTCTATCACTCAGAGGCAGGTCTGCAGCATCGGAAAAGGAG	5760
353	CAATTAAAGGAAGTATCCTTCTATCACTCAGAGGCAGGTCTGCAGCATCGGAAAAGGAG	5760

DAH	CGCAAGGAACCGAATCAGCTAGATGCATTGCGGGCAGTCCCCTGGATCAAGATCTGCAG	5744
29B	CGCAAGGAACCGAATCAGCTAGATGCATTGCGGGCAGTCCCCTGGATCAAGATCTGCAG	5820
e19	CGCAAGGAACCGAATCAGCTAGATGCATTGCGGGCAGTCCCCTGGATCAAGATCTGCAG	5820
74e19	CGCAAGGAACCGAATCAGCTAGATGCATTGCGGGCAGTCCCCTGGATCAAGATCTGCAG	5820
74	CGCAAGGAACCGAATCAGCTAGATGCATTGCGGGCAGTCCCCTGGATCAAGATCTGCAG	5820
211	CGCAAGGAACCGAATCAGCTAGATGCATTGCGGGCAGTCCCCTGGATCAAGATCTGCAG	5820
246	CGCAAGGAACCGAATCAGCTAGATGCATTGCGGGCAGTCCCCTGGATCAAGATCTGCAG	5820
353	CGCAAGGAACCGAATCAGCTAGATGCATTGCGGGCAGTCCCCTGGATCAAGATCTGCAG	5820

DAH	GATCGGGAACAGCAGGAGGATGCTACCACACCTTACGAATGGCGATTATCAGCAGAAC	5804
29B	GATCGGGAACAGCAGGAGGATGCTACCACACCTTACGAATGGCGATTATCAGCAGAAC	5880
e19	GATCGGGAACAGCAGGAGGATGCTACCACACCTTACGAATGGCGATTATCAGCAGAAC	5880
74e19	GATCGGGAACAGCAGGAGGATGCTACCACACCTTACGAATGGCGATTATCAGCAGAAC	5880
74	GATCGGGAACAGCAGGAGGATGCTACCACACCTTACGAATGGCGATTATCAGCAGAAC	5880
211	GATCGGGAACAGCAGGAGGATGCTACCACACCTTACGAATGGCGATTATCAGCAGAAC	5880
246	GATCGGGAACAGCAGGAGGATGCTACCACACCTTACGAATGGCGATTATCAGCAGAAC	5880
353	GATCGGGAACAGCAGGAGGATGCTACCACACCTTACGAATGGCGATTATCAGCAGAAC	5880

DAH	TCCTCGTTGGATCAACC GCCCGAAAGCCCCATGCCATGGTGATGATCAGGGTTCTCAC	5864
29B	TCCTCGTTGGATCAACC GCCCGAAAGCCCCATGCCATGGTGATGATCAGGGTTCTCAC	5940
e19	TCCTCGTTGGATCAACC GCCCGAAAGCCCCATGCCATGGTGATGATCAGGGTTCTCAC	5940
74e19	TCCTCGTTGGATCAACC GCCCGAAAGCCCCATGCCATGGTGATGATCAGGGTTCTCAC	5940
74	TCCTCGTTGGATCAACC GCCCGAAAGCCCCATGCCATGGTGATGATCAGGGTTCTCAC	5940
211	TCCTCGTTGGATCAACC GCCCGAAAGCCCCATGCCATGGTGATGATCAGGGTTCTCAC	5940
246	TCCTCGTTGGATCAACC GCCCGAAAGCCCCATGCCATGGTGATGATCAGGGTTCTCAC	5940
353	TCCTCGTTGGATCAACC GCCCGAAAGCCCCATGCCATGGTGATGATCAGGGTTCTCAC	5940

DAH	TTGCCATTAGCCTGCCATCCGGATTCATTGCGAGCAGTACTCCTGATGGGCAGACTGTA	5924
29B	TTGCCATTAGCCTGCCATCCGGATTCATTGCGAGCAGTACTCCTGATGGGCAGACTGTA	6000
e19	TTGCCATTAGCCTGCCATCCGGATTCATTGCGAGCAGTACTCCTGATGGGCAGACTGTA	6000
74e19	TTGCCATTAGCCTGCCATCCGGATTCATTGCGAGCAGTACTCCTGATGGGCAGACTGTA	6000
74	TTGCCATTAGCCTGCCATCCGGATTCATTGCGAGCAGTACTCCTGATGGGCAGACTGTA	6000
211	TTGCCATTAGCCTGCCATCCGGATTCATTGCGAGCAGTACTCCTGATGGGCAGACTGTA	6000
246	TTGCCATTAGCCTGCCATCCGGATTCATTGCGAGCAGTACTCCTGATGGGCAGACTGTA	6000
353	TTGCCATTAGCCTGCCATCCGGATTCATTGCGAGCAGTACTCCTGATGGGCAGACTGTA	6000

DAH	ATGGCTACTGCTTCCAGAATATTCCAGCAGCGCAAGGTGATATTCGGCGACTTACGTT	5984
29B	ATGGCTACTGCTTCCAGAATATTCCAGCAGCGCAAGGTGATATTCGGCGACTTACGTT	6060
e19	ATGGCTACTGCTTCCAGAATATTCCAGCAGCGCAAGGTGATATTCGGCGACTTACGTT	6060
74e19	ATGGCTACTGCTTCCAGAATATTCCAGCAGCGCAAGGTGATATTCGGCGACTTACGTT	6060
74	ATGGCTACTGCTTCCAGAATATTCCAGCAGCGCAAGGTGATATTCGGCGACTTACGTT	6060
211	ATGGCTACTGCTTCCAGAATATTCCAGCAGCGCAAGGTGATATTCGGCGACTTACGTT	6060
246	ATGGCTACTGCTTCCAGAATATTCCAGCAGCGCAAGGTGATATTCGGCGACTTACGTT	6060
353	ATGGCTACTGCTTCCAGAATATTCCAGCAGCGCAAGGTGATATTCGGCGACTTACGTT	6060

DAH	GTGCCTGATGCAGACGCTTGGACGGCGACAGGGGATATGAGATCTACGATCCCAGTCCG	6044
29B	GTGCCTGATGCAGACGCTTGGACGGCGACAGGGGATATGAGATCTACGATCCCAGTCCG	6120
e19	GTGCCTGATGCAGACGCTTGGACGGCGACAGGGGATATGAGATCTACGATCCCAGTCCG	6120
74e19	GTGCCTGATGCAGACGCTTGGACGGCGACAGGGGATATGAGATCTACGATCCCAGTCCG	6120
74	GTGCCTGATGCAGACGCTTGGACGGCGACAGGGGATATGAGATCTACGATCCCAGTCCG	6120
211	GTGCCTGATGCAGACGCTTGGACGGCGACAGGGGATATGAGATCTACGATCCCAGTCCG	6120
246	GTGCCTGATGCAGACGCTTGGACGGCGACAGGGGATATGAGATCTACGATCCCAGTCCG	6120
353	GTGCCTGATGCAGACGCTTGGACGGCGACAGGGGATATGAGATCTACGATCCCAGTCCG	6120

DAH	AAATGTGCAGAGCTGCCGACGAGCAGAACGTGGCAGTACTGGCGGTGGAAACTCAGCGGA	6104
29B	AAATGTGCAGAGCTGCCGACGAGCAGAACGTGGCAGTACTGGCGGTGGAAACTCAGCGGA	6180
e19	AAATGTGCAGAGCTGCCGACGAGCAGAACGTGGCAGTACTGGCGGTGGAAACTCAGCGGA	6180
74e19	AAATGTGCAGAGCTGCCGACGAGCAGAACGTGGCAGTACTGGCGGTGGAAACTCAGCGGA	6180
74	AAATGTGCAGAGCTGCCGACGAGCAGAACGTGGCAGTACTGGCGGTGGAAACTCAGCGGA	6180
211	AAATGTGCAGAGCTGCCGACGAGCAGAACGTGGCAGTACTGGCGGTGGAAACTCAGCGGA	6180
246	AAATGTGCAGAGCTGCCGACGAGCAGAACGTGGCAGTACTGGCGGTGGAAACTCAGCGGA	6180
353	AAATGTGCAGAGCTGCCGACGAGCAGAACGTGGCAGTACTGGCGGTGGAAACTCAGCGGA	6180

DAH	GAACACATTGCTGCCAAGAAAGGGTCGCCAGCCTACCATCATGAGCAGCTCGATGCCA	6164
29B	GAACACATTGCTGCCAAGAAAGGGTCGCCAGCCTACCATCATGAGCAGCTCGATGCCA	6240
e19	GAACACATTGCTGCCAAGAAAGGGTCGCCAGCCTACCATCATGAGCAGCTCGATGCCA	6240
74e19	GAACACATTGCTGCCAAGAAAGGGTCGCCAGCCTACCATCATGAGCAGCTCGATGCCA	6240
74	GAACACATTGCTGCCAAGAAAGGGTCGCCAGCCTACCATCATGAGCAGCTCGATGCCA	6240
211	GAACACATTGCTGCCAAGAAAGGGTCGCCAGCCTACCATCATGAGCAGCTCGATGCCA	6240
246	GAACACATTGCTGCCAAGAAAGGGTCGCCAGCCTACCATCATGAGCAGCTCGATGCCA	6240
353	GAACACATTGCTGCCAAGAAAGGGTCGCCAGCCTACCATCATGAGCAGCTCGATGCCA	6240

DAH	GATGATGTCATCGTGGGTCTCACTGCAACCCTCGACTGCGTCAGCAGCCAGTTCTAAT	6224
29B	GATGATGTCATCGTGGGTCTCACTGCAACCCTCGACTGCGTCAGCAGCCAGTTCTAAT	6300
e19	GATGATGTCATCGTGGGTCTCACTGCAACCCTCGACTGCGTCAGCAGCCAGTTCTAAT	6300
74e19	GATGATGTCATCGTGGGTCTCACTGCAACCCTCGACTGCGTCAGCAGCCAGTTCTAAT	6300
74	GATGATGTCATCGTGGGTCTCACTGCAACCCTCGACTGCGTCAGCAGCCAGTTCTAAT	6300
211	GATGATGTCATCGTGGGTCTCA G TGCAACCCTCGACTGCGTCAGCAGCCAGTTCTAAT	6300
246	GATGATGTCATCGTGGGTCTCACTGCAACCCTCGACTGCGTCAGCAGCCAGTTCTAAT	6300
353	GATGATGTCATCGTGGGTCTCACTGCAACCCTCGACTGCGTCAGCAGCCAGTTCTAAT	6300

DAH	GCCAGTTCGCACACAGGACGCCAAGTCTGAAGAAAACAGTGGCGGATTCGGTCGCAAT	6284
29B	GCCAGTTCGCACACAGGACGCCAAGTCTGAAGAAAACAGTGGCGGATTCGGTCGCAAT	6360
e19	GCCAGTTCGCACACAGGACGCCAAGTCTGAAGAAAACAGTGGCGGATTCGGTCGCAAT	6360
74e19	GCCAGTTCGCACACAGGACGCCAAGTCTGAAGAAAACAGTGGCGGATTCGGTCGCAAT	6360
74	GCCAGTTCGCACACAGGACGCCAAGTCTGAAGAAAACAGTGGCGGATTCGGTCGCAAT	6360
211	GCCAGTTCGCACACAGGACGCCAAGTCTGAAGAAAACAGTGGCGGATTCGGTCGCAAT	6360
246	GCCAGTTCGCACACAGGACGCCAAGTCTGAAGAAAACAGTGGCGGATTCGGTCGCAAT	6360
353	GCCAGTTCGCACACAGGACGCCAAGTCTGAAGAAAACAGTGGCGGATTCGGTCGCAAT	6360

DAH	AAGGCAAACTTCATTAATGCCACCTATTTAACCAAGCGAACGGCAGCAATGCCAGC	6344
29B	AAGGCAAACTTCATTAATGCCACCTATTTAACCAAGCGAACGGCAGCAATGCCAGC	6420
e19	AAGGCAAACTTCATTAATGCCACCTATTTAACCAAGCGAACGGCAGCAATGCCAGC	6420
74e19	AAGGCAAACTTCATTAATGCCACCTATTTAACCAAGCGAACGGCAGCAATGCCAGC	6420
74	AAGGCAAACTTCATTAATGCCACCTATTTAACCAAGCGAACGGCAGCAATGCCAGC	6420
211	AAGGCAAACTTCATTAATGCCACCTATTTAACCAAGCGAACGGCAGCAATGCCAGC	6420
246	AAGGCAAACTTCATTAATGCCACCTATTTAACCAAGCGAACGGCAGCAATGCCAGC	6420
353	AAGGCAAACTTCATTAATGCCACCTATTTAACCAAGCGAACGGCAGCAATGCCAGC	6420

DAH	CACAAGAGCAATGCCTCCAATGCTCGAGTACCAGCAGTAACACCAACTTGACAAGTCAC	6404
29B	CACAAGAGCAATGCCTCCAATGCTCGAGTACCAGCAGTAACACCAACTTGACAAGTCAC	6480
e19	CACAAGAGCAATGCCTCCAATGCTCGAGTACCAGCAGTAACACCAACTTGACAAGTCAC	6480
74e19	CACAAGAGCAATGCCTCCAATGCTCGAGTACCAGCAGTAACACCAACTTGACAAGTCAC	6480
74	CACAAGAGCAATGCCTCCAATGCTCGAGTACCAGCAGTAACACCAACTTGACAAGTCAC	6480
211	CACAAGAGCAATGCCTCCAATGCTCGAGTACCAGCAGTAACACCAACTTGACAAGTCAC	6480
246	CACAAGAGCAATGCCTCCAATGCTCGAGTACCAGCAGTAACACCAACTTGACAAGTCAC	6480
353	CACAAGAGCAATGCCTCCAATGCTCGAGTACCAGCAGTAACACCAACTTGACAAGTCAC	6480

DAH	CCAGTGGCTATGGGCAATCTTGAACATCGAGAGTGGTGGCAGTGGTTGGCTGGTAGT	6464
29B	CCAGTGGCTATGGGCAATCTTGAACATCGAGAGTGGTGGCAGTGGTTGGCTGGTAGT	6540
e19	CCAGTGGCTATGGGCAATCTTGAACATCGAGAGTGGTGGCAGTGGTTGGCTGGTAGT	6540
74e19	CCAGTGGCTATGGGCAATCTTGAACATCGAGAGTGGTGGCAGTGGTTGGCTGGTAGT	6540
74	CCAGTGGCTATGGGCAATCTTGAACATCGAGAGTGGTGGCAGTGGTTGGCTGGTAGT	6540
211	CCAGTGGCTATGGGCAATCTTGAACATCGAGAGTGGTGGCAGTGGTTGGCTGGTAGT	6540
246	CCAGTGGCTATGGGCAATCTTGAACATCGAGAGTGGTGGCAGTGGTTGGCTGGTAGT	6540
353	CCAGTGGCTATGGGCAATCTTGAACATCGAGAGTGGTGGCAGTGGTTGGCTGGTAGT	6540

DAH	TATACTGGAACACCCCGCTTCTATACTCCATCAGCGACGCCCTGGAGGAGGCAGCGGTATG	6524
29B	TATACTGGAACACCCCGCTTCTATACTCCATCAGCGACGCCCTGGAGGAGGCAGCGGTATG	6600
e19	TATACTGGAACACCCCGCTTCTATACTCCATCAGCGACGCCCTGGAGGAGGCAGCGGTATG	6600
74e19	TATACTGGAACACCCCGCTTCTATACTCCATCAGCGACGCCCTGGAGGAGGCAGCGGTATG	6600
74	TATACTGGAACACCCCGCTTCTATACTCCATCAGCGACGCCCTGGAGGAGGCAGCGGTATG	6600
211	TATACTGGAACACCCCGCTTCTATACTCCATCAGCGACGCCCTGGAGGAGGCAGCGGTATG	6600
246	TATACTGGAACACCCCGCTTCTATACTCCATCAGCGACGCCCTGGAGGAGGCAGCGGTATG	6600
353	TATACTGGAACACCCCGCTTCTATACTCCATCAGCGACGCCCTGGAGGAGGCAGCGGTATG	6600

DAH	GCCATTAGCGACAATCTTAACACTACAGACTACTAGACGAGTCATAGCCAGCGAACAGGCC	6584
29B	GCCATTAGCGACAATCTTAACACTACAGACTACTAGACGAGTCATAGCCAGCGAACAGGCC	6660
e19	GCCATTAGCGACAATCTTAACACTACAGACTACTAGACGAGTCATAGCCAGCGAACAGGCC	6660
74e19	GCCATTAGCGACAATCTTAACACTACAGACTACTAGACGAGTCATAGCCAGCGAACAGGCC	6660
74	GCCATTAGCGACAATCTTAACACTACAGACTACTAGACGAGTCATAGCCAGCGAACAGGCC	6660
211	GCCATTAGCGACAATCTTAACACTACAGACTACTAGACGAGTCATAGCCAGCGAACAGGCC	6660
246	GCCATTAGCGACAATCTTAACACTACAGACTACTAGACGAGTCATAGCCAGCGAACAGGCC	6660
353	GCCATTAGCGACAATCTTAACACTACAGACTACTAGACGAGTCATAGCCAGCGAACAGGCC	6660

DAH	ACCATCCTAACGACTAGCAGCCCCAATCCAACTACGAGATGATGCATCCACCAACCAGT	6644
29B	ACCATCCTAACGACTAGCAGCCCCAATCCAACTACGAGATGATGCATCCACCAACCAGT	6720
e19	ACCATCCTAACGACTAGCAGCCCCAATCCAACTACGAGATGATGCATCCACCAACCAGT	6720
74e19	ACCATCCTAACGACTAGCAGCCCCAATCCAACTACGAGATGATGCATCCACCAACCAGT	6720
74	ACCATCCTAACGACTAGCAGCCCCAATCCAACTACGAGATGATGCATCCACCAACCAGT	6720
211	ACCATCCTAACGACTAGCAGCCCCAATCCAACTACGAGATGATGCATCCACCAACCAGT	6720
246	ACCATCCTAACGACTAGCAGCCCCAATCCAACTACGAGATGATGCATCCACCAACCAGT	6720
353	ACCATCCTAACGACTAGCAGCCCCAATCCAACTACGAGATGATGCATCCACCAACCAGT	6720

DAH	CTGGTCAGTACCAATCCGAACTATATGCCCATGAATGAGACTCCAGTGCAGATGGCGGGT	6704
29B	CTGGTCAGTACCAATCCGAACTATATGCCCATGAATGAGACTCCAGTGCAGATGGCGGGT	6780
e19	CTGGTCAGTACCAATCCGAACTATATGCCCATGAATGAGACTCCAGTGCAGATGGCGGGT	6780
74e19	CTGGTCAGTACCAATCCGAACTATATGCCCATGAATGAGACTCCAGTGCAGATGGCGGGT	6780
74	CTGGTCAGTACCAATCCGAACTATATGCCCATGAATGAGACTCCAGTGCAGATGGCGGGT	6780
211	CTGGTCAGTACCAATCCGAACTATATGCCCATGAATGAGACTCCAGTGCAGATGGCGGGT	6780
246	CTGGTCAGTACCAATCCGAACTATATGCCCATGAATGAGACTCCAGTGCAGATGGCGGGT	6780
353	CTGGTCAGTACCAATCCGAACTATATGCCCATGAATGAGACTCCAGTGCAGATGGCGGGT	6780

DAH	GTGACCATTAGCCATAATCGAATTACCAGCCATGCAGGCGCCGTTGAATGCACGCCAA	6764
29B	GTGACCATTAGCCATAATCGAATTACCAGCCATGCAGGCGCCGTTGAATGCACGCCAA	6840
e19	GTGACCATTAGCCATAATCGAATTACCAGCCATGCAGGCGCCGTTGAATGCACGCCAA	6840
74e19	GTGACCATTAGCCATAATCGAATTACCAGCCATGCAGGCGCCGTTGAATGCACGCCAA	6840
74	GTGACCATTAGCCATAATCGAATTACCAGCCATGCAGGCGCCGTTGAATGCACGCCAA	6840
211	GTGACCATTAGCCATAATCGAATTACCAGCCATGCAGGCGCCGTTGAATGCACGCCAA	6840
246	GTGACCATTAGCCATAATCGAATTACCAGCCATGCAGGCGCCGTTGAATGCACGCCAA	6840
353	GTGACCATTAGCCATAATCGAATTACCAGCCATGCAGGCGCCGTTGAATGCACGCCAA	6840

DAH	AGCCAAAGTAGCTCGACGGGACAACGAGCAGGAGGAGCAGTGAGGATGAGGACGAC	6824
29B	AGCCAAAGTAGCTCGACGGGACAACGAGCAGGAGGAGCAGTGAGGATGAGGACGAC	6900
e19	AGCCAAAGTAGCTCGACGGGACAACGAGCAGGAGGAGCAGTGAGGATGAGGACGAC	6900
74e19	AGCCAAAGTAGCTCGACGGGACAACGAGCAGGAGGAGCAGTGAGGATGAGGACGAC	6900
74	AGCCAAAGTAGCTCGACGGGACAACGAGCAGGAGGAGCAGTGAGGATGAGGACGAC	6900
211	AGCCAAAGTAGCTCGACGGGACAACGAGCAGGAGGAGCAGTGAGGATGAGGACGAC	6900
246	AGCCAAAGTAGCTCGACGGGACAACGAGCAGGAGGAGCAGTGAGGATGAGGACGAC	6900
353	AGCCAAAGTAGCTCGACGGGACAACGAGCAGGAGGAGCAGTGAGGATGAGGACGAC	6900

DAH	GACGTGGACGATGAGCATGTGGAGCACATCAAGATGGAGCGCATGCCATTGAGTCGGCCC	6884
29B	GACGTGGACGATGAGCATGTGGAGCACATCAAGATGGAGCGCATGCCATTGAGTCGGCCC	6960
e19	GACGTGGACGATGAGCATGTGGAGCACATCAAGATGGAGCGCATGCCATTGAGTCGGCCC	6960
74e19	GACGTGGACGATGAGCATGTGGAGCACATCAAGATGGAGCGCATGCCATTGAGTCGGCCC	6960
74	GACGTGGACGATGAGCATGTGGAGCACATCAAGATGGAGCGCATGCCATTGAGTCGGCCC	6960
211	GACGTGGACGATGAGCATGTGGAGCACATCAAGATGGAGCGCATGCCATTGAGTCGGCCC	6960
246	GACGTGGACGATGAGCATGTGGAGCACATCAAGATGGAGCGCATGCCATTGAGTCGGCCC	6960
353	GACGTGGACGATGAGCATGTGGAGCACATCAAGATGGAGCGCATGCCATTGAGTCGGCCC	6960

DAH	AGGCAAAGAGCGTTGCCAGCAAGACGCAGCCCTCGCAGTCGCAGCGTCAGCCAAACG	6944
29B	AGGCAAAGAGCGTTGCCAGCAAGACGCAGCCCTCGCAGTCGCAGCGTCAGCCAAACG	7020
e19	AGGCAAAGAGCGTTGCCAGCAAGACGCAGCCCTCGCAGTCGCAGCGTCAGCCAAACG	7020
74e19	AGGCAAAGAGCGTTGCCAGCAAGACGCAGCCCTCGCAGTCGCAGCGTCAGCCAAACG	7020
74	AGGCAAAGAGCGTTGCCAGCAAGACGCAGCCCTCGCAGTCGCAGCGTCAGCCAAACG	7020
211	AGGCAAAGAGCGTTGCCAGCAAGACGCAGCCCTCGCAGTCGCAGCGTCAGCCAAACG	7020
246	AGGCAAAGAGCGTTGCCAGCAAGACGCAGCCCTCGCAGTCGCAGCGTCAGCCAAACG	7020
353	AGGCAAAGAGCGTTGCCAGCAAGACGCAGCCCTCGCAGTCGCAGCGTCAGCCAAACG	7020

69		
DAH	AGAAAATCTCCTACGAATCCAACCTCGGAATCGGAGCGACAGGAGCCGGAAACCGATCC	7004
29B	AGAAAATCTCCTACGAATCCAACCTCGGAATCGGAGCGACAGGAGCCGGAAACCGATCC	7080
e19	AGAAAATCTCCTACGAATCCAACCTCGGAATCGGAGCGACAGGAGCCGGAAACCGATCC	7080
74e19	AGAAAATCTCCTACGAATCCAACCTCGGAATCGGAGCGACAGGAGCCGGAAACCGATCC	7080
74	AGAAAATCTCCTACGAATCCAACCTCGGAATCGGAGCGACAGGAGCCGGAAACCGATCC	7080
211	AGAAAATCTCCTACGAATCCAACCTCGGAATCGGAGCGACAGGAGCCGGAAACCGATCC	7080
246	AGAAAATCTCCTACGAATCCAACCTCGGAATCGGAGCGACAGGAGCCGGAAACCGATCC	7080
353	AGAAAATCTCCTACGAATCCAACCTCGGAATCGGAGCGACAGGAGCCGGAAACCGATCC	7080

DAH	AACTTGCTTAAAGAGAACCTGGCTCGGACCGGGCAGTACGCCAAGGCCTCCACCACCAAT	7064
29B	AACTTGCTTAAAGAGAACCTGGCTCGGACCGGGCAGTACGCCAAGGCCTCCACCACCAAT	7140
e19	AACTTGCTTAAAGAGAACCTGGCTCGGACCGGGCAGTACGCCAAGGCCTCCACCACCAAT	7140
74e19	AACTTGCTTAAAGAGAACCTGGCTCGGACCGGGCAGTACGCCAAGGCCTCCACCACCAAT	7140
74	AACTTGCTTAAAGAGAACCTGGCTCGGACCGGGCAGTACGCCAAGGCCTCCACCACCAAT	7140
211	AACTTGCTTAAAGAGAACCTGGCTCGGACCGGGCAGTACGCCAAGGCCTCCACCACCAAT	7140
246	AACTTGCTTAAAGAGAACCTGGCTCGGACCGGGCAGTACGCCAAGGCCTCCACCACCAAT	7140
353	AACTTGCTTAAAGAGAACCTGGCTCGGACCGGGCAGTACGCCAAGGCCTCCACCACCAAT	7140

DAH	GGATTTCATCGGAAGGGAGGC Gtaatcgttacgaactgttagttctgtagaaaaacatgtaa	7124
29B	GGATTTCATCGGAAGGGAGGC Gtaatcgttacgaactgttagttctgtagaaaaacatgtaa	7200
e19	GGATTTCATCGGAAGGGAGGC Gtaatcgttacgaactgttagttctgtagaaaaacatgtaa	7200
74e19	GGATTTCATCGGAAGGGAGGC Gtaatcgttacgaactgttagttctgtagaaaaacatgtaa	7200
74	GGATTTCATCGGAAGGGAGGC Gtaatcgttacgaactgttagttctgtagaaaaacatgtaa	7200
211	GGATTTCATCGGAAGGGAGGC Gtaatcgttacgaactgttagttctgtagaaaaacatgtaa	7200
246	GGATTTCATCGGAAGGGAGGC Gtaatcgttacgaactgttagttctgtagaaaaacatgtaa	7200
353	GGATTTCATCGGAAGGGAGGC Gtaatcgttacgaactgttagttctgtagaaaaacatgtaa	7200

DAH	atagatgaggagaaggcagcatggatatagataagtagcaagtcttgcgcagccaag	7184
29B	atagatgaggagaaggcagcatggatatagataagtagcaagtcttgcgcagccaag	7260
e19	atagatgaggagaaggcagcatggatatagataagtagcaagtcttgcgcagccaag	7260
74e19	atagatgaggagaaggcagcatggatatagataagtagcaagtcttgcgcagccaag	7260
74	atagatgaggagaaggcagcatggatatagataagtagcaagtcttgcgcagccaag	7260
211	atagatgaggagaaggcagcatggatatagataagtagcaagtcttgcgcagccaag	7260
246	atagatgaggagaaggcagcatggatatagataagtagcaagtcttgcgcagccaag	7260
353	atagatgaggagaaggcagcatggatatagataagtagcaagtcttgcgcagccaag	7260

DAH	cttcggaaatttgactctctatgcctgcaaatttagggtgaggagatgttagtcca	7244
29B	cttcggaaatttgactctctatgcctgcaaatttagggtgaggagatgttagtcca	7320
e19	cttcggaaatttgactctctatgcctgcaaatttagggtgaggagatgttagtcca	7320
74e19	cttcggaaatttgactctctatgcctgcaaatttagggtgaggagatgttagtcca	7320
74	cttcggaaatttgactctctatgcctgcaaatttagggtgaggagatgttagtcca	7320
211	cttcggaaatttgactctctatgcctgcaaatttagggtgaggagatgttagtcca	7320
246	cttcggaaatttgactctctatgcctgcaaatttagggtgaggagatgttagtcca	7320
353	cttcggaaatttgactctctatgcctgcaaatttagggtgaggagatgttagtcca	7320

DAH	aagatatttgatatttcaatatgactggctagcaatccattgtaacattctagcgat	7304
29B	aagatatttgatatttcaatatgactggctagcaatccattgtaacattctagcgat	7380
e19	aagatatttgatatttcaatatgactggctagcaatccattgtaacattctagcgat	7380
74e19	aagatatttgatatttcaatatgactggctagcaatccattgtaacattctagcgat	7380
74	aagatatttgatatttcaatatgactggctagcaatccattgtaacattctagcgat	7380
211	aagatatttgatatttcaatatgactggctagcaatccattgtaacattctagcgat	7380
246	aagatatttgatatttcaatatgactggctagcaatccattgtaacattctagcgat	7380
353	aagatatttgatatttcaatatgactggctagcaatccattgtaacattctagcgat	7380

DAH	gaagcttcaaacaagaaggattgtttagttaaaatttgcattaatagaaaaattaagc	7364
29B	gaagcttcaaacaagaaggattgtttagttaaaatttgcattaatagaaaaattaagc	7440
e19	gaagcttcaaacaagaaggattgtttagttaaaatttgcattaatagaaaaattaagc	7440
74e19	gaagcttcaaacaagaaggattgtttagttaaaatttgcattaatagaaaaattaagc	7440
74	gaagcttcaaacaagaaggattgtttagttaaaatttgcattaatagaaaaattaagc	7440
211	gaagcttcaaacaagaaggattgtttagttaaaatttgcattaatagaaaaattaagc	7440
246	gaagcttcaaacaagaaggattgtttagttaaaatttgcattaatagaaaaattaagc	7440
353	gaagcttcaaacaagaaggattgtttagttaaaatttgcattaatagaaaaattaagc	7440

DAH	acattgtgtcagtctcgtaaaataaacacaccccccgtagatggatgacacccttctt	7424
29B	acattgtgtcagtctcgtaaaataaacacaccccccgtagatggatgacacccttctt	7500
e19	acattgtgtcagtctcgtaaaataaacacaccccccgtagatggatgacacccttctt	7500
74e19	acattgtgtcagtctcgtaaaataaacacaccccccgtagatggatgacacccttctt	7500
74	acattgtgtcagtctcgtaaaataaacacaccccccgtagatggatgacacccttctt	7500
211	acattgtgtcagtctcgtaaaataaacacaccccccgtagatggatgacacccttctt	7500
246	acattgtgtcagtctcgtaaaataaacacaccccccgtagatggatgacacccttctt	7500
353	acattgtgtcagtctcgtaaaataaacacaccccccgtagatggatgacacccttctt	7500

DAH	cttgggtgtcgaacccaaaattgtaaattttagattcatagagcctaatttcattctaa	7484
29B	cttgggtgtcgaacccaaaattgtaaattttagattcatagagcctaatttcattctaa	7560
e19	cttgggtgtcgaacccaaaattgtaaattttagattcatagagcctaatttcattctaa	7560
74e19	cttgggtgtcgaacccaaaattgtaaattttagattcatagagcctaatttcattctaa	7560
74	cttgggtgtcgaacccaaaattgtaaattttagattcatagagcctaatttcattctaa	7560
211	cttgggtgtcgaacccaaaattgtaaattttagattcatagagcctaatttcattctaa	7560
246	cttgggtgtcgaacccaaaattgtaaattttagattcatagagcctaatttcattctaa	7560
353	cttgggtgtcgaacccaaaattgtaaattttagattcatagagcctaatttcattctaa	7560

DAH	ttgttcgacgattaggcaaacgactatatggagtcgaagatatggaaacaaaggcttaat	7544
29B	ttgttcgacgattaggcaaacgactatatggagtcgaagatatggaaacaaaggcttaat	7620
e19	ttgttcgacgattaggcaaacgactatatggagtcgaagatatggaaacaaaggcttaat	7620
74e19	ttgttcgacgattaggcaaacgactatatggagtcgaagatatggaaacaaaggcttaat	7620
74	ttgttcgacgattaggcaaacgactatatggagtcgaagatatggaaacaaaggcttaat	7620
211	ttgttcgacgattaggcaaacgactatatggagtcgaagatatggaaacaaaggcttaat	7620
246	ttgttcgacgattaggcaaacgactatatggagtcgaagatatggaaacaaaggcttaat	7620
353	ttgttcgacgattaggcaaacgactatatggagtcgaagatatggaaacaaaggcttaat	7620

DAH	ttgctaagaacgagccaaaaagattgctaaccgccttcacgattttgttcaagccatc	7604
29B	ttgctaagaacgagccaaaaagattgctaaccgccttcacgattttgttcaagccatc	7680
e19	ttgctaagaacgagccaaaaagattgctaaccgccttcacgattttgttcaagccatc	7680
74e19	ttgctaagaacgagccaaaaagattgctaaccgccttcacgattttgttcaagccatc	7680
74	ttgctaagaacgagccaaaaagattgctaaccgccttcacgattttgttcaagccatc	7680
211	ttgctaagaacgagccaaaaagattgctaaccgccttcacgattttgttcaagccatc	7680
246	ttgctaagaacgagccaaaaagattgctaaccgccttcacgattttgttcaagccatc	7680
353	ttgctaagaacgagccaaaaagattgctaaccgccttcacgattttgttcaagccatc	7680

DAH	gaaccaccctaaccgcgcatgtataataccaatagttcgatttttcattttatgttttta	7664
29B	gaaccaccctaaccgcgcatgtataataccaatagttcgatttttcattttatgttttta	7740
e19	gaaccaccctaaccgcgcatgtataataccaatagttcgatttttcattttatgttttta	7740
74e19	gaaccaccctaaccgcgcatgtataataccaatagttcgatttttcattttatgttttta	7740
74	gaaccaccctaaccgcgcatgtataataccaatagttcgatttttcattttatgttttta	7740
211	gaaccaccctaaccgcgcatgtataataccaatagttcgatttttcattttatgttttta	7740
246	gaaccaccctaaccgcgcatgtataataccaatagttcgatttttcattttatgttttta	7740
353	gaaccaccctaaccgcgcatgtataataccaatagttcgatttttcattttatgttttta	7740

DAH	cgtatt	7670
29B	cgtatt	7746
e19	cgtatt	7746
74e19	cgtatt	7746
74	cgtatt	7746
211	cgtatt	7746
246	cgtatt	7746
353	cgtatt	7746

Fig S4. Master PROT alignment HR alleles with Wildtypes wDAH and 29B (HR)

DAH	MFNMPRGVTKS SKS KRG KIKMENDMAAATTTACTLGHICVLCRQEMLLDTCCRQAVEAV	60
29B	MFNMPRGVTKS SKS KRG KIKMENDMAAATTTACTLGHICVLCRQEMLLDTCCRQAVEAV	60
e19	MFNMPRGVTKS SKS KRG KIKMENDMAAATTTACTLGHICVLCRQEMLLDTCCRQAVEAV	60
74e19	MFNMPRGVTKS SKS KRG KIKMENDMAAATTTACTLGHICVLCRQEMLLDTCCRQAVEAV	60
74	MFNMPRGVTKS SKS KRG KIKMENDMAAATTTACTLGHICVLCRQEMLLDTCCRQAVEAV	60
211	MFNMPRGVTKS SKS KRG KIKMENDMAAATTTACTLGHICVLCRQEMLLDTCCRQAVEAV	60
246	MFNMPRGVTKS SKS KRG KIKMENDMAAATTTACTLGHICVLCRQEMLLDTCCRQAVEAV	60
353	MFNMPRGVTKS SKS KRG KIKMENDMAAATTTACTLGHICVLCRQEMLLDTCCRQAVEAV	60

DAH	DSPASSEEAYSSSNSSCQASSEISAAEVWFLSHDDIVLCRRPKFDEVETTGKKRDKVKCS	120
29B	DSPASSEEAYSSSNSSCQASSEISAAEVWFLSHDDIVLCRRPKFDEVETTGKKRDKVKCS	120
e19	DSPASSEEAYSSSNSSCQASSEISAAEVWFLSHDDIVLCRRPKFDEVETTGKKRDKVKCS	120
74e19	DSPASSEEAYSSSNSSCQASSEISAAEVWFLSHDDIVLCRRPKFDEVETTGKKRDKVKCS	120
74	DSPASSEEAYSSSNSSCQASSEISAAEVWFLSHDDIVLCRRPKFDEVETTGKKRDKVKCS	120
211	DSPASSEEAYSSSNSSCQASSEISAAEVWFLSHDDIVLCRRPKFDEVETTGKKRDKVKCS	120
246	DSPASSEEAYSSSNSSCQASSEISAAEVWFLSHDDIVLCRRPKFDEVETTGKKRDKVKCS	120
353	DSPASSEEAYSSSNSSCQASSEISAAEVWFLSHDDIVLCRRPKFDEVETTGKKRDKVKCS	120

DAH	GHQCSNECDDGSKNNRQQRENFNIFSNCHNILRTLQSLLLLMFNC G IFNKRRRRQHQQQ	180
29B	GHQCSNECDDGSKNNRQQRENFNIFSNCHNILRTLQSLLLLMFNC G IFNKRRRRQHQQQ	180
e19	GHQCSNECDDGSKNNRQQRENFNIFSNCHNILRTLQSLLLLMFNC G IFNKRRRRQHQQQ	180
74e19	GHQCSNECDDGSKNNRQQRENFNIFSNCHNILRTLQSLLLLMFNC G IFNKRRRRQHQQQ	180
74	GHQCSNECDDGSKNNRQQRENFNIFSNCHNILRTLQSLLLLMFNC G IFNKRRRRQHQQQ	180
211	GHQCSNECDDGSKNNRQQRENFNIFSNCHNILRTLQSLLLLMFNC G IFNKRRRRQHQQQ	180
246	GHQCSNECDDGSKNNRQQRENFNIFSNCHNILRTLQSLLLLMFNC G IFNKRRRRQHQQQ	180
353	GHQCSNECDDGSKNNRQQRENFNIFSNCHNILRTLQSLLLLMFNC G IFNKRRRRQHQQQ	180

DAH	HHHYQHHQHHHQHLQRQQANVS YTKFLLL LQT LAA ATTRLSLSPK NYK QQQQLQHNQQ	240
29B	HHHYQHHQHHHQHLQRQQANVS YTKFLLL LQT LAA ATTRLSLSPK NYK QQQQLQHNQQ	240
e19	HHHYQHHQHHHQHLQRQQANVS YTKFLLL LQT LAA ATTRLSLSPK NYK QQQQLQHNQQ	240
74e19	HHHYQHHQHHHQHLQRQQANVS YTKFLLL LQT LAA ATTRLSLSPK NYK QQQQLQHNQQ	240
74	HHHYQHHQHHHQHLQRQQANVS YTKFLLL LQT LAA ATTRLSLSPK NYK QQQQLQHNQQ	240
211	HHHYQHHQHHHQHLQRQQANVS YTKFLLL LQT LAA ATTRLSLSPK NYK QQQQLQHNQQ	240
246	HHHYQHHQHHHQHLQRQQANVS YTKFLLL LQT LAA ATTRLSLSPK NYK QQQQLQHNQQ	240
353	HHHYQHHQHHHQHLQRQQANVS YTKFLLL LQT LAA ATTRLSLSPK NYK QQQQLQHNQQ	240

DAH	LPRATPQQKQQEKDRHKCFHYKHNSYSPG I SLLL FILLANTLAIQAVVLP AHQ QHLLHN	300
29B	LPRATPQQKQQEKDRHKCFHYKHNSYSPG I SLLL FILLANTLAIQAVVLP AHQ QHLLHN	300
e19	LPRATPQQKQQEKDRHKCFHYKHNSYSPG I SLLL FILLANTLAIQAVVLP AHQ QHLLHN	300
74e19	LPRATPQQKQQEKDRHKCFHYKHNSYSPG I SLLL FILLANTLAIQAVVLP AHQ QHLLHN	300
74	LPRATPQQKQQEKDRHKCFHYKHNSYSPG I SLLL FILLANTLAIQAVVLP AHQ QHLLHN	300
211	LPRATPQQKQQEKDRHKCFHYKHNSYSPG I SLLL FILLANTLAIQAVVLP AHQ QHLLHN	300
246	LPRATPQQKQQEKDRHKCFHYKHNSYSPG I SLLL FILLANTLAIQAVVLP AHQ QHLLHN	300
353	LPRATPQQKQQEKDRHKCFHYKHNSYSPG I SLLL FILLANTLAIQAVVLP AHQ QHLLHN	300

DAH	DIADGLDKTALSVSGTQSRWTRSESNPTMRLSQNVKPCSKMDIRNMVSHFNQLENCTVIE	360
29B	DIADGLDKTALSVSGTQSRWTRSESNPTMRLSQNVKPCSKMDIRNMVSHFNQLENCTVIE	360
e19	DIADGLDKTALSVSGTQSRWTRSESNPTMRLSQNVKPCSKMDIRNMVSHFNQLENCTVIE	360
74e19	DIADGLDKTALSVSGTQSRWTRSESNPTMRLSQNVKPCSKMDIRNMVSHFNQLENCTVIE	360
74	DIADGLDKTALSVSGTQSRWTRSESNPTMRLSQNVKPCSKMDIRNMVSHFNQLENCTVIE	360
211	DIADGLDKTALSVSGTQSRWTRSESNPTMRLSQNVKPCSKMDIRNMVSHFNQLENCTVIE	360
246	DIADGLDKTALSVSGTQSRWTRSESNPTMRLSQNVKPCSKMDIRNMVSHFNQLENCTVIE	360
353	DIADGLDKTALSVSGTQSRWTRSESNPTMRLSQNVKPCSKMDIRNMVSHFNQLENCTVIE	360

DAH	GFLLIDLINDASPLNRSFPKLTEVTDYIIYRTGLHSLSKIFPNLSVIRGNKLF DGYAL	420
29B	GFLLIDLINDASPLNRSFPKLTEVTDYIIYRTGLHSLSKIFPNLSVIRGNKLF DGYAL	420
e19	GFLLIDLINDASPLNRSFPKLTEVTDYIIYRTGLHSLSKIFPNLSVIRGNKLF DGYAL	420
74e19	GFLLIDLINDASPLNRSFPKLTEVTDYIIYRTGLHSLSKIFPNLSVIRGNKLF DGYAL	420
74	GFLLIDLINDASPLNRSFPKLTEVTDYIIYRTGLHSLSKIFPNLSVIRGNKLF DGYAL	420
211	GFLLIDLINDASPLNRSFPKLTEVTDYIIYRTGLHSLSKIFPNLSVIRGNKLF DGYAL	420
246	GFLLIDLINDASPLNRSFPKLTEVTDYIIYRTGLHSLSKIFPNLSVIRGNKLF DGYAL	420
353	GFLLIDLINDASPLNRSFPKLTEVTDYIIYRTGLHSLSKIFPNLSVIRGNKLF DGYAL	420

DAH	VVYSNFDLMDLGLHKLRSITRGGVRIEKNHKLCYDRTIDWLEILAENETQLVVLTENGKE	480
29B	VVYSNFDLMDLGLHKLRSITRGGVRIEKNHKLCYDRTIDWLEILAENETQLVVLTENGKE	480
e19	VVYSNFDLMDLGLHKLRSITRGGVRIEKNHKLCYDRTIDWLEILAENETQLVVLTENGKE	480
74e19	VVYSNFDLMDLGLHKLRSITRGGVRIEKNHKLCYDRTIDWLEILAENETQLVVLTENGKE	480
74	VVYSNFDLMDLGLHKLRSITRGGVRIEKNHKLCYDRTIDWLEILAENETQLVVLTENGKE	480
211	VVYSNFDLMDLGLHKLRSITRGGVRIEKNHKLCYDRTIDWLEILAENETQLVVLTENGKE	480
246	VVYSNFDLMDLGLHKLRSITRGGVRIEKNHKLCYDRTIDWLEILAENETQLVVLTENGKE	480
353	VVYSNFDLMDLGLHKLRSITRGGVRIEKNHKLCYDRTIDWLEILAENETQLVVLTENGKE	480

DAH	KECRLSKCPGEIRIEEGHDTTAIEGELNASCQLHNNRRLCWNSKLCQTKCPEKCRNNCID	540
29B	KECRLSKCPGEIRIEEGHDTTAIEGELNASCQLHNNRRLCWNSKLCQTKCPEKCRNNCID	540
e19	KECRLSKCPGEIRIEEGHDTTAIEGELNASCQLHNNRRLCWNSKLCQTKCPEKCRNNCID	540
74e19	KECRLSKCPGEIRIEEGHDTTAIEGELNASCQLHNNRRLCWNSKLCQTKCPEKCRNNCID	540
74	KECRLSKCPGEIRIEEGHDTTAIEGELNASCQLHNNRRLCWNSKLCQTKCPEKCRNNCID	540
211	KECRLSKCPGEIRIEEGHDTTAIEGELNASCQLHNNRRLCWNSKLCQTKCPEKCRNNCID	540
246	KECRLSKCPGEIRIEEGHDTTAIEGELNASCQLHNNRRLCWNSKLCQTKCPEKCRNNCID	540
353	KECRLSKCPGEIRIEEGHDTTAIEGELNASCQLHNNRRLCWNSKLCQTKCPEKCRNNCID	540

DAH	EHTCCSQDCLGGCVIDKNGNESCISCRNSFNNICMDSCPCKGYQFDSRCVTANECITLT	600
29B	EHTCCSQDCLGGCVIDKNGNESCISCRNSFNNICMDSCPCKGYQFDSRCVTANECITLT	600
e19	EHTCCSQDCLGGCVIDKNGNESCISCRNSFNNICMDSCPCKGYQFDSRCVTANECITLT	600
74e19	EHTCCSQDCLGGCVIDKNGNESCISCRNSFNNICMDSCPCKGYQFDSRCVTANECITLT	600
74	EHTCCSQDCLGGCVIDKNGNESCISCRNSFNNICMDSCPCKGYQFDSRCVTANECITLT	600
211	EHTCCSQDCLGGCVIDKNGNESCISCRNSFNNICMDSCPCKGYQFDSRCVTANECITLT	600
246	EHTCCSQDCLGGCVIDKNGNESCISCRNSFNNICMDSCPCKGYQFDSRCVTANECITLT	600
353	EHTCCSQDCLGGCVIDKNGNESCISCRNSFNNICMDSCPCKGYQFDSRCVTANECITLT	600

DAH	KFETNSVYSGIPYNGQCITHCPTGYQKSENKRMCEPCPGKCDKECSSGLIDSLERAREF	660
29B	KFETNSVYSGIPYNGQCITHCPTGYQKSENKRMCEPCPGKCDKECSSGLIDSLERAREF	660
e19	KFETNSVYSGIPYNGQCITHCPTGYQKSENKRMCEPCPGKCDKECSSGLIDSLERAREF	660
74e19	KFETNSVYSGIPYNGQCITHCPTGYQKSENKRMCEPCPGKCDKECSSGLIDSLERAREF	660
74	KFETNSVYSGIPYNGQCITHCPTGYQKSENKRMCEPCPGKCDKECSSGLIDSLERAREF	660
211	KFETNSVYSGIPYNGQCITHCPTGYQKSENKRMCEPCPGKCDKECSSGLIDSLERAREF	660
246	KFETNSVYSGIPYNGQCITHCPTGYQKSENKRMCEPCPGKCDKECSSGLIDSLERAREF	660
353	KFETNSVYSGIPYNGQCITHCPTGYQKSENKRMCEPCPGKCDKECSSGLIDSLERAREF	660

DAH	HGCTIITGTEPLTISIKRESGAHVMDELKYGLAAVHKIQSSLMVHLTYGLKSLKFQSLT	720
29B	HGCTIITGTEPLTISIKRESGAHVMDELKYGLAAVHKIQSSLMVHLTYGLKSLKFQSLT	720
e19	HGCTIITGTEPLTISIKRESGAHVMDELKYGLAAVHKIQSSLMVHLTYGLKSLKFQSLT	720
74e19	HGCTIITGTEPLTISIKRESGAHVMDELKYGLAAVHKIQSSLMVHLTYGLKSLKFQSLT	720
74	HGCTIITGTEPLTISIKRESGAHVMDELKYGLAAVHKIQSSLMVHLTYGLKSLKFQSLT	720
211	HGCTIITGTEPLTISIKRESGAHVMDELKYGLAAVHKIQSSLMVHLTYGLKSLKFQSLT	720
246	HGCTIITGTEPLTISIKRESGAHVMDELKYGLAAVHKIQSSLMVHLTYGLKSLKFQSLT	720
353	HGCTIITGTEPLTISIKRESGAHVMDELKYGLAAVHKIQSSLMVHLTYGLKSLKFQSLT	720

DAH	EISGDPPMDADKYALYVLDNRDLDELWGPQNQTVFIRKGFFHFNPKLCVSTINQLLPML	780
29B	EISGDPPMDADKYALYVLDNRDLDELWGPQNQTVFIRKGFFHFNPKLCVSTINQLLPML	780
e19	EISGDPPMDADKYALYVLDNRDLDELWGPQNQTVFIRKGFFHFNPKLCVSTINQLLPML	780
74e19	EISGDPPMDADKYALYVLDNRDLDELWGPQNQTVFIRKGFFHFNPKLCVSTINQLLPML	780
74	EISGDPPMDADKYALYVLDNRDLDELWGPQNQTVFIRKGFFHFNPKLCVSTINQLLPML	780
211	EISGDPPMDADKYALYVLDNRDLDELWGPQNQTVFIRKGFFHFNPKLCVSTINQLLPML	780
246	EISGDPPMDADKYALYVLDNRDLDELWGPQNQTVFIRKGFFHFNPKLCVSTINQLLPML	780
353	EISGDPPMDADKYALYVLDNRDLDELWGPQNQTVFIRKGFFHFNPKLCVSTINQLLPML	780

DAH	ASKPKFFEKSVDVGADSNGNRGSCGTAVLNVTLQSVGANSAMLNVTTKVEIGEPQKPSNAT	840
29B	ASKPKFFEKSVDVGADSNGNRGSCGTAVLNVTLQSVGANSAMLNVTTKVEIGEPQKPSNAT	840
e19	ASKPKFFEKSVDVGADSNGNRGSCGTAVLN D TLQSVGANSAMLNVTTKVEIGEPQKPSNAT	840
74e19	ASKPKFFEKSVDVGADSNGNRGSCGTAVLN D TLQSVGANSAMLNVTTKVEIGEPQKPSNAT	840
74	ASKPKFFEKSVDVGADSNGNRGSCGTAVLNVTLQSVGANSAMLNVTTKVEIGEPQKPSNAT	840
211	ASKPKFFEKSVDVGADSNGNRGSCGTAVLNVTLQSVGANSAMLNVTTKVEIGEPQKPSNAT	840
246	ASKPKFFEKSVDVGADSNGNRGSCGTAVLNVTLQSVGANSAMLNVTTKVEIGEPQKPSNAT	840
353	ASKPKFFEKSVDVGADSNGNRGSCGTAVLNVTLQSVGANSAMLNVTTKVEIGEPQKPSNAT	840

DAH	IVFKDPRAFIGVFYHMIDPYGNSTKSSDDPCDDRWKVSSPEKSGVMVLSNLIPYTNYSY	900
29B	IVFKDPRAFIGVFYHMIDPYGNSTKSSDDPCDDRWKVSSPEKSGVMVLSNLIPYTNYSY	900
e19	IVFKDPRAFIGVFYHMIDPYGNSTKSSDDPCDDRWKVSSPEKSGVMVLSNLIPYTNYSY	900
74e19	IVFKDPRAFIGVFYHMIDPYGNSTKSSDDPCDDRWKVSSPEKSGVMVLSNLIPYTNYSY	900
74	IVFKDPRAFIGVFYHMIDPYGNSTKSSDDPCDDRWKVSSPEKSGVMVLSNLIPYTNYSY	900
211	IVFKDPRAFIGVFYHMIDPYGNSTKSSDDPCDDRWKVSSPEKSGVMVLSNLIPYTNYSY	900
246	IVFKDPRAFIGVFYHMIDPYGNSTKSSDDPCDDRWKVSSPEKSGVMVLSNLIPYTNYSY	900
353	IVFKDPRAFIGVFYHMIDPYGNSTKSSDDPCDDRWKVSSPEKSGVMVLSNLIPYTNYSY	900

DAH	YVRTMAISSELTN AESDVKNFR TNPGRPSKVTEVVATAISDSKINVTWSYLDKPYGV LTR	960
29B	YVRTMAISSELTN AESDVKNFR TNPGRPSKVTEVVATAISDSKINVTWSYLDKPYGV LTR	960
e19	YVRTMAISSELTN AESDVKNFR TNPGRPSKVTEVVATAISDSKINVTWSYLDKPYGV LTR	960
74e19	YVRTMAISSELTN AESDVKNFR TNPGRPSKVTEVVATAISDSKINVTWSYLDKPYGV LTR	960
74	YVRTMAISSELTN AESDVKNFR TNPGRPSKVTEVVATAISDSKINVTWSYLDKPYGV LTR	960
211	YVRTMAISSELTN AESDVKNFR TNPGRPSKVTEVVATAISDSKINVTWSYLDKPYGV LTR	960
246	YVRTMAISSELTN AESDVKNFR TNPGRPSKVTEVVATAISDSKINVTWSYLDKPYGV LTR	960
353	YVRTMAISSELTN AESDVKNFR TNPGRPSKVTEVVATAISDSKINVTWSYLDKPYGV LTR	960

DAH	YFIKAKLINRPTRN NN RDY CTEPLVKAMENDLPATTPTKKISDPLAGDCKC VEGSKKTSS	1020
29B	YFIKAKLINRPTRN NN RDY CTEPLVKAMENDLPATTPTKKISDPLAGDCKC VEGSKKTSS	1020
e19	YFIKAKLINRPTRN NN RDY CTEPLVKAMENDLPATTPTKKISDPLAGDCKC VEGSKKTSS	1020
74e19	YFIKAKLINRPTRN NN RDY CTEPLVKAMENDLPATTPTKKISDPLAGDCKC VEGSKKTSS	1020
74	YFIKAKLINRPTRN NN RDY CTEPLVKAMENDLPATTPTKKISDPLAGDCKC VEGSKKTSS	1020
211	YFIKAKLINRPTRN NN RDY CTEPLVKAMENDLPATTPTKKISDPLAGDCKC VEGSKKTSS	1020
246	YFIKAKLINRPTRN NN RDY CTEPLVKAMENDLPATTPTKKISDPLAGDCKC VEGSKKTSS	1020
353	YFIKAKLINRPTRN NN RDY CTEPLVKAMENDLPATTPTKKISDPLAGDCKC VEGSKKTSS	1020

75		
DAH	QEYDDRKVQAGMЕFENALQNFI FVPNIRKSKNGSSDKSDGAEGAALDSNAIPNGGATNPS	1080
29B	QEYDDRKVQAGMЕFENALQNFI FVPNIRKSKNGSSDKSDGAEGAALDSNAIPNGGATNPS	1080
e19	QEYDDRKVQAGMЕFENALQNFI FVPNIRKSKNGSSDKSDGAEGAALDSNAIPNGGATNPS	1080
74e19	QEYDDRKVQAGMЕFENALQNFI FVPNIRKSKNGSSDKSDGAEGAALDSNAIPNGGATNPS	1080
74	QEYDDRKVQAGMЕFENALQNFI FVPNIRKSKNGSSDKSDGAEGAALDSNAIPNGGATNPS	1080
211	QEYDDRKVQAGMЕFENALQNFI FVPNIRKSKNGSSDKSDGAEGAALDSNAIPNGGATNPS	1080
246	QEYDDRKVQAGMЕFENALQNFI FVPNIRKSKNGSSDKSDGAEGAALDSNAIPNGGATNPS	1080
353	QEYDDRKVQAGMЕFENALQNFI FVPNIRKSKNGSSDKSDGAEGAALDSNAIPNGGATNPS	1080

DAH	RRRRDALEPELDDVEGSVLLRHVR SITDDTDAFFEKDENTYKDEEDLSSNKQFYEVFA	1140
29B	RRRRDALEPELDDVEGSVLLRHVR SITDDTDAFFEKDENTYKDEEDLSSNKQFYEVFA	1140
e19	RRRRDALEPELDDVEGSVLLRHVR SITDDTDAFFEKDENTYKDEEDLSSNKQFYEVFA	1140
74e19	RRRRDALEPELDDVEGSVLLRHVR SITDDTDAFFEKDENTYKDEEDLSSNKQFYEVFA	1140
74	RRRRDALEPELDDVEGSVLLRHVR SITDDTDAFFEKDENTYKDEEDLSSNKQFYEVFA	1140
211	RRRRDALEPELDDVEGSVLLRHVR SITDDTDAFFEKDENTYKDEEDLSSNKQFYEVFA	1140
246	RRRRDALEPELDDVEGSVLLRHVR SITDDTDAFFEKDENTYKDEEDLSSNKQFYEVFA	1140
353	RRRRDALEPELDDVEGSVLLRHVR SITDDTDAFFEKDENTYKDEEDLSSNKQFYEVFA	1140

DAH	KELPPNQTHFVFEKLHRHFTRYAIFVVACREEIPSEKLRDTSFKSLCSDYDTVFQTTKRK	1200
29B	KELPPNQTHFVFEKLHRHFTRYAIFVVACREEIPSEKLRDTSFKSLCSDYDTVFQTTKRK	1200
e19	KELPPNQTHFVFEKLHRHFTRYAIFVVACREEIPSEKLRDTSFKSLCSDYDTVFQTTKRK	1200
74e19	KELPPNQTHFVFEKLHRHFTRYAIFVVACREEIPSEKLRDTSFKSLCSDYDTVFQTTKRK	1200
74	KELPPNQTHFVFEKLHRHFTRYAIFVVACREEIPSEKLRDTSFKSLCSDYDTVFQTTKRK	1200
211	KELPPNQTHFVFEKLHRHFTRYAIFVVACREEIPSEKLRDTSFKSLCSDYDTVFQTTKRK	1200
246	KELPPNQTHFVFEKLHRHFTRYAIFVVACREEIPSEKLRDTSFKSLCSDYDTVFQTTKRK	1200
353	KELPPNQTHFVFEKLHRHFTRYAIFVVACREEIPSEKLRDTSFKSLCSDYDTVFQTTKRK	1200

DAH	KFADIVMDLKVDLEHANNTESPVRVRWTTPVDPNGEIVTYEVAYKLQKPDQVEEKKCIPA	1260
29B	KFADIVMDLKVDLEHANNTESPVRVRWTTPVDPNGEIVTYEVAYKLQKPDQVEEKKCIPA	1260
e19	KFADIVMDLKVDLEHANNTESPVRVRWTTPVDPNGEIVTYEVAYKLQKPDQVEEKKCIPA	1260
74e19	KFADIVMDLKVDLEHANNTESPVRVRWTTPVDPNGEIVTYEVAYKLQKPDQVEEKKCIPA	1260
74	KFADIVMDLKVDLEHANNTESPVRVRWTTPVDPNGEIVTYEVAYKLQKPDQVEEKKCIPA	1260
211	KFADIVMDLKVDLEHANNTESPVRVRWTTPVDPNGEIVTYEVAYKLQKPDQVEEKKCIPA	1260
246	KFADIVMDLKVDLEHANNTESPVRVRWTTPVDPNGEIVTYEVAYKLQKPDQVEEKKCIPA	1260
353	KFADIVMDLKVDLEHANNTESPVRVRWTTPVDPNGEIVTYEVAYKLQKPDQVEEKKCIPA	1260

DAH	ADFNQTAGYLIKLN EGLYSFRVRANSIAGYGDFTEVEHIKEPPP SYAKVFFWLLGIGLA	1320
29B	ADFNQTAGYLIKLN EGLYSFRVRANSIAGYGDFTEVEHIKEPPP SYAKVFFWLLGIGLA	1320
e19	ADFNQTAGYLIKLN EGLYSFRVRANSIAGYGDFTEVEHIKEPPP SYAKVFFWLLGIGLA	1320
74e19	ADFNQTAGYLIKLN EGLYSFRVRANSIAGYGDFTEVEHIKEPPP SYAKVFFWLLGIGLA	1320
74	ADFNQTAGYLIKLN EGLYSFRVRANSIAGYGDFTEVEHIKEPPP SYAKVFFWLLGIGLA	1320
211	ADFNQTAGYLIKLN EGLYSFRVRANSIAGYGDFTEVEHIKEPPP SYAKVFFWLLGIGLA	1320
246	ADFNQTAGYLIKLN EGLYSFRVRANSIAGYGDFTEVEHIKEPPP SYAKVFFWLLGIGLA	1320
353	ADFNQTAGYLIKLN EGLYSFRVRANSIAGYGDFTEVEHIKEPPP SYAKVFFWLLGIGLA	1320

DAH	FLIVSLFGYVCYLHKRKVPSDLHMNTEVNPFYASMQYIPDDWEVLRENIIQLAPLGQGS	1380
29B	FLIVSLFGYVCYLHKRKVPSDLHMNTEVNPFYASMQYIPDDWEVLRENIIQLAPLGQGS	1380
e19	FLIVSLFGYVCYLHKRKVPSDLHMNTEVNPFYASMQYIPDDWEVLRENIIQLAPLGQGS	1380
74e19	FLIVSLFGYVCYLHKRKVPSDLHMNTEVNPFYASMQYIPDDWEVLRENIIQLAPLGQGS	1380
74	FLIVSLFGYVCYLHKRKVPSDLHMNTEVNPFYASMQYIPDDWEVLRENIIQLAPLGQGS	1380
211	FLIVSLFGYVCYLHKRKVPSDLHMNTEVNPFYASMQYIPDDWEVLRENIIQLAPLGQGS	1380
246	FLIVSLFGYVCYLHKRKVPSDLHMNTEVNPFYASMQYIPDDWEVLRENIIQLAPLGQGS	1380
353	FLIVSLFGYVCYLHKRKVPSDLHMNTEVNPFYASMQYIPDDWEVLRENIIQLAPLGQGS	1380

	FGMVYEGILKSFPNGVDERCAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGVC	1440
DAH	FGMVYEGILKSFPNGVDERCAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGVC	1440
29B	FGMVYEGILKSFPNGVDERCAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGVC	1440
e19	FGMVYEGILKSFPNGVDERCAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGVC	1440
74e19	FGMVYEGILKSFPNGVDERCAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGVC	1440
74	FGMVYEGILKSFPNGVDERCAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGVC	1440
211	FGMVYEGILKSFPNGVDERCAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGVC	1440
246	FGMVYEGILKSFPNGVDERCAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGVC	1440
353	FGMVYEGILKSFPNGVDERCAIKTVNENATDRERTNFLSEASVMKEFDTYHVVRLLGVC	1440
	*****	*****
DAH	SRGPALVVMELMKGDLKSYLRAHRPEERDEAMMTYLNRIGVTGNVPQPPTYGRIVQMAI	1500
29B	SRGPALVVMELMKGDLKSYLRAHRPEERDEAMMTYLNRIGVTGNVPQPPTYGRIVQMAI	1500
e19	SRGPALVVMELMKGDLKSYLRAHRPEERDEAMMTYLNRIGVTGNVPQPPTYGRIVQMAI	1500
74e19	SRGPALVVMELMKGDLKSYLRAHRPEERDEAMMTYLNRIGVTGNVPQPPTYGRIVQMAI	1500
74	SRGPALVVMELMKGDLKSYLRAHRPEERDEAMMTYLNRIGVTGNVPQPPTYGRIVQMAI	1500
211	SRGPALVVMELMKGDLKSYLRAHRPEERDEAMMTYLNRIGVTGNVPQPPTYGRIVQMAI	1500
246	SRGPALVVMELMKGDLKSYLRAHRPEERDEAMMTYLNRIGVTGNVPQPPTYGRIVQMAI	1500
353	SRGPALVVMELMKGDLKSYLRAHCPEERDEAMMTYLNRIGVTGNVPQPPTYGRIVQMAI	1500
	*****	*****
DAH	EIADGMAYLAACKFVHDLAARNCMVADDLTVKIGDFGMRDIFYETDYYRKGTGLLPVR	1560
29B	EIADGMAYLAACKFVHDLAARNCMVADDLTVKIGDFGMRDIFYETDYYRKGTGLLPVR	1560
e19	EIADGMAYLAACKFVHDLAARNCMVADDLTVKIGDFGMRDIFYETDYYRKGTGLLPVR	1560
74e19	EIADGMAYLAACKFVHDLAARNCMVADDLTVKIGDFGMRDIFYETDYYRKGTGLLPVR	1560
74	EIADGMAYLAACKFVHDLAARNCMVADDLTVKIGDFGMRDIFYETDYYRKGTGLLPVR	1560
211	EIADGMAYLAACKFVHDLAARNCMVADDLTVKIGDFGMRDIFYETDYYRKGTGLLPVR	1560
246	EIADGMAYLAACKFVHDLAARNCMVADDLTVKIGDFGMRDIFYETDYYRKGTGLLPVR	1560
353	EIADGMAYLAACKFVHDLAARNCMVADDLTVKIGDFGMRDIFYETDYYRKGTGLLPVR	1560
	*****	*****
DAH	WMPPESLRDGVYSSASDVFSFGVVLWEMATLAAQPYQGLSNEQLRYVIDGGVMERPENC	1620
29B	WMPPESLRDGVYSSASDVFSFGVVLWEMATLAAQPYQGLSNEQLRYVIDGGVMERPENC	1620
e19	WMPPESLRDGVYSSASDVFSFGVVLWEMATLAAQPYQGLSNEQLRYVIDGGVMERPENC	1620
74e19	WMPPESLRDGVYSSASDVFSFGVVLWEMATLAAQPYQGLSNEQLRYVIDGGVMERPENC	1620
74	WMPPESLRDGVYSSASDVFSFGVVLWEMATLAAQPYQGLSNEQLRYVIDGGVMERPENC	1620
211	WMPPESLRDGVYSSASDVFSFGVVLWEMATLAAQPYQGLSNEQLRYVIDGGVMERPENC	1620
246	WMPPESLRDGVYSSASDVFSFGVVLWEMATLAAQPYQGLSNEQLRYVIDGGVMERPENC	1620
353	WMPPESLRDGVYSSASDVFSFGVVLWEMATLAAQPYQGLSNEQLRYVIDGGVMERPENC	1620
	*****	*****
DAH	PDFLHKLMQRWCWHRSSARPSFLDIAYLEPQCPNSQFKEVSFYHSEAGLQHREKERKER	1680
29B	PDFLHKLMQRWCWHRSSARPSFLDIAYLEPQCPNSQFKEVSFYHSEAGLQHREKERKER	1680
e19	PDFLHKLMQRWCWHRSSARPSFLDIAYLEPQCPNSQFKEVSFYHSEAGLQHREKERKER	1680
74e19	PDFLHKLMQRWCWHRSSARPSFLDIAYLEPQCPNSQFKEVSFYHSEAGLQHREKERKER	1680
74	PDFLHKLMQRWCWHRSSARPSFLDIAYLEPQCPNSQFKEVSFYHSEAGLQHREKERKER	1680
211	PDFLHKLMQRWCWHRSSARPSFLDIAYLEPQCPNSQFKEVSFYHSEAGLQHREKERKER	1680
246	PDFLHKLMQRWCWHRSSARPSFLDIAYLEPQCPNSQFKEVSFYHSEAGLQHREKERKER	1680
353	PDFLHKLMQRWCWHRSSARPSFLDIAYLEPQCPNSQFKEVSFYHSEAGLQHREKERKER	1680
	*****	*****
DAH	NQLDAFAAVPLDQDLQDREQQEDATTPLRMGDYQQNSSLQPPESPIAMVDDQGSHPFS	1740
29B	NQLDAFAAVPLDQDLQDREQQEDATTPLRMGDYQQNSSLQPPESPIAMVDDQGSHPFS	1740
e19	NQLDAFAAVPLDODLQDREQQEDATTPLRMGDYQQNSSLQPPESPIAMVDDQGSHPFS	1740
74e19	NQLDAFAAVPLDQDLQDREQQEDATTPLRMGDYQQNSSLQPPESPIAMVDDQGSHPFS	1740
74	NQLDAFAAVPLDQDLQDREQQEDATTPLRMGDYQQNSSLQPPESPIAMVDDQGSHPFS	1740
211	NQLDAFAAVPLDQDLQDREQQEDATTPLRMGDYQQNSSLQPPESPIAMVDDQGSHPFS	1740
246	NQLDAFAAVPLDQDLQDREQQEDATTPLRMGDYQQNSSLQPPESPIAMVDDQGSHPFS	1740
353	NQLDAFAAVPLDQDLQDREQQEDATTPLRMGDYQQNSSLQPPESPIAMVDDQGSHPFS	1740
	*****	*****

	LPSGFIASSTPDGQTVMATAFQNIPAAQGDISATYVVPADALDGDGRGYEIYDPSPKAE	1800
29B	LPSGFIASSTPDGQTVMATAFQNIPAAQGDISATYVVPADALDGDGRGYEIYDPSPKAE	1800
e19	LPSGFIASSTPDGQTVMATAFQNIPAAQGDISATYVVPADALDGDGRGYEIYDPSPKAE	1800
74e19	LPSGFIASSTPDGQTVMATAFQNIPAAQGDISATYVVPADALDGDGRGYEIYDPSPKAE	1800
74	LPSGFIASSTPDGQTVMATAFQNIPAAQGDISATYVVPADALDGDGRGYEIYDPSPKAE	1800
211	LPSGFIASSTPDGQTVMATAFQNIPAAQGDISATYVVPADALDGDGRGYEIYDPSPKAE	1800
246	LPSGFIASSTPDGQTVMATAFQNIPAAQGDISATYVVPADALDGDGRGYEIYDPSPKAE	1800
353	LPSGFIASSTPDGQTVMATAFQNIPAAQGDISATYVVPADALDGDGRGYEIYDPSPKAE	1800

	LPTSRSGSTGGGKLSGEQHLLPRKGRQPTIMSSMPDDVIGSSLQPSSTAASSNASSH	1860
29B	LPTSRSGSTGGGKLSGEQHLLPRKGRQPTIMSSMPDDVIGSSLQPSSTAASSNASSH	1860
e19	LPTSRSGSTGGGKLSGEQHLLPRKGRQPTIMSSMPDDVIGSSLQPSSTAASSNASSH	1860
74e19	LPTSRSGSTGGGKLSGEQHLLPRKGRQPTIMSSMPDDVIGSSLQPSSTAASSNASSH	1860
74	LPTSRSGSTGGGKLSGEQHLLPRKGRQPTIMSSMPDDVIGSSLQPSSTAASSNASSH	1860
211	LPTSRSGSTGGGKLSGEQHLLPRKGRQPTIMSSMPDDVIGGSSVQPSSTAASSNASSH	1860
246	LPTSRSGSTGGGKLSGEQHLLPRKGRQPTIMSSMPDDVIGSSLQPSSTAASSNASSH	1860
353	LPTSRSGSTGGGKLSGEQHLLPRKGRQPTIMSSMPDDVIGSSLQPSSTAASSNASSH	1860
	*****:*****	
	TGRPSLKKTVADSVRNKANFINRHLFNHKRTGSNASHKSNASNAPSTSNTNLSPVAM	1920
29B	TGRPSLKKTVADSVRNKANFINRHLFNHKRTGSNASHKSNASNAPSTSNTNLSPVAM	1920
e19	TGRPSLKKTVADSVRNKANFINRHLFNHKRTGSNASHKSNASNAPSTSNTNLSPVAM	1920
74e19	TGRPSLKKTVADSVRNKANFINRHLFNHKRTGSNASHKSNASNAPSTSNTNLSPVAM	1920
74	TGRPSLKKTVADSVRNKANFINRHLFNHKRTGSNASHKSNASNAPSTSNTNLSPVAM	1920
211	TGRPSLKKTVADSVRNKANFINRHLFNHKRTGSNASHKSNASNAPSTSNTNLSPVAM	1920
246	TGRPSLKKTVADSVRNKANFINRHLFNHKRTGSNASHKSNASNAPSTSNTNLSPVAM	1920
353	TGRPSLKKTVADSVRNKANFINRHLFNHKRTGSNASHKSNASNAPSTSNTNLSPVAM	1920
	*****:*****	
	GNLGTIESGGSGSAGSYTGPFRYTPSATPGGGSGMAISDNPYRLDESIASEQATILT	1980
29B	GNLGTIESGGSGSAGSYTGPFRYTPSATPGGGSGMAISDNPYRLDESIASEQATILT	1980
e19	GNLGTIESGGSGSAGSYTGPFRYTPSATPGGGSGMAISDNPYRLDESIASEQATILT	1980
74e19	GNLGTIESGGSGSAGSYTGPFRYTPSATPGGGSGMAISDNPYRLDESIASEQATILT	1980
74	GNLGTIESGGSGSAGSYTGPFRYTPSATPGGGSGMAISDNPYRLDESIASEQATILT	1980
211	GNLGTIESGGSGSAGSYTGPFRYTPSATPGGGSGMAISDNPYRLDESIASEQATILT	1980
246	GNLGTIESGGSGSAGSYTGPFRYTPSATPGGGSGMAISDNPYRLDESIASEQATILT	1980
353	GNLGTIESGGSGSAGSYTGPFRYTPSATPGGGSGMAISDNPYRLDESIASEQATILT	1980

	TSSPNPNEYEMMHPPTSLVSTNPYMPMNETPVQMAGVTISHNPNEYQPMQAPLNARQSQSS	2040
29B	TSSPNPNEYEMMHPPTSLVSTNPYMPMNETPVQMAGVTISHNPNEYQPMQAPLNARQSQSS	2040
e19	TSSPNPNEYEMMHPPTSLVSTNPYMPMNETPVQMAGVTISHNPNEYQPMQAPLNARQSQSS	2040
74e19	TSSPNPNEYEMMHPPTSLVSTNPYMPMNETPVQMAGVTISHNPNEYQPMQAPLNARQSQSS	2040
74	TSSPNPNEYEMMHPPTSLVSTNPYMPMNETPVQMAGVTISHNPNEYQPMQAPLNARQSQSS	2040
211	TSSPNPNEYEMMHPPTSLVSTNPYMPMNETPVQMAGVTISHNPNEYQPMQAPLNARQSQSS	2040
246	TSSPNPNEYEMMHPPTSLVSTNPYMPMNETPVQMAGVTISHNPNEYQPMQAPLNARQSQSS	2040
353	TSSPNPNEYEMMHPPTSLVSTNPYMPMNETPVQMAGVTISHNPNEYQPMQAPLNARQSQSS	2040

	SDEDNEQEEDDEDDEDVVDEHVEHIKMERMPLSRPQRALPSKTQPPRSRSVSQTRKSP	2100
29B	SDEDNEQEEDDEDDEDVVDEHVEHIKMERMPLSRPQRALPSKTQPPRSRSVSQTRKSP	2100
e19	SDEDNEQEEDDEDDEDVVDEHVEHIKMERMPLSRPQRALPSKTQPPRSRSVSQTRKSP	2100
74e19	SDEDNEQEEDDEDDEDVVDEHVEHIKMERMPLSRPQRALPSKTQPPRSRSVSQTRKSP	2100
74	SDEDNEQEEDDEDDEDVVDEHVEHIKMERMPLSRPQRALPSKTQPPRSRSVSQTRKSP	2100
211	SDEDNEQEEDDEDDEDVVDEHVEHIKMERMPLSRPQRALPSKTQPPRSRSVSQTRKSP	2100
246	SDEDNEQEEDDEDDEDVVDEHVEHIKMERMPLSRPQRALPSKTQPPRSRSVSQTRKSP	2100
353	SDEDNEQEEDDEDDEDVVDEHVEHIKMERMPLSRPQRALPSKTQPPRSRSVSQTRKSP	2100

DAH	TNPNSGIGATGAGNRSNLLKENWLRPASTPRPPPPNGFIGREA	2143
29B	TNPNSGIGATGAGNRSNLLKENWLRPASTPRPPPPNGFIGREA	2143
e19	TNPNSGIGATGAGNRSNLLKENWLRPASTPRPPPPNGFIGREA	2143
74e19	TNPNSGIGATGAGNRSNLLKENWLRPASTPRPPPPNGFIGREA	2143
74	TNPNSGIGATGAGNRSNLLKENWLRPASTPRPPPPNGFIGREA	2143
211	TNPNSGIGATGAGNRSNLLKENWLRPASTPRPPPPNGFIGREA	2143
246	TNPNSGIGATGAGNRSNLLKENWLRPASTPRPPPPNGFIGREA	2143
353	TNPNSGIGATGAGNRSNLLKENWLRPASTPRPPPPNGFIGREA	2143

Table S1. Survival statistics of alleles in Quantitative Complementation Test. Male and female cohorts. Wildtype test alleles are *InR*⁺ on extracted third chromosomes (NC series) sampled from the Raleigh Farmer's Market by T.F. Mackay. Mutant *InR* test alleles are archival EMS-treated third chromosomes with lesions mapped over a deficiency, reported in Fernandez (59) provided by M. Frasch. N(0): initial cohort size, *InR*^{allele} / *InR*^{E19} (*InR*^{allele} / E19) and *InR*^{allele} / TM3, *InR*⁺ *Sb* (*InR*^{allele} / TM3) combined. * : mutant *InR* alleles were tested in two independent blocks; data shows combined sample size and average (after block adjustment) survival parameter estimates.

Survival parameters are adjusted across blocks. In each of five periods used to construct life tables (blocks), the relative hazard coefficient β was estimated for a universal wildtype chromosome (NC442) as NC442/E19 relative to NC442/TM3 (Table S2). To adjust for period effects among blocks, the β_{NC442} expressed in the block was subtracted from each β estimated for wildtype and *InR* mutants tested within the corresponding block. Proportional Hazard is the ratio of estimated mortality among compared cohorts after adjusting for their common time dependent underlying hazard $h_0(t)$, estimated as $\exp(|\beta|)$; unity indicates no overall mortality difference. **Bold** values indicate test allele decreases mortality (*InR*^{allele} / TM3: *InR*^{allele} / E19). Normal values indicate test allele elevates mortality (*InR*^{allele} / E19: *InR*^{allele} / TM3). Estimates are used to visualize the distribution of allelic effects (fold-change) among naturally segregating polymorphisms of *InR* (Figure 2, main text). Life Expectancy is observed difference between the Kaplan-Meier estimated median life span of the *InR*^{allele} / E19 cohort and the *InR*^{allele} / TM3 cohort, adjusted by subtracting the difference in life expectancy between NC442/E19 relative to NC442/TM3 of the block.

Block adjusted β for each *InR* EMS allele is evaluated by z-test relative to the distribution of β among NC alleles where mean value is -0.15 (both sexes) and SD is 0.44 and 0.48 for males and females respectively. Adjusting for multiple comparisons with Bonferroni correction ($p < 0.0055$), the alleles *InR*⁷⁴ and *InR*²¹¹ in both sexes decrease mortality more than expected; *InR*³⁵³ reduces mortality about 2.5-fold in females, but not significantly by our criteria; *InR*²⁶² modestly decreases mortality in both sexes, but no more than many wildtype *InR* alleles; *InR*³²⁷ is strongly adult deleterious in both sexes.

Table S2. Block survival statistics of the universal wildtype test chromosome (*InR⁺*) NC442 in Quantitative Complementation Test as NC442/E19 relative to NC442/TM3.

	Block	N(0)	β	s.e.	Life expectancy (difference, days) NC442/E19 – NC442/TM3
Males					
NC442	1	500	-0.258	0.091	4.9
NC442	2	395	-1.18	0.114	14
NC442	3	433	-0.504	0.099	5
NC442	4	665	-0.999	0.084	17.9
NC442	5	452	-0.825	0.101	8.1
Females					
NC442	1	463	-0.095	0.094	1.92
NC442	2	457	-1.279	0.102	16.5
NC442	3	571	-0.33	0.085	4.4
NC442	4	424	-0.973	0.106	14.49
NC442	5	445	-0.596	0.097	8.2

Table S3. Summary life table statistics for *InR* hemizygotes and *InR* wildtype homozygotes. Data of Figure 5A (Trial 1) and 5B (Trial 2) in main text, females. Wildtype *InR* included three independent accessions: *InR*^{+(HR)29B} (29+), *InR*^{+(HR)13A} (13+) and wDah. Null *InR* alleles were derived from the accessions used to produce wildtype *InR*^{+(HR)29B} (29+) and *InR*^{+(HR)13A} (13+). Trials were conducted in independent periods, with contemporary cohorts of indicated genotypes.

Genotype class	Genotype	N(0)	Median survival (d)	Median 95% c.i.	Mean survival (d)	Mean s.e.
Trial 1						
Wildtype homozygote						
	29+/29+	351	44	42-44	41.7	0.52
	wDah/29+	352	40	38-42	39.6	0.55
	29+/wDah	361	44	44-46	42.2	0.57
	wDah/wDah	348	48	46-48	45.9	0.62
Hemizygote						
	29+/29null	366	48	48-50	46.9	0.52
	wDah/29null	348	44	42-46	44.1	0.59
Trial 2						
Wildtype homozygote						
	13+/13+	443	34	36-36	33.7	0.51
	13+/29+	451	36	36-38	35.1	0.44
	29+/13+	342	38	38-40	37.99	0.42
	29+/29+	475	46	44-46	43.4	0.45
	wDah/13+	492	44	44-44	43.6	0.32
	wDah/29+	476	40	40-40	39.7	0.38
	wDah/wDah	461	42	42-42	41.2	0.37
Hemizygote						
	13+/13null	446	42	40-42	39.8	0.54
	29+/29null	484	50	48-50	48.8	0.41
	wDah/13null	465	44	44-46	44	0.38
	wDah/29null	467	42	40-44	42.3	0.47

Relative mRNA abundance of *InR* from whole body lysates of 20 females per biological replicate. Three biological replicates each genotype, each with three technical replicates. *InR* mRNA CT normalized to Rp49 of each sample, 2(-ddCT) estimation method. Means with s.e. and difference confidence interval, $t= 3.12$, $p = 0.036$.

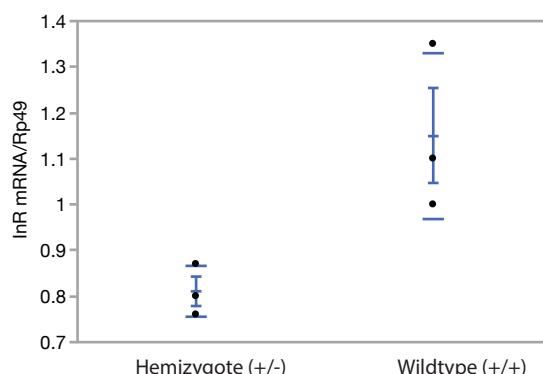


Table S4. Summary survival statistics of homologous recombination genotypes presented in main text figure 6. Subfigures with controls (+/+; +/−) in blue or green were contemporary cohorts. Proportional hazard analysis within each subfigure with relative mortality risk compared to wildtype or hemizygote, log-likelihood test. Accessions are independently generated stocks of indicated HR alleles.

FIG.	HR GENOTYPE & ACCESSION	N ₀	MEDIAN SURVIVAL (95% C.I.)	RISK RATIO TO WILDTYPE	P < χ ²	RISK RATIO TO HEMIZYGOTE	P < χ ²
6A	+/ +/ <i>InR</i> (E19)-2C +/ <i>InR</i> (E19)-14B +/ <i>InR</i> (E19)-22A	328 337 331 351	44 (42, 44) 40 (40, 42) 42 (40, 44) 48 (46, 52)	0.828 0.895 1.495	0.0154 0.155 <0.0001		
6B	+/ +/ <i>InR</i> (74) +/ <i>InR</i> (211) +/ <i>InR</i> (74, E19)	486 473 487 468	40 (38, 40) 44 (44, 44) 38 (38, 38) 40 (40, 40)	0.523 1.06 0.810	<0.0001 0.34 0.0012		
6C	+/ +/- (hemizygote) +/ <i>InR</i> (74) +/ <i>InR</i> (211) +/ <i>InR</i> (246) +/ <i>InR</i> (353) +/ <i>InR</i> (74, E19)	334 346 291 347 332 347 343	48 (46, 48) 50 (50, 52) 50 (48, 50) 44 (42, 44) 50 (48, 50) 60 (58, 62) 48 (48, 50)	0.648 0.660 1.60 0.743 0.215 0.881	<0.0001 <0.0001 1.02 <0.0001 <0.0001 0.331 0.0996	1.54 1.02 0.816 2.47 1.15 0.076 1.36	<0.0001 0.816 <0.0001 0.076 <0.0001 0.0001 <0.0001
6D	+/ +/ <i>InR</i> (353)-8.1 +/ <i>InR</i> (353)-15.4 +/ <i>InR</i> (353)-20.1	348 359 357 362	46 (44, 48) 62 (62, 64) 56 (54, 58) 58 (56, 58)	0.247 0.261 0.275	<0.0001 <0.0001 <0.0001		
6E	+/ <i>InR</i> (E19)/ <i>InR</i> (74) <i>InR</i> (E19)/ <i>InR</i> (211) <i>InR</i> (211)/ <i>InR</i> (74, E19) <i>InR</i> (74)/ <i>InR</i> (211)	486 446 471 456 350	40 (38, 40) 54 (52, 56) 52 (50, 52) 48 (46, 48) 46 (46, 48)	0.179 0.238 0.350 0.334	<0.0001 <0.0001 <0.0001 <0.0001		
6F	+/ +/- (hemizygote) <i>InR</i> (E19)/ <i>InR</i> (74) <i>InR</i> (E19)/ <i>InR</i> (211) <i>InR</i> (74)/ <i>InR</i> (211)	334 346 324 351 339	48 (46, 48) 50 (50, 52) 62 (62, 64) 58 (56, 58) 58 (58, 60)	0.138 0.259 0.234	<0.0001 <0.0001 0.215 <0.0001 0.405 <0.0001 0.366 <0.0001	0.215 0.405 0.366	<0.0001 <0.0001 <0.0001
6G	+/ +/- (hemizygote) <i>InR</i> (E19)/ <i>InR</i> (353) <i>InR</i> (E19)/ <i>InR</i> (246) <i>InR</i> (74)/ <i>InR</i> (353)	334 346 330 320 348	48 (46, 48) 50 (50, 52) 62 (58, 66) 46 (44, 48) 70 (68, 70)	0.154 0.75 0.142	<0.0001 0.0004 0.216 <0.0001 1.05 0.199	0.216 1.05 0.529 0.199	<0.0001 <0.0001 <0.0001
6H	+/ <i>InR</i> (E19)/ <i>InR</i> (246) <i>InR</i> (E19)/ <i>InR</i> (353) <i>InR</i> (74)/ <i>InR</i> (353)	359 347 360 354	46 (44, 48) 48 (46, 50) 61 (60, 64) 67 (66, 68)	0.679 0.213 0.189	<0.0001 <0.0001 <0.0001		