
Compilation of tRNA sequences and sequences of tRNA genes

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INTRODUCTION

This compilation contains 455 tRNA sequences and 981 sequences of tRNA genes published thus far covering the literature up to the end of 1988. A survey of published sequences is given in Table 1. The numbering system derived from yeast tRNA-Phe as adopted in the Cold Spring Harbour Symposium on tRNA in 1979 is used (Fig. 1, Ref. 1). Secondary structure of tRNAs are indicated by region-specific underlining of sequences. In the case of most mitochondrial tRNAs the alignment according to the accepted numbering system (Fig. 1) is not possible, therefore for D- and T-regions an arbitrary alignment was adopted. For the nomenclature of modified nucleosides see Table 2 and Fig. 2.

Each sequence is identified by one letter code for the particular amino acid depicting the specificity of tRNA, a three digit number, the anticodon sequence in unmodified form and the name of the organism from which the tRNA originates. The numbers for tRNAs are used in such way that each biological species has the same number throughout the compilation. In the case of tRNA genes the nucleotide preceding nucleotide residue 1 and the nucleotide following residue 76, as well as the intervening sequences, have been excluded from the compilation. Some tRNA-His and tRNA-His genes possess a nucleoside in position -1. The CCA sequence is included in the gene sequence only if it is coded for in the gene. The occurrence of intervening sequences after a particular nucleotide is indicated by an asterik and defined in the footnote.

Sequences of tRNA-precursors and tRNA-mutants are included in the compilation only when there are no mature and wild type tRNA-sequences, respectively, available. These cases are indicated in the footnote. The sequence originating from transformed or differentiated cells are considered as a separate entry only in the case of nucleoside replacement (i.e. different genes).

Footnotes are numbered according to the coordinates of the corresponding nucleoside and are indicated in the sequence by asteriks. The references and footnotes are listed at the end of the particular compilation with the corresponding sequence number. The references are restricted to one citation unless additional specific sequence information was obtained in a later work.

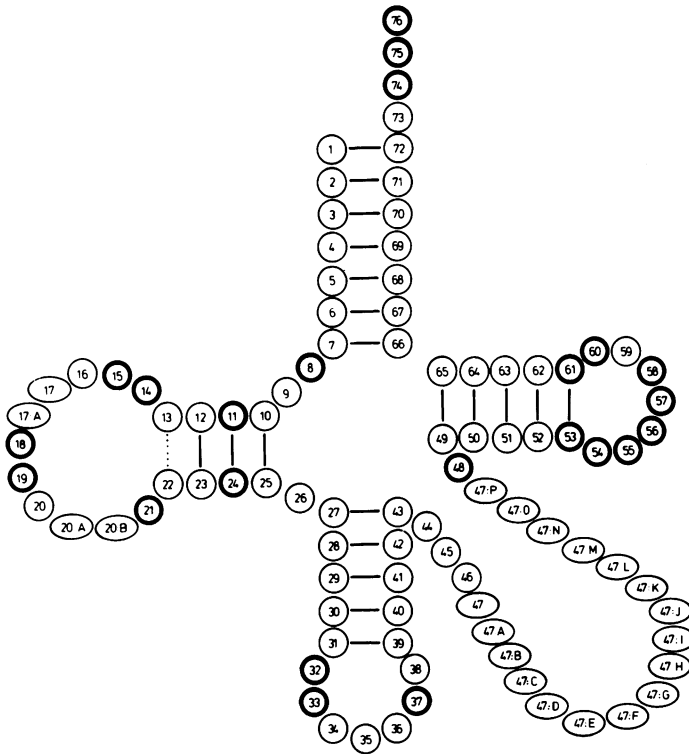


Fig. 1: Numbering of nucleotides in tRNAs. Circles represent nucleotides which are always present; among these, the thick-edged circles denote invariant or semiinvariant nucleotides. Ovals represent nucleotides which are not present in each sequence: these are the nucleotides before the two constant GMP residues (18,19) in the D-loop, the nucleotides after these GMP residues and the nucleotides in the variable loop which may be up to 17 nucleotides.

A nucleotide to be added at a given site is indicated by the number of the preceding nucleotide followed by a colon and a capital letter in alphabetical order. Thus, e.g. 20:A and 20:B mean the first and the second nucleotide after position 20. The absence of a nucleotide is indicated by the absence of a number, e.g. if no residue is found in position 17, the sequence then reads C16-C18. The numbering for the D-loop when one, two or three nucleotides are present, each between 15 and 18 or between 19 and 21, is then 16 and 16, 17 and 16, 17 and 20 and 20, 20:A and 20, 20:A, 20:B respectively. When the variable loop is five-membered, the numbering is 44, 45, 46, 47, 48. 47 is eliminated when the variable loop is four-membered. For large variable loops, capital letters are added onto 47, e.g. for thirteen nucleotides 44, 45, 46, 47, 47:A, 47:B, 47:C, 47:D, 47:E, 47:F, 47:G, 47:H, 47:I.

Table 1. Sequences included in the compilation

name (1)	numberrange	tRNA (2)	tRNA genes (2)
virus			
Avian Onco-Virus	000-099		
Chicken ASV/AMV/RSV	010	M	
House M-Mulv.	012	P, P	
Phage T4	013..014	R, G, G, I, L, P, S, T	R, Q, G, I, L, P, S, T
Phage T5	020	N, D, Q, H, L, P	A, D, Q, G, H, I, L, K, M, X, P, S, S, T, V
	040..041		
archaeobacteria			
Archaeoglobus fulg.	100-199		
Halobacterium cut.	108		A, C
Halobacterium hal.	110..112	A, R, N, Q, G, H, X, S, T, V, V, V	A, C
Halobacterium med.	113		A
Halobacterium vol.	115		M
	120..124		C, M
		A, A, A, R, R, R, N, D, C, Q, E, E, G, G, G, H, I, L, L, L, L, L, L, K, K, R, X, F, P, P, P, S, S, S, T, T, M, Y, V, V, X	A
Halococcus morrhuae	130		A
Methanobac. formi.	135		A, R, N, D, Q, E, H, I, L, K, F, P, T, T, Y, V
Methanobac. therm.	140	N, G	E
Methanococ. vani.	145..146		D, K, P, T, Y
Ruminobacter amylo.	147		A
Methanococ. voltae	148		G, L, X, F, S, V
Methanospir. hung.	149		A
Sulfolobus acido.	150	X	
Sulfolobus solfa.	160	M, X	
Thermoplasma acido.	180		
Thermococcus celer.	185		
Thermoprot. tenax	190..191		A, A, L, L, X
eubacteria			
Mycoplasma capric.	200-299		
Mycoplasma mycoid.	200..201	F, T, I	Q, L, L, K, T, M, W, Y
Mycoplasma P650	203..204	A, G, I, X, P, S, T, V	A, R, R, N, D, E, G, I, M, X, F, P, S, T, V
Spiroplasma melif.	205		L, X
Streptomyces gris.	206		A, K, D, C, I, M, X, F, P, S
Streptomyces rim.	215	X	X, X
Staphylococ. epid.	218..219		
Mycobac. smeg.	220..221		
Bacillus stearo.	227	G, G	
Lactobac. bulg.	230	X	
Bacillus subtilis	232	L, F, Y, V	
	235..238	A, R, G, L, K, K, M, X, F, P, T, Y, V, V, I, X, X	S
Thermus thermophi.	240..241		A, A, A, R, R, N, M, D, C, Q, E, G, G, H, H, I, I, L, L, L, L, K, M, X, F, F, P, S, S, T, T, T, M, Y, V, G

Thiobacillus ferro.	243			
E. coli	250..254	A, A, A, R, R, R, R, R, N, D, C, Q, Q, E, E, G, G, G, H, I, I, I, L, L, L, K, M, X, X, F, P, P, S, S, S, S, S, T, W, Y, Y, V, V, V, G, H, L, P, P, P	A, I A, R, R, R, D, Q, Q, E, G, G, H, I, I, L, L, L, K, M, X, X, F, P, P, S, S, S, T, T, T, W, Y, Y, V, V, V, R, H, L, P H, P R, H, L, P R, G, I, T, T, Y A, I A, I L L	
Salmonella typhi.	255..257			
Photobact. phosph.	258			
Aeromonas hydroph.	259			
Pseudomonas aer.	260..261			
Campylobac. jejuni	270			
Caulobacter cres.	272			
Rhizobium meliloti	275			
Bordetella pertus.	277			
Rhodospiril. rub.	280			
Agmenellium quad.	285			
Anacystis nidulans	290..291	L, F F, L, X E	A, I A, E, G, I, L, S A, I	
Synechocystis sp.	293			
Cyanophora parad.	295			
Pyraliella littora.	297			
Chloroplasts	300-399			
Chlamydomo. reinh.	300			
Chlorella elipso.	302			
Euglena gracilis	304..306	F	A, I A, I A, R, N, D, C, Q, E, G, G, H, I, L, L, L, K, M, X, F, P, S, S, T, W, Y, V	
Scenedesmus obliq.	308	M, X, Y	A, R, R, R, N, D, C, Q, E, G, G, H, I, I, L, L, L, K, M, X, F, P, P, S, S, T, W, Y, V, V, C, G, M, X, S, T, V, P, S, S, T, W, Y, V, V, R, D, C, E, G, G, M, X, P, S, T, W, Y A, R, N, C, H, I, L, L, M, F, P, S, S, T, W, V, V, M L, I, M, V H A, I, M, V	
Marchantia polym.	310..312			
Hordeum vulgare	313..314			
Triticum aestivum	315..316			
Zea mays	320..322	Q E I	A, R, R, R, N, D, C, Q, E, G, G, H, I, I, L, L, L, K, M, X, F, P, P, S, S, T, W, Y, V, V, C, G, M, X, S, T, V, P, S, S, T, W, Y, V, V, R, D, C, E, G, G, M, X, P, S, T, W, Y A, R, N, C, H, I, L, L, M, F, P, S, S, T, W, V, V, M L, I, M, V H A, I, M, V	
Arabidopsis thal.	324			
Brassica oleracea	325			
Glycine max	327..329	L, L, L	A, R, R, N, D, C, Q, E, G, G, H, I, I, I, L, L, L, K, M, X, F, P, P, S, S, S, T, T, W, Y, V, V, H P, H R I H	
Medicago sativa	333			
Nicotiana tabacum	335..337			
Nicotiana debneyi	340			
Oenothera sp.	342			
Pelargonium zonale	345			
Pennisetum america.	347			
Petunia hybrida	348			
Phaseolus vulgaris	350..352			
Pisum sativum	355..356	L, L, L, X, F, W	R, R, N, D, E, G, H, L, K, X, P, S, T, W, Y, V K, V A, R, D, C, E, H, I, I, L, M, S, S, T, Y, V, P, R, N E, L, L, F, T, Y L	
Sinapis alba	360			
Spinacia oleracea	365..366			
Spirodelia oligorn.	370..371	I, I, L, M, X, F, P, T, M, V		
Vicia faba	375..376			
Sorghum bicolor	378			

mitochondria	400-499		
single cell organisms and fungi 400-429			
Chlamydomo. reinh.	402		
Trypanosoma brucei	403..404		
Paramecium prim.	405		
Paramecium tetra.	406	F, Y	
Tetrahymena pyrif.	407		
Tetrahymena therm.	408		
Aspergillus nidul.	410..411		
Neurospora crassa	412..413	A, L, L, X, T, W, Y, V	
Podospora anserina	414	R, R, G, I, L, K, M, M, F, P,	
Saccharomyces cer.	417..419	S, S, T, W, Y	
Schizosaccha. pom.	421		
Torulopsis glab.	425..426		
Yeast	428	H, X	
plants 430-449			
Arabidopsis thal.	430		
Glycine max	432		
Lupinus luteus	434		
Onoothera sp.	436		
Phaseolus vulgaris	438..440		
Triticum aestivum	442..443	L, L, L, M, X, F, P, W, Y	
Zea mays	445		
animals 450-499			
Ascaris suum	450..451		
Caenorhabdi. eleg.	452		
Artemia sp.	453		
Locusta migratoria	454..455		
Aedes albopictus	456		
Drosophila melano.	457	R, D, Q, E, G, I, K, X, S, V	
Drosophila yakuba	459..460		
Drosophila virilis	461		
Asterina pectini.	462..463		
Paracentrotus liv.	464..465		
Strongylocen. purp.	466..467		
Xenopus laevis	468..469		

W, A, A
 X, Y
 W, Y
 H, L, X, F, W
 L, X, Y
 A, R, N, D, C, C, Q, E, G, G, H, I, L, L, K,
 M, W, Y, F, P, S, T, W, Y, V
 A, R, C, V
 D, S, W
 A, R, R, N, D, C, Q, E, G, H, I, K, M
 F, P, S, S, T, T, Y, Y, V, V
 G, H, L
 A, R, N, D, C, C, Q, E, G, H, I, L, K, M, X,
 F, P, S, S, T, T, W, Y, V
 L, X, W

 M, M, X
 E, M, X
 M, G, X
 X, S, W
 S
 D, Q, X, P, S, S, W, Y
 D, C, H, M, X, P, S, W

 A, R, N, D, C, C, Q, E, G, H, I, L, L, K, X,
 F, P, S, S, T, W, Y, V
 D, C, G, H, X, T
 E, F
 D, G, L, L, K, S
 A, R, N, E, G, L, F, S, V
 D, C, G, L, K, W, Y
 A, R, N, D, C, C, Q, E, G, H, I, L, L, K, X,
 F, P, S, S, T, W, Y, V
 Q, T, Y
 H, S, S
 A, R, N, D, C, C, Q, E, G, H, I, L, L, X,
 F, P, S, S, T, W, Y, V
 A, R, N, D, C, C, Q, E, G, H, I, L, L, K, X,
 F, P, S, S, T, W, Y, V
 A, R, N, D, C, C, Q, E, G, H, I, L, L, K, X,
 F, P, S, S, T, W, Y, V
 F, F, P, S, S, T, W, Y, V

Rana catesbeiana	470		A, N, C, Q, I, L, X, F, P, T, M, Y
Cephalorhyn.com	471		F, P, T
Hamster	473		
Rat	474..476	R, D, K, S R, D, L, L, L, K, F, M, V, V	A, R, M, N, M, D, D, C, C, Q, Q, E, G, H, I, L, L, K, K, Y, X, F, P, P, S, S, S, T, I, M, W, Y, V, A, R, N, D, C, Q, E, G, H, I, L, L, K, X, F, P, S, S, T, M, Y, V
Mouse	477..478		F, P, S, S, T, M, Y, V
Bovine	480..481	R, E, G, I, L, L, K, X, S, T, M, V,	F, P, S, S, T, M, Y, V
Green Monkey	482		H, L, S
Macaca fuscata	483		H, L, S
Macaca mulatta	484		H, L, S
Macaca fasciula.	485		H, L, S
Macaca sylvanus	486		H, L, S
Saimiri sciureus	487		H, L, S
Tarsius syrichta	488		H, L, S
Lemur catta	489		H, L, S
Chimpanzee	490		H, L, S
Gibbon	491		H, L, S
Gorilla	492		H, L, S
Orang Utan	493		H, L, S
Human	494..495	S	A, R, N, D, C, Q, E, G, H, I, L, L, K, X, F, P, S, S, T, M, Y, V
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eukaryotes	500-999		

single cell organisms and fungi	500-599		
Euglena gracilis	510	D, F	Q
Tetrahymena therm.	530..532	Q, Q, Q, X X, F, Y	E, M, V, V
Scenedesmus oblig.	540		L, F
Dictyostelium oblig.	555..556	X, F	S, S
Neurospora crassa	560		K, K, M, X, P, S, S, M, Y, V, R, R, D, E, H, K, X, X, F, S, S
Podospora anserina	565..566		I, F, F, S, S, S
Saccharomyces cer.	570..572	G, G, H, H, I, L, L, K, M, X, P, S, M E, F, Y A, I, L, X, P, Y, V A, R, R, R, N, D, C, E, K, F, F, S, S, T, Y, V, V, V	
Schizosaccha.pom.	575..577		
Torulopsis utilis	580		
Yeast	590..592		

plants	600-699		
Arabidopsis thal.	615		Y
Hordeum vulgare	620..621	E, E, F	
Triticum aestivum	625..626	R, G, M, X, F, M, Y, Y	
Brassica napus	630		D, M, X
Glycine max	635		L, P, P
Lupinus luteus	640	M, E, G, H, I, M, X, F, P, S, Y, V L, L, L, X	
Phaseolus vulgare	645..647		

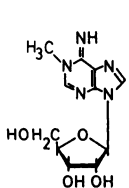
Pisum sativum	650								
Nicotiana rustica	655..656	F	Y, Y						Y
Petunia sp.	660								N
Sorghum bicolor	665								G
Oryza sativa	670								G

animals 700-999									
Caenorhabdi eleg.	700	L							D, L, K, P, W
Asterina amurensis	750	X							A, A, E, G, K
Bombax mori	770..771	A, A, G, G, F, F							A, R, R, N, D, E, E, E, G, G, H, I, L,
Drosophila melano.	780..782	E, H, K, K, X, F, S, S, Y, V, V, V							L, K, K, M, X, F, P, P, S, T, Y, V, V
									S
Drosophila simul.	785	X							A, N, L, K, X, X, F, Y, Y, V
Euphausia sperba	790	D, X, F							
Xenopus laevis	830..831	X							
Salmon	850								
Chicken	870	W							K, S, W
Mouse	950..952	R, R, Q, Q, E, I, K, K, M, X, F, F, S, V							A, D, E, G, H, I, L, K, K, X, P, P
Rat	955..957	N, N, D, D, Q, E, L, L, K, K, K, S,							D, D, E, E, E, G, G, L, L, K, F, P, P
		S, S, V, V							
Rabbit	965..967	D, X, K, K, M, X, F, V							
Bovine	970..972	R, K, R, N, D, Q, L, F, F, S, S, T, M, Y							
Calf	974	F							
Cow	975..976	L, L							
Sheep	980	H, X							
Human	995..997	N, E, G, G, H, L, M, X, F, S, S, Y, Y, V							N, N, Q, E, G, G, L, L, K, X, X, P, S, S, S,
									T, Y, Y, V, V
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(1) abbreviated names of organisms
 (2) specificity of the tRNA is defined by amino acid one-letter-code

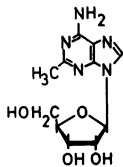
Table 2. ABBREVIATIONS FOR MODIFIED NUCLEOSIDES

A1 = M1A	= 1-METHYLADENOSINE
A2 = M2A	= 2-METHYLADENOSINE
A3 = M1AM	= 2'-O-METHYL-1-METHYLADENOSINE
A4 = I6A	= N6-ISOPENTENYLADENOSINE
A5 = MS2I6A	= 2-METHYLTHIO-N6-ISOPENTENYLADENOSINE
A6 = M6A	= N6-METHYLADENOSINE
A7 = T6A	= N-((9-BETA-D-RIBOFURANOSYLPURINE-6-YL)CARBAMOYL)-THREONINE
A8 = MT6A	= N-((9-BETA-D-RIBOFURANOSYLPURINE-6-YL)N-METHYL-CARBAMOYL)-THREONINE
A9 = MS2T6A	= N-((9-BETA-D-RIBOFURANOSYL-2-METHYLTHIOPURINE-2-YL)-CARBAMOYL)THREONINE
C2 = S2C	= 2-THIOCYTIDINE
C3 = CM	= 2'-O-METHYLCYTIDINE
C4 = AC4C	= N4-ACETYLCYTIDINE
C5 = M5C	= 5-METHYLCYTIDINE
C6 = M3C	= 3-METHYLCYTIDINE
C7 = ACP2C	= N2-(5-AMINO-5-CARBOXYPENTYL)CYTIDINE
D = D	= DIHYDROURIDINE
F = F	= PSEUDOURIDINE
F1 = M1F	= 1-METHYLPSEUDOURIDINE
F3 = FM	= 2'-O-METHYLPSEUDOURIDINE
G1 = M1G	= 1-METHYLGUANOSINE
G2 = M2G	= N2-METHYLGUANOSINE
G3 = GM	= 2'-O-METHYLGUANOSINE
G4 = M22G	= N2, N2-DIMETHYLGUANOSINE
G5 = M22GM	= 2'-O-METHYL-N2, N2-DIMETHYLGUANOSINE
G7 = M7G	= 7-METHYLGUANOSINE
I = I	= INOSINE
I1 = M1I	= 1-METHYLINOSINE
Q = Q	= QUEUOSINE
Q1 = MANQ	= BETA, D-MANNOSYLQUEUOSINE
Q2 = GALQ	= BETA, D-GALACTOSYLQUEUOSINE
T1 = T	= 5-METHYLURIDINE
T2 = S2T	= 5-METHYL-2-THIOURIDINE
T3 = TM	= 2'-O-METHYL-5-METHYLURIDINE
U1 = MAM5U	= 5-METHYLAMINOMETHYLURIDINE
U2 = S2U	= 2-THIOURIDINE
U3 = UM	= 2'-O-METHYLURIDINE
U4 = S4U	= 4-THIOURIDINE
U6 = CM5U	= 5-CARBAMOYLMETHYLURIDINE
U7 = MCM5U	= 5-METHOXYCARBONYLMETHYLURIDINE
U8 = MAM5S2U	= 5-METHYLAMINOMETHYL-2-THIOURIDINE
U9 = MCM5S2U	= 5-METHOXYCARBONYLMETHYL-2-THIOURIDINE
V1 = O5U	= URIDINE-5-OXYACETIC ACID
V2 = MO5U	= 5-METHOXYURIDINE
V3 = MV	= URIDINE-5-OXOACETIC ACID METHYL ESTER
V4 = CMNM5U	= 5-CARBOXYMETHYLAMINOMETHYLURIDINE
V5 = CMNM5S2U	= 5-CARBOXYMETHYLAMINOMETHYL-2-THIOURIDINE
V6 = CHM5U	= 5-(CARBOXYHYDROXYMETHYL)URIDINE
V7 = ACP3U	= 3-(3-AMINO-3-CARBOXYPROPYL)URIDINE
V8 = MCHM5U	= 5-(CARBOXYHYDROXYMETHYL)URIDINE METHYL ESTER
Y1 = YW	= WYBUTOSINE
Y2 = O2YW	= WYBUTOSINE
Z1 = CZ	= CIS-ZEATIN



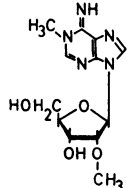
A1 = M1A

1-Methyladenosine



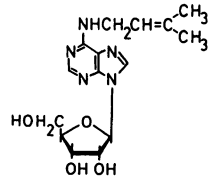
A2 = M2A

2-Methyladenosine



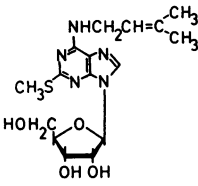
A3 = M1AM

2'-O-Methyl-1-methyladenosine



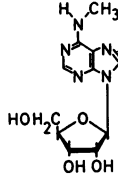
A4 = I6A

N6-Isopentenyladenosine



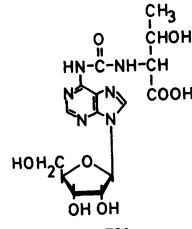
A5 = MS2I6A

2-Methylthio-N6-isopentenyl-
-adenosine



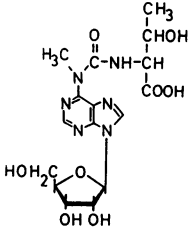
A6 = M6A

N6-Methyladenosine



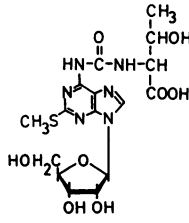
A7 = T6A

N-((9-B-D-Ribofuranosylpurine-6-yl)-
-carbamoyl)-threonine



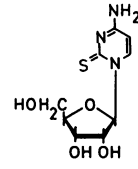
A8 = MT6A

N-((9-B-D-Ribofuranosylpurine-6-yl)-
-N-methyl-carbamoyl)threonine



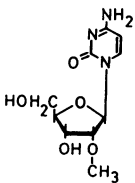
A9 = MS2T6A

N-((9-B-D-Ribofuranosyl-2-methylthio-
-purine-6-yl)carbamoyl)threonine



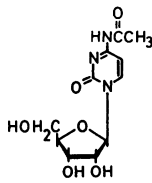
C2 = S2C

2-Thiocytidine



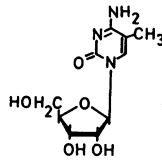
C3 = CM

2'-O-Methylcytidine



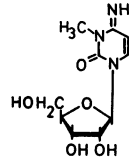
C4 = AC4C

N4-Acetylcytidine



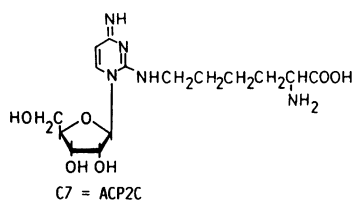
C5 = MSC

5-Methylcytidine

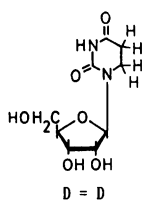


C6 = M3C

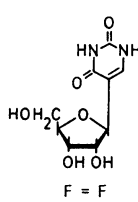
3-Methylcytidine



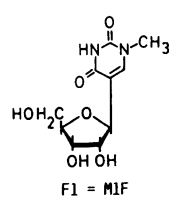
N2-(5-Amino-5-carboxypentyl)cytidine



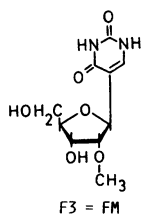
Dihydrouridine



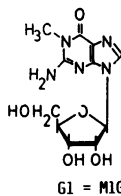
Pseudouridine



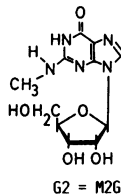
1-Methylpseudouridine



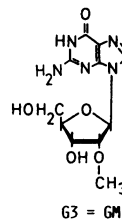
2'-O-Methylpseudouridine



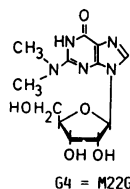
1-Methylguanosine



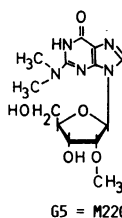
N2-Methylguanosine



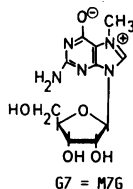
2'-O-Methylguanosine



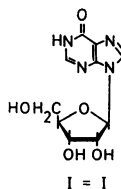
N2,N2-Dimethylguanosine



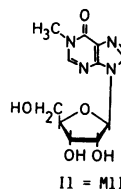
2'-O-Methyl-N2,N2-dimethylguanosine



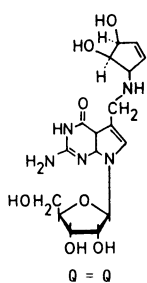
7-Methylguanosine



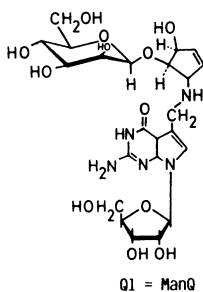
Inosine



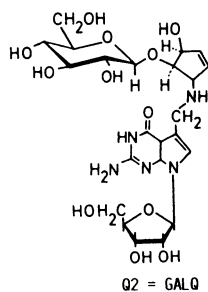
1-Methylinosine



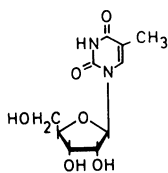
Queuosine



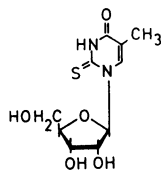
Beta, D-Mannosylqueuosine



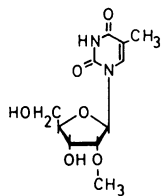
Beta, D-Galactosylqueuosine



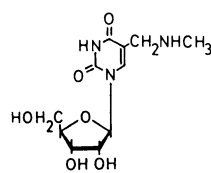
5-Methyluridine



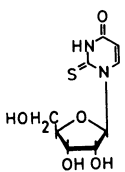
5-Methyl-2-thiouridine



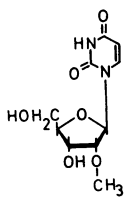
2'-O-Methyl-5-methyluridine



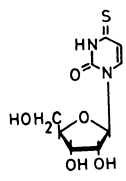
5-Methylaminomethyluridine



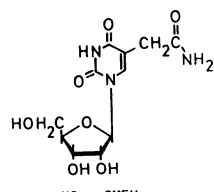
2-Thiouridine



2'-O-Methyluridine



4-Thiouridine



5-Carbamoylmethyluridine

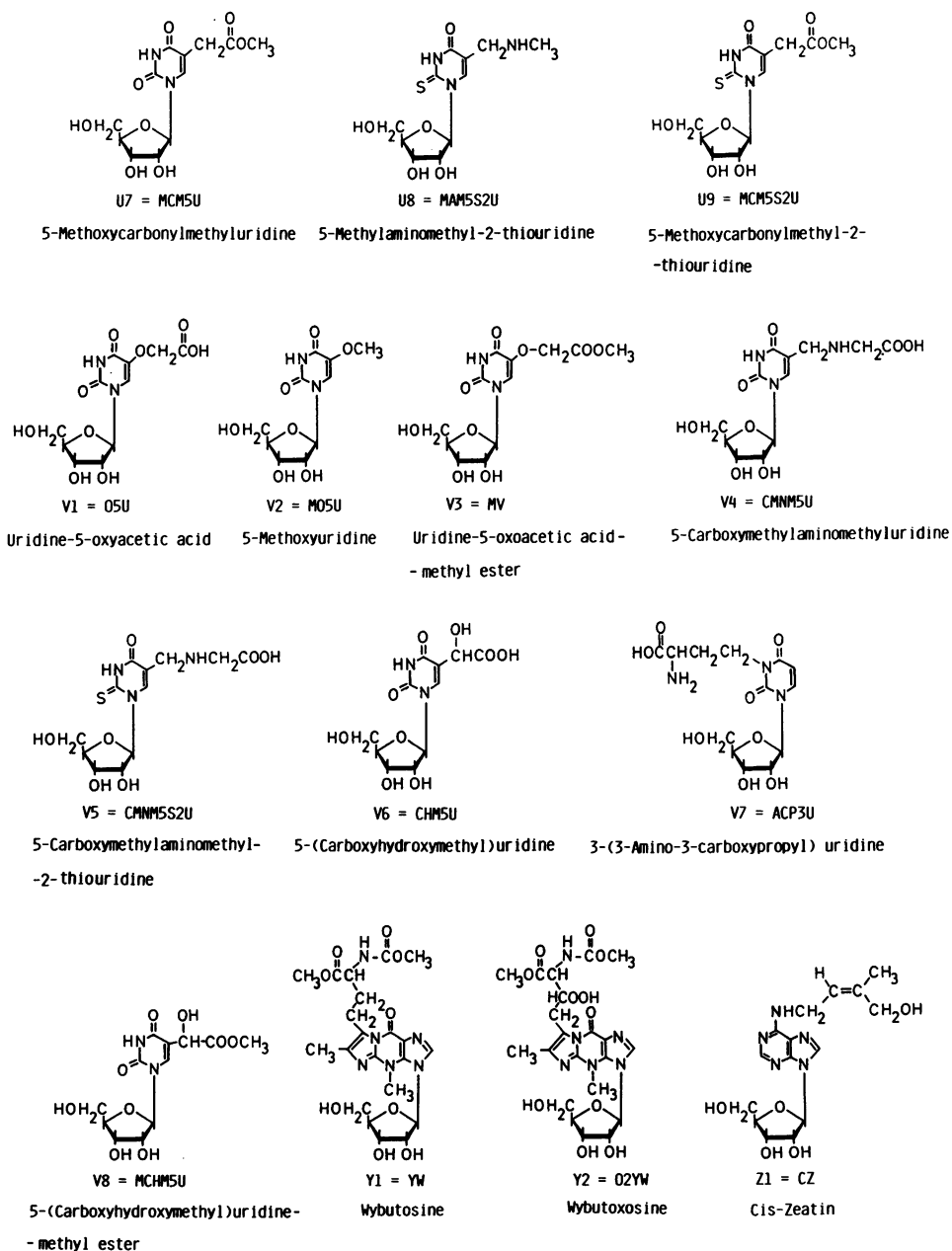


Fig. 2: Structure of modified nucleosides found in tRNAs

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Ref. 1: Transfer-RNA: Structure, Properties and Recognition. P.R.Schimmel, D.Soell, J.N.Abelson, Eds., 1979, Cold Spring Harbour Laboratory, N.Y., pp. 518-519

	AMINOACYL STEM	D STEM	D LOOP	D STEM	D STEM	ANTIC. STEM	ANTIC. LOOP	ANTIC. STEM
	1 2 3 4 5 6 7 8 9 10 12 14 16 A 19 A 21 23 25 27 29 31 33 35 37 39 41 43	11 13 15 17 18 20 B 22 24 26 28 30 32 34 36 38 40 42						
A L A N I N E								
A110	CGC HALOBACTERIUM CUT.	G G C U C G U A G A U C A G C	GGU	A G A U C G A C U U C C U U C G C A A G G A A G				
A120	CGC HALOBACTERIUM VOL.	G G C U C G U A G A U C A G U	GGC	A G A U C G A C U U C C U U C G C A A G G A A G				
A121	GGC HALOBACTERIUM VOL.	G G C U C G U A G A U C A G G	GGU	A G A U C A C U C C C U U G G C A U G G A G				
A122	UGC HALOBACTERIUM VOL.	G G C C C A U A G C U C A G U	GGU	A G A G U G C C U C C U U G C A A G G A G G				
A203	UGC MYCOPLASMA MYCOID.	G G C C C U U A G C U C A G C D	GGG	A G A G C A C C U G C C U U G C A G C G C A G G				
A235	UGC BACILLUS SUBTILIS	G G A S C C U U A G C U C A G C D	GGG	A G A G C C C U G C U U V 2 G C A G C G C A G G				
A250	GGC E. COLI	G G G C C U A U A G C U C A G C D	GGG	A G A G C C U U G C A U G G C A U G C A A G				
A251	UGC E. COLI	G G G C C A U A G C U C A G C D	GGG	A G A G C C C U G C U U V 1 G C A C G C A G G				
A252	UGC E. COLI	G G G C C U A U A G C U C A G C D	GGG	A G A G C C C U G C U U V 1 G C A C G C A G G				
A412	UGC NEUROSPORA CRASSA MITO	G G G G G U A U A G U A U A A D U	GGD	A G U A C A G C A A U C U U G C U C A U U G C				
A560	AGC TORULOPSIS UTILIS	G G G C G U G U G T G C G U A G D D	GGD	A G C G C G A F U C G C U I G C I F G C G A A				
A590	AGC YEAST	G G G C G U G U G T G C G U A G D C	GGD	A G C G C G A C U C C C U I G C I F G G G A G				
A770	AGC BOMBYX MORI	G G G C G U A G G C U C A G A D	GGU	A G A G C G A C U C G C U 3 U I G C I F G 3 F G A G				
A771	AGC BOMBYX MORI	G G G G C C G U A G G C U C A G A D	GGU	A G A G C G A C U C G C U 3 U I G C I F G 3 C G A G				
A R G I N I N E								
R020	UCU PHAGE T4	G U C C C G C U G G U G U A U	G3GAD	A G C A U A C G A U C C U C U A A G F U U G				
R110	CGC HALOBACTERIUM CUT.	G U C C G G A U A G A G F A G U	GGACUAU	A U C C U U G G C U U G G C U G G G I A G C C A G				
R120	CCG HALOBACTERIUM VOL.	G G C C C G U A G C U C A A U	GGAC	A G A G U G A C U U G G U U C C G G I A C C A A G				
R121	CGC HALOBACTERIUM VOL.	G U C C U G A U A G A G F A G U	GGACUAU	A U C C U C U G G C U U G C G G I A G C C A G				
R122	UCS HALOBACTERIUM VOL.	G G G C C C U A G C U C A U C U G G A C	AGAGU	A U C U G G C U C U G G C U C U G G I A C C A A G				
R235	AGC BACILLUS SUBTILIS	G C G C C C G U A G C U C A U	GGAD	A G A G C G U U U G A C U I C G G I A U C A A A				

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R252	CCU E. COLI	GUC	CU	U	A	G	U	A	A	U	G	G	A	G	A	A	A	C	G	A	G	C	C	C	C	C	C	U	C	C	U	A	A	G	G	G	C	U	G	G	C	U			
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R417	ACG SACCHAROMYCES CER. MITO	AUA	UCU	U	A	A	U	A	A	D	G	G	D	A	A	A	A	A	F	A	G	A	A	C	U	A	A	A	A	C	G	A	A	A	C	G	A	A	A	A	A	A	A		
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R456	UCG AEDES ALBOPICTUS MITO	AAA	UA	U	G	A	A	16	C	G	A	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	
R473	UCG HAMSTER MITO	UGG	U	G	A	U	A	16	U	U	A	C	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	
R474	UCG MT. MORRIS HEPATOMA MITO	UAG	U	U	U	A	A	A	A	U	A	A	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	
R480	UCG BOVINE LIVER MITO	UGU	A	C	U	U	A	A	A	A	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	
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R253	G	G	C																									
R254	A	G	G	U																								
R417	U	U	A																									
R418	A	U	A	U																								
R456	C	U																										
R473	A	G																										
R474	A	G																										
R480	A	G																										
R590	A	G	A	D																								
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R970	G	G	A	D																								
R971	G	G	A	D																								
R972	A	G	G	U																								

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M120	GUJ	HALOBACTERIUM	VOL.	G	C	G	C	C	G	U	A	G	C	U	C	A	G	U	G	U	A	G	A	G	A	C	G	C	G	G	U	U	A	T	C	C	A	C	G								
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M995	GUJ	HUMAN	LIVER/PLAC.	G	U	C	U	C	U	G	U	G	G	C	A	A	D	C	G	D	V	A	G	C	G	C	U	U	A	T	C	C	G	G													
A S P A R T I C A C I D																																															
D040	GUC	PHAGE	T5	G	C	G	A	C	C	G	G	G	C	U	G	G	C	U	U	A	G	G	U	A	C	U	C	C	C	C	U	G	C	A	C	G	G	G	A	G							
D120	GUC	HALOBACTERIUM	VOL.	G	C	C	C	G	G	G	U	G	A	U	G	F	A	G	U	G	G	C	C	A	U	C	A	C	G	A	C	C	C	U	G	C	A	C	G	G	U	C	G				
D250	GUC	E. COLI		G	G	A	G	C	G	G	A	G	U	C	A	G	D	C	G	D	A	G	A	A	C	C	U	C	G	C	C	U	U	C	A	Z	C	G	A	G	G						
D456	GUC	AEDIS	ALBOPICTUS	A	A	A	A	A	U	A	G	U	U	U	A	A	U	C	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
D473	GUC	HAMSTER	MITO	A	A	G	A	U	A	U	A	G	U	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
D474	GUC	RAT	LIVER	G	A	G	A	U	A	U	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
D475	GUC	RAT	MORRIS	G	A	G	A	U	A	U	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
D510	GUC	EUGLENA	GRACILIS	U	C	U	C	G	G	U	A	G	U	A	F	A	G	D	G	3	G	D	A	A	G	U	A	G	F	C	C	G	C	C	U	G	C	A	C	G	G	A					
D590	GUC	YEAST		U	C	C	U	G	A	A	A	G	U	F	A	A	D	G	D	C	A	G	A	A	G	G	G	C	C	F	U	G	C	G	G	C	C	F	U	G	C	G	C				
D850	GUC	XENOPUS	LAEVIS	U	C	C	U	C	G	U	A	F	A	G	U	A	F	A	G	U	G	G	D	G	A	G	U	A	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C

45 47 B D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
 44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76

ASPARAGINE

N040 G G G T U U G C U G G T F C G A A U C C A G C A G G A C C G C C A
 N110 U U G U C S C A G G F I F C S G G A C C C U G G U G G C G G C G A A C
 N120 U U G U C S C A G G F I F C S G A G U C C U G C C G U G G C G C C A
 N140 A G G C C C C A G G F C S 11 A A U C C G C C G G C G C A C C A
 N250 A U G 7 U C A C U G T F C G A G U C C A G U C A G A G G A G C C A
 N590 A G A D C S G U G A G T F C A A T C C C U C A C U G G G U C G C C A
 N640 A G G 7 D C S G U A G G T F C G A 1 G C C C U A C U G G G G C G C C A
 N955 A G G 7 D U G U G G N F C G A 1 G C C C A C C F A G G A C G C C A
 N956 A G G 7 D U G U G G N F C G A 1 G C C C A C C F A G G A C G C C A
 N970 A G G 7 D U G U G G N F C G A 1 G C C C A C C C A G G A C G C C A
 N995 A G G 7 D U G U G G N F C G A 1 G C C C A C C F A G G A C G C C A

ASPARTIC ACID

D040 A G A A U G U G G G T F C A A A U C C C A F C G G U C G G C G C C A
 D120 U G A C G G G F I F C S 11 A A U C C G C C U C G G G C G C C A
 D250 G G G 7 U C G G G T F C G A G U C C C G F C C G U C C G C C A
 D456 A A A A U U A G A U C A U C U A A U F F U U U A C C A
 D473 A A U U A U A G A C U A A U C U C A U A U A U C U A C C A
 D474 A G U U A U A G A C U U A A A U C U A U A U C U A C C A
 D475 A G U U A U A G A C U U A A A U C U A U A U C U A C C A
 D510 A G A C C G G G T F C A A U U C C C G C C G G A G A G C C A
 D590 A G A U C S G G G G T F C A A U U C C C G U C G G G A G C C A
 D830 A G A C S G G G G T F C A A U U C C C G A C G G G G A G C C A

45 47 B D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
 44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76

D955 A G A C5C5G G G G T F C G A U U C C C C G A C G G G G A G C C A
 D956 A G A C5C5G G G G T F C G A U U C C C C G A C G G G G A G C C A
 D965 A G G A C5C5G G G G T F C G A I A U C C C C G A C G G G A G C C A
 D970 G G G T A C5N U G A G T F C G A I A C U C A C G G C A C C G C C A

 C Y S T E I N E
 C120 A C C C C C5G C C G G F I F C3I1A A U C C G G C C U U G G C U C C A
 C250 C U A G U C C G G T F C G A C U C C G G A C G C C C U C C A
 C590 U G G 70 C5C U U A G T F C G A I U C C U G G U G C G A G C U C C A

 G L U T A M I N E
 Q020 G A U G C A A G G T F C G A G U C C U U U A U U C C C A G C C A
 Q040 G A U C A U G G T F C A A A U C C A U A U C C C U G C C A
 Q110 C G A C G A A G F C3G A U C U U C C C G G G A C U A C C A
 Q120 C G A C C A G G F C3G A A U C C U G G U G G A C U A C C A
 Q250 C A U U C C G A G G T F C G A A U C C U C G U A C C C A G C C A
 Q251 C A U U C C U G G T F C G A U C C A G G U A C C C A G C C A
 Q313 U A U D C G G G G T F C G A A U C C U U C C G U C C C A G C C A
 Q456 U A G A A U G U U A A U A U C U A U A U A U A A C C A
 Q530 U G A C5U G G G U F C G A I A U C C C A G U A G G A C C U C C A
 Q531 U G A C5U G G G U F C G A I A U C C C A G U G G G A C C U C C A
 Q532 U G A C5U G G G U F C G A I A U C C C A G U A C G A C C U C C A
 Q950 C G A U C5C5G A G F F C A A I A U C U C G G U G G A A C C U C C A
 Q951 C G A U C5C5G A G F F C A A I A U C U C G G U G G A A C C U C C A

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43										
0955	CUG	RAT	LIVER	G	G	U	U	C	C	A	U	G	T	G	U	G	F	A	A	D	G	S	G	D	D	A	G	C	A	C	U	C	U	G	G	A	C	S	U	C	U	G	A	A	F	C	C	A	G				
0970	CUG	BOVINE	LIVER	G	G	U	U	C	C	A	U	G	T	G	U	G	F	A	A	D	G	S	G	D	D	A	G	C	A	C	U	C	U	G	G	A	C	S	U	C	U	G	A	A	F	C	C	A	G				
G L U T A M I C A C I D																																																					
E120	CUC	HALOBLACTERIUM	VOL.	G	C	U	C	U	G	U	G	G	A	U	G	A	U	G	C	C	G	G	C	C	A	U	C	A	U	A	U	C	A	C	C	C	U	C	A	C	C	U	C	A	C	C	G	G	A				
E121	UUC	HALOBLACTERIUM	VOL.	G	C	U	C	G	G	U	G	G	A	U	G	A	U	G	C	C	G	G	C	C	A	U	C	A	U	C	U	C	A	C	C	C	U	C	A	C	C	U	C	G	A	C	C	C	G	A			
E250	UUC	E. COLI		G	U	C	C	C	U	C	G	U	C	F	A	G	A	G	G	C	C	C	A	G	A	C	A	C	C	G	C	C	C	U	U	8	C	U	U	8	C	A	C	2	C	8	C	G	G				
E251	UUC	E. COLI		G	U	C	C	C	U	C	G	U	C	F	A	G	A	G	G	C	C	C	A	G	A	C	A	C	C	G	C	C	U	U	8	C	U	U	8	C	A	C	2	C	8	C	G	G					
E252	UUC	E. COLI		G	U	C	C	C	U	C	G	U	C	F	A	G	A	G	G	C	C	C	A	G	A	C	A	C	C	G	C	C	U	U	8	C	U	U	8	C	A	C	2	C	8	C	G	G					
E293	UUC	SYNECHOCYSTIS	SP.*	G	C	C	C	C	A	U	C	G	U	C	U	A	G	A	G	G	C	C	D	A	G	A	C	A	C	C	U	C	C	C	C	U	C	U	C	C	C	U	U	8	C	A	C	2	C	8	C	G	G
E315	UUC	TRITICUM AESTIVUM	CHLORO	G	C	C	C	C	A	U	C	G	U	C	A	U	A	G	A	G	G	D	D	C	A	G	A	C	A	U	C	U	C	F	C	U	U	8	C	A	A	G	G	G	G	G	G	G	G	G			
E456	UUC	AEDIS ALBOPICTUS	MITO	A	U	U	A	U	A	U	A	T	G	U	U	A	A	A	A	A	A	A	C	A	A	C	A	F	F	A	C	A	U	U	U	U	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
E480	UUC	BOVINE LIVER		G	U	C	U	G	U	A	T	G	U	U	G	A	U	G	A	A	G	A	C	A	C	G	A	F	G	G	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U		
E575	UUC	SCHIZOSACCHARIA	POM.	U	C	C	G	U	U	G	U	G	U	C	C	A	C	A	C	G	G	D	A	G	G	A	U	C	G	U	C	G	C	U	U	9	U	9	U	9	U	9	U	9	U	9	U	9	U	9	U	9	U
E590	UUC	YEAST		U	C	C	G	A	U	A	G	U	G	F	A	C	A	C	G	G	D	A	U	C	A	F	A	C	G	C	G	C	U	U	9	U	9	U	9	U	9	U	9	U	9	U	9	U	9	U	9	U	
E620	UUC	HORDEUM VULGARE		U	C	G	U	C	G	U	A	G	U	C	F	A	G	G	D	A	G	G	D	F	A	G	G	A	U	C	U	C	G	G	C	U	U	F	C	A	C	C	C	G	G	G	G	G	G	G	G		
E621	CUC	HORDEUM VULGARE		U	C	G	U	C	G	U	A	G	U	C	F	A	G	G	D	A	G	G	D	F	A	G	G	A	U	C	U	C	G	G	C	U	U	F	C	A	C	C	C	G	G	G	G	G	G	G	G		
E640	CUC	LUPINUS LUTEUS		U	C	C	U	U	G	U	A	G	U	C	F	A	G	D	D	A	G	G	D	C	A	G	G	A	U	U	C	G	G	C	U	U	C	A	C	C	C	G	G	A	A	A	A	A	A	A			
E780	UUC	DROSOPHILA	MELANO.	U	C	C	A	U	A	G	U	C	F	A	G	D	A	G	G	D	A	G	G	D	C	A	G	G	A	U	U	C	U	G	G	C	U	U	9	U	9	U	9	U	9	U	9	U	9	U	9	U	
E950	CUC	MOUSE		U	C	C	U	G	U	G	U	C	F	A	G	U	A	G	C	A	C	C	C	A	G	G	A	U	U	C	C	G	G	C	U	C	C	A	C	C	C	C	C	C	C	C	C	C	C	C			
E955	UUC	RAT LIVER		U	C	C	A	U	G	U	C	F	A	G	C	A	A	G	C	G	D	A	G	G	A	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U			
E995	CUC	HUMAN		U	C	C	U	G	U	G	U	C	F	A	G	U	A	G	C	G	D	A	G	G	A	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U			

	45	47	B	D	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75																					
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76																				
Q955	C	G	A								U	C	S	G	A	G	F	F	C	A	A	U	C	U	C	G	G	U	G	G	A	A	C	C	U	C	C	A							
Q970	C	G	A								U	C	S	G	A	G	F	F	C	A	A	U	C	U	C	G	G	U	G	G	A	A	C	C	U	C	C	A							
GLUTAMIC ACID																																													
E120	U	G	A								C	S	A	G	G	G	F	F	C	S	G	A	A	U	C	C	U	G	A	C	G	G	A	C	C	A	C	C	A						
E121	G	G	A								C	S	A	G	G	G	F	F	C	S	11	A	U	C	C	U	G	A	C	G	G	A	C	C	A	C	C	A	C	C	A				
E250	U	A	A								C	A	G	G	G	T	F	C	G	A	A	U	C	C	C	U	G	G	G	A	C	G	C	C	A	C	C	A	C	C	A				
E251	U	A	A								C	A	G	G	G	T	F	C	G	A	A	U	C	C	C	U	A	G	G	G	A	C	G	C	C	A	C	C	A	C	C	A			
E252	U	A	A								C	A	G	G	G	T	F	C	G	A	A	U	C	C	C	U	A	G	G	G	A	C	G	C	C	A	C	C	A	C	C	A			
E293	C	G	A								A	G	G	G	A	T	F	C	G	A	A	U	C	C	C	U	G	G	G	G	U	A	C	C	A	C	C	A	C	C	A				
E315	C	A	G								C	S	G	G	G	A	T	F	C	G	A	C	U	U	C	C	C	F	G	G	G	U	A	C	C	A	C	C	A	C	C	A			
E456	A	A	A								U	A	A	A	U	U	A						U	U	U	U	F	A	U	A	A	U	A	C	C	A	C	C	A	C	C	A			
E480	U	A	G								U	S	A	U	G	G	J	F	A	G	A	U	U	C	C	A	U	G	A	G	A	U	A	C	C	A	C	C	A	C	C	A			
E575	G	G	A								G	S	G	G	G	T	F	C	G	A	C	U	U	C	C	C	G	A	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A		
E590	A	G	A								C	S	G	G	G	T	F	C	G	A	C	U	U	C	C	C	G	U	A	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A	
E620	A	G	A								C	S	C	G	G	T	F	C	A	A	G	U	U	C	C	C	G	C	G	A	C	G	G	A	C	C	A	C	C	A	C	C	A		
E621	A	G	A								C	S	C	G	G	T	F	C	G	A	G	U	U	C	C	C	G	C	G	A	C	G	G	A	C	C	A	C	C	A	C	C	A		
E640	A	G	A								C	S	C	G	G	T	F	C	A	A	G	U	U	C	C	C	G	C	G	A	C	G	G	A	C	C	A	C	C	A	C	C	A		
E780	A	G	G								C	G	G	G	T	F	C	G	A	U	U	U	C	C	C	G	U	A	G	G	G	A	C	C	A	C	C	A	C	C	A	C	C	A	
E950	C	G	G								C	S	C	S	G	G	T	F	C	G	A	U	U	C	C	C	G	U	A	G	G	G	A	C	C	A	C	C	A	C	C	A	C	C	A
E955	C	G	G								C	S	C	S	G	G	T	F	C	G	A	C	U	U	C	C	C	G	U	G	G	A	C	C	A	C	C	A	C	C	A	C	C	A	
E995	C	G	G								C	S	C	S	G	G	T	F	C	G	A	U	U	C	C	C	G	U	C	A	G	G	A	C	C	A	C	C	A	C	C	A	C	C	A

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44					
	G L Y C I N E																																																
G020	UCC PHAGE T4	GCGGAUAUCGUAUAU	G36D	AUUAAC	UACA	GA	CUUCCA	AFCUGA																																									
G110	GCC HALOBACTERIUM CUT.	GCGUGUA	62U	GFAGU	GGU	AUAC	GUAGCC	UUGCCA	UGUCA																																								
G120	CCC HALOBACTERIUM VOL.	GCGCGAUG	62U	CCAGU	GGU	AGGA	CA	CGAGC	UUGCCA	GCUCG																																							
G121	GCC HALOBACTERIUM VOL.	GCGUGGUA	62U	GFAGU	GGU	AUCA	CG	UGAGC	UUGCCA	GGUCA																																							
G122	GCC HALOBACTERIUM VOL.	GCACCGUG	62U	CUAAU	GGU	AACA	UUGC	CUNCCA	GCACA																																								
G140	GCC METHANOBAC. THERM.	GCGCGUA	U	6U	CCU	GU	UA	GA	CA	CUGC	CUGCC	CAC	GCCAG																																				
G203	UCC MYCOPLASMA MYCOID.	GCAGGUG	U4A	GUUAAU	GGC	AGAC	U	CAGC	CUUCC	A6A	GCUGA																																						
G220	UCC STAPHYLOCOCC. EPID.*	GCGGAG	U4A	UUUAAU	UUD	AGAA	UA	CGUU	CUUCC	CGAACG																																							
G221	UCC STAPHYLOCOCC. EPID.*	GCGGAG	U4A	GUUAAU	UUD	AGAA	UA	CGUU	CUUCC	CGAACG																																							
G235	UCC BACILLUS SUBTILIS	GCGGUGUA	GUU	AGU	GGD	AAC	C	UAGC	CUV4C	CAAGCUGA																																							
G250	CCC E-COLI	GCGGCGUA	GUU	CAU	GGD	AGAC	U	AGC	CUUCC	CAAGCUGA																																							
G251	GCC E-COLI	GCGGAAUA	GUU	CAU	GGD	AGAC	U	AGC	CUUCC	CAAGCUGA																																							
G252	UCC E-COLI	GCGGCAUC	GUU	AAU	GGCU	AUA	C	UAGC	CUUCC	CAAGCUGA																																							
G255	CCC SALMONELLA TYPHI.	GCGGCGUA	GUU	CAU	GGD	AGAC	U	AGC	CUUCC	CAAGCUGA																																							
G417	UCC SACCCHAROMYCES CER. MITO	AUAUAUA	U	16U	AAU	AUD	GGD	AAC	U	6AG	AUG	F	CUUCC	A4A	ACAAU																																		
G456	UCC AEDIS ALBOPICTUS MITO	AUUUAUA	U	16U	AAU	AUD	GGD	AAC	U	6AG	AUG	F	CUUCC	A4A	ACAAU																																		
G480	UCC BOVINE LIVER MITO	AUUUUUA	U	16U	AAU	AUD	GGD	AAC	U	6AG	AUG	F	CUUCC	A4A	ACAAU																																		
G570	GCC SACCCHAROMYCES CER. MITO	GCGCAAGU	U	16U	U	F	AGD	GGD	AAA	U	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A			
G571	UCC SACCCHAROMYCES CER. MITO	GCGGCGUA	GUU	CAU	GGD	AGAC	U	AGC	CUUCC	CAAGCUGA																																							
G625	GCC WHEAT GERM	GCA	C3C	A	U	6U	U	C	F	A	G	D	GGU	AGAA	UA	GA	CA	C	U	G	C	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A				
G640	GCC LUPINUS LUTEUS	GCA	C3C	A	U	6U	U	C	F	A	G	D	GGU	AGAA	UA	GA	CA	C	U	G	C	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A				

45 47 8 D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
 44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76

GLYCINE

6020	UGA	UGAGG	UFCGAUU	UCUAU	UUCGCGU	UCCA
6110	CAA	CSCSU	GGGF	C311A	AUCCCA	GCAAGCGCA
6120	GAG	CSC	GGGF	C311A	AUCCCGG	UCCGCGCA
6121	CAA	C5U	GGGF	C311A	AUCCAG	CAGCGCA
6122	CAA	C5U	GGGF	C311A	AUCCAG	CAGCGCA
6123	UUA	U5U	GGGF	C3G	AUCCAG	CAGCGCA
6140	CGU	C	CGGF	F	C311A	AUCCCGG
6203	UUG	UG	GGG	UFCGAUU	CCUU	CACCGU
6220	AGA	UA	AGG	UGCAA	AUCCUA	CUCCGU
6221	AGG	UA	AGG	UGCAA	AUCCUA	CUCCGU
6235	UGU	UG	AGG	UFCGAUU	CCUA	CACCGU
6250	AUA	C	AGG	UFCGAUU	CCUU	CACCGU
6251	G G 67U	C	CGG	UFCGAUU	CCGU	UCCCGU
6252	UGA	UG	GGG	UFCGAUU	CCCG	UCCCGU
6255	AUA	C	AGG	UFCGAUU	CCUU	CACCGU
6417	GAA	UG	CGG	UFCGAUU	CCCG	UCCCGU
6456	AGG	A	CUAA	JAA	U	UUAGU
6480	UAG	UU	CGG	UFCGAUU	CCGA	AAGAUA
6570	GG	C	CGG	UFCGAUU	CCGG	UCCCGU
6571	GGA	C	AGG	UFCGAUU	CCGU	UCCCGU
6625	AGA	C5C5G	GG	UFCGAUU	CCCG	UCCCGU
6640	AGA	C5C5G	GG	UFCGAUU	CCCG	UCCCGU

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44									
G770	GCC	BOMBYX	MORI																																																		
G771	UCC	BOMBYX	MORI																																																		
G995	CCC	HUMAN	PLACENTA																																																		
G996	GCA	HUMAN	PLACENTA																																																		
H I S T I D I N E																																																					
H040	GUG	PHAGE	T5																																																		
H110	GUG	HALOBACTERIUM	CUT. G																																																		
H120	GUG	HALOBACTERIUM	VOL. G																																																		
H250	GUG	E. COLI																																																			
H255	GUG	SALMONELLA	TYPHI.																																																		
H428	GUG	YEAST																																																			
H570	GUG	SACCHAROMYCES	CER. G																																																		
H571	GUG	SACCHAROMYCES	CER. G																																																		
H640	GUG	LUPTINUS	LUTEUS																																																		
H780	GUG	DROSOPHILA	MELANO. G																																																		
H980	GUG	SHEEP	LIVER																																																		
H995	GUG	HUMAN	HELA CELLS ⁸																																																		
I S O L E U C I N E																																																					
I020	MAU	PHAGE	T4																																																		
I120	GAU	HALOBACTERIUM	VOL.																																																		
I121	UAU	HALOBACTERIUM	VOL.																																																		
I203	GAU	MYCOPLASMA	MYCOID.																																																		

		45	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75						
		44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76					

6770	CGG	C5C5G	G	G	T	F	G	A	U	U	C	C	C	G	G	C	G	A	U	G	C	A	C	C	A	C	C	A			
6771	UGA	U	C5C5G	G	G	T	F	G	A	U	U	C	C	C	G	C	C	A	C	G	C	A	C	C	A	C	C	A			
6995	CGA	C	C5C5G	G	G	T	F	G	A	U	U	C	C	C	G	G	C	G	G	C	C	A	C	C	A	C	C	A			
6996	AGG	C5C5G	G	G	T	F	G	A	U	U	C	C	C	G	G	C	C	A	U	G	C	A	C	C	A	C	C	A			
		=====																													
		H I S T I D I N E																													
H040	CCUA	U	G	G	G	A	T	F	G	A	U	U	C	A	U	C	A	U	A	G	C	C	A	C	C	A	C	C	A		
H110	AGA	C	G	G	G	T	F	C	S	G	A	U	U	C	C	G	C	C	C	U	G	G	A	C	C	C	C	C	A		
H120	AGA	C	G	G	G	T	F	C	S	A	U	U	C	C	G	C	A	C	C	U	G	G	A	C	C	C	C	C	A		
H250	UUGU	C	G	U	G	G	T	F	G	A	U	U	C	C	C	A	U	A	G	C	C	A	C	C	C	C	C	C	A		
H255	UUGU	C	G	U	G	G	T	F	G	A	U	U	C	C	C	A	U	C	C	A	U	A	G	C	C	A	C	C	A		
H428	AAA	U	C	G	A	G	T	F	G	A	U	U	C	C	A	U	C	C	A	G	U	A	U	C	A	C	C	C	A		
H570	AAA	C	C	U	G	T	F	G	A	U	U	C	U	A	G	A	G	G	A	G	G	C	A	C	C	C	C	C	A		
H571	AAA	C	C	U	G	T	F	G	A	U	U	C	U	A	G	A	G	A	G	G	A	G	G	C	A	C	C	C	A		
H640	AAA	C	C	U	G	G	T	F	C	G	A	U	U	C	C	A	U	C	C	A	G	A	G	G	C	C	A	C	C	A	
H780	UAA	C	C	C	S	A	G	G	T	F	C	G	A	U	U	C	C	G	C	A	G	C	C	A	C	C	C	C	A		
H980	CAA	C	C	U	C	G	G	T	F	C	G	A	U	U	C	C	G	A	U	C	C	G	U	C	A	C	G	C	C	A	
H995	CAA	C	C	U	C	G	G	T	F	C	G	A	U	U	C	C	G	A	U	C	C	G	U	C	A	C	G	C	C	A	
		=====																													
		I S O L E U C I N E																													
I020	AGGU	U	A	C	C	A	G	T	F	C	A	A	U	U	C	U	G	G	U	C	U	G	G	U	C	A	C	C	A		
I120	AGGC	C	C	S	G	C	G	T	F	C	S	I	A	U	U	C	C	G	G	U	G	C	C	C	A	C	C	C	A		
I121	UGGU	C	S	A	U	G	G	T	F	C	S	G	A	C	C	C	A	U	G	G	C	C	C	C	C	C	C	C	A	C	A
I203	AGGU	C	G	U	G	G	T	F	C	A	A	G	U	U	C	C	A	A	U	A	U	C	C	C	A	C	C	C	A	C	A
		=====																													

45 47 B D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
 44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76

1240 A G G7U C G U G G T2F C A A1G U C C A C C A U C G C C C A C C A
 1250 A G G7V7 C G U G G T F C A A G U C C A C F C A G G C C U A C C A
 1251 A G G7V7 C G U G G T F C A A G U C C A C F C A G G C C U A C C A
 1252 U G G7V7 C G U G G T F C A A G U C C A G C A G G G C C A C C A
 1320 A G G7V7 C U U G G T F C A A G U C C A G G A U G G C C A C C A
 1365 A G G7V7 C U C U G G T F C A A G U C C A G G A U G G C C A C C A
 1366 A A U U C G U A G G T F C A A U U C C U A C U G G A U G C A C C A
 1417 A A A U A U A G G T F C A A U C C C U G U A G U U C A C C A
 1456 A U U A U A A G A U U A A U A C U U F A U C A U U A C C A
 1480 A U A A U A G A G G U C A A A C C C U C U A U U U C U A C C A
 1570 G G A D C S A G C G G T F C G A1U C C C G C U A G A G C C A C C A
 1580 A G A D C S A G C A G T F C G A1U C C U G C U A G G A C C A C C A
 1640 A A G D C S A G C A G T F C G A1U C C U G C U A U G G G C C A C C A
 1950 A G G7D C S G C G G G T F C G A1U C C C C G U A C G G G C C A C C A

LEUCINE

L020 C G G A U G A U U U C C U U G U G G G T F C G A G U C C C A C U U C U G C C A C C A
 L040 A G C U A A A U G C G U G G A G T F C G A G U C U C C U A G C C C A C C A
 L120 U C C G U A G G G G U C C S G U G G G F I F C31A U C C C U C C C U G C C A C C A
 L121 U C C A U A G G A G U C C S G C A G G F I F C31A A U C C U G U C C C U G C A C C A
 L122 U C C U G A G A G G U C C S G C C G G F I F C31A A U C C G G U C C C A C C A C C A
 L123 U C C G U A G G G G U C C G C G A G F I F C3G A A U C U C G U C C C C G C A C C A
 L124 U G G U G A G A C C U U C S G C A G G F I F C3G A A C C C U G U C C C G G C A C C A
 L230 U G G G C U U U G C C C G U G U G G G T F C G A C U C C C A C U G G C A C C A

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44				
L235	CAG	BACILLUS	SUBTILIS	G	C	C	G	G	U	G	U	G	C	G	G	A	A	U	D	G	G	D	A	G	A	C	C	G	C	U	A	G	A	U	C	A	G	G	A	F	C	U	A	G				
L250	AAA	E. COLI	G	C	C	G	G	A	U	G	U	G	G	A	A	D	C	G	3	6	D	A	G	A	C	C	A	G	G	G	A	F	U	N	A	A	5	A	F	C	C	U						
L251	CAG	E. COLI	G	C	G	A	G	G	U	G	C	G	G	A	A	D	D	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	G	F	G	F	U	A	G					
L252	GAG	E. COLI	G	C	C	G	A	G	U	G	U	G	C	G	A	A	D	D	3	6	D	A	G	A	C	C	G	C	U	A	C	C	U	U	G	A	G	N	F	G	U	A	G					
L255	CAG	SALMONELLA	TYPHI.	G	C	G	A	A	G	G	C	G	A	A	D	D	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	U	A	G	N	F	G	U	A	G					
L280	CAA	RHODOSPIRIL	RUB.	G	C	C	U	U	U	G	A	G	C	G	A	A	D	G	D	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	G	C		
L290	CAA	AMACYSTIS	NIDULANS	G	C	G	C	A	G	U	G	C	G	A	A	D	D	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	G	C				
L291	CAG	AMACYSTIS	NIDULANS	G	C	G	A	A	C	U	G	C	G	A	A	D	D	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G				
L327	CAA	GLYCINE	MAX	G	C	C	U	G	U	G	U	G	A	A	U	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G						
L328	UAA	GLYCINE	MAX	G	C	G	G	A	U	A	G	C	G	A	A	U	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G					
L329	UAG	GLYCINE	MAX	G	C	C	G	C	U	A	G	U	G	A	A	U	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G					
L350	CAA	PHASEOLUS	VULGARIS	G	C	U	U	C	A	U	G	U	G	A	A	U	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G					
L351	UAA	PHASEOLUS	VULGARIS	G	C	G	G	A	U	A	G	C	G	A	A	U	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G					
L352	UAG	PHASEOLUS	VULGARIS	G	C	C	G	C	U	A	U	G	U	G	A	A	U	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G				
L365	UAG	SPINACEA	OLERACEA	G	C	C	G	C	U	A	U	G	U	G	A	A	U	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G				
L412	UAA	NEUROSPORA	CRASSA	A	U	C	C	G	A	G	U	A	U	G	A	A	D	3	6	D	A	G	A	C	A	U	A	C	A	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G		
L413	UAG	NEUROSPORA	CRASSA	A	U	A	G	G	U	U	C	U	G	A	A	D	3	6	D	A	G	A	C	A	U	A	C	A	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G			
L417	UAG	SALICORNIA	CER.	G	C	U	A	U	U	U	G	U	G	A	A	D	3	6	D	A	G	A	C	C	G	C	U	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G					
L438	CAA	PHASEOLUS	VULGARIS*	G	U	C	A	G	A	U	G	C	C	5	6	A	G	D	3	6	D	A	G	A	C	C	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G						
L439	CAA	PHASEOLUS	VULGARIS*	G	U	C	A	G	A	U	G	C	C	5	6	A	G	D	3	6	D	A	G	A	C	C	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G						
L440	MAG	PHASEOLUS	VULGARIS*	G	A	U	A	G	U	U	G	C	C	4	6	A	G	D	3	6	D	A	G	A	C	C	A	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G						
L474	UAG	RAT	LIVER	A	C	U	U	U	A	U	A	G	2	6	A	U	A	G	A	3	6	D	A	A	U	C	C	A	F	U	G	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G		
L475	UAG	RAT	MORRIS	A	C	U	U	U	A	U	A	G	2	6	A	U	A	G	A	3	6	D	A	A	U	C	C	A	F	U	G	G	C	U	A	G	C	A	A	N	A	F	C	U	A	G		

45	47	B	D	F	H	J	L	M	P	K9	51	53	55	57	59	61	63	65	67	69	71	73	75	76
44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76
L235		GGUCUU	AUGGACC							UGAGGGT	FCAAGU	CCUU	CACCCGC	ACCA										
L250		CGCGGU	CGCGCUG							UCGGGT	FCAAGU	CCGC	UCCGGGU	ACCA										
L251		UGCCUA	CGGACG							UGGGGT	FCAAGU	CCCC	CCUCGC	ACCA										
L252		UGCCCA	AUGGGCU							UCGGGT	FCAAGU	CCGU	CCUCGGU	ACCA										
L255		UGUCUA	CGGACG							UGGGGT	FCAAGU	CCCC	CCUCGC	ACCA										
L280		UUUGUA	CGGACG							UGUAGT	FCGACU	UCCC	CAAAGGC	ACCA										
L290		CGUA	CGGUAUG							UGGGGT	FCGAGU	CCAC	CUGCC	ACCA										
L291		UGGU	UCAGACUG							UCGGGT	FCAAGU	CCGG	GUCCGC	ACCA										
L327		UGCUA	CAGAGCG							UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L328		CGACUA	AGAAUCA							UGAGGT	FCAAGU	CCUC	UCAAAGC	ACCA										
L329		UGCUA	GAGCA							UCGGT	FCGAGU	CCG	UA	CGGC	ACCA									
L350		UGCUA	AGAGCG							UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L351		CGACUA	AGAAUCA							UCGGT	FCGAGU	CCG	UA	CGGC	ACCA									
L352		UGCUA	GAGCA							UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L365		UGCGA	GAGCA							UCGGT	FCGAGU	CCG	UA	CGGC	ACCA									
L412		GGCU	UAGCUG							UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L413		UGUU	UAACUG							UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L417		UA	CUUACAGUA							UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L438		U3C	UCGAGAGGGG							UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L439		U3C	UCGAGAGGGG							UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L440		U3C	CGAANGG							UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L474		AAA								UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										
L475		AAA								UGAGGT	FCGAGU	CCUC	UCAAAGC	ACCA										

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42					
L476	UAG RAT MORRIS HEPATOMA MITO	A	C	U	U	G	U	A	U	A	G	26	A	U	G	A	G	A	G	D	A	U	C	C	A	F	U	G	G	U	C	U	A	G	G	U	A	G	G	A	C	C	A	A			
L480	UAA BOVINE LIVER	G	U	A	A	G	G	G	G	G	A	G	G	A	G	C	C	C	G	G	D	A	U	G	G	C	A	F	A	A	A	C	U	N	A	A	A	C	F	U	U	A					
L481	UAG BOVINE LIVER MITO	A	C	U	U	U	A	A	A	A	G	26	A	U	G	A	G	A	G	D	A	U	C	C	A	F	U	G	G	U	C	U	A	G	G	U	A	G	G	A	C	C	A	A			
L570	CAA SACCHAROMYCES CER.	G	G	U	G	U	U	U	G	G	2C	C	4G	A	G	C																															
L571	UAG SACCHAROMYCES CER.	G	G	A	G	U	U	U	G	G	2C	C	4G	A	G	D																															
L580	CAA TORULOPSIS UTILIS	G	G	A	U	U	U	U	G	G	2C	C	4G	A	G	C																															
L645	CAA PHASEOLUS VULGARIS	G	U	C	A	G	G	A	U	G	G	2C	C	5G	A	G	D																														
L646	CAA PHASEOLUS VULGARIS	G	U	C	A	G	G	A	U	G	G	2C	C	5G	A	G	D																														
L647	MAG PHASEOLUS VULGARIS	G	A	U	A	G	U	U	U	G	G	2C	C	4G	A	G	D																														
L700	AAG CAEORHABDI. ELEG.	G	G	A	G	A	G	A	U	G	G	C	C	4G	A	G	C																														
L955	CAA RAT MORRIS HEPATOMA	G	U	C	A	G	G	A	U	G	G	2C	C	4G	A	G	D																														
L956	CAA RAT MORRIS HEPATOMA	G	U	C	A	G	G	A	U	G	G	2C	C	4G	A	G	D																														
L970	AAG BOVINE LIVER	G	G	U	A	G	C	G	U	G	G	2C	C	4G	A	G	C																														
L975	AAG COM MAMMARY GLAND	G	G	U	A	G	C	G	U	G	G	2C	C	4G	A	G	C																														
L976	CAG COM MAMMARY GLAND	G	U	C	A	G	G	A	U	G	G	2C	C	4G	A	G	C																														
L995	MAG HUMAN HELA-CELLS	G	U	C	A	G	G	A	U	G	G	2C	C	4G	A	G	D																														
	LYSINE																																														
K120	UUU HALOBACTERIUM VOL.	G	G	G	C	U	G	G	U	A	G	C	U	C	A	N	U	A	G	G	C	A	G	G	C	A	G	G	C	U	G	G	C	U	A	G	G	C	U	A	G	G	C	C	A	G	
K121	CUU HALOBACTERIUM VOL.	G	G	G	C	C	G	G	U	A	G	C	U	C	A	N	U	A	G	G	C	A	G	G	C	A	G	G	C	U	G	G	C	U	A	G	G	C	U	A	G	G	C	C	A	G	
K235	UUU BACILLUS SUBTILIS	G	A	G	C	C	A	U	A	G	C	U	C	A	G	U	D	G	G	D																											
K236	UUU BACILLUS SUBTILIS	G	A	G	C	C	A	U	A	G	C	U	C	A	G	U	D	G	G	D																											
K250	UUU E.COLI	G	G	U	C	C	U	U	A	G	C	U	C	A	G	U	D	G	G	D																											
K417	UUU SACCHAROMYCES CER. MITO	G	A	G	A	A	U	A	U	G	U	U	A	A	D	G	G	D																													

	45	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75															
	44	46	A	C	E	G	I	K	L	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76														
L476	AAA	C5U	U	G	G	J	G	C	A	C	U	C	C	A	A	C	C	A	A	A	A	A	A	A	G	U	A	C	C	A									
L480	UAUC	CSA	G	A	G	A	U	U	C	A	A	U	C	C	U	C	C	U	C	C	U	A	A	C	C	A	C	C	A	C	C	A							
L481	AA	A	U	U	G	G	J	G	C	A	C	U	C	C	A	A	U	C	C	A	A	A	A	A	A	A	G	U	A	C	C	A							
L570	UAUCGUAAGAUG	CSA	A	G	A	G	T	F	C	G	A	U	C	C	U	A	G	C	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C	A					
L571	UAUCUCGGAUG	CSA	A	G	A	G	T	F	C	G	A	U	C	C	U	A	G	C	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C	A					
L580	UAUCGUAAGAUG	CSA	U	G	A	G	T	F	C	G	A	U	C	C	U	A	G	C	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C	A					
L645	U3CUUCGAGAGGG	CS6	U	G	G	T	F	C	A	A	U	C	C	A	C	U	C	U	C	U	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A				
L646	U3CUUCGAAAGAGGG	CS6	U	G	G	T	F	C	A	A	U	C	C	A	C	U	C	U	C	U	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A				
L647	U3CGAANGGG	CS6	U	G	G	T	F	C	A	A	U	C	C	A	C	U	C	U	C	U	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A				
L700	UCCUUCGGGG	C	G	U	G	G	T	U	C	G	A	U	C	C	A	C	U	C	U	C	U	C	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C		
L955	N ⁺ FCGGAUGGAG	CS6	U	G	G	T	F	C	G	A	U	C	C	A	C	U	C	U	C	U	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C		
L956	N ⁺ FCGUAUGGAG	CS6	U	G	G	T	F	C	G	A	U	C	C	A	C	U	C	U	C	U	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C		
L970	UCFCFCGCGGG	CS6	U	G	G	T	F	C	G	A	U	C	C	A	C	U	C	U	C	U	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C		
L975	UCFCFCGCGGG	CS6	U	G	G	T	F	C	G	A	U	C	C	A	C	U	C	U	C	U	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C		
L976	U3CFCFCUGGAGG	CS6	U	G	G	T	F	C	G	A	U	C	C	A	C	U	C	U	C	U	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C		
L995	U3CFCFCGGAUGGAG	CS6	U	G	G	T	F	C	G	A	U	C	C	A	C	U	C	U	C	U	C	G	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C		
L Y S I N E																																							
K120	CGGU	CS6	G	G	G	F	F	C	S	I	A	G	U	C	C	U	C	C	A	G	C	C	C	C	G	C	C	A	C	C	A	C	C	A	C	C			
K121	CGGU	CS6	C	G	F	G	F	F	C	S	I	A	U	C	G	C	G	U	C	C	G	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	
K235	GGG7U	C	A	A	G	T	F	C	G	A	U	C	C	U	C	U	C	U	C	U	C	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C
K236	GGG7U	C	A	A	G	T	F	C	G	A	U	C	C	U	C	U	C	U	C	U	C	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C	A	C	C
K250	UGG7U7	C	G	A	G	T	F	C	G	A	U	C	C	U	C	U	C	U	C	U	C	A	C	C	A	C	C	C	C	C	C	C	C	C	C	C	C	C	
K417	CCA	U	G	C	U	G	G	T	F	C	A	C	U	C	C	A	C	U	C	C	A	G	C	U	A	U	C	U	C	C	A	C	C	A	C	C	A	C	C

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44							
K456	CUU	AEDS	ALBOP	ICTUS	C	U	U	A	G	A	U	G	A	G	A	A	A	G	G	C	A	A	A	G	G	A	F	G	A	U	C	U	C	U	U	A	A	U	A	A	U	A	U	A	U						
K473	UUU	HAN	STER	MITO	C	A	C	A	U	G	A	G	U	C	U	A	G	G	C	A	A	G	G	C	A	A	G	G	C	A	C	U	U	A	A	C	U	U	U	A	A	A	A	A	A	A	A				
K474	UUU	RAT	LIVER	MITO	C	A	U	G	C	G	A	G	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U			
K480	UUU	BOVINE	LIVER	MITO	C	A	C	A	A	G	A	G	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U		
K570	UUU	SACC	HARON	CYCES	G	C	U	U	G	U	J	G	U	G	U	A	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G		
K590	UUU	YEAST			F	C	C	U	G	U	A	G	U	C	A	G	D	D	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G		
K780	CUU	DROSOPHILA	MELANO.		G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G		
K781	UUU	DROSOPHILA	MELANO.		G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	
K950	CUU	MOUSE	FIBROS.	SV 40	G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	
K951	CUU	MOUSE			G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	
K955	CUU	RAT	LIVER		G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	
K956	CUU	RAT	LIVER		G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	
K957	UUU	RAT	LIVER		G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G
K965	CUU	RABBIT	LIVER		G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G
K966	CUU	RABBIT	LIVER		G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G
K967	UUU	RABBIT	LIVER		G	C	C	G	G	C	A	G	U	C	A	G	D	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	
METHIONINE																																																			
M010	CAU	AVIAN	OMCO.-VIRUS		G	C	C	C	C	U	A	G	U	C	A	G	D	A	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G		
M120	CAU	HALO	BACTERIUM	VOL.	G	C	C	G	G	G	U	G	C	U	F	A	N	C	U	G	G	A	C	A	F	A	G	G	C	C	G	C	A	C	U	C	U	A	A	A	A	A	A	A	A	A	A	A	A	A	A
M180	CAU	THERMOP	PLASMA	ACIDO.	G	C	C	G	G	G	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U
M235	CAU	BACILLUS	SUBTILIS		G	C	C	G	G	U	A	G	U	C	A	G	C	A	G	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G
M250	CAU	E. COLI			G	C	U	A	C	G	U	A	G	U	C	A	G	D	A	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	
M308	CAU	SCENEDES	MUS	OB	G	C	C	U	G	C	U	A	G	U	C	A	G	D	A	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	
CHLORO																																																			

	45 47 B D F H J L N P	49 51 53	55 57 59	61 63 65	67 69 71	73 75
	44 46 A C E G I K M O	48 50 52	54 56 58 60	62 64 66 68 70 72 74 76		

K456	A A U A	U A G U A A A	U U A G C A Q U	U A C F F C U A A U G A C C A		
K473	A A U	U G A G A G A	C U U C U A	G U C U C A U G G A U G A C C A		
K474	A G U U	A G A G C A C A A	A U C U C A C A U G A C C A			
K480	A G A U	U G A G A G C A U A U A	C U C U C U U G G U G A C C A			
K570	A G G U	U A G G G G T F C G A G C	C C C C U A C A G G G C U C C A			
K590	A U G U	C S A G G G G T F C G A G C	C C C C U A F G A G G A G C C A			
K780	G G G U	C G U G G G N F C G A G C	C C C A C G U G G G C G C C A			
K781	G G G U	C S C A G G G T F C A A G U	C C C U G U U C G C G C G C C A			
K950	G G G U	C S G U G G G T F C G A G C	C C C A C G U G G G C G C C A			
K951	G G G U	C S G U G G G T F C G A G C	C C C A C G U U G G G C G C C A			
K955	G G G U	C S G U G G G T F C G A G C	C C C A C G U G G G C G C C A			
K956	G G G U	C S G U G G G T F C G A G C	C C C A C G U G G G C G C C A			
K957	G G G U	C S C S A G G G T F C A A G U	C C C U G U C G G G C G C C A			
K965	G G G U	C S G U G G G T F C G A G C	C C C A C G U G G G C G C C A			
K966	G G G U	C S G U G G G T F C G A G C	C C C A C G U G G G C G C C A			
K967	G G G U	C S C S A G G G T F C A A G U	C C C U G U C G G G C G C C A			

		M E T H I O N I N E				
M010	A G G U	C S C S U G A G T F C G A A C	C U C A G A G G G G C C A C C A			
M120	A G A U	C S G U G G G T F C S G A G	C C C A C C C G G C A C C A			
M180	A G G U	C U C G G G F C S G A U C	C C C G A U C C C G G C A C C A			
M235	A G G U	C G G G G G T F C G A U C	C C C U C C G C C G C U A C C A			
M250	G G G U	C A C A G G T F C G A A U	C C C U C G U A G C C A C C A			
M308	A A G U	C A C U A G T F C G A A U	C U A G U A G C A G G C A C C A			

	45	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76		
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76		
M365	G A G 7 7										C A U U G G T	F C A A A U	C C A A U A	G U A G G U A C C A													
M417	U A A U										G J A A G G T	F C A A U U	C C U U C U	A C A A G U A C C A													
M418	U A A										U C U A A G G T	F C A A U U	C C U U C U	A C A A G U A C C A													
M438	G A G 7 7										C A U U G G T	F C A A A U	C C A A U A	G U A G G U A C C A													
M570	A G 6 7 0										C S G A G A G T	F C G A 1 A C	C U C U C	C U G A G C A C C A													
M625	A G 6 7 0										C S G A G A G T	F C G A 1 G C	C U C U C	C A C C C C A C C A													
M640	A G 6 7 0										C S G A G A G T	F C G A 1 G C	C U C U C U	C A C C C C A C C A													
M950	A G 6 7 0										C S G U G A G T	F C G A 1 U C	C U C A C A	C G G G C A C C A													
M965	A G 6 7 0										C S G U G A G T	F C G A 1 U C	C U C A C A	C G G G C A C C A													
M995	A G 6 7 0										C S G U G A G T	F C G A 1 U C	C U C A C A	C G G G C A C C A													
M E T H I O N I N E - I N I T I A T O R																											
X110	A G A U										C G G U A G F I F	C 3 1 1 A U	C U A C C U	C C C G C U A C C A													
X120	A G A U										C G G U A G F I F	C 3 1 1 A U	C U A C C U	C C C G C U A C C A													
X130	A G A U										C A G U A G M	F 1 C 3 1 1 A U	C U A C U	U C C G C U A C C A													
X150	A G G U										C S C C U G G U 3 U	C 3 1 1 A 1 U	C C A G G C	C G C G C U A C C A													
X180	A G A U										C A U U G G F	C 3 N 1 A 1 U	C C A U C	C C C G C U A C C A													
X203	A G G C										C S C A G G U	F C G A G U	C C U G C	C C C G C A C C A													
X215	A G G U										C S C A G G U	F C A 1 A U	C C U G U	C C C G C U A C C A													
X227	A G 6 7 0										C S C A G G U	F C G A 1 U	C C U G U	C C C G C U A C C A													
X235	A G G U										C S C A G G T	F C A A U	C C U G C	C C C G C A C C A													
X240	A G 6 7 0										C G C C G G T 2 F	C A 1 A U	C C G G C	C C C G C A C C A													
X241	A G 6 7 0										C G C C G G T 2 F	C A 1 A U	C C C G C	C C C G C A C C A													
X250	A G A U										C S U C G G T	F C A A U	C C G G C	C C C G C A C C A													

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	27	29	31	33	35	37	39	41	43							
X251	CAU E. COLI	C	G	G	G	G	G	U	A	G	G	C	A	G	C	C	U	G	G	G	A	G	C	U	C	G	U	C	G	G	G	C	U	C	A	U	A	C	C	C	G
X290	CAU AMACYSTIS NIDULANS	C	G	C	G	G	G	G	U	A	G	A	G	C	A	G	C	C	U	G	G	A	G	C	U	C	G	U	C	G	G	C	U	C	A	U	A	C	C	C	G
X308	CAU SCENEDESMUS OBLIQ. CHILORO	C	G	C	A	G	G	A	U	A	G	A	G	C	A	G	C	A	G	C	U	G	36	D	A	G	C	U	C	G	G	C	U	C	A	U	A	F	C	C	A
X355	CAU PHASEOLUS VULGARIS	C	G	C	G	G	A	U	A	G	A	G	C	A	G	C	A	C	U	G	36	D	A	G	C	U	C	G	C	A	G	C	U	C	A	U	A	C	C	U	G
X365	CAU PHASEOLUS VULGARIS CHILORO	C	G	C	G	G	A	U	A	G	A	G	C	A	G	C	A	C	U	G	36	D	A	G	C	U	C	G	C	A	G	C	U	C	A	U	A	C	C	U	G
X412	CAU PHASEOLUS VULGARIS CHILORO	U	G	C	G	G	A	U	A	U	U	G	G	A	A	D	A	G	D																						
X428	CAU NEUROSPORA CRASSA	U	G	C	A	U	A	G	A	U	G	A	G	U	A	A	D	A	G	D																					
X438	CAU YEAST	U	G	C	A	U	A	G	A	U	G	A	G	U	A	A	D	A	G	D																					
X438	CAU PHASEOLUS VULG. "	A	G	C	G	G	G	U	A	G	A	G	U	A	G	U	A	G	D																						
X456	CAU AEDES ALBOPICTUS "	A	A	A	A	G	A	U	A	A	G	C	U	A	A	U																									
X480	CAU BOVINE LIVER "	A	G	U	A	G	C	U	C	A	G	C	U	A	U																										
X530	CAU TETRAHYMENA THERM.	A	G	C	A	G	G	U	G	C	G	A	A	D																											
X540	CAU SCENEDESMUS OBLIQ.	A	G	C	U	G	A	G	U	G	2C	G	C	A	G	D																									
X560	CAU NEUROSPORA CRASSA	A	G	C	U	G	C	A	U	G	C	G	C	A	G	C																									
X570	CAU SACCHAROMYCES CER.	A	G	C	G	C	G	U	G	2C	G	C	A	G	D																										
X580	CAU TORULOPSIS UTILIS	A	G	C	G	U	C	U	U	G	2C	G	C	A	G	D																									
X625	CAU WHEAT GERM	A	U	C	A	G	U	G	2C	G	C	A	G	C																											
X640	CAU LUPINUS LUTEUS	A	U	C	A	G	U	G	2C	G	C	A	G	C																											
X645	CAU PHASEOLUS VULGARIS	A	U	C	A	G	U	G	2C	G	C	A	G	C																											
X750	CAU ASTERINA ANURENSIS	A	G	C	A	G	A	G	U	G	2C	G	C	A	G	U																									
X780	CAU DROSOPHILA MELANO.	A	G	C	A	G	A	G	U	G	2C	G	C	A	G	U																									
X790	CAU EUPHASTIA SPERBA	A	G	C	A	G	A	G	U	G	2C	G	C	A	G	U																									
X830	CAU XENOPUS LAEVIS	A	G	C	A	G	U	G	2C	G	C	A	G	C																											
X850	CAU SALMON LIVER	A	G	C	A	G	U	G	2C	G	C	A	G	C																											

	45	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76															
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76															

X251	A	G	G7U	C	U	C	G	G	G	T	F	C	A	A	U	C	G	G	C	C	C	C	C	C	C	A	C	C	A											
X290	A	G	G7U	C	A	G	G	G	T	F	C	A	A	U	C	C	U	C	C	C	C	C	C	C	C	C	C	C	A	C	A									
X308	A	U	G7D	C	G	C	A	G	G	T	F	C	A	A	U	C	C	U	C	C	U	G	C	A	C	C	C	C	A	C	C	A								
X355	A	A	G7V7	U	A	C	G	G	G	T	F	C	A	A	U	C	C	G	G	U	C	C	G	C	A	C	C	C	A	C	C	A								
X365	A	G	G7U	C	A	C	G	G	G	T	F	C	A	A	U	C	C	U	G	C	C	C	C	C	C	C	C	C	A	C	C	A								
X412	U	G	A	C	A	U	A	G	G	J	G	C	A	A	U	C	C	G	U	A	U	C	C	G	C	A	C	C	C	A	C	C	A							
X428	U	U	A	U	A	C	C	T	F	C	A	A	U	C	C	G	U	A	U	U	G	C	F	A	C	C	C	A	C	C	A	C	C	A						
X438	A	G	A	D	G	A	G	G	T	F	G	A	A	U	C	C	U	G	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	A					
X456	U	U	A	U	A	A	G	G	J	A	U	A	A	U	C	C	U	F	F	C	U	U	U	A	C	C	C	C	A	C	C	C	C	C	C	A				
X480	A	A	A	U	G	U	G	G	J	U	A	U	A	U	C	C	U	C	C	G	A	C	U	A	C	C	C	C	A	C	C	C	C	C	C	C	A			
X530	A	A	G7U	C	S	A	G	A	G	A	F	C	G	A	A	C	C	U	C	U	C	U	C	U	C	U	C	C	C	A	C	C	C	C	C	C	C	A		
X540	A	G	G7D	C	S	A	C	A	G	A	U	C	G	A	A	C	C	U	U	C	C	A	C	C	U	C	C	C	C	A	C	C	C	C	C	C	C	A		
X560	A	G	G7*	C	A	C	U	C	G	A	U	C	G	A	A	C	G	A	U	G	C	A	G	C	A	C	C	C	C	C	C	C	C	C	C	C	C	C	A	
X570	A	U	G7D	C	S	C	S	U	C	G	A	U	C	G	A	A	C	C	G	N	C	G	G	C	U	A	C	C	C	C	C	C	C	C	C	C	C	C	A	
X580	A	U	G7D	C	S	C	S	C	U	G	A	U	C	G	A	A	C	C	A	G	A	G	A	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	A	
X625	A	G	G7D	C	S	C	S	C	A	G	A	F	C	G	A	A	C	C	N	G	C	U	C	U	G	A	U	C	C	C	C	C	C	C	C	C	C	C	A	
X640	A	G	G7D	C	S	N	C	A	G	A	F	C	G	A	A	C	C	U	G	C	U	C	U	G	A	U	C	C	C	C	C	C	C	C	C	C	C	C	A	
X645	A	G	G7D	C	S	C	C	A	G	A	F	C	G	A	A	C	C	U	G	S	G	C	U	C	U	A	U	C	C	C	C	C	C	C	C	C	C	C	C	A
X750	A	G	G7D	C	S	C	G	A	G	A	F	C	G	A	A	C	C	U	C	G	C	U	C	U	G	A	U	C	C	C	C	C	C	C	C	C	C	C	C	A
X780	A	G	G7D	C	S	C	G	A	G	A	U	C	G	A	A	C	C	U	G	C	U	C	U	G	A	U	C	C	C	C	C	C	C	C	C	C	C	C	C	A
X790	A	G	G7U	C	G	G	A	G	A	F	C	G	A	A	C	U	A	C	U	C	U	C	U	G	A	U	C	C	C	C	C	C	C	C	C	C	C	C	C	A
X830	A	G	G7D	C	S	G	A	U	G	A	U	C	G	A	A	C	C	A	U	C	C	U	C	U	G	A	U	C	C	C	C	C	C	C	C	C	C	C	C	A
X850	A	G	G7D	C	S	G	A	U	G	A	U	C	G	A	A	C	C	A	U	C	C	U	C	U	G	A	U	C	C	C	C	C	C	C	C	C	C	C	C	A

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44											
X950	CAU	HOUSE	MYELOMA	A	G	C	A	G	A	G	U	G	G	C	G	C	G	C	A	G	G	A	A	G	C	G	U	G	C	U	G	G	C	C	A	U	A	T	A	C	C	C	A	G											
X955	CAU	RABBIT	LIVER	A	G	C	A	G	A	G	U	G	G	C	G	C	A	G	C	G	G	A	A	G	C	G	U	G	C	U	G	G	C	C	A	U	A	T	A	C	C	C	A	G											
X980	CAU	SHEEP	MAMMARY GLAND	A	G	C	A	G	A	G	U	G	G	C	G	C	A	G	C	G	G	A	A	G	C	G	U	G	C	U	G	G	C	C	A	U	A	T	A	C	C	C	A	G											
X995	CAU	HUMAN	PLACENTA	A	G	C	A	G	A	G	U	G	G	C	G	C	A	G	C	G	G	A	A	G	C	G	U	G	C	U	G	G	C	C	A	U	A	T	A	C	C	C	A	G											
P H E N Y L A L A N I N E																																																							
F120	GAA	HALOBACTERIUM	VOL.	G	C	C	G	C	C	U	A	G	C	U	C	A	A	C	U	G	G	G	A	G	A	C	U	C	A	C	U	G	A	A	C	U	G	A	A	G	A	A	F	C	C	A	G								
F200	GAA	MYCOPLASMA	CAPRIC.	G	G	U	C	G	U	G	A	G	C	U	C	A	G	U	C	G	G	D	A	G	A	C	A	G	C	A	G	C	U	G	A	A	G	A	G	A	G	A	G	A	G	A	F	C	U	G	C				
F230	GAA	BACILLUS	STEARO.	G	G	C	U	C	G	G	U	A	G	C	U	C	A	G	U	C	G	D	A	G	A	C	A	G	C	A	G	C	U	G	A	A	G	A	G	A	G	A	G	A	A	A	F	C	C	U	U				
F235	GAA	BACILLUS	SUBTILIS	G	G	C	U	C	G	G	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	G	A	G	A	A	A	A	A	A	A	F	C	C	U	U				
F250	GAA	E-COLI		G	C	C	C	G	G	A	U	A	G	C	U	C	A	G	D	C	G	D	A	G	A	C	A	G	C	U	G	A	A	G	A	A	G	A	A	A	A	A	A	A	A	A	A	F	C	C	C				
F280	GAA	RHODOSPIRIL	RUB.	G	C	C	C	G	G	U	A	G	C	U	C	A	G	D	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	F	C	A	C	G		
F285	GAA	AGHENEILLUM	QUADR.	G	C	C	A	G	A	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	C		
F304	GAA	EUGLEMA	GRACILIS	G	C	U	G	G	A	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	C	
F350	GAA	PHASEOLUS	VULGARIS	G	U	C	G	G	A	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	C	
F365	GAA	SPINACIA	OLERACEA	G	U	C	G	G	A	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	C
F407	GAA	TETRAHYMENA	PYRIF-U	G	C	U	A	A	G	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	C	
F417	GAA	SACCHAROMYCES	CER.	G	C	U	U	A	U	A	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	U	
F438	GAA	PHASEOLUS	VULGARIS	G	U	U	A	G	A	G	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	U		
F474	GAA	RAT MORRIS	HEPATOMA	G	U	U	A	U	A	G	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	U		
F510	GAA	EUGLEMA	GRACILIS	G	C	C	G	A	C	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	U		
F540	GAA	SCENEDESMUS	OBLIQ.	G	G	C	U	U	G	A	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	F	C	C	U	U		
F560	GAA	NEUROSPORA	CRASSA	G	C	C	G	G	U	A	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	F	C	S	U	G	A	
F575	GAA	SCHIZOSACCHARIA	POM.	G	U	C	G	C	A	U	A	G	C	U	C	A	G	U	C	G	D	G	A	G	A	C	A	G	C	U	G	A	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	F	C	S	U	G	U	

	44	45	46	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	
X950	A	G	G	T																C	S	G	A	U	G	G	A	U	C	G	A	A	C	C	A	U	C	C	U	C	U	G	G	C	U	A	C	C	A
X955	A	G	G	T																C	S	G	A	U	G	G	A	U	C	G	A	A	C	C	A	U	C	C	U	C	U	G	G	C	U	A	C	C	A
X980	A	G	G	T																C	S	G	A	U	G	G	A	U	C	G	A	A	C	C	A	U	C	C	U	C	U	G	G	C	U	A	C	C	A
X995	A	G	G	T																C	S	G	A	U	G	G	A	U	C	G	A	A	C	C	A	U	C	C	U	C	U	G	G	C	U	A	C	C	A
P H E N Y L A L A N I N E																																																	
F120	C	U	G	U																C	S	C	C	G	G	F	F	C	S	I	A	U	C	C	G	G	A	G	G	C	G	G	C	A	C	C	A		
F200	G	U	G	U																C	G	G	G	U	F	C	A	A	U	C	C	G	G	C	C	A	C	G	A	C	C	A	C	C	A	C	C	A	
F230	G	U	G	U																C	G	G	G	T	F	C	G	A	U	C	C	G	G	C	C	G	A	G	C	C	A	C	C	A	C	C	A		
F235	G	U	G	U																C	G	C	G	T	F	C	G	A	U	C	C	G	C	C	G	A	G	C	C	A	C	C	A	C	C	A			
F250	G	U	G	U																C	U	U	G	G	T	F	C	G	A	U	C	C	G	A	G	U	C	C	G	G	C	A	C	C	A				
F280	G	U	G	U																C	G	G	G	T	F	C	G	A	U	C	C	G	C	C	C	C	G	G	G	C	A	C	C	A					
F285	G	U	G	U																C	G	C	G	T	F	C	A	A	U	C	C	G	C	U	C	C	G	G	C	A	C	C	A						
F304	G	U	G	U																C	A	C	C	A	G	T	F	C	A	A	U	C	G	G	U	C	C	U	A	G	C	A	C	C	A				
F350	G	U	G	U																C	A	C	C	A	G	T	F	C	A	A	U	C	G	G	U	C	C	U	A	G	C	A	C	C	A				
F365	G	U	G	U																C	A	C	C	A	G	T	F	C	A	A	U	C	G	G	U	C	C	U	A	G	C	A	C	C	A				
F407	A	G	G	U																C	A	U	G	G	U	J	C	C	G	A	U	C	C	A	U	C	U	C	U	A	G	G	C	A	C	C	A		
F417	U	U	A																	C	A	U	G	A	G	T	F	C	G	A	U	C	C	A	U	A	A	G	G	C	A	C	C	A					
F438	G	U	G	U																C	A	G	U	G	T	F	C	G	A	U	C	C	A	U	C	U	C	U	A	A	G	C	C	A					
F474	A	G	A																	C	U	G	A	U	C	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
F510	A	G	U																	C	A	C	C	A	G	T	F	C	G	A	U	C	C	G	G	A	G	C	G	C	A	C	C	A					
F560	A	G	G	U																C	S	C	C	A	G	T	F	C	G	A	U	C	C	S	U	G	G	U	C	A	G	C	C	A	C	C	A		
F560	A	G	G	U																C	S	G	U	G	T	F	C	G	A	U	C	C	A	C	A	C	A	A	A	C	C	G	C	A	C	C	A		
F575	U	G	G	U																C	A	U	C	G	T	F	C	G	A	U	C	C	C	G	G	U	U	G	G	A	C	A	C	C	A				


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45 47 B D F H J L M P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
44 46 A C E G I K O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76
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F590  A G G7U      C C S U G U G T F C G A I U C C A C A G A A U U C G C C A C C A
F591  A G G7U      C C S U G U G T F C G A I U C C A C A G A G U U C G C C A C C A
F620  A G G7D      C G C G U G T F C G A I U C C A C G C U C A C G C C A C C A
F625  A G G7D      C G C G U G T F C G A I U C C A C C C U C A C G C C A C C A
F630  A G G7D      C G C G U G T F C G A I U C C A C G C U C A C G C C A C C A
F640  A G G7D      C A C G U G T F C G A I U C C A C G U U C A C C G C A C C A
F650  A G G7D      C G C G U G T F C G A I U C C A C G C U C A C G C C A C C A
F770  A G G7Dn      C C C U G G T F C A A I U C C C G G G U U C G G C A C C A
F771  A G G7Dn      C C C U G G T F C G A I U C C C G G G U U C G G C A C C A
F780  A G G7Dn      C C C G G T F C A A I U C C C G G G U U C G G C A C C A
F830  A G G7D      C C S C U G G T F C G A I U C C C G G G U U C G G C A C C A
F950  A G G7U      C C S C U G G T F C G A I U C C C G G G U U C G G C A C C A
F951  A G G7U      C C S C U G G T F C G A I U C C C G G G U U C G G C A C C A
F952  A G G7U      C C S C U G G T F C G A I U C C C G G G U U C G G C A C C A
F965  A G G7Dn      C C S C U G G T F C G A I U C C C G G G U U C G G C A C C A
F970  A G G7D      C C S C U G G T F C G A I U C C C G G G U U C G G C A C C A
F971  A G G7D      C C S C U G G T F C A A I U C C C G G G U U C G G C A C C A
F974  A G G7D      C C S C U G G T F C G A I U C C C G G G U U C G G C A C C A
F995  A G G7Dn      C C S C U G G T F C G A I U C C C G G G U U C G G C A C C A
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P R O L I N E
P013  A G G7D      C C S C G G G F F C A A I A U C C C C G G A C G A G C C C C C A
P014  A G G7D      C C S C G G G F F C A A I A U C C C C G G A C G A G C C C C C A
P020  A G G7U      C A A G G T F C A A A U C C U U G A U G G A G A C C A

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P040	UGG PHAGE T5	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43
P120	CGG HALOBACTERIUM VOL.	U A G C U C A A U U G G C D A G A G U A C A C C G U U U G G G T G C G G U G
P121	GGG HALOBACTERIUM VOL.	G G G C C G G U G G A G F A N C U U G G U A U C C U U C G G C C U U C A G G G T F G C C G
P122	GGG HALOBACTERIUM VOL.	G G G A C C G U G G A G F A G U G G U A U C C U C G C C A U G G G T U C G G U A
P203	UGG MYCOPLASMA MYCOID.	G G A C C G U G G A G F A G C C U G G U A A C U U C G G C C U U G G G T U G C C C G
P235	UGG BACILLUS SUBTILIS	C G G G A G G U G C U C A G U U G G D A G A G C A U C G G U U U G G G T A C C G A A
P255	CGG SALMONELLA TYPHI.	C G G G A G U A G C U C A G C U U G G D A G A G C A C A U G G F U V 2 6 G G T A C C A U G
P256	GGG SALMONELLA TYPHI.	C G G U G A U U A G G C G C A G C C U G G D A G C G C A C U U C G U 3 U C G G G T A C G A A G
P257	UGG SALMONELLA TYPHI.	C G G C A C G U A G C G C A G C C U G G D A G C G C A C C G U C C 3 U G G G T U F G C G G
P365	UGG SPINACIA OLERACEA	C G C G A G U A G C G C A G C U U G G D A G C G C A C U G G U 3 U V 1 6 G G T A C C A G U
P417	UGG SACCHAROMYCES CER. CHILDO MITO	A G G A U G U A G C G A G C U U G 3 G D A G C C C F U U G U F U N G G N F A C A A A
P438	UGG PHASEOLUS VULGARIS MITO	C A G A U A G A G C C A A A G G D C A G G C G C U U U C F U U G G G T A G A A A G
P570	UGG SACCHAROMYCES CER.	C G A G G U G U A G C G C A G U C D G 3 G D C A G C G C A U C U G U U U U G G G T F A C A G A
P571	UGG SACCHAROMYCES CER.	G G G C 3 G U G U G U C F A G D G G D A U G A U U C U C G C F U N G G G T F G C G A G
P580	UGG TORULOPSIS UTILIS	G G G C 3 G U G U G U C F A G D G G D A U G A U U C U C G C U U G G G T F G F G A G
P640	AGG LUPTINUS LUTEUS	G G G A C A U G G U C F A G D G G D A U G A U U C U C G C U U I G G G T F G C G A G
S E R I N E		
S020	UGA PHAGE T4	G G A G G C G U A G G C A G A G U G 3 G D D U A U G C A C C G G U C 3 U N G A A 5 A A C C G G
S110	CGA HALOBACTERIUM CUT.	G C G A G G U A G C A F A N C U U G C C A U G C G A G U U G C C U U C 4 G A G 1 A C C A A C
S120	GCU HALOBACTERIUM VOL.	G U U G C G U A G C C A N C C U G G C C A A G G C G A C U G G G U U G G C U A 7 A C C A G
S121	CGA HALOBACTERIUM VOL.	G C C G A G G U A G C C F A N C C C G G C C A A G G C G A G U A G A U U C 4 G A A F C U A C
S122	GGG HALOBACTERIUM VOL.	G C C A G G A U G C C G A N C G G U A A G G C G A C A C C G C U G G A A A G C G U G
S203	UGA MYCOPLASMA MYCOID.	G G A A G A U A C C C A A G U C C G G C D G A N G G G A U C G G U C U U G A A 6 A A C C G A

	45 47 B D F H J L N P	49 51 53 55 57 59 61 63 65	67 69 71 73 75
	44 46 A C E G I K M O	48 50 52 54 56 58 60 62 64	66 68 70 72 74 76
P040	GG GTU	UGAAGGTCFCGAGUCCUUC	AUUGGAGACCA
P120	UAA	CSCUCAGFIFC3GAAUCUGAG	CCGGCCCA
P121	GGA	C5UCAGGFC3GACUCUCAG	GGUCCCA
P122	UGA	C5CCGGFIFC311AAUC	CGGGGCGUCCCA
P203	GGGU	CGAGGUCAAAUCUUGU	CUUCCGACCA
P235	GGGTU	CGAGGTCFCAAUCUUGU	CUUCCGACCA
P255	GGGTU	CGAGGTCFCAAUCUCU	AUACCGACCA
P256	GGGTU	CGAGGTCFCAAUCUCU	CGUGCCGACCA
P257	GGGTU	CGAGGTCFCAAUCUCF	CUCCGACCA
P365	AUGTU	CACGGTCFAAUCUUGU	CAUCCUACCA
P417	ACCU	AGUAGTCFAGAUCAUCU	CUAUCUGACCA
P438	GGGTC	CAUAGGTCFCAAUCUGU	CACCUUGACCA
P570	AGGTC	CUGGGTCFCAUUCUCCAG	CUCGCCCA
P571	AGGTC	NUGGGTCFCAUUCUCCAG	CUCGCCCA
P580	UGGTU	C5CAAGGTCFCAUUCUCCU	GUCCGACCA
P640	AGGTU	C5CAGTCFCAUUCUCCG	AUGUCCCA
S E R I N E			
S020	CAGUCGUCGGCGACU	CAUAGGTCFAAUCU	CUAUCGCCUCCGCA
S110	GUCCACA CGGACU	C5AAGGFC311AAUCUCU	CCUGGCGCA
S120	UGGCGUCA GCCC	C5CGGGFIFC3GAAUC	CCCGCGCAACGCA
S121	UGUCCAUUCGGACA	C5UAGGFC311AAUCU	CACCGCGCGCA
S122	UUCUUCUGGGAU	C5GGGGFIFC311AAUC	CCUCCUGGCGCA
S203	GAGUCGGGAACCCGAG	CGGGGUUCGAAUC	CCCUAUCUCCGCA

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45				
S250	CGA	E.	COLI	U	G	C	C	G	G	A	C	C	G	G	A	C	G	G	A	C	G	G	A	C	C	G	G	A	C	C	G	G	A	C	C	G	G	A	C	C	G	G	A	C	C	G	G		
S251	GGU	E.	COLI	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G		
S252	GGG	E.	COLI	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G		
S253	GGG	E.	COLI	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G		
S254	UGA	E.	COLI	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G	C	C	G	A	G	G		
S417	GGU	SACCHAROMYCES CER.	G	G	A	A	A	U	A	C	U	A	D	A	A	A	G	A	U	A	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F			
S418	UGA	SACCHAROMYCES CER.	G	G	A	U	G	A	C	U	A	G	D	A	A	A	G	A	U	A	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F			
S419	UGA	SACCHAROMYCES CER.	G	G	A	U	G	A	C	U	A	G	D	A	A	A	G	A	U	A	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F			
S456	GGU	AEDES ALBOPICTUS	G	G	A	U	G	A	C	U	A	G	D	A	A	A	G	A	U	A	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F	F			
S473	GGU	HAMSTER	G	G	A	A	U	G	A	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	
S480	GGU	BOWTIE HEART	G	G	A	A	A	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	
S481	GGU	BOWTIE LIVER	G	G	A	A	A	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	U	
S494	GGU	HUMAN	G	G	A	A	A	G	C	U	C	U	C	U	C	A	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	
S570	UGA	SACCHAROMYCES CER.	G	G	C	A	C	U	A	G	C	C	A	G	D	A	A	G	G	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G
S590	AGA	YEAST	G	G	C	A	C	U	A	G	C	C	A	G	D	A	A	G	G	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G
S591	AGA	YEAST	G	G	C	A	C	U	A	G	C	C	A	G	D	A	A	G	G	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G
S640	GGU	LUPINUS LUTEUS	G	U	C	A	G	U	A	G	C	C	A	G	D	A	A	G	G	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G
S780	GGU	DROSOPHILLA MELANO.	G	A	C	G	A	G	G	U	G	C	C	A	G	D	A	A	G	G	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A
S781	AGA	DROSOPHILLA MELANO.	G	C	A	G	S	C	U	A	G	C	C	A	G	D	A	A	G	G	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	
S782	CGA	DROSOPHILLA MELANO.	G	C	A	G	S	C	U	A	G	C	C	A	G	D	A	A	G	G	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	
S950	UGA	MOUSE LIVER	G	C	C	C	G	A	U	G	A	U	C	C	A	G	U	C	A	G	U	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	
S955	AGA	RAT LIVER	G	U	A	G	U	C	C	A	G	A	G	D	A	A	G	G	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	
S956	GGU	RAT LIVER	G	A	C	G	A	G	G	U	G	C	C	A	G	D	A	A	G	G	C	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	

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45 47 B D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76
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S250  A G U A G G G G C A A C U C U A C   C G G G G T F C A A U U   C C C C C U C U C U C C   G C C A
S251  U A U G C G C U C A A A G C U G C A U C   C G G G G T F G A A U   C C C C G C U C A C C C   G C C A
S252  U A U A C G C A C G U A U   : : : : : C G G G G T F G A A U   C C C C C C U C A C C C   G C C A
S253  U A U A C G G C A C G U A U   C G G G G T F G A A U   C C C C C C U C A C C C   G C C A
S254  C G A C C G A A G G G U U   C A G A G T F G A A U   C U C U G C G C U U C C   G C C A
S417  U G A A U U G U A A U U C U   : : : : : U A U G A G T F G A A U   C U C A U A U U U U C C   G C C A
S418  U A G U C U U U A U U G G C U A   : : : : : C G U A G G T F C A A U   C C U A C A U C A U C C   G C C A
S419  U U G U C U U U A U U G C U A   : : : : : C G U A G G T F C A A U   C C U A C A U C A U C C   G C C A
S456  U C U U   : : : : : U A U G G U U U A A U U   C C A U F A A U U U C U   J C C A
S473  G C U   A C C A U G U A A A U A A C   A U G G C U U C U U A C C A
S480  U G C U   C C A U A U C U A A U A G   A U G G C U U U U U C C   G C C A
S481  U G C U   C C A U A U C U A A U A G   A U G G C U U U U U C C   G C C A
S494  G C C   C C A U G U C U A C A A C   A U G G C U U U C U C A C C A
S570  U36 G G C U C U G C C C G   C56 C U G G T F C A A U   C C U G C U G G U G C   G C C A
S590  U36 G G C U C U G C C C G   C56 C A G G T F C A A U   C C U G C A G U G U C   G C C A
S591  U36 G G C U U G C C C G   C56 C A G G T F G A A U   C C U G C A G U U G U C   G C C A
S640  U36 A G A U U G C C U G G   C56 G G G G T F C G A I U   C C G C U A C U G A C   G C C A
S780  U36 U G C U C60 G C A C G   C56 U G G G T F G G A I U   C C C A U C C U C U C   G C C A
S781  U30 C C C U C60 G G G A G   C56 U A G G T F C G A I U   C C U A C C G A C U G C   G C C A
S782  U30 C C C U C60 G G G A G   C56 U A G G T F G G A I U   C C U A C C G G C U G C   G C C A
S950  A G C U G U U A G C G A C A   G A G U G G U F C A A I U   C C A C C U U C G G G C   G C C A
S955  U36 G G G U C60 C C C C G   C56 C A G G T F G G A I U   C C U G C C G A C U A C   G C C A
S956  U36 F G C U C60 G C A C G   : : : : : C56 U G G G T F C G A I U   C C C A U C C U C G U C   G C C A

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	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43				
S957	AGA	RAT	MORRIS	HEPATOMA	G	U	A	G	U	C	G	U	G	C	C	A	G	D	G	G	D	A	G	G	C	A	F	G	G	A	C	S	U	I	G	A	A	A	F	C	C	A	U				
S970	CCA	BOVINE	LIVER	G	C	C	C	G	G	A	U	G	A	U	C	C	U	C	A	G	U	G	G	C	C	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G	G		
S971	NCA	BOVINE	LIVER	G	C	C	C	G	A	U	G	A	U	C	C	U	A	G	U	G	G	U	G	G	G	C	A	G	C	A	G	C	U	N	C	A	A	A	C	C	U	G	U	G	U		
S995	UCA	HUMAN	HELA	C	C	C	G	A	U	G	A	U	C	C	U	C	A	G	U	G	G	U	G	G	G	C	A	G	C	U	N	C	A	A	A	C	C	U	G	U	G	U	G	U	G		
S996	UCA	HUMAN		G	U	A	G	U	C	G	U	G	C	C	A	G	D	G	S	G	D	A	G	G	C	A	F	G	G	A	C	U	G	A	A	A	F	C	C	A	U	G	U	G	U		
T H R E O N I N E																																															
T020	UGU	PHAGE	T4	G	C	U	G	A	U	U	A	G	C	U	C	A	G	D	A	G	G	A	G	G	C	A	C	C	U	C	A	C	U	H	G	U	N	A	F	G	A	G	G	G			
T110	GGU	HALOBACTERIUM	CUT.	G	C	C	G	G	A	U	G	C	U	F	A	G	C	G	G	U	A	A	A	G	C	G	A	C	C	G	U	C	C	U	C	C	U	G	G	U	A	T	A	G	A	C	
T120	CGU	HALOBACTERIUM	VOL.	G	C	C	G	U	G	A	G	C	U	C	A	N	U	U	G	G	C	A	G	A	G	C	A	U	U	C	C	U	C	G	U	A	T	A	G	A	G	A	G	A	U	G	
T121	GGU	HALOBACTERIUM	VOL.	G	C	C	U	G	G	A	G	C	U	F	A	G	C	G	G	U	A	A	A	G	C	G	U	C	C	U	C	G	U	C	C	U	G	G	U	A	T	A	G	A	C		
T200	AGU	MYCOPLASMA	CAPRIC.	G	C	U	G	A	C	U	N	A	G	C	U	C	A	G	U	D	G	G	A	G	G	C	A	U	U	G	A	C	U	A	G	A	C	U	A	G	U	A	T	A	F	C	
T201	UGU	MYCOPLASMA	CAPRIC.	G	C	U	G	A	C	U	N	A	G	C	U	C	A	G	C	A	G	C	A	G	G	C	A	G	G	C	A	C	U	G	A	C	U	G	A	C	U	G	U	A	T	A	F
T203	UGU	MYCOPLASMA	MYCOID.	G	C	U	G	A	C	U	A	G	C	U	C	A	G	C	A	G	C	A	G	C	A	G	G	C	A	C	U	G	A	C	U	G	A	C	U	G	U	A	T	A	F	C	
T235	UGU	BACILLUS	SUBTILIS	G	C	C	G	U	G	A	G	C	U	C	A	A	U	D	G	G	D	A	G	G	C	A	C	U	G	A	C	U	G	A	C	U	V	2	G	U	A	T	A	F			
T250	GGU	E. COLI		G	C	U	G	A	U	A	G	C	U	C	A	G	D	D	G	D	A	G	G	C	C	A	C	C	C	U	G	G	U	A	B	A	G	G	G	U	G	G	U	G	G		
T365	GGU	SPINACIA	OLERACEA	G	C	C	C	U	U	A	A	C	U	C	A	G	U	G	S	G	D	A	G	A	G	U	A	C	G	C	A	C	C	A	U	G	G	U	A	B	A	G	G	C	G		
T412	UGU	NEUROSPORA	CRASSA	G	C	C	U	G	U	A	G	C	U	A	A	A	A	A	G	D	A	A	U	G	C	A	U	U	F	J	U	U	G	A	T	A	F	C	A	A	U	G	U	A	T		
T417	UAG	SACCHAROMYCES	CER.	G	U	A	A	U	A	A	A	U	U	A	A	D	G	D	A	A	A	U	G	F	A	U	G	F	J	U	U	A	G	G	T	F	C	A	A	U	A	G	G	U			
T480	UGU	MITO	LIVER	G	U	C	U	U	G	A	G	U	A	C	A	U	C	U	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
T590	AGU	YEAST		G	C	U	C	U	A	U	G	G	C	A	A	G	D	D	G	D	A	G	G	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	
T591	AGU	YEAST		G	C	U	C	U	A	U	G	G	C	A	A	G	D	D	G	D	A	G	G	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A	C	A		
T970	GGU	BOVINE	LIVER	G	C	C	C	U	G	U	G	C	C	U	A	G	C	D	G	D	C	A	A	G	G	C	C	U	G	U	A	T	A	G	C	C	U	G	U	A	T	A	F	C			

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44
T R Y P T O P H A M																																												
W012	CCA	CHICKEN	ASV/MNV/RSV	G A C C U C G U	U G162C	G C A A C	G3G D	A G C G C	64F C U G A	C3U C3C A G1A F3C A G A																																		
W120	CCA	MALOBACTERIUM	VOL.	G G G C U G U	U G C C A	A N C C C G G C	A U G G C	64A C U G A	C3U C3C A G1A U3C A G U																																			
W250	CCA	E.COLI		A G G G G C G U A A	G U U C A	A A D D G G D	A G A G C	A C C G G U	C3U C C A A5A A C C G G																																			
W350	CCA	PHASEOLUS	VULGARIS	A C G C U C U	U A G U U C	A G U U C G3G C	A G A A C	G G G F	C U C3C A N A A C C C G																																			
W365	CCA	SPINACIA	OLERACEA	G C G C U C U	U A G U U C	A G U U C G3G D	A G A A C	64F G G G F	C U C C A N A A C C C G																																			
W412	UCA	NEUROSPORA	CRASSA	A A G A G U A	U A G U U U	A A D G G D	A A A C A	G A A G C	U N C A N C U U U A																																			
W417	UCA	MITOCHONDRION	CER.	A A G A U A	U A G U U U	A A D G G D	A A A C A	G U U G A	F U N C A N A F C A A U																																			
W438	CCA	PHASEOLUS	VULGARIS	G C G C U C U	U A G U U C	A G U U C G3G D	A G A A C	64C G G G F	C U C3C A N A A C C C G																																			
W474	UCA	RAT MORRIS	HEPATOMA	A G A A G U U	U A G C G A	U A U A C	A G U C C A	A G A G C	C U N C A A5A G C C C U																																			
W480	UCA	BOVINE	LIVER	A G G A A U U	U A G26	U U A A C	A G A C C A	A G A G C	C U N C A A5A G C C C U																																			
W570	CCA	SACCHAROMYCES	CER.	G A A G C G G U	U G162C	U C A A D G3G D	A G A G C	F F C G A	C3U C3C A A A F C G A A																																			
W625	CCA	WHEAT	GERM	G G A N C C G U	U G162C	G C A A D D G3G D	A G C G C	64F C U G A	C3U C3C A G1A F3C A G A																																			
W870	CCA	CHICKEN		G A C C U C G U	U G162C	G C A A C G3G D	A G C G C	64F C U G A	C3U C3C A G1A F3C A G A																																			
W970	CCA	BOVINE	LIVER	G A C C U C G U	U G162C	G C A A D G3G D	A G C G C	64F C U G A	C3U C3C A G1A F3C A G A																																			
T Y R O S I N E																																												
Y120	GUA	MALOBACTERIUM	VOL.	C C G C U C U	U A G C U C A	N C C U G G C	A G A G C A	G C C G A	C3U G U A G1A F C G G C																																			
Y230	GUA	BACILLUS	STEARO.	G G A G G G U A A	U G C G A	A G U G3G C U A	A1C G C G	G C C G A	C U Q U A A5A F C C G C																																			
Y235	GUA	BACILLUS	SUBTILIS	G G A G G G U A A	U G C G A	A G U G3G C U A	A1C G C G	G C C G A	C U Q U A A4A F C C G C																																			
Y236	GUA	BACILLUS	SUBTILIS	G G A G G G G U A A	U G C G A	A G U G3G C U A	A1C G C G	G C C G A	C U Q U A A5A F C C G C																																			
Y250	GUA	E.COLI		G G U G G G G U A U	U C C C G A	G C G3G C C A	A A G G G A	G C A G A	C U Q U A A5A F C U G C																																			
Y251	GUA	E.COLI		G G U G G G G U A U	U C C C G A	G C G3G C C A	A A G G G A	G C A G A	C U Q U A A5A F C U G C																																			
Y308	GUA	SENEDESMUS	OBLIQ.	G G G C C G A	U G C C G A	A G U N G3G D U	A U G G G G	G C C G A	F U G U A A4A F C C G C																																			

45 47 B D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
 44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76

TRYPTOPHAN

W012 A G GTC U G C G U G F C G A I A U C A C G U C G G G G U C A C C A
 W120 C G A U C S G G G G F I F C S I I A A U C C C U C G G C C C C A C C A
 W250 G U G7U U G G A G T F C G A G U C U C U C G C C C C U G C C A
 W350 A U G7U C U A G G T F C A A A U C U A C A G A G C G U G C C A
 W365 A U G N C U A G G T F C A A G U C U A C A G A G C G U G C C A
 W412 A A U U C U U A G T F C G A G U C U A A G U A C U C U U G C C A
 W417 C A U U A G G A G T F C G A A U C U C U U A U C C U U G C C A
 W438 A U G7N C U A G G T F C A A A U C C U A C A G A G C G U G C C A
 W474 U A G A A A C A A C A C A G U U A C U C U C U G C C A
 W480 A A G C A G U A C A A U U A C U U A A U U C C U G C C A
 W570 G G G7D U G C A G G T F C A A I U U C C U G F C C G U U C A C C A
 W625 A G G D U G C G U G T F C G A I U U C A C G F C G G U C C A C C A
 W870 A G G7C U G C G U G F C G A I A U C A C G U C G G G G U C A C C A
 W970 A G G7D U G C G U G F C G A I A U C A C G U C G G G G U C A C C A

TYROSINE

Y120 U U G U C S C C C G G F I F C S I I A A U C G G G G A G A G C G G A C C A
 Y230 U C C U U G G G U U C G C G G T F C G A A U C G U C C C C U C C A C C A
 Y235 U C C U C A G G G U U C G C A G T F C G A A U C U G C C C C U C C A C C A
 Y236 U C C U C A G G G U U C G C A G T F C G A A U C U G C C C C U C C A C C A
 Y250 C G U C A U C G A C U C A A G G T F C G A A U C U U C F C C C A C C A C C A
 Y251 C G U C A C A G A C U U C A A G G T F C G A A U C U U C C C C A C C A C C A
 Y308 U G G U U A C G C C U A C G U G G T F C G A A U C C G A C U C G G C C C A C C A

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43												
Y407	GUA	TETRAMYEMA	PYRIF.																																																				
Y412	GUA	HELIOSPORA	CRASSA																																																				
Y417	GUA	SACCHAROMYCES	CER.																																																				
Y438	MUA	PHASEOLUS	WULG.																																																				
Y540	GUA	SCENEDESMUS	ORLIG.																																																				
Y575	GUA	SCHIZOSACCHA.	POM.																																																				
Y580	GUA	TORULOPSIS	UTILIS																																																				
Y590	GUA	YEAST																																																					
Y625	GUA	WHEAT	GERM																																																				
Y626	GUA	WHEAT	LEAVES																																																				
Y640	GUA	LUPINUS	LUTEUS																																																				
Y655	GUA	NICOTIANA	RUSTICA																																																				
Y656	GUA	NICOTIANA	RUSTICA																																																				
Y780	GUA	DROSOPHILA	MELANO.																																																				
Y970	GUA	BOVINE	LIVER																																																				
Y995	GUA	HUMAN	PLACENTA																																																				
Y996	GUA	HUMAN	PLACENTA																																																				
V A L I N E																																																							
V110	CAC	HALOBACTERIUM	CUT.																																																				
V111	CAC	HALOBACTERIUM	CUT.																																																				
V112	GAC	HALOBACTERIUM	CUT.																																																				
V120	CAC	HALOBACTERIUM	VOL.																																																				
V121	GAC	HALOBACTERIUM	VOL.																																																				

	45	47	B	D	F	H	J	L	N	O	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75		
	44	46	A	C	E	G	I	K	M	O	P	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76	
Y407	U3G	G	G	U	A	U	C	C	G	U		C	G	U	C	G	G	A	U	C	C	G	A	U	C	C	A
Y412	U	G	A	C	U	A	A	U	A	G	U	C	S	G	U		C	G	A	A	U	C	C	U	U	C	C
Y417	U	G	A	C	U	A	A	G	U	C	U	U															
Y438	U	G	A	A	G	U	U	U	C	A		C	U	A	G	G	T	F	C	G	A	A	U	C	C	G	A
Y540	A	G	G	A								C	A	C	U	G	T	F	C	G	A	A	U	C	C	G	A
Y575	U	G	G	T								C	S	G	C	A	G	T	F	C	G	A	U	C	G	G	A
Y580	A	C	A	D								C	S	G	G	C	T	F	C	G	A	A	U	C	G	C	C
Y590	A	G	A	D								C	S	G	G	C	T	F	C	G	A	A	U	C	G	G	A
Y625	A	G	G	T								C	A	C	U	G	J	F	C	G	A	A	U	C	C	G	A
Y626	A	G	G	T								C	A	C	U	G	J	F	C	G	A	A	U	C	C	G	A
Y640	A	G	G	T								C	A	C	U	G	J	F	C	G	A	A	U	C	C	G	A
Y655	A	G	G	T								C	A	C	U	G	J	F	C	G	A	A	U	C	C	G	A
Y656	A	G	G	T								C	G	C	U	J	F	C	G	A	A	U	C	C	G	G	A
Y780	A	G	G	T								C	G	C	U	J	F	C	G	A	A	U	C	C	G	G	A
Y970	A	G	G	T								C	S	G	C	U	G	T	F	C	G	A	U	C	C	G	A
Y995	A	G	G	T								C	S	G	C	U	G	T	F	C	G	A	U	C	C	G	A
Y996	A	G	G	T								C	S	G	C	U	G	T	F	C	G	A	U	C	C	G	A
V A L I N E																											
V110	A	G	G	U								C	G	G	C	G	F	G	S	I	A	U	C	C	G	C	C
V111	A	G	G	U								C	G	C	S	G	F	F	C	S	I	A	U	C	C	C	C
V112	A	G	G	U								C	G	C	S	G	F	F	C	S	I	A	U	C	C	C	C
V120	A	G	G	C								C	S	G	C	G	F	F	C	S	G	A	U	C	C	C	C
V121	A	G	G	C								C	S	G	C	A	G	F	F	C	S	I	A	U	C	C	C

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44								
V203	UAC	MYCOPLASMA	MYCOID.																																																	
V230	GAC	BACILLUS	STEARO.																																																	
V235	UAC	BACILLUS	SUBTILIS																																																	
V250	GAC	E. COLI																																																		
V251	GAC	E. COLI																																																		
V252	UAC	E. COLI																																																		
V365	UAC	SPINACIA	OLERACEA																																																	
V412	UAC	CHILORO																																																		
V456	UAC	AEDES	ALBOPICTUS																																																	
V474	UAC	RAT	LIVER																																																	
V475	UAC	RAT	MORRIS	HEPATOMA																																																
V480	UAC	RYNINE	LIVER																																																	
V580	AAC	TORULOPSIS	UTILIS																																																	
V590	AAC	YEAST																																																		
V591	CAC	YEAST																																																		
V592	UAC	YEAST																																																		
V640	UAC	LUPTINUS	LUTEUS																																																	
V780	AAC	DROSOPHILA	MELANO.																																																	
V782	CAC	DROSOPHILA	MELANO.																																																	
V950	AAC	MOUSE	MYELOMA																																																	
V955	AAC	RAT	ASCIT. HEPATOMA																																																	
V956	MAC	RAT	LIVER																																																	

	45	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76
V203	C	G	G	G	U	C	A	U	G	G	U	C	A	U	A	G	C	C	U	A	A	C	A	C	C
V230	A	G	G	U	C	G	U	G	G	T	F	C	G	A	G	C	C	A	G	U	C	G	G	A	A
V235	G	G	G	U	C	G	G	G	T	F	C	G	A	G	C	C	C	G	U	C	A	C	C	C	C
V250	G	G	G	U	C	G	G	G	T	F	C	G	A	G	C	C	C	A	U	C	G	G	A	C	C
V251	G	G	G	U	C	G	G	G	T	F	C	G	A	G	C	C	C	A	U	C	G	G	A	C	C
V252	G	G	G	U	C	G	G	G	T	F	C	G	A	G	C	C	C	C	U	A	C	C	C	C	C
V365	A	A	G	U	C	J	A	C	G	G	T	F	C	G	A	G	C	C	G	U	A	G	C	C	C
V412	A	G	G	U	U	G	G	G	T	F	C	G	A	U	C	A	C	C	C	A	U	U	C	C	C
V456	A	G	A	A	U	U	G	G	A	A	C	A	A	F	A	A	A	U	G	A	C	C	A	C	C
V474	A	G	A	A	U	C	A	U	A	A	A	U	G	A	C	A	C	A	U	U	G	A	C	C	C
V475	A	G	A	A	U	C	A	U	A	A	A	U	G	A	C	A	C	A	U	U	G	A	C	C	C
V480	A	G	A	A	U	C	A	U	U	A	A	U	G	A	C	A	C	A	U	U	G	A	C	C	C
V580	A	C	G	U	C	C	S	C	A	G	T	F	C	G	A	U	C	C	U	G	G	C	G	A	A
V590	A	C	G	U	C	C	S	C	A	G	T	F	C	G	A	U	C	C	U	G	G	C	G	A	A
V591	A	G	G	U	C	C	S	C	A	G	T	F	C	G	A	U	C	C	U	G	G	U	G	G	A
V592	A	G	A	D	C	S	C	A	G	T	F	C	G	A	U	C	C	U	G	G	U	G	G	A	C
V640	A	G	G	U	C	C	C	G	G	T	F	C	G	A	G	U	C	C	C	G	G	C	G	A	C
V780	A	G	G	U	C	C	C	G	G	T	F	C	G	A	G	U	C	C	C	G	G	C	G	A	C
V781	A	G	G	U	C	C	C	G	G	T	F	C	G	A	G	U	C	C	C	G	G	C	G	A	C
V782	A	G	G	U	C	C	C	G	G	T	F	C	G	A	G	U	C	C	C	G	G	U	G	G	A
V950	A	G	G	U	C	C	S	C	C	G	J	F	C	G	A	A	C	C	C	G	G	C	G	G	A
V955	A	G	G	U	C	C	S	C	C	G	J	F	C	G	A	A	C	C	C	G	G	C	G	G	A
V956	A	G	G	U	C	C	S	C	C	G	J	F	C	G	A	A	C	C	C	G	G	C	G	G	A

K121/34	PARTIALLY C	K121/34	PARTIALLY C
K121/39	PARTIALLY U	K121/39	PARTIALLY U
K235/37	N IS A7 OR A9	K235/37	N IS A7 OR A9
K236/32	N IS A MODIFIED C	K236/32	N IS A MODIFIED C
K417/34	N IS A DERIVATIVE OF U	K417/34	N IS A DERIVATIVE OF U
K474/34	N IS A DERIVATIVE OF U	K474/34	N IS A DERIVATIVE OF U
K480/34	N IS A MODIFIED U	K480/34	N IS A MODIFIED U
K550/37	HARLOT	K550/37	HARLOT
K570/9	PARTIALLY G	K570/9	PARTIALLY G
K590/34	N IS A DERIVATIVE OF U9	K590/34	N IS A DERIVATIVE OF U9
K780/54	N IS PROBABLY T3	K780/54	N IS PROBABLY T3
K781/34	N IS PROBABLY U9	K781/34	N IS PROBABLY U9
K950/27	PARTIALLY MODIFIED	K950/27	PARTIALLY MODIFIED
K950/27	N IS PROBABLY A PRECURSOR OF A7	K950/27	N IS PROBABLY A PRECURSOR OF A7
K951/34	N IS A MODIFIED U	K951/34	N IS A MODIFIED U
K951/37	N IS EITHER A7 OR A DERIVATIVE OF A7	K951/37	N IS EITHER A7 OR A DERIVATIVE OF A7
K956/36	N IS PROBABLY A MODIFIED U	K956/36	N IS PROBABLY A MODIFIED U
M010/20	N IS PROBABLY A MODIFIED U	M010/20	N IS PROBABLY A MODIFIED U
M120/15	N IS A MODIFIED G	M120/15	N IS A MODIFIED G
M235/46	PARTIALLY G	M235/46	PARTIALLY G
M235/47	PARTIALLY U	M235/47	PARTIALLY U
M250/16	PARTIALLY U	M250/16	PARTIALLY U
M250/16	N IS PROBABLY U	M250/16	N IS PROBABLY U
M438/0	ELONGATION- AND INITIATION-trRNA MAY ORIGINATE	M438/0	ELONGATION- AND INITIATION-trRNA MAY ORIGINATE
M570/67	PARTIALLY U MODIFIED	M570/67	PARTIALLY U MODIFIED
M945/34	N IS A MODIFIED G	M945/34	N IS A MODIFIED G
X110/15	N IS A MODIFIED G	X110/15	N IS A MODIFIED G
X120/15	N IS A MODIFIED G	X120/15	N IS A MODIFIED G
X130/57	Z-YAMAIZUMI ET AL. (1982) NUCL. ACIDS SYMP. SE	X130/57	Z-YAMAIZUMI ET AL. (1982) NUCL. ACIDS SYMP. SE
X150/9	N IS A MODIFIED NUCLEOTIDE	X150/9	N IS A MODIFIED NUCLEOTIDE
X150/26	N IS PROBABLY G5	X150/26	N IS PROBABLY G5
X150/57	Z-YAMAIZUMI ET AL. (1982) NUCL. ACIDS SYMP. SE	X150/57	Z-YAMAIZUMI ET AL. (1982) NUCL. ACIDS SYMP. SE
X180/52	PARTIALLY G	X180/52	PARTIALLY G
X203/17	PARTIALLY U	X203/17	PARTIALLY U
X412/38	N IS PROBABLY F	X412/38	N IS PROBABLY F
X428/72	PARTIALLY U	X428/72	PARTIALLY U
X438/0	ELONGATION- AND INITIATION-trRNA MAY ORIGINATE	X438/0	ELONGATION- AND INITIATION-trRNA MAY ORIGINATE
X456/0	ELONGATION- AND INITIATION-trRNA MAY ORIGINATE	X456/0	ELONGATION- AND INITIATION-trRNA MAY ORIGINATE
X480/0	ELONGATION- AND INITIATION-trRNA MAY ORIGINATE	X480/0	ELONGATION- AND INITIATION-trRNA MAY ORIGINATE
X540/64	N IS PROBABLY G3	X540/64	N IS PROBABLY G3
X560/28	N IS A MODIFIED PYRIMIDINE	X560/28	N IS A MODIFIED PYRIMIDINE
X560/47	PARTIALLY U	X560/47	PARTIALLY U
T366/34	N IS PROBABLY A MODIFIED C	T366/34	N IS PROBABLY A MODIFIED C
L030/24	N IS A MODIFIED U	L030/24	N IS A MODIFIED U
L120/15	N IS A MODIFIED G	L120/15	N IS A MODIFIED G
L121/15	N IS A MODIFIED C	L121/15	N IS A MODIFIED C
L121/57	PARTIALLY A	L121/57	PARTIALLY A
L122/15	N IS A MODIFIED G	L122/15	N IS A MODIFIED G
L123/15	N IS A MODIFIED G	L123/15	N IS A MODIFIED G
L124/15	N IS A MODIFIED G	L124/15	N IS A MODIFIED G
L124/34	N IS PROBABLY V2	L124/34	N IS PROBABLY V2
L230/8	PROBABLY U4	L230/8	PROBABLY U4
L250/24	N IS A MODIFIED A	L250/24	N IS A MODIFIED A
L250/27	N IS A MODIFIED G	L250/27	N IS A MODIFIED G
L251/37	N IS A MODIFIED U	L251/37	N IS A MODIFIED U
L255/37	N IS A DERIVATIVE OF G	L255/37	N IS A DERIVATIVE OF G
L260/37	N IS A MODIFIED A	L260/37	N IS A MODIFIED A
L290/57	N IS A MODIFIED A	L290/57	N IS A MODIFIED A
L328/37	N IS A MODIFIED A	L328/37	N IS A MODIFIED A
L350/37	N IS A4 OR A5	L350/37	N IS A4 OR A5
L351/34	N IS A MODIFIED U	L351/34	N IS A MODIFIED U
L351/37	N IS A4 OR ZEATIN	L351/37	N IS A4 OR ZEATIN
L412/34	N IS A MODIFIED U	L412/34	N IS A MODIFIED U
L412/34	N IS A MODIFIED U	L412/34	N IS A MODIFIED U
L438/0	IMPORTED FROM CYTOPLASMA	L438/0	IMPORTED FROM CYTOPLASMA
L438/34	N IS AN MODIFIED NUCLEOTIDE	L438/34	N IS AN MODIFIED NUCLEOTIDE
L439/0	IMPORTED FROM CYTOPLASMA	L439/0	IMPORTED FROM CYTOPLASMA
L439/34	N IS AN MODIFIED NUCLEOTIDE	L439/34	N IS AN MODIFIED NUCLEOTIDE
L440/0	IMPORTED FROM CYTOPLASM	L440/0	IMPORTED FROM CYTOPLASM
L440/34	N IS A MODIFIED NUCLEOTIDE	L440/34	N IS A MODIFIED NUCLEOTIDE
L480/34	N IS A MODIFIED U	L480/34	N IS A MODIFIED U
L645/34	N IS A MODIFIED NUCLEOTIDE	L645/34	N IS A MODIFIED NUCLEOTIDE
L649/34	N IS A MODIFIED NUCLEOTIDE	L649/34	N IS A MODIFIED NUCLEOTIDE
L647/47	N IS A MODIFIED NUCLEOTIDE	L647/47	N IS A MODIFIED NUCLEOTIDE
L955/10	PARTIALLY C	L955/10	PARTIALLY C
L955/12	PARTIALLY G	L955/12	PARTIALLY G
L955/34	N IS A MODIFIED C	L955/34	N IS A MODIFIED C
L955/37	N IS A MODIFIED G	L955/37	N IS A MODIFIED G
L955/44	N IS A MODIFIED U	L955/44	N IS A MODIFIED U
L955/45	PARTIALLY U	L955/45	PARTIALLY U
L956/10	PARTIALLY G	L956/10	PARTIALLY G
L956/12	PARTIALLY C	L956/12	PARTIALLY C
L956/34	N IS A MODIFIED C	L956/34	N IS A MODIFIED C
L956/37	N IS A MODIFIED U	L956/37	N IS A MODIFIED U
L956/45	PARTIALLY U	L956/45	PARTIALLY U
L995/0	PRECURSOR	L995/0	PRECURSOR
L995/34	N IS A MODIFIED NUCLEOTIDE	L995/34	N IS A MODIFIED NUCLEOTIDE
K120/15	N IS A MODIFIED G	K120/15	N IS A MODIFIED G
K120/34	N IS A MODIFIED U	K120/34	N IS A MODIFIED U
K121/15	N IS A MODIFIED G	K121/15	N IS A MODIFIED G

X540/64	N IS A MODIFIED G
X570/64	N IS A MODIFIED A
X570/65	N IS A MODIFIED G
X580/64	N IS A DERIVATIVE OF G
X625/64	N IS A MODIFIED G
X640/28	PARTIALLY U
X640/49	N IS A MODIFIED C
X630/48	R.A.KOSKI ET AL. ONLY FIND C
X650/26	PARTIALLY G2
X995/26	PARTIALLY DEFICIENT
F120/15	N IS A MODIFIED G
F260/12	N IS MODIFIED U
F304/12	N IS PROBABLY A
F407/37	N IS A DERIVATIVE OF X
F430/37	N IS A4 OR A5
F510/49	N IS A MODIFIED C
F540/47	N IS A MODIFIED A
F560/37	N IS A DERIVATIVE OF Y
F575/9	N IS A MODIFIED G
F575/10	N IS PROBABLY G2
F575/25	N IS PROBABLY G2
F625/37	N IS PROBABLY G2
F630/37	N IS Y1 OR Y2
F630/37	N IS PROBABLY Y2
F770/32	PARTIALLY C
F770/47	PARTIALLY U
F770/48	MODIFICATION EITHER IN POSITION 48 OR 49
F771/32	PARTIALLY C
F771/47	PARTIALLY U
F771/48	MODIFICATION EITHER IN POSITION 48 OR 49
F807/32	PARTIALLY U
F830/37	N IS PROBABLY C
F950/32	PARTIALLY C
F950/34	PARTIALLY G
F950/37	N IS A UNDERMODIFIED WYBUTOSINE
F952/32	PARTIALLY C
F952/34	PARTIALLY G
F965/47	PARTIALLY U
F965/54	PARTIALLY U
F995/37	PARTIALLY MODIFIED
P020/8	PARTIALLY MODIFIED U
P020/34	N IS A MODIFIED U
P120/15	N IS A MODIFIED G
P120/34	PARTIALLY C
P120/38	PARTIALLY U
P120/48	PARTIALLY C
P120/55	PARTIALLY U
P121/55	PARTIALLY U
P122/15	PARTIALLY MODIFIED
P122/18	PARTIALLY C
P122/49	PARTIALLY C
P255/32	PARTIALLY U
P257/32	PARTIALLY U
P257/65	PARTIALLY U
P365/10	N IS PROBABLY A MODIFIED G
P365/34	N IS A MODIFIED U
P365/37	N IS A MODIFIED A
P570/35	N IS PROBABLY A MODIFIED U
P571/26	N IS C OR C5
P571/34	N IS A MODIFIED U
P571/48	N IS C OR C5
P571/49	N IS C OR C5
P580/34	K.OGAWA, T.KONDO, M.KAWAKAMI, S.TAKEMURA (1983) NUCL. ACIDS SYMP. SER. 12, 131-132
S020/34	N IS A MODIFIED U
S110/15	N IS A MODIFIED G
S120/15	N IS A MODIFIED G
S121/15	N IS A MODIFIED G
S122/15	N IS A MODIFIED G
S205/23	N IS A MODIFIED A
S254/33	Y.YAMADA, H.ISHIKURA (1975) BIOCHIM. BIOPHYS.
S481/49	PARTIALLY MODIFIED
S570/32	N IS PROBABLY C6
S570/34	N IS A MODIFIED U
S950/34	N IS PROBABLY A MODIFIED U
S970/0	CAN BE PHOSPHORYLATED TO PHOSPHOSERYL-TRNA; UG
S971/0	CAN BE PHOSPHORYLATED TO PHOSPHOSERYL-TRNA; UG
T075/34	N IS PROBABLY A
T080/37	N IS A MODIFIED A
T120/15	N IS A MODIFIED A
T121/15	PARTIALLY MODIFIED
T200/8	N IS A MODIFIED U
T201/8	N IS A MODIFIED U
T203/37	N IS A MODIFIED A
T235/20	PARTIALLY U
T590/48	PARTIALLY C
T591/48	PARTIALLY C
W120/15	N IS A MODIFIED G
W225/37	N IS A4 OR A5
W412/34	N IS A MODIFIED U
W412/37	N IS A MODIFIED A
W417/34	N IS A MODIFIED U
W417/37	N IS A4 OR A5
W438/37	N IS A4 OR A5
W438/47	N IS A MODIFIED NUCLEOTIDE

V955/48 PARTIALLY C
 V955/49 PARTIALLY
 V955/34 N T S I OR C

W674/74 N IS A MODIFIED U
 W680/74 N IS A MODIFIED U
 W670/75 PARTIALLY U
 W625/76 N T S A MODIFIED U
 W625/58 PARTIALLY A
 W970/77 PARTIALLY G
 W970/16 PARTIALLY C
 W970/34 PARTIALLY C
 W970/46 PARTIALLY G
 W970/47 PARTIALLY C
 W970/57 PARTIALLY A
 Y120/15 N IS A MODIFIED G
 Y120/16 PARTIALLY G
 Y120/75 N T S A MODIFIED U
 Y308/37 PARTIALLY A
 Y407/37 N IS A MODIFIED A
 Y412/37 N IS PROBABLY A4
 Y412/47 PROBABLY MODIFIED
 Y412/48 PROBABLY MODIFIED
 Y417/17 A PARTIALLY U
 Y438/18 PARTIALLY MODIFIED
 Y438/34 N IS A MODIFIED NUCLEOTIDE
 Y438/77 N IS A4 OR A5
 Y507/27 N IS A MODIFIED NUCLEOTIDE
 Y528/27 PARTIALLY G
 Y625/34 PARTIALLY G
 Y626/47 PROBABLY X
 Y655/47 PROBABLY X
 Y656/47 PROBABLY X
 Y780/34 PARTIALLY MODIFIED
 Y780/54 PARTIALLY MODIFIED
 Y955/17 PARTIALLY MODIFIED
 Y120/12 N IS A MODIFIED G
 Y263/23 PROBABLY U
 Y565/34 N IS A MODIFIED U
 Y656/37 PARTIALLY A
 Y591/10 PARTIALLY G
 Y592/34 N.YAMAMOTO ET AL. (1985) J. BIOCHEM. 97, 361-364
 Y780/20 PARTIALLY MODIFIED
 Y780/49 PROBABLY C5
 Y782/20 N IS A MODIFIED U
 Y782/26 PARTIALLY MODIFIED
 Y782/34 N IS A MODIFIED U
 W526/49 PARTIALLY MODIFIED
 W526/52 PARTIALLY C
 W550/35 PARTIALLY C
 W550/38 PARTIALLY C
 W550/43 PARTIALLY G
 W565/32 PARTIALLY C
 W565/38 PARTIALLY C

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1123	R. GUPTA (1986) SYSTEM. APPL. MICROBIOL. 7, 102-105	L481	S.-H. CHANG, S.-K. KO, E. HAN, T.S. CHEN, N.R. MITLETSER, D. CO
1124	R. GUPTA (1986) SYSTEM. APPL. MICROBIOL. 7, 102-105	L481	BIOCHEM. BIOPHYS. RES. COMM. 51, 951-955 (1975)
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L290	B. LARUE, N. NEMHUSE, K. NITOGHOSIAN, R.J. CEDERGREEN (1981)	L955	E. RANDEPATH, R.C. GUPTA, H.P. MORRIS, K. RANDEPATH
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L328	M. L. OSORIO-ALMEIDA ET AL. (1980)	L976	I.G. VASILJEVA, M.A. TUKALO, T.A. KRIVILYI, G.K. MA
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	AMINOACYL STEM	D STEM	D LOOP	D STEM	ANTIC. STEM	ANTIC. LOOP	ANTIC. STEM	ANTIC. LOOP	ANTIC. STEM
	1 2 3 4 5 6 7 8 9 10 12 14 16 A 19 A	11 13 15 17 18 20 B	21 23 25 27 29 31 33 35 37 39 41 43	22 24 26 28 30 32 34 36 38 40 42					
A L A N I N E									
A040	UGC PHAGE T5 G G G C G G A A T A G T G T C A G C G G G **** ** *****				A G C A C A C C A G A C T T G C A A T C T G G				
A108	UGC ARCHAELOBUS FULG. G G G C T C G T A G C T C A G C G G G **** *****				A G A G C G C C G C C T T G C G A G G C G G				
A110	UGC HALOBACTERIUM CUT. G G G C C C A T A G C T C A G T G G T **** *****				A G A G T G C C T C C T T G C A A G G A G G				
A113	UGC HALOBACTERIUM HAL. G G G C C C A T A G C T C A G T G G T **** *****				A G A G T G C C T C C T T G C A A G G A G G				
A135	UGC METHANOBAC. FORMI. G G G C C C G T A G C T C A G A C T G G G **** *****				A G A G C G C C G C C C T T G C A A G G C G G				
A140	UGC METHANOBAC. THERM. G G G C C C G T A G C T C A G A C T G G G **** *****				A G A G C G C C G C C C T T G C A A G G C G G				
A145	UGC METHANOCOC. VANI. G G G C C C G T A G C T C A G T T G G G **** *****				A G A G C G C T G C C C T T G C A A G G C A G				
A149	UGC METHANOSPIR. HUNG. G G G C T C G T A G C T C A G C T G G A **** *****				A G A G C G C G G C G T T G C A C G C C G				
A185	UGC THERMOCOCCUS CELER G G G C C C C T A G C T C A G C C T G G G **** *****				A G A G C G T C G G C T T G C A A G C C G A				
A190	UGC THERMOPROT. TENAX G G G C C G G T A G T C T A G C G G A **** *****				A G A C G C C G C C T T G C G C G C G G G				
A191	CGC THERMOPROT. TENAX G G G C C G G T A G T C T A G C G G A **** *****				A G A C G C C G C C T T G C G C G C G G G				
A203	UGC MYCOPLASMA MYCOID. G G G C C C T T A G C T C A G C T G G G **** *****				A G A G C A C C T G C C T T G C A C G C A G G				
A206	UGC SPIROPLASMA MELIF. G G G C C C G T A G C T C A G C T G G G **** *****				A G A G C A C C T G C C T T G C A C G C A G G				
A235	UGC BACILLUS SUBTILIS G G G C C C T T A G C T C A G C T G G G **** *****				A G A G C G C C T G C C T T G C A C G C A G G				
A236	UGC BACILLUS SUBTILIS G G G C C T A G C T C A G C T G G G **** *****				A G A G C G C C T G C C T T G C A C G C A G G				
A237	UGC BACILLUS SUBTILIS G G G C C C T A G C T C A G C T G G G **** *****				A G A G C G C C T G C C T T G C A C G C A G G				
A238	UGC BACILLUS SUBTILIS G G G C C C T A G C T C A G C T G G G **** *****				A G A G C G C C T G C C T T G C A C G C A G G				
A243	UGC THIOBACILLUS FERRO. G G G G C T G T A G C T C A G T T G G G **** *****				A G A G C A C C T G C C T T T G C A G C A G G				
A250	UGC E. COLI G G G G C T A T A G C T C A G C T G G G **** *****				A G A G C G C C T G C C T T T G C A C G C A G G				
A260	UGC PSEUDOMONAS AER. G G G G C C C T A G C T C A G C T G G G **** *****				A G A G C G C C T G C C T T T G C A C G C A G G				
A270	UGC CAMPYLOBAC. JEJUNI G G G G C A T A G C T C A G C T G G G **** ** *****				A G A G C G C C T G C C T T T G C A C G C A G G				

EXTRA ARM		TF STEM	TF LOOP	TF STEM	AMINOACYL STEM
45 47 B D F H J L N P 44 46 A C E G I K M O		49 51 53 50 52 54 56 58 60	55 57 59 54 56 58 60	61 63 65 62 64 66 68 70 72	67 69 71 73 75 76
ALANINE					
A040	T A G	G G A G G G T	T C G A G T	C C C T C T	T T G T C C A C C A
A108	A G G C	C G C G G G T	T C A A A T	C C C G C	C G A G T C C A
A110	A T G C	C T G G G T	T C G A A T	C C C A G T	G G G T C C A
A113	A T G C	C T G G G T	T G G A A T	C C C A G T	G G G T C C A
A135	A G G C	C C G C C T	T C A A A T	C C C G G T	G G G T C C A
A140	A G G C	C C G G G T	T C A A A T	C C C G G T	G G G T C C A
A145	A G G C	C G T G G G T	T C A A A T	C C C G C	G G G T C C A
A149	A G G C	C T G G G G T	T C A A A T	C C C C A	C G G G T C C A
A185	A G G C	C C G G G T	T C G A A T	C C C G G C	C G G T C C A C C A
A190	A G A T	C C G G G T	T C G A A T	C C C G G C	C G G T C C A
A191	A G A T	C C G G G T	T C G A A T	C C C G G C	C G G T C C A
A203	G G G T	C G A C G G T	T C G A T C	C C G T	A G G G T C C A C C A
A206	G G G T	C G A C G G T	T C G A T C	C C G T	C G G G T C C A C C A
A235	A G G T	C A G C G G T	T C G A T C	C C G C T	A G G C T C C A
A236	A G G T	C A G C G G T	T C G A T C	C C G C T	A G G C T C C A C C A
A237	A G G T	C A G C G G T	T C G A T C	C C G C T	A G G C T C C A
A238	A G G T	C A G C G G T	T C G A T C	C C G C T	A G G C T C C A C C A
A243	G G G T	C A T C G G T	T C G A G A	C C G G T	C A G C T C C A C C A
A250	A G G T	C T G C G G T	T C G A T C	C C G C A	T A G C T C C A C C A
A260	A G G T	C A G G A G T	T C G A T C	C T C C T	T G G C T C C A C C A
A270	A G G T	C A G C G G T	T C G A T C	C C G C T	A T C T C C A C C A

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	27	29	31	33	35	37	39	41	43																						
A272	UGC CAULOBACTER CRES.	G	G	G	G	C	C	C	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	G	C	C	T	G	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G									
A290	UGC ANACYSTIS NIDULANS	G	G	G	G	G	G	T	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	T	G	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G									
A295	UGC CYANOPHORA PARAD.	G	G	G	G	G	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	T	G	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G									
A297	UGC PYLAIELLA LITTORA.	G	G	G	G	G	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	T	G	C	T	T	G	C	A	T	A	G	C	A	G	C	A	G	C	A	G								
A300	UGC CHLAMYDOMONAS REINH.	G	G	G	G	G	A	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	T	G	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G								
A302	UGC CHLORELLA ELLIPSO.	G	G	G	A	G	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	T	G	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G									
A304	UGC EUGLENA GRACILLIS	G	G	G	G	T	A	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	T	G	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G								
A305	UGC EUGLENA GRACILLIS	G	G	G	G	T	A	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	T	G	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G								
A310	UGC MARCHANTIA POLYM.	G	G	G	G	A	T	A	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G									
A320	UGC ZEA MAYS	G	G	G	G	A	T	A	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	C	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G								
A327	UGC GLYCINE MAX	G	G	G	G	A	T	A	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G	C	A	G						
A335	UGC LECTIANA TABACUM	G	G	G	G	A	T	A	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G	C	A	G						
A365	UGC SPINACIA OLERACEA	G	G	G	G	A	T	A	T	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	A	G	C	C	C	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G	C	A	G						
A403	UGC TRYPANOSOMA BRUCEI	A	A	A	T	A	T	A	T	A	T	A	T	A	T	T	T	C	T	C	T	A	A	G	T	G	C	A	A	T	G	C	T	T	C	G	A	A	A	T	G	C	T	T	C	G	A	A	T	G	C	A	A			
A404	UGC TRYPANOSOMA BRUCEI	A	A	A	T	A	T	A	T	A	T	A	T	A	T	T	T	C	T	A	A	A	A	T	G	A	A	T	G	C	A	A	T	G	C	T	T	C	G	A	A	A	T	G	C	T	T	C	G	A	A	T	G	C	A	A
A410	UGC ASPERGILLUS NIDUL.	G	G	G	G	A	T	A	T	A	T	A	G	C	T	T	A	A	C	T	G	G	T	A	A	A	A	C	G	C	G	A	T	T	T	G	C	A	T	A	T	C	G	T	T	G	C	A	T	A	T	C	G	T	A	T
A412	UGC NEUROSPORA CRASSA	G	G	G	G	A	T	A	T	A	T	A	G	C	T	T	A	A	A	T	G	G	T	A	A	A	A	A	G	C	A	T	C	T	T	G	C	T	A	T	T	G	C	A	T	T	C	G	T	A	T	C	G	T	A	T
A417	UGC SACCHAROMYCES CER.	G	G	G	G	A	T	A	T	A	T	A	G	C	T	T	A	A	A	T	T	G	G	T	A	A	A	A	A	A	T	T	G	C	A	T	T	G	C	A	T	T	G	C	A	T	T	C	G	T	A	T				
A425	UGC TORULOPSIS GLAB.	G	G	G	G	A	T	A	T	A	T	A	G	C	T	T	A	A	A	T	T	G	G	T	A	A	A	A	A	A	T	T	G	C	A	T	T	G	C	A	T	T	G	C	A	T	T	C	G	T	A	T				
A450	UGC ASCARIS SUUM	G	G	G	G	T	A	T	A	T	A	T	A	G	C	T	T	A	A	G	T	T	A	A	A	A	A	A	T	T	G	C	A	T	T	G	C	A	A	G	C	A	G	C	A	G	C	A	G	C	A	G				
A456	UGC AEDES ALBOPICTUS	A	G	G	G	A	T	A	T	A	T	A	T	A	T	A	T	T	A	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A					
A459	UGC DROSOPHILA YAKUBA	A	G	G	G	T	A	T	A	T	A	T	A	T	A	T	T	A	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A					
A464	UGC PARACENTROTUS LIV.	G	G	G	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A				

	45	47	B	D	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76												
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76												
A272	T	G	T								C	G	T	C	G	G	A	T	C	G	G	A	T	C	G	G	T	C	C	A	C	C	A				
A290	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	C	C	T	C	C	A	C	C	A
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A297	A	C	G	T							C	A	G	C	A	G	T	C	G	A	G	T	C	T	G	C	T	A	T	C	T	C	C	A	C	C	A
A300	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	T	C	T	C	C	A	C	C	A
A302	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	T	C	T	C	C	A	C	C	A
A304	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	T	C	T	C	C	A	C	C	A
A305	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	T	C	T	C	C	A	C	C	A
A310	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	T	C	T	C	C	A	C	C	A
A320	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	T	C	T	C	C	A	C	C	A
A327	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	T	C	T	C	C	A	C	C	A
A335	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	T	C	T	C	C	A	C	C	A
A365	A	T	G	T							C	A	G	C	G	G	T	C	G	A	G	T	C	C	G	G	T	A	T	C	T	C	C	A	C	C	A
A403	T	T	G	A	A	A	A	A	A	T	T	A	A	A	T	A	A	T	T	A	A	T	T	A	A	T	T	T	T	T	T	T	T	T	T	T	
A404	T	T	G	A	A	A	A	A	A	T	T	A	A	A	T	A	A	T	T	A	A	T	T	A	A	T	T	T	T	T	T	T	T	T	T	T	
A410	T	A	T								T	C	A	G	G	A	T	C	G	A	G	T	C	T	G	A	T	A	A	C	T	C	C	A	C	C	A
A412	T	T	G								T	C	A	G	G	T	C	A	A	G	T	C	C	T	T	G	A	T	C	T	C	C	A	C	C	A	
A417	T	A	A								T	A	G	A	G	T	C	A	A	G	T	C	C	A	T	C	A	T	T	C	C	A	C	C	A	C	C
A425	A	G	A								T	A	G	A	G	T	C	A	A	G	T	C	C	A	T	C	A	T	T	C	C	A	C	C	A	C	C
A450	T	G	A	T	T	C					T	A	G	A	G	T	C	A	A	G	T	C	C	A	T	C	A	T	T	C	C	A	C	C	A	C	C
A456	A	A	G								T	A	T	G	A	A	T	T	A	A	T	T	C	A	A	T	T	A	T	T	A	C	C	T	T	A	
A459	A	A	G								T	A	T	G	A	A	T	T	A	A	T	T	C	A	A	T	T	A	T	T	A	C	C	T	T	A	
A464	A	A	T								C	T	A	A	G	T	A	A	T	T	A	T	T	C	A	A	T	T	A	T	T	A	C	C	T	T	A

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42							
R255	CCG	SALMONELLA	TYPHI.	G	G	C	C	C	G	T	A	G	C	T	A	G	C	T	G	G	A	T	A	G	A	G	C	G	C	T	G	C	C	T	C	C	G	G	A	G	G	C	A	G					
R259	CCG	AEROMONAS	HYDROPH.	G	C	C	C	G	T	A	G	C	T	A	G	C	T	A	G	C	T	G	G	A	T	A	G	A	G	C	G	C	T	C	C	G	G	A	G	G	C	A	G	G	C	A	G		
R304	ACG	EUGLENA	GRACILLIS	G	G	C	T	T	A	G	C	T	C	A	G	T	G	G	A	C	T	A	G	A	G	A	G	C	G	C	T	C	C	G	G	A	G	G	C	C	T	C	C	G	G	A	G		
R310	ACG	MARCHANTIA	POLYM.	G	G	C	T	T	A	G	C	T	C	A	G	T	G	G	A	C	T	A	G	A	G	A	G	C	G	C	T	C	C	G	G	A	G	G	C	C	T	C	C	G	G	A	G		
R311	CCG	MARCHANTIA	POLYM.	G	G	G	T	T	G	T	A	G	C	T	C	A	G	T	G	G	A	C	T	A	G	A	G	C	G	C	T	C	C	G	G	A	G	G	T	T	C	C	G	A	T	C	A	T	
R312	UCU	MARCHANTIA	POLYM.	G	C	G	T	C	C	A	T	G	T	C	T	A	A	G	G	A	T	A	G	A	G	A	G	A	G	G	T	T	T	C	T	A	A	C	C	T	C	T	A	A	C	C	T		
R315	UCU	TRITICUM	AESTIVUM	G	C	G	T	C	C	A	T	G	T	C	T	A	A	G	G	A	T	A	G	A	G	A	G	A	G	G	T	T	T	C	T	A	A	C	C	T	C	T	A	A	C	C	T		
R320	ACG	ZELONIA	CHLORO	G	G	C	T	G	T	A	G	C	T	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	G	T	T	C	C	G	G	A	G	G	C	T	C	A	G	C	C	G	
R335	ACG	NICOTIANA	TABACUM	G	G	C	C	T	G	T	A	G	C	T	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	G	T	T	C	C	G	G	A	G	G	C	T	C	A	G	C	C	G
R336	UCU	NICOTIANA	TABACUM	G	C	G	T	C	C	A	T	G	T	C	T	A	A	G	G	A	T	A	G	A	G	A	G	A	G	A	G	G	T	T	C	C	G	G	A	G	G	C	T	C	A	G	C	C	G
R345	ACG	PELARGONIUM	ZONALE	G	G	C	C	T	G	T	A	G	C	T	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	G	T	T	C	C	G	G	A	G	G	C	T	C	A	G	C	C	G
R355	ACG	PISUM	SATIVUM	G	G	C	C	T	C	G	T	A	G	C	T	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	G	T	T	C	C	G	G	A	G	G	C	T	C	A	G	C	C	G	
R356	UCU	PISUM	SATIVUM	G	C	T	C	C	A	T	G	T	C	T	A	A	G	G	A	T	A	G	A	G	A	G	A	G	A	G	G	T	T	C	C	G	G	A	G	G	C	T	C	A	G	C	C	G	
R365	UCU	SPINACIA	OLERACEA	G	C	G	T	C	C	A	T	G	T	C	T	A	A	G	G	A	T	A	G	A	G	A	G	A	G	A	G	G	T	T	C	C	G	G	A	G	G	C	T	C	A	G	C	C	G
R370	ACG	SPIRODELA	OLIGORH.	G	G	C	C	T	G	T	A	G	C	T	C	A	G	A	G	A	G	A	G	A	G	A	G	A	G	A	G	G	T	T	C	C	G	G	A	G	G	C	T	C	A	G	C	C	G
R371	UCU	SPIRODELA	OLIGORH.	G	C	G	T	C	C	A	T	G	T	C	T	A	A	G	G	A	T	A	G	A	G	A	G	A	G	A	G	G	T	T	C	C	G	G	A	G	G	C	T	C	A	G	C	C	G
R410	UCU	ASPERGILLUS	NIDUL.	T	T	C	T	A	T	A	G	C	T	C	A	A	G	A	G	A	T	A	G	A	G	A	G	A	G	A	A	A	T	A	A	C	T	T	C	T	A	A	T	A	T	T	T	T	
R412	UCU	NEUROSPORA	CRASSA	T	T	C	C	A	A	T	A	G	C	T	C	A	A	G	A	G	A	T	A	G	A	G	A	G	A	G	A	G	A	T	A	A	C	T	T	C	T	A	A	T	A	T	T	T	
R417	UCU	SACCHAROMYCES	CER.	G	C	T	C	T	A	G	C	T	C	A	A	G	A	G	A	T	A	A	G	A	G	A	G	A	G	A	A	A	C	T	T	C	T	A	A	T	A	T	A	T	A	T	A		
R418	ACG	SACCHAROMYCES	CER.	A	T	A	C	T	T	A	A	T	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	T	A	G	A	T	A	A	C	T	T	A	A	T	A	T	A	T	A		
R425	UCU	TORULOPSIS	GLAB.	G	C	T	C	T	C	T	A	G	C	T	T	A	A	G	A	G	A	T	A	A	G	A	G	A	T	A	A	T	A	A	C	T	T	A	A	T	A	T	A	T	A	T	A		
R450	ACG	ASCARIS	SUUM	A	G	A	C	G	T	A	A	T	A	G	A	T	A	A	G	A	T	A	A	G	A	G	A	T	A	A	T	A	A	C	T	A	T	A	T	A	T	A	T	A	T	A	T		
R456	UCG	AEDIS	ALBOPICTUS	A	A	A	T	A	T	A	G	C	G	A	T	T	A	A	T	A	A	G	A	T	A	A	G	A	T	A	A	T	A	A	C	T	A	T	A	T	A	T	A	T	A	T	A		

	45 47 B D F H J L M P	49 51 53	55 57 59	61 63 65	67 69 71	73 75
	44 46 A C E G I K N O	48	50 52	54 56 58 60	62 64 66 68 70 72 74 76	
R255	A G G T	C T C A G G G	T C G A A T	C C T G T	C G G G C G T	A C C A
R259	A G G T	C A C A G G T	T C G A A T	C C T G T	C G G G C G C	A C C A
R304	G A G T	A G G G C C T	T C G A A T	C C C T C T T	G C C C G	
R310	G T G T	C G G G G G T	T C G A A T	C C C T C C T	G C C C A	
R311	A A G T	C A A G G G T	T C G A A T	C C C T C T A	A C C C T	
R312	C A G	T A T A G G T	T C G A A T	C C T A T T G G A	C G T A	
R315	T G G	T A T A G G T	T C A A A T	C C T A T T G G A	C G C A	
R320	G T G T	C G G G G G T	T C G A A T	C C C T C C T	C C C C A	
R335	G T G T	C G G G G T	T C G A A T	C C C T C C T	C G C C C A	
R336	T G G	T A T A G G T	T C A A A T	C C T A T T G G A	C G C A	
R345	A T G T	C G G G G G T	T C G A A T	C C C T C C T	A G C C A	
R355	G T G T	C G G G G G T	T C G A A T	C C C T C C T	C G C C C A	
R356	T G G	T A T A G G T	T C A A A T	C C T A T T G G A	C G C A	
R365	T G G	T A T A G G T	T C A A A A	C C T A T T G G A	C G C A	
R370	G T G T	C G G G G G T	T C G A A T	C C C T C C T	C G C C C A	
R371	T G G	T A T A G G T	T C A A G T	C C T A T T G G A	C G C A	
R410	T G A	T C T A A G T	T C G A G T	C T T A G A T	A A G A A G	
R412	C G A	T T T T A G T	T C G A A T	C T A A A T	T G A G A A T	
R417	A T A T	T C C A T G T	T C A A A T	C A T G G A	A A G C T A	
R418	T T A	T A T A G G T	T C A A A T	C C T A T A	A G A T A T	
R425	A T A T	T C C A T G T	T C A A A T	C A T G G A	G A G A G T A	
R450	A A G A G A G					
R456	C T T	A G G T G A A A T	T C A C C C	C A T A T T T		

	45	47	B	D	F	H	J	K	L	M	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75		
	46	48	A	C	E	G	I	K	M	O			50	52	54	56	58	60	62	64	66	68	70	72	74	76		
N335	TGGT		C	G	T	A	G	G	T	T	C	G	T	A	A	C	C	T	A	C	T	T	G	G	G	A	G	
N355	TGGT		C	G	T	A	G	G	T	T	C	A	A	A	T	C	C	T	A	T	T	T	G	C	G	A	G	
N370	TGGT		C	G	T	A	G	G	T	T	C	G	G	A	A	T	C	C	T	A	C	T	T	G	G	G	A	G
M410	TG A		T	A	G	A	T	G	T	T	C	A	A	T	C	A	T	C	A	T	T	A	A	G	G	C	T	
M417	A T A		G	A	T	G	G	T	T	C	A	A	T	C	C	T	T	A	A	G	G	C	G					
M425	A T A		C	A	T	A	G	G	T	T	C	A	A	T	C	C	T	T	G	A	A	G	A	C	G			
M434	TGGT		C	G	T	A	G	G	T	T	C	A	A	T	C	C	T	A	T	T	G	A	A	G	A	C	G	
M450	GGT G A T																											
M456	T A A T		T	G	A	A	T	T	T	A	T	T	C	A	A	T	T	A	A	G								
M459	T A A T		T	G	A	G	T	A	A	C	T	C	A	A	T	T	A	A	G									
M464	A A A T		A	G	T	A	G	G	T	A	A	A	A	A	A	C	T	A	T	C	T	C	C	G	A	G		
M466	A A A T		A	A	T	A	G	G	T	A	A	A	T	A	A	C	T	A	T	C	C	C	A	G				
M468	A T G T		T	G	C	G	G	A	T	C	G	N	G	C	C	G	T	T	C	T	A	G						
M470	A T A T		A	C	G	G	A	T	C	G	A	G	G	C	C	G	T	C	A	T	C	A	T	A	G			
M474	T T T T		C	G	T	A	G	G	T	T	G	A	T	C	T	C	A	A	T	C	T	A	G					
M475	T T T T		C	G	T	A	G	G	T	T	G	A	A	T	C	T	C	A	A	T	C	T	A	G				
M476	T T T T		C	G	T	A	G	G	T	T	G	A	A	T	C	T	C	A	A	T	C	T	A	G				
M477	T T T T		C	G	T	A	G	G	T	T	G	A	A	T	C	T	C	A	A	T	C	T	A	G				
M480	G T T T		C	G	T	G	G	G	A	T	T	A	A	T	C	T	G	C	A	A	T	C	A	G				
M494	T G T T		T	G	G	T	T	A	A	G	T	C	C	A	T	T	G	G	C	T	A	G						
M660	T G G T		C	G	T	A	G	G	T	T	C	G	A	A	T	C	T	A	T	T	G	G	G	A	G			
N780	A G G T		T	G	G	T	G	G	A	T	C	G	A	G	T	C	A	C	C	C	G	G	G	G	C	G		
N830	A G G T		T	G	G	T	G	G	A	T	C	G	A	G	T	C	A	C	C	C	A	G	G	A	G	C	G	

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43				
D452	GUC CAENORHABDI. ELEG.	A	A	G	T	T	T	A	G	T	A	A	A	A	A	A	A	A	A	A	A	G	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T		
D454	GUC LOCUSTA MIGRATORIA MITO	A	A	G	G	A	G	T	A	G	T	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D457	GUC DROSOPHILA MELANO. MITO	A	A	A	A	A	A	T	A	G	T	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T
D459	GUC DROSOPHILA YAKUBA MITO	A	A	A	A	A	A	T	A	G	T	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T
D464	GUC PLOCEUS CENTROTUS LIV. MITO	A	G	A	G	T	C	T	A	G	T	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D466	GUC STRONGYLOCEUS PURP. MITO	A	G	A	G	G	C	C	T	A	G	T	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D468	GUC XENOPUS LAEVIS MITO	G	A	G	A	T	G	T	A	G	T	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D474	GUC RAT MITO	G	A	G	A	T	A	G	T	A	A	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D475	GUC RAT MITO	G	A	G	A	T	A	G	T	A	A	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D477	GUC MOUSE MITO	A	A	G	A	A	A	T	A	G	T	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T
D480	GUC BOVINE MITO	G	A	G	T	T	A	G	T	A	A	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T
D494	GUC HUMAN MITO	A	A	G	T	A	T	A	G	A	A	A	A	A	A	A	A	A	A	A	A	T	A	C	A	T	T	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D570	GUC SACCHAROMYCES CER. MITO	T	C	C	G	T	G	A	T	A	G	T	T	T	T	T	T	T	T	T	T	G	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T		
D575	GUC SCHIZOSACCHA. POM. MITO	T	C	T	C	T	T	A	G	T	A	T	A	G	G	G	G	G	G	G	G	T	A	C	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D635	GUC GLYCINE MAX MITO	G	T	C	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D700	GUC CAENORHABDI. ELEG. MITO	T	C	C	T	C	G	T	A	G	T	A	T	A	G	T	A	T	A	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D780	GUC DROSOPHILA MELANO. MITO	T	C	C	T	C	G	A	T	A	G	T	A	T	A	G	T	A	T	A	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D950	GUC MOUSE MITO	T	C	C	T	C	G	T	A	G	T	A	T	A	G	T	A	T	A	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
D955	GUC RAT MITO	T	C	C	T	C	G	T	A	G	T	A	T	A	G	T	A	T	A	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T		
D956	GUC RAT MITO	T	C	C	T	C	G	T	A	G	T	A	T	A	G	T	A	T	A	G	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T		

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44
C Y S T E I N E																																												
C110	GCA	HALOBACTERIUM	CUT.																																									
C120	GCA	HALOBACTERIUM	VOL.																																									
C206	GCA	SPIROPLASMA	MELIF.																																									
C235	GCA	BACILLUS	SUBTILIS																																									
C304	GCA	EUGLENA	GRACILIS																																									
C310	GCA	MICROSPORIDIUM	PHOENICIA POLYM.																																									
C315	GCA	TRITICUM	AESTIVUM																																									
C320	GCA	ZEA	MAYS																																									
C335	GCA	NICOTIANA	TABACUM																																									
C365	GCA	SPINACIA	OLERACEA																																									
C410	GCA	ASPERGILLUS	NIDUL.																																									
C411	GCA	ASPERGILLUS	NIDUL.																																									
C412	GCA	NEUROSPORA	CRASSA																																									
C417	GCA	SACCHAROMYCES	CER.																																									
C425	GCA	TORULOPSIS	GLAB.																																									
C445	GCA	ZEA	MAYS																																									
C450	GCA	ASCARIS	SUUM																																									
C452	GCA	CAENORHABDI.	ELEG.																																									
C457	GCA	DROSOPHILA	MELANO.																																									
C459	GCA	DROSOPHILA	YAKUBA																																									
C464	GCA	PARACENTROTUS	LIV.																																									

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42									
C466	GCA	STRONGYLOGEN. PURP. MITO	G G T	T	I	G	T	A	G	T	G	T	A	T	C	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G							
C468	GCA	XEMPOUS LAEVIS MITO	A	A	G	C	C	T	G	C	G	T	G	T	G	A	C	A	T	T	C	C	A	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G				
C470	GCA	RAMA CATESBETANA MITO	G	G	C	T	G	G	T	G	T	C	A	C	A	T	T	C	G	G	G	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G		
C474	GCA	RAT MITO	A	G	C	T	T	A	G	T	G	A	T	T	C	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	
C475	GCA	RAT MITO	A	G	C	T	T	A	G	T	G	A	T	T	C	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	
C477	GCA	MUSE MITO	G	G	T	T	A	G	T	G	A	T	T	C	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G		
C480	GCA	BOVINE MITO	A	G	C	C	T	G	T	G	A	T	T	C	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G		
C494	GCA	HUMAN MITO	A	G	C	T	C	G	A	G	T	G	A	T	T	C	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G
C570	GCA	SACCHAROMYCES CER. MITO	G	C	T	C	G	T	G	C	C	A	G	T	G	G	T	A	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G	G	A	T	T	C	G
G L U T A M I N E																																																			
0020	UUG	PHAGE T4	I	G	G	G	A	T	A	G	C	C	A	G	T	G	G	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	
0040	UUG	PHAGE T5	T	G	G	G	G	A	T	A	G	C	T	A	G	C	T	A	A	G	C	T	A	A	G	C	T	A	A	G	C	T	A	A	G	C	T	A	A	G	C	T	A	A	G	C	T	A	A		
0145	UUG	METHANOCOCC. VANI.	A	G	C	C	A	G	T	A	G	T	G	T	A	G	T	G	G	C	C	A	T	C	A	T	C	A	T	A	G	G	T	A	G	G	C	T	T	G	G	A	C	C	T	G	G	A	C	C	
0200	UUG	MYCOPLASMA CAPRIC.	T	G	G	G	C	T	A	G	C	C	A	G	C	G	G	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A		
0235	UUG	BACILLUS SUBTILIS	T	G	G	G	C	T	A	G	C	C	A	G	C	G	G	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A		
0250	UUG	E. COLI	T	G	G	G	T	A	C	G	C	A	G	C	G	G	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A			
0251	CUG	E. COLI	T	G	G	G	T	A	C	G	C	A	G	C	G	G	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A			
0304	UUG	EUGLENA GRACILIS CHLORO	T	G	A	G	G	C	C	A	G	T	G	G	C	A	G	T	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A			
0310	UUG	MARCHANTIA POLYM. CHLORO	T	G	G	G	C	C	T	A	G	C	C	A	G	T	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G				
0335	UUG	NICOTIANA TABACUM CHLORO	T	G	G	G	C	C	T	A	G	C	C	A	G	T	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G	G	C	A	T	A	G				
0410	UUG	ASPERGILLUS NIDUL. MITO	T	A	T	G	T	T	G	T	C	A	T	A	T	C	G	A	T	A	T	C	G	A	T	A	T	C	A	T	A	T	C	A	T	A	T	C	A	T	A	T	C	A	T	A	T				
0417	UUG	SACCHAROMYCES CER. MITO	I	G	A	T	C	C	T	A	G	C	A	T	A	G	T	A	C	A	T	A	G	T	A	C	A	T	A	T	C	A	T	A	T	C	A	T	A	T	C	A	T	A	T	C	A				
0425	UUG	ICHELLOPSIS GLAB. MITO	I	G	A	T	C	C	T	A	G	C	A	T	A	G	T	A	C	A	T	A	G	T	A	C	A	T	A	T	C	A	T	A	T	C	A	T	A	T	C	A	T	A	T	C	A				

		45 47 B D F H J L N P	49 51 53	55 57 59	61 63 65	67 69 71	73 75
		44 46 A C E G I K M O	48	50 52 54 56 58 60	62 64 66 68	70 72	74 76
C466	GA	TGCAAC	TAGGAC	GIIGCC	CAAAGCT		
C468	AGA	AGCAA	CGAAG	GTIIGC	CGGGCTTC		
C470	AGA	TGAGT	ATAATA	CCTGC	CGGGCTT		
C474	AGG	TGTAGA	GAACTCCT	CTACT	TAAAGGCTT		
C475	AGG	TGTAGA	GAACTCCT	CTACT	TAAAGGCTT		
C477	AGG	TGTAGA	GAAAT	CTACT	TAAAGCTT		
C480	AGA	AGAGCT	TCAA	TCTIGC	CGGGCTT		
C494	AGA	AGAGCT	TCAA	ACTIGC	CGGGGCTT		
C570	TGGT	CCTTAG	TCGATCTGAG	TGCGAGCT			
Q020	GATG	CAAAGG	TCGAGT	CCTTT	ATGCCAAG		
Q040	GAT	CATTGG	TCAAAT	CCAA	ATCCCTGCCA		
Q145	GGA	CGGGT	TCGAAT	CCGG	CTGGGCTA		
Q200	CATG	CCCGG	TCGAAT	CCTGC	TAGCCCAACA		
Q235	CATG	CATTGG	TCGAAT	CCAGC	TAGCCCAAG		
Q250	CATT	CCTGG	TCGAAT	CCAGG	TACCAGCCA		
Q251	CATT	CAGGG	TCGAAT	CCTCG	TACCAGCCA		
Q304	CATT	CAGG	TCGAAT	CCTCC	CGCCICAG		
Q310	TATT	CAGGG	TCGAAT	CCTTC	CGTCCCAAG		
Q335	TATT	CAGGG	TCGAAT	CCTTC	CGTCCCAAG		
Q410	TAT	TGGTG	TCGAGT	CGCCC	CAACATAA		
Q417	AGT	CCTTGT	TCGAAT	CAAAG	CGATTCAA		
Q425	AGT	GTTTGT	TCGAAT	CAACC	CGATTCAA		

GLUTAMINE


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45 47 B D F H J L M P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76
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0442 C A T G C A A G G G T T C G A A T C C T T T T A C T C C A G
0450 A G G G T T G T A G T A G T A G T A G T A G T A G T A G
0459 T A G T A G T A G T A G T A G T A G T A G T A G T A G
0461 T A G T A G T A G T A G T A G T A G T A G T A G T A G
0464 T G T G G G T C A A T T C C C A T C C C T T A A
0466 C T T G G G T C C A T T C C C A C T C C T T A A
0468 T G C A G G T T C A A T T C C T G T C T T C T A A
0470 T A G G G T T C A A T T C C T A T C T C T I I A G
0474 T G A G G T T C A A T T C C I A T T G I C C T A G
0475 T G T A G G T T C A A T T C C I A T T G I C C T A G
0477 T G T A G G T T C A A T T C C T A T T G T C C T A G
0480 A G T A G G T T C G A T T C C T A T A G T T C T A G
0494 G A T G G G T T C G A T T C T C A T A G C C C A G
0530 C T T G C G T T C G A A T C C C A G T G G G A C C T
0570 C C C G G T T C G A A T C C G G G T A G G A C C T
0571 C C C A G T T C G A A T C C G G G T G G G A C C
0995 T C C G A G T T C A A A T C T C G G T G G A A C C T
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GLUTAMIC ACID
E145 C G A C T C G G G T T C A A A T C C C G G C C G G A G C A
E147 T A A C A C G G G T T C G A A C C C C G T A G G G A C C G C C A
E203 G A C A C A C G G G T T C G A A C C C C G T A C A G G C T A C C A
E235 T A A C A C G G G T T C G A A T C C C G T A C G G G T C A
E250 T A A C A G G G G T T C G A A T C C C C T A G G G G A C G C C A
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	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43													
E295	UUC	CYANOPIORA	PARAD.	**																																																				
E304	UUC	EUGLEMA	GRACILIS																																																					
E310	UUC	MARCHANTIA	POLYM.																																																					
E315	UUC	TRITICUM	AESTIVUM																																																					
E335	UUC	NICOTIANA	TABACUM																																																					
E355	UUC	PIZUN	SATIVUM																																																					
E365	UUC	SPUNGLIA	OLERACEA																																																					
E375	UUC	VICIA	FABA																																																					
E410	UUC	ASPERGILLUS	NIDUL.																																																					
E417	UUC	SACCHAROMYCES	CER.																																																					
E425	UUC	TORULOPSIS	GLAB.																																																					
E432	UUC	MITCINE	MAX																																																					
E450	UUC	ASCARIS	SUUM																																																					
E453	UUC	ARTEMIA	SP.																																																					
E456	UUC	AEDIS	ALBOPICTUS																																																					
E459	UUC	DROSOPHILA	YAKUBA																																																					
E464	UUC	MARCENTROTUS	LIV.																																																					
E466	UUC	STRONGYLOGEN.	PURP.																																																					
E468	UUC	XENOPUS	LAEVIS																																																					
E474	UUC	RAT																																																						
E477	UUC	MOUSE																																																						
E480	UUC	BOVINE																																																						
E494	UUC	HUMAN																																																						

	45	47	B	D	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75		
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76	
E295	C	G	A								C	G	G	G	A	T	C	C	C	T	G	G	G	G	G	A
E304	C	A	A								C	G	G	G	A	T	C	C	C	T	G	G	G	G	G	A
E310	C	G	A								C	G	G	G	A	T	C	C	C	T	G	G	G	G	G	A
E315	C	A	G								C	G	G	A	T	C	C	C	T	G	G	G	G	G	G	A
E335	C	A	G								C	G	G	G	A	T	C	C	C	T	G	G	G	G	G	A
E355	C	A	A								C	G	G	G	A	T	C	C	C	T	G	G	G	G	G	A
E365	C	A	A								C	G	G	G	A	T	C	C	C	T	G	G	G	G	G	A
E375	C	A	A								C	G	G	G	A	T	C	C	C	T	G	G	G	G	G	A
E410	G	T	G								C	G	G	A	T	C	C	C	T	G	G	G	G	G	G	A
E417	T	A	A								T	A	C	G	T	C	G	A	T	C	C	G	A	T	A	A
E425	A	A	A								T	G	T	G	T	C	A	T	C	C	A	A	C	T	A	A
E432	A	G	A								C	A	C	G	T	C	G	A	T	C	C	G	T	A	G	G
E450	A	G	G	T	T	A					T	G	G	A	T	C	A	A								
E453	A	A	A								T	A	C	C	T	A	A									
E456	A	A	A								T	A	A	A	T	T	T	T	T	T	T	T	T	T	T	A
E459	T	A	A								T	A	A	A	T	T	A	T	T	T	T	T	T	T	T	A
E464	A	A	G								T	T	G	A	T	A	A	A	T	C	T	C	A	T	A	G
E466	T	A	A	G							T	T	G	A	T	A	A	A	T	C	T	C	A	T	A	G
E468	A	A	G								T	T	G	A	T	A	A	A	T	C	T	C	A	T	A	G
E474	T	A	G								T	C	T	G	T	A	G	A	T	C	C	T	G	T	G	G
E477	T	G	G								T	C	A	C	A	T	A	T	G	C	C	T	G	T	A	C
E480	T	A	G								T	C	A	T	A	T	C	C	A	T	C	C	A	T	A	G
E494	T	G	G								T	C	G	T	T	G	T	A	G	T	C	C	T	G	C	G

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	
E555	UUC	DICTYOSTELIUM	DIS.	T	C	C	T	C	A	T	G	T	G	T	A	G	T	C	A	C	A	T	A	C	A	C	T	A	G	T	C	T	T	C	A	C	A	C	T	G	G		
E570	UUC	SACCHAROMYCES	CER.	T	C	C	G	A	T	A	G	T	A	A	C	G	G	C	T	A	C	A	T	C	A	C	A	T	C	A	C	C	G	T	G	G	G	G	G	G	G	G	G
E571	CUC	SACCHAROMYCES	CER.	T	C	C	G	A	T	A	G	T	A	A	C	G	G	C	T	A	C	A	T	C	A	C	A	T	C	A	C	C	G	T	G	G	G	G	G	G	G	G	G
E575	UUC	SCHIZOSACCHA.	POM.	T	C	C	G	T	G	T	G	T	C	C	A	A	C	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	G	T	C	G	C	T	T	C	A	C	G
E770	UUC	BOMBYX	MORI	T	C	C	C	G	T	A	T	G	G	T	C	T	A	G	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	T	G	C	T	T	C	A	C
E780	CUC	DROSOPHILA	MELANO.	T	C	C	T	A	T	T	G	T	C	T	A	G	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	C	G	G	C	T	C	A	C	C	G
E781	CUC	DROSOPHILA	MELANO.	T	C	C	A	T	T	G	T	C	T	A	G	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	C	G	G	C	T	C	A	C	C	G	
E782	UUC	DROSOPHILA	MELANO.	T	C	C	C	A	T	A	G	T	C	T	A	G	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	T	G	C	T	T	C	A	C	C	
E950	CUC	MOUSE		T	C	C	T	G	T	G	T	C	T	A	G	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	G	G	C	C	T	C	A	C	G	C	
E955	CUC	RAT		T	C	C	T	G	T	G	T	C	T	A	G	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	G	G	C	C	T	C	A	C	G	C	
E956	CUC	RAT		T	C	C	T	G	T	G	T	C	T	A	G	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	G	G	C	C	T	C	A	C	G	C	
E957	CUC	RAT		T	C	C	T	G	T	G	T	C	T	A	G	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	G	G	C	C	T	C	A	C	G	C	
E995	UUC	HUMAN		T	C	C	T	G	T	G	T	C	T	A	G	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	C	G	G	C	C	T	C	A	C	G	C	
G L Y C I N E																																											
G020	UCC	PHAGE T4		G	C	G	G	A	T	C	G	T	A	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T
G040	UCC	PHAGE T5		G	C	G	G	T	A	A	G	C	A	C	G	G	G	T	G	T	C	G	A	G	G	C	C	A	G	C	C	T	C	C	A	A	G	C	C	T	C	C	
G160	CCC	SULFOLOBUS	SOLFA.	G	C	G	C	C	T	C	G	T	A	G	T	G	G	A	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A	T	A	G	G	A
G203	UCC	MYCOPLASMA	MYCOID.	G	C	A	G	G	T	G	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A
G235	UCC	BACILLUS	SUBTILIS	G	C	G	G	T	G	T	A	G	T	T	A	G	T	A	A	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	
G236	GCC	BACILLUS	SUBTILIS	G	C	G	G	A	G	T	A	G	T	T	A	G	T	A	A	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	
G240	GCC	THERMUS	THERMOPHI.	G	C	G	G	G	A	G	T	A	G	T	C	A	G	T	C	A	G	T	C	A	G	T	C	A	G	T	C	A	G	T	C	A	G	T	C	A	G	T	
G250	UCC	E. COLI		G	C	G	G	C	A	T	C	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	
G251	GCC	E. COLI		G	C	G	G	C	A	T	A	C	T	C	A	G	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	A	T	A	

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42													
G260	UCC	PSEUDONUMAS	AER.																																																				
G295	GCC	CYANOPHORA	PARAD.*																																																				
G304	GCC	EUGLENA	GRACILIS																																																				
G305	UCC	EUGLENA	GRACILIS																																																				
G310	GCC	MARCHANTIA	POLYM.																																																				
G311	UCC	MARCHANTIA	POLYM.																																																				
G313	GCC	HORDEUM	VULGARE																																																				
G314	UCC	HORDEUM	VULGARE																																																				
G315	GCC	TRITICUM	AESTIVUM																																																				
G316	UCC	TRITICUM	AESTIVUM																																																				
G335	UCC	NICOTIANA	TABACUM																																																				
G336	GCC	NICOTIANA	TABACUM																																																				
G355	GCC	PISUM	SATIVUM																																																				
G410	ACC	ASPERGILLUS	NIDUL.																																																				
G411	UCC	ASPERGILLUS	NIDUL.																																																				
G417	UCC	SACCHAROMYCES	CER.																																																				
G421	UCC	NEUROSPORA	CRASSA																																																				
G425	UCC	TORULOPSIS	GLAB.																																																				
G434	GCC	LUPINUS	LUTEUS																																																				
G450	UCC	ASCARIS	SUUM																																																				
G452	UCC	CAENORHABD.	ELEG.																																																				
G454	UCC	LOCUSTA	MIGRATORIA																																																				
G456	UCC	MEGALOPTERIS	ALBOPUNCTUS																																																				

	45	47	B	D	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75				
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76			

6260	T	G	A	T	G	C	C	G	G	T	T	C	G	G	G	T	C	C	C	G	C	T	A	C	C	C	G	C
6295	A	T	G	T	C	G	C	G	T	T	C	G	A	T	C	G	C	G	T	C	A	C	C	C	G	C	T	
6304	A	T	G	C	C	A	T	G	G	G	T	C	G	A	G	T	C	C	C	A	T	A	T	C	T	G	C	T
6305	A	T	G	T	T	G	C	G	T	G	T	C	G	A	T	C	A	C	G	T	A	C	C	C	G	C	T	
6310	A	T	A	T	G	C	G	G	G	T	C	G	A	T	C	C	C	G	C	T	A	C	C	C	G	C		
6311	C	G	A	T	G	C	G	G	G	T	C	G	A	T	C	C	C	G	C	T	A	C	C	C	G	C		
6313	A	G	A	T	A	C	G	G	T	T	C	G	A	T	C	C	C	G	C	T	G	C	C	G	C	T		
6314	C	G	A	T	G	C	G	G	T	T	C	G	A	T	C	C	C	G	C	T	A	C	C	C	G	C		
6315	A	G	A	T	A	C	G	G	G	T	C	G	A	T	C	C	C	G	C	T	G	C	C	G	C	T		
6316	C	G	A	T	G	C	G	G	G	T	C	G	A	T	C	C	C	G	C	T	A	C	C	C	G	C		
6335	C	G	A	T	G	C	G	G	G	T	C	G	A	T	C	C	C	G	C	T	A	C	C	C	G	C		
6336	A	G	A	T	G	C	G	G	G	T	C	G	A	T	C	C	C	G	C	T	A	C	C	C	G	C		
6355	A	G	A	T	C	C	G	G	T	T	C	A	A	T	C	C	C	G	C	T	A	C	C	C	G	C		
6410	A	T	G	T	G	T	C	G	A	T	T	C	G	A	T	T	C	G	A	C	T	A	G	C	C	G	C	
6411	G	T	G	T	G	T	C	G	A	T	T	C	G	A	T	T	C	G	A	C	T	A	G	T	C	A	T	
6417	G	A	A	T	G	C	G	A	G	T	T	C	G	A	T	T	C	G	C	T	A	T	C	T	A	T		
6421	T	T	A	T	G	T	C	G	A	T	T	C	G	A	T	T	C	G	A	C	T	A	T	C	G	C		
6425	G	A	T	T	G	T	C	G	A	T	T	C	G	A	T	T	C	A	C	T	A	T	C	A	G			
6434	A	G	G	T	T	G	A	G	G	T	T	C	A	A	G	T	T	C	C	T	C	T	C	C	G	C		
6450	T	A	G	T	T	T	T																					
6452	A	G	G	T	T	T																						
6454	A	G	A	T	T	G	A	A	T																			
6456	A	G	G	T	T	A	A	T	A																			

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44										
6457	UCC	DROSOPHILA	MELANO.																																																			
6459	UCC	DROSOPHILA	YAKUBA																																																			
6464	UCC	PARACENTROTUS	LIV.																																																			
6466	UCC	STRONGYLOCEAN	PURP.																																																			
6468	UCC	AEROPUS	LAEVIS																																																			
6474	UCC	MOUSE	MITO																																																			
6477	UCC	MOUSE	MITO																																																			
6480	UCC	BOVINE	MITO																																																			
6494	UCC	HUMAN	MITO																																																			
6570	GCC	SACCHAROMYCES	CER.																																																			
6665	GCC	SORGHUM	BICOLOR																																																			
6670	GCC	ORYZA	SATIVA																																																			
6770	GCC	BOMBYX	MORI																																																			
6780	GCC	DROSOPHILA	MELANO.																																																			
6781	GCC	DROSOPHILA	MELANO.																																																			
6950	UCC	MOUSE																																																				
6955	UCC	RAT																																																				
6956	UCC	RAT																																																				
6995	GCC	HUMAN																																																				
6996	GCC	HUMAN																																																				
1040	GUG	PHAGE T5																																																				

	45	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76	
	44	46	A	C	E	G	I	K	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76		
6457	AGG										T	C	T	A	T	A	A	T	A	A	T	A	A	T	A	A
6459	AGG										T	C	T	A	T	A	A	T	A	A	T	A	A	T	A	A
6464	TAGC										T	C	T	A	G	G	A	A	A	T	C	T	A	G	A	T
6466	CAG										T	C	T	T	G	G	T	G	A	A	A	T	T	A	A	G
6468	AAG										T	C	T	T	A	G	T	A	G	A	T	C	T	A	G	A
6474	TAA										T	C	T	G	A	A	A	A	C	T	C	A	G	A	A	G
6477	AGA										T	C	T	G	A	T	A	A	C	C	C	A	G	A	A	G
6480	TAG										T	T	C	C	G	T	C	A	G	T	C	C	G	A	A	G
6494	TAG										T	T	T	G	A	C	A	C	A	T	T	C	A	A	A	A
6570	GGC										C	C	C	G	G	T	C	G	A	T	C	C	G	G	C	T
6665	AGA										C	C	G	G	G	T	C	G	A	T	C	C	G	G	C	T
6670	AGA										C	C	G	G	G	T	C	G	A	T	C	C	G	G	C	T
6770	CGG										C	C	G	G	G	T	C	G	A	T	C	C	G	G	C	T
6780	CGG										C	C	G	G	G	T	C	G	A	T	C	C	G	G	C	T
6781	CGG										C	C	G	G	G	T	C	G	A	T	C	C	C	G	A	T
6950	TGA										C	C	G	G	G	T	C	G	A	T	C	C	C	G	A	T
6955	TGA										C	C	G	G	G	T	C	G	A	T	C	C	C	G	A	T
6956	TGA										C	C	G	G	G	T	C	G	A	T	C	C	C	G	A	T
6995	AGG										C	C	G	G	G	T	C	G	A	T	C	C	C	G	A	T
6996	AGA										C	C	G	G	G	T	C	A	A	T	C	C	C	G	A	T
1040	CCTA										T	G	T	G	G	A	T	C	G	A	T	T	C	C	A	C

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42		
H I S T I D I N E																																												
H145	GUG	METHANOC.	VANI.	G	G	C	C	G	A	G	G	T	A	G	G	G	T	A	G	G	G	T	A	T	C	C	T	G	A	G	G	A	C	T	G	T	G	G	A	T	C	C	T	
H235	GUG	BACILLUS	SUBTILIS	G	G	C	G	G	T	T	G	T	G	C	G	A	A	G	T	G	G	T	A	A	C	G	C	A	G	A	T	T	G	T	G	G	C	T	C	T	G	G		
H236	GUG	BACILLUS	SUBTILIS	G	G	C	G	G	T	T	G	T	G	C	G	A	A	G	T	G	G	T	A	A	C	G	C	A	G	A	T	T	G	T	G	G	C	T	C	T	G	G		
H250	GUG	È.	COLI	G	G	T	G	C	T	A	T	A	G	C	T	C	A	G	T	G	T	G	A	A	G	A	G	C	C	T	G	G	A	T	T	G	T	G	A	T	C	C	A	G
H255	GUG	SALMONELLA	TYPHI.	G	G	T	G	C	T	A	T	A	G	C	T	C	A	G	T	G	T	G	A	A	G	A	G	C	C	T	G	G	A	T	T	G	T	G	A	T	C	C	A	G
H258	GUG	PHOTOBACT.	PHOSPH.	G	G	T	G	C	T	A	T	A	G	C	T	C	A	G	T	G	T	G	A	A	G	A	G	C	C	G	G	A	T	T	G	T	G	A	T	C	C	G	G	
H259	GUG	AEROMONAS	HYDROPH.	G	G	T	G	C	T	G	T	A	G	C	T	C	A	G	T	G	T	G	A	A	G	A	G	C	C	G	G	A	T	T	G	T	G	A	T	C	C	G	G	
H304	GUG	EUGLENA	GRACILIS	G	G	T	G	G	T	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	A	G	G	A	T	T	G	T	G	A	T	C	C	T	
H310	GUG	CHLORELLA	ELABORATA	G	G	C	G	A	C	T	A	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	A	T	T	G	T	G	A	T	C	C	T		
H320	GUG	ZEA	MAYS	G	G	C	G	A	T	G	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	T	G	T	G	A	T	C	C	A	G	T	C	C	
H333	GUG	MEDICAGO	SATIVA	G	G	C	G	A	T	G	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	T	G	T	G	A	T	C	C	A	G	T	C	C	
H335	GUG	NICOTIANA	TABACUM	G	G	C	G	A	T	G	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	T	G	T	G	A	T	C	C	A	G	T	C	C	
H340	GUG	NICOTIANA	DEBNEYI	G	G	C	G	A	T	G	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	T	G	T	G	A	T	C	C	A	G	T	C	C	
H348	GUG	CEJUNIA	HYBRIDA	G	G	C	G	A	T	G	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	T	G	T	G	A	T	C	C	A	G	T	C	C	
H355	GUG	PISUM	SATIVUM	G	G	C	G	A	T	G	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	T	G	T	G	A	T	C	C	A	G	T	C	C	
H365	GUG	SPINACIA	OLERACEA	G	G	C	G	A	T	G	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	T	G	T	G	A	T	C	C	A	G	T	C	C	
H407	GUG	TETRAHYMENA	PYRIF.	G	G	T	G	G	A	T	A	T	A	G	C	T	C	A	A	T	C	A	T	C	A	T	C	A	T	C	A	T	C	A	T	C	A	T	C	A	T	C		
H410	GUG	ASPERGILLUS	MIDUL.	G	G	T	G	G	T	G	T	A	G	T	T	C	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
H417	GUG	SACCAROMYCES	CER.	G	G	T	G	A	T	A	T	T	T	C	A	A	T	A	T	T	C	A	A	T	T	T	C	A	A	T	T	C	A	A	T	T	C	A	A	T	T	C		
H421	GUG	MITO	ZOSACCHIA.	POM.	G	G	T	C	A	T	T	T	C	A	A	T	T	C	A	A	T	T	C	A	A	T	T	C	A	A	T	T	C	A	A	T	T	C	A	A	T	T		
H425	GUG	TORULOPSIS	GLAB.	G	C	T	A	A	T	A	T	T	C	A	A	T	T	C	A	A	T	T	C	A	A	T	T	C	A	A	T	T	C	A	A	T	T	C	A	A	T	T		
H445	GUG	ZEA	MAYS	G	G	C	G	A	T	G	T	A	G	C	C	A	A	G	T	G	T	G	A	A	G	A	G	C	A	G	T	G	T	G	A	T	C	C	A	G	T	C		

	45	47	B	D	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75						
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76					

H450	T	G	A	A	A	T																		T	G	G	A	G	C	T
H452	A	G	A	A	T	A																		T	A	G	A	G	C	T
H459	T	G	A																											
H462	A	A	T																											
H464	A	G	T																											
H466	A	G	A																											
H468	A	G	T																											
H474	C	A	A																											
H477	C	A	A																											
H480	C	A	A																											
H483	C	C	A																											
H484	C	C	A																											
H485	C	T	A																											
H486	C	T	A																											
H487	T	A	A																											
H488	T	A	A																											
H489	A	A	A																											
H490	C	A	A																											
H491	C	A	A																											
H492	T	A	A																											
H493	T	A	A																											
H494	C	A	A																											
H570	A	A	A																											

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	
1575	GUG SCHIZOSACCHA. POM.	GCT	C	A	C	A	T	G	T	C	C	A	G	T	G	G	T	A	G	A	G	A	C	T	C	A	T	C	A	T	C	C	G	T	T	G	T	G	G	C	C	G	A	T	G
1780	GUG DIROSPHILLA MELANO.	G	C	C	G	T	G	A	T	C	T	C	T	A	G	T	G	G	T	A	G	A	C	C	C	A	C	G	T	T	G	T	G	G	C	C	G	T	G	G	C	C	T	G	G
1950	GUG MOUSE	G	C	C	G	T	G	A	T	C	T	C	T	A	G	T	A	G	G	T	A	G	T	A	C	T	C	T	C	T	G	C	G	T	T	G	T	G	T	G	G	C	C	A	G
I S O L E U C I N E																																													
1020	CAU PHAGE T4	G	G	C	C	C	T	G	T	A	G	C	T	C	A	A	T	G	G	T	A	G	A	G	C	A	G	T	C	C	C	T	C	A	T	A	G	G	G	A	A	G	G	A	
1040	CAU PHAGE T5	A	C	T	C	G	T	A	G	C	T	A	G	C	T	A	A	G	C	T	A	A	A	G	C	A	C	T	C	G	C	T	G	A	T	A	A	C	C	G	A	G	A		
1145	CAU METHANOCOC. VANI. ^m	G	G	G	C	T	C	G	T	A	G	C	T	C	A	G	G	C	T	G	G	T	A	G	A	G	T	G	C	T	C	G	G	C	T	C	A	T	A	C	C	G	A		
1203	CAU MYCOPLASMA MYCOID.	G	G	A	C	C	T	T	A	G	C	T	C	A	G	T	G	T	A	G	A	G	C	A	T	C	C	G	G	C	T	C	A	T	A	C	C	G	A						
1206	CAU SPIROPLASMA MELIF.	G	G	A	C	C	T	T	A	G	C	T	C	A	G	T	G	T	A	G	A	G	C	A	T	C	C	G	G	C	T	C	A	T	A	C	C	G	A						
1235	GAU BACILLUS SUBTILIS	G	G	G	C	C	T	G	T	A	G	C	T	C	A	G	C	T	G	G	T	A	G	A	G	C	G	C	A	C	C	C	T	G	A	T	A	G	C	G	T	G			
1236	GAU BACILLUS SUBTILIS	G	G	G	C	C	T	G	T	A	G	C	T	C	A	G	C	T	G	G	T	A	G	A	G	C	G	C	A	C	C	C	T	G	A	T	A	G	C	G	T	G			
1237	CAU BACILLUS SUBTILIS	G	G	A	C	C	T	T	A	G	C	T	C	A	G	T	G	T	A	G	A	G	C	A	T	C	C	G	G	C	T	C	A	T	A	C	C	G	A						
1243	GAU THIOBACILLUS FERRO.	G	G	G	C	C	T	A	G	C	T	C	A	G	C	T	G	G	T	A	G	A	G	C	A	T	C	C	G	C	T	G	A	T	A	G	C	G	T	G					
1250	GAU E. COLI	A	G	G	C	T	G	T	A	G	C	T	C	A	G	G	T	G	T	A	G	A	G	C	G	C	A	C	C	C	T	G	A	T	A	G	G	G	T	G					
1251	CAU E. COLI	G	G	C	C	C	T	A	G	C	T	C	A	G	T	G	T	A	G	A	G	A	G	C	A	G	G	C	A	C	C	T	G	A	T	A	C	C	T	G	C	T			
1260	GAU PSEUDOMONAS AER.	G	G	G	T	C	T	G	T	A	G	C	T	C	A	G	T	G	T	A	G	A	G	C	G	C	A	C	C	C	T	G	A	T	A	G	G	G	T	G					
1270	GAU CAMPYLOBAC. JEJUNI	G	G	C	C	A	T	A	G	C	T	C	A	G	C	T	A	G	C	T	A	G	C	T	A	G	C	T	A	G	C	T	G	A	T	A	G	G	T	G					
1272	GAU CAULOBACTER CRES.	A	G	C	C	T	G	T	A	G	C	T	C	A	G	G	T	G	T	A	G	A	G	C	G	T	A	C	C	C	T	G	A	T	A	G	C	T	G	A					
1290	GAU ANACYSTIS NIDULANS	G	G	G	C	T	A	T	A	G	C	T	C	A	G	G	T	G	T	A	G	A	G	C	G	T	A	C	C	C	T	G	A	T	A	G	C	T	G	A	A				
1295	GAU CYANOPHORA PARAD. ^m	G	G	G	C	T	A	T	A	G	C	T	C	A	G	G	T	G	T	A	G	A	G	C	G	T	A	C	C	C	T	G	A	T	A	G	C	T	G	A	A				
1297	GAU PYLAIELLA LITTORA. ^m	G	G	G	C	T	A	T	A	G	C	T	C	A	G	G	T	G	T	A	G	A	G	C	G	T	A	C	C	C	T	G	A	T	A	G	C	T	G	A	A				
1300	GAU CHLAMYDOMONAS REINH.	G	G	G	C	C	A	T	A	G	C	T	C	A	G	G	T	G	T	A	G	A	G	C	G	T	A	C	C	C	T	G	A	T	A	G	C	T	G	A	A				
1302	GAU CHLORELLA ELLIPSO. CHLORO	G	G	G	C	C	A	T	A	G	C	T	C	A	G	G	T	G	T	A	G	A	G	C	G	T	A	C	C	C	T	G	A	T	A	G	C	T	G	A	A				

45 47 B D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
 44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76

H575 CGA
 H780 TAA
 H950 CAA

ISOLEUCINE

1020 A G G T
 1040 A G A T
 1145 T G G T
 1203 C G G T
 1206 T G G T
 1235 A G G T
 1236 A G G T
 1237 C G G T
 1243 A G G T
 1250 A G G T
 1251 T G G T
 1260 A G G T
 1270 A G G T
 1272 A G G T
 1290 A G G T
 1295 A G G C
 1297 A G G G
 1300 A A G T
 1302 A G G T

CCAGGTTCGATTCCITGGTGGGGCA
 CCAGGTTCGAATCCITGGTCAAGGCA
 CTCGGTTCGAATCCGAGTCAAGGCA
 TACCAGTTCAAATCTGGTCTGGGTCA
 CCGGGTTTAAATCCCTCCGGAGTACCA
 CATGGGTTCAAATCCCATCGGGCCA
 CATGGTTCAGTCCAATAGGTCCACCA
 CACTGGTTCAGTCCAGTAGGTCCACCA
 CGTGGTTCGAGTCCACTCAGGCCACCA
 CATGGTTCGAGTCCATTAGGCCACCA
 CATGGTTCGAGTCCATCAGGTCCACCA
 CAGTGGTTCGAGTCCACTGGGCCACCA
 CGTGGTTCAGTCCACTCAGGCCACCA
 CGTGGTTCAGTCCAGTAGGTCCACCA
 CGGAGTTCGAATCTGCCAGACCCACCA
 CCAAGTTCAGTCTTGTAGGCCACCA
 CGCAGTTCGAGTCTGCCTAGGCCACCA
 CCTGGTTCAGTCCAGTAGGCCACCA
 CCTGGTTCGAATCCAGGTAGGCCCA
 CCTGGTTCAAATCCAGGTAGGCCCA
 CAAAGTTCAAATCTTCA TAGGCCA
 CACTGATTCGAATTCAGCATAGGCCA

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	
1480	GAU BOVINE	A	G	A	A	A	T	A	T	G	T	C	T	G	C	A					A	A	G	A	G	T	T	A	C	T	T	G	A	T	A	G	A	G	T	A	A			
1494	GAU HITO	A	G	A	A	A	T	A	T	G	T	C	T	G	A					A	A	G	A	G	T	T	A	C	T	T	G	A	T	A	G	A	G	T	A	A				
1570	AU SACCHAROMYCES CER.	G	G	T	C	T	C	A	G	T	G	C	C	C	A	G	T	G	G	T	A	G	G	C	A	C	T	C	T	A	T	A	A	C	G	C	G							
1590	AU YEAST	G	C	T	C	T	C	A	G	T	G	C	C	C	A	G	T	G	G	T	A	G	G	C	A	C	T	C	T	A	T	A	A	C	G	C	G							
1780	AU DROSOPHILA MELANO.	G	G	C	C	A	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	C	G	T	C	C	T	G	C	T	A	T	A	A	C	G	C	G					
1950	AU MOUSE	G	G	C	C	G	G	T	A	G	C	T	C	A	G	T	G	G	T	A	G	A	G	C	G	T	C	C	T	G	C	T	A	T	A	A	C	G	C	G				
L E U C I N E																																												
1020	UAA PHAGE T4	G	C	G	G	A	T	G	T	C	A	A	A	T	G	G	T	A	G	G	C	A	C	A	G	C	C	A	C	T	T	A	A	A	T	G	C	T	G					
1040	UAG PHAGE T5	G	G	G	C	T	A	T	G	T	G	A	A	C	T	G	A	C	T	A	G	C	A	T	A	C	G	G	C	T	T	A	G	T	C	C	G							
1145	UAG METHANOCOCC. VANI.	G	C	A	G	G	G	T	G	T	C	G	A	G	C	T	G	C	C	A	A	G	A	T	G	C	A	G	G	A	T	C	C	T	G									
1160	CAG SULFOLOBUS SOLFA.	G	C	G	G	G	G	T	G	C	C	G	A	G	C	A	G	T	C	A	A	G	G	G	T	C	G	G	C	T	C	A	G	G	C	C	C	G						
1190	CAA THERMOPROT. TENAX	G	C	G	G	G	G	T	G	C	C	G	A	G	C	C	A	G	T	A	A	G	G	G	T	C	A	G	G	C	T	C	A	G	A	C	C	T	G					
1191	CAG THERMOPROT. TENAX	C	C	C	C	A	G	T	G	C	C	G	A	T	A	G	G	T	A	A	G	G	C	A	T	G	G	A	T	C	C	A												
1200	UAA MYCOPLASMA CAPRIC.	G	G	G	A	T	G	G	G	A	T	T	G	G	A	A	T	G	G	C	A	G	C	G	A	C	T	A	G	G	A	T	C	C	A									
1201	UAG MYCOPLASMA CAPRIC.	G	G	G	A	T	G	G	G	A	T	T	G	G	A	A	T	G	G	C	A	G	C	G	A	C	T	A	G	G	A	T	C	C	A									
1205	UAG MYCOPLASMA PG50	G	G	G	G	A	T	G	C	C	G	A	A	T	G	G	C	A	G	C	A	G	C	A	C	T	A	G	G	A	T	C	T	A	G									
1235	CAG BACILLUS SUBTILIS	G	C	G	G	A	T	G	C	C	G	A	A	T	G	G	C	A	G	C	C	A	T	A	G	C	T	A	G	A	T	C	A	G	G	C	T	A	G					
1236	CAG BACILLUS SUBTILIS	G	C	G	C	C	T	G	T	G	G	A	A	T	G	G	C	A	G	C	C	A	T	A	G	C	T	A	G	A	T	C	A	G	G	C	T	A	G					
1237	UAA BACILLUS SUBTILIS	G	C	C	G	G	T	G	T	G	G	A	A	T	G	G	C	A	G	C	C	A	T	A	G	C	T	A	G	A	T	C	C	A										
1238	CAA BACILLUS SUBTILIS	G	C	C	G	G	T	G	T	G	G	A	A	T	G	G	C	A	G	C	C	A	T	A	G	C	T	A	G	A	T	C	C	A										
1250	CAG E. COLI	G	C	G	A	G	T	G	C	C	G	A	A	T	G	G	T	A	G	C	G	C	T	A	G	C	T	A	G	G	T	C	A	G										
1251	UAG E. COLI	G	C	G	G	A	G	T	G	C	C	G	A	A	T	G	G	T	A	G	C	G	C	A	G	C	C	A	G	A	T	T	A	G	G	T	C	T	G					
1252	CAA E. COLI	G	C	C	G	A	G	T	G	C	C	G	A	A	T	G	G	T	A	G	C	G	C	A	G	T	T	C	A	A	A	T	C	A	A									

	45 47 B D F H J L N P	49 51 53	55 57 59	61 63 65	67 69 71	73 75
	44 46 A C E G I K M O	48 50 52	54 56 58 60	62 64	66 68 70 72	74 76
1480	A T A A	T A G A G C T	T C A A A C	C C T C T	A A I I C T A	
1494	A T A A	T A G G C T	T T A A A C	C C C C T	A T I I C T A	
1570	G G A T	C A R C G G T	T C G A T C	C G C C T	A G A G A C C A	
1590	C C G T	C G T G G G T	T C A A T C	C C C A C C T	C G A G C A	
1780	A G G T	C G C G G G T	T C G A T C	C C C T C A T	G G G C C A	
1950	A G G T	C G C G G G T	T C G A T C	C C C G T A C G	G G C C A	
	LEUCINE					
L020	C G G A A T G A T T I C C T	T G T G G G T	T C G A G T	C C C A C T	T C T C G C A C C A	
L040	A G C T A A A T G C G	T G G G A G T	T C G A G T	C T C C C T	A G C C C C A C C A	
L145	T C C A G T A G T G G T T	C A G G G T	T C A A A T	C C C T G E	C C C T G C A	
L160	T G G T G T A G C C C T G	C G T G G G T	T C A A A T	C C C A C C	C C C C G C A	
L190	T G G C G T A G C C C T G	C G T G G G T	T C G A A T	C C C A C C	C C C G C A	
L191	T G G C G T A G C C C T G	C G T G G G T	T C A A A T	C C C A C C	C C C C G C A	
L200	C G G G C T A A T A T C C T G	T G C C G G T	T C A A G T	C G G G C T	T G G G G A C C A	
L201	C G T C T T G A C G	T A A G G G T	T C A A G T	C C C T T A T	C C C C C A C C A	
L205	C G T C T T G A C G	T A A G G G T	T C A A G T	C C C T T A T	C C C C C C A C C A	
L235	T G T C T T A C A G A C G	T G G G G G T	T C A A G T	C C C T T C A T	C C G C C A C C A	
L236	C G T A G G T G A C T A C C G	T G C G G G T	T C A A G T	C G G C C C T	C G G C A	
L237	C G G T A G G T G A C T A C C G	T G C C G G T	T C A A G T	C G G C C C T	C G G C A C C A	
L238	T T C C T T C T G G A G	T G T C G G T	T C G A C C	C G G A C C A C	C C G G T A	
L250	T G T C C T T A C G G A C G	T G G G G G T	T C A A G T	C C C C C C T	C C C A C C A C C A	
L251	C G C C G C A A G G T G	T G C G A G T	T C A A G T	C T C G C T	C C C G C A C C A	
L252	C G T A G A A T A C G	T G C C G G T	T C G A G T	C C G G C T	T C G G C A C C A	

	45	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75		
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76	
L255	T	G	T	C	T	A	C	G	G	A	C	G														
L259	T	G	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C
L275	T	A	C	T	C	G	A	A	G	G	G	T														
L277	T	G	C	C	G	C	A	A	G	G	T															
L295	C	G	A	T	T	G	T	A	T	A	G	A	T	C	G											
L304	T	G	T	C	T	T	A	T	A	G	T	G														
L305	T	G	C	T	T	A	A	A	G	C	G															
L306	T	G	C	A	T	A	A	G	C	T																
L310	T	G	C	T	A	A	G	C	A																	
L311	T	G	G	C	T	T	A	A	G	C	C	G														
L312	T	G	C	T	A	A	G	G	C	T																
L320	C	G	A	C	T	T	A	A	G	T	C	G														
L321	T	G	C	T	A	A	A	G	C	C	G															
L325	T	G	C	T	A	A	A	A	G	C	G															
L335	T	G	C	T	A	A	T	G	C	A																
L336	C	G	A	C	T	T	A	A	A	A	T	C	G													
L337	T	G	C	T	A	A	T	A	G	C	G															
L355	T	G	C	T	A	A	C	A	G	C	G															
L365	T	G	C	T	A	A	A	A	G	C	G															
L375	T	G	C	T	A	A	C	A	G	C	G															
L376	C	G	A	C	T	T	A	A	A	A	T	C	G													
L378	C	G	A	C	T	T	A	A	G	T	C	G														
L407	T	T	C	C	A	T	A	G	G	A	G															

	45 47 B D F H J L N P	49 51 53	55 57 59	61 63 65	67 69 71	73 75
	44 46 A C E G I K M O	48 50 52	54 56 58 60	62 64 66	68 70 72	74 76
	T T C C T A C G G G A G	T G C C G G	T T C G A C C C	C C G G T	C A T A A G T A	
L408	T G A C G C A A G T C G	T A A C G T	T C G A A T	C G T T T	C T C G G A T A	
L410	T A G T C A A A C T T	T G C A A G	T T C A A G T	C T T G T	T A C C C G T A	
L411	C G C T T C G A G C G	T G T T G G	T C G A G T	C C A A C	T T A G G T A	
L421	T A C T T T A T A G T A	T G A G G T	T C A A A T	C C T T C	A A A T A G C A	
L425	T A C T T T A C A G T A	T G A G G T	T C A A G T	C C T T T	A A A T A G C A	
L428	T G T A T A C T	
L430	A G A T G G T	A G A T A G			G G A C A A C T	
L451		T G T A G A	T T T T T T	T C T A C A	A A T A G T A	
L454	T T A	T A A G T	T T G A C	C T T T A	A A T A G A A	
L455	A A A	T A T A T	T T A A T T A	A T T A T	A A A T A G T A	
L456	T T A	T A A G T	A T T T	T A C T T T	A T T A G A A	
L457	A T A	T A A G T	A T T T	T A C T T T	T A T T A G A A	
L459	A T A	T G T A A T	T T T	T A T T A C A	A A T A G T A	
L460	A T A	C A A G G	T T C A A C T	C C T T C	T T T C A G C T	
L464	C A T	C A A G G	T T C A A C T	C C T T T	T T T A C C T	
L465	C T A T	C A G G G	T T C A A C T	C C T T C	T T T C A G C T	
L466	T A A	C A G G G	T T C A A C	T C C T T	C T T T A G C	
L467	C T A T	C A G G G	T T C A A A T	C C C C T	C G C T A A C T	
L468	T T A T	T C T T G G	T G C A A A T	C C A A G T	A A A A G C T	
L469	A A A C	T C T T G G	T G C A A G T	C C A A G T	A A A A G C T	
L470	C A T C	C T T G G	T G C A A G T	C C A A G T	A A A A G C T	
L474	A A A	C T T G G	T G C A A C T	C C A A A	A A A A G T A	
L475	T T C C	C A G A G G	T T C A A A T	C C T C T	C C C T A A T A	

	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43			
L477	UAG	MOUSE	ACTTTTAA	A	GGAT	A	ATAGT	A	TCCA	T	GGT	C	T	T	AGGA	A	CCAA																				
L478	UAA	MOUSE	ATTAGGT	G	CAG	A	GCCAGGAA	A	TTCG	T	AAG	C	T	T	AAAC	C	CTTG																				
L480	UAA	BOVINE	GTTAAGGT	G	CAG	A	GCCCGGTA	A	TGCA	T	AAA	C	T	T	AACT	A	TTTTA																				
L481	UAG	BOVINE	ACTTTTAA	A	GGAT	A	GTAGT	T	TCCG	T	GGT	C	T	T	AGGA	A	CCAA																				
L483	UAG	MACACA FUSCATA	ACTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGC	C	T	T	AGG	A	CCAA																				
L484	UAG	MACACA MULATTA	ACTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGC	C	T	T	AGG	A	CCAA																				
L485	UAG	MACACA FASCICULA	ACTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGC	C	T	T	AGG	A	CCAA																				
L486	UAG	MACACA SYLVANUS	ACTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGC	C	T	T	AGG	A	CCAA																				
L487	UAG	SAIMIRI SCIUREUS	ACTTTTAA	A	GGAT	A	GTAGT	T	TCCA	T	GGT	C	T	T	AGG	A	CCAA																				
L488	UAG	TARSIUS SYRICHITA	ACTTTTAA	A	GGAT	A	GTAGT	T	TCCA	T	GGT	C	T	T	AGG	A	CCAA																				
L489	UAG	LEPOR CATTA	ACTTTTAA	A	GGAT	A	GTAGT	T	TCCA	T	GGT	C	T	T	AGG	A	CCAA																				
L490	UAG	CHIMPANZEE	ACTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGT	C	T	T	AGG	A	CCAA																				
L491	UAG	GIBBON	ACTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGT	C	T	T	AGG	A	CCAA																				
L492	UAG	GORILLA	ACTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGT	C	T	T	AGG	A	CCAA																				
L493	UAG	ORANG UTAN	GCTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGT	C	T	T	AGG	A	CCAA																				
L494	UAG	HITO	ACTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGT	C	T	T	AGG	A	CCAA																				
L495	UAG	HUMAN	ACTTTTAA	A	GGAT	A	ACAGC	T	TCCA	T	GGT	C	T	T	AGG	A	CCAA																				
L560	AAG	NEUROSPORA CRASSA	GGCAAGAT	G	CCG	A	GC	AGC	G	GTCTA	A	GGC	G	C	CA	C	G	T	A	A	G	G	C	C	G	T	G	G									
L570	CAA	SACCHAROMYCES CER.	GTTGT	T	G	CCG	A	GC	G	GTCTA	A	GGC	G	C	CA	G	T	A	A	G	C	T	CA	A	G	C	T	C	A	G	T	C	A	G			
L645	CAA	PHASEOLUS VULGARIS	GTCAGGAT	G	CCG	A	GC	AGT	G	GTCTA	A	GGC	G	C	CA	G	A	C	T	CA	A	G	T	C	T	G	G										
L700	AAG	CAENORHABDI. ELEG.	GAGAGAT	G	CCG	A	GC	AGC	G	GTCTA	A	GGC	G	C	CA	G	A	C	T	CA	A	G	T	C	T	G	G										
L780	CAA	DROSOPHILA MELANO.	GTCAGGAT	G	CCG	A	GC	AGC	G	GTCTA	A	GGC	G	C	CA	G	A	C	T	CA	A	G	T	C	T	G	G										
L781	CAG	DROSOPHILA MELANO.	GTCAGGAT	G	CCG	A	GC	AGT	G	GTCTA	A	GGC	G	C	CA	G	T	CA	A	G	T	C	A	G	T	C	G	C	A	G	T	C	A	G	T	C	G

	45	47	B	D	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76
L477	AAA										C	T	T	G	G	T	G	C	A	A	T	C	C	A	A
L478	TTCC										C	A	G	G	T	T	C	A	A	A	T	C	C	T	C
L480	TATC										C	A	G	A	T	T	C	A	A	A	T	C	C	T	C
L481	AA										A	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L483	AAA										C	A	T	T	G	T	G	C	A	A	A	T	C	C	A
L484	AAA										T	A	T	T	G	T	G	C	A	A	A	T	C	C	A
L485	AAA										C	A	T	T	G	T	G	C	A	A	A	T	C	C	A
L486	AAA										T	A	T	T	G	T	G	C	A	A	A	T	C	C	A
L487	AAA										C	A	T	T	G	T	G	C	A	A	A	T	C	C	A
L488	AAA										A	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L489	AAA										A	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L490	AAA										T	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L491	AAA										T	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L492	AAA										T	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L493	AAA										T	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L494	AAA										T	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L495	AAA										T	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L560	TCCGAAAGGG										T	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L570	TATCGTAAAGATG										C	T	T	G	T	G	C	A	A	A	T	C	C	A	A
L645	TCTTCGAGAGAGGG										C	A	G	A	T	T	C	G	A	A	T	C	C	A	A
L700	TCCCTCGG66G										C	T	G	G	T	T	C	A	A	A	T	C	C	A	A
L780	TCTCTCTGA66G										C	T	G	G	T	T	C	G	A	A	T	C	C	A	A
L781	TCTACTCTGTAGG										C	T	G	G	T	T	C	G	A	A	T	C	C	A	A

	45	47	B	D	E	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76	
	44	46	A	C	E	G	I	K	M	O		50	52	54	56	58	60	62	64	66	68	70	72	74	76		
L830	T	C	C	C	C	C	T	G	G	G	T	T	C	G	A	A	T	C	C	A	C	T	C	T	G	A	
L950	T	C	C	C	C	C	T	G	G	G	T	T	C	G	A	A	T	C	C	A	C	T	C	T	G	A	
L955	T	C	C	C	C	T	A	G	A	G	G	T	T	C	G	A	A	T	C	C	A	C	T	C	T	G	
L956	T	C	C	C	C	T	G	A	G	G	T	T	C	G	A	A	T	C	C	A	C	T	C	T	G	A	
L957	T	C	C	C	C	T	G	A	G	G	T	T	C	G	A	A	T	C	C	A	C	T	C	T	G	A	
L995	T	C	T	C	T	C	G	G	A	G	G	T	T	C	G	A	A	T	C	C	A	C	T	C	T	G	
L996	T	C	T	C	T	C	G	G	G	G	T	T	C	G	A	A	T	C	C	A	C	T	C	T	G	A	
LYSINE																											
K040	A	G	G	T	T	C	A	G	G	T	T	C	G	A	G	A	T	C	C	A	C	C	C	C	C	C	
K145	C	G	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	
K148	C	G	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	
K200	G	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	C	
K205	G	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	C	
K235	G	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	C	
K250	T	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	C	
K304	C	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	C	
K310	G	A	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	
K335	T	A	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	
K355	A	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	C	
K360	T	A	G	T	T	C	G	A	A	T	T	C	G	G	G	C	C	C	C	C	C	C	C	C	C	C	
K410	C	T	A	C	T	T	A	G	T	T	T	C	G	A	A	T	T	C	C	A	C	C	C	C	C	C	
K417	C	C	A	T	T	C	A	A	C	T	T	C	G	A	A	T	T	C	C	A	C	C	C	C	C	C	
K425	C	C	A	T	T	C	A	A	C	T	T	C	G	A	A	T	T	C	C	A	C	C	C	C	C	C	

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42													
K450	UUU	ASCARIS	SULM																																																				
K454	UUU	LOCUSTA	MIGRATORIA																																																				
K457	UUU	DROSOPHILA	MELANO.																																																				
K459	UUU	DROSOPHILA	YAKUBA																																																				
K466	UUU	STRONGYLOEN	PURP.																																																				
K468	UUU	XENOPUS	LAEVIS																																																				
K474	UUU	RAT																																																					
K475	UUU	RAT																																																					
K477	UUU	MOUSE																																																					
K480	UUU	BOVINE																																																					
K480	UUU	BOVINE																																																					
K494	UUU	HUMAN																																																					
K370	UUU	SACCHAROMYCES	CER.																																																				
K371	UUU	SACCHAROMYCES	CER.																																																				
K375	UUU	SCHIZOSACCHA.	POM.																																																				
K700	UUU	CAENORHABDI.	ELEG.																																																				
K770	UUU	BOMBYX	MORI																																																				
K780	UUU	DROSOPHILA	MELANO.																																																				
K781	UUU	DROSOPHILA	MELANO.																																																				
K830	UUU	XENOPUS	LAEVIS																																																				
K870	UUU	CHICKEN																																																					
K950	UUU	MOUSE																																																					
K951	UUU	MOUSE																																																					
K955	UUU	RAT																																																					

	45	47	B	D	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76			
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76			

K450	A	A	A	T	G	G	G	T	T	G	T	C																
K454	A	A	A	A																								
K457	T	T	A	A																								
K459	T	T	T	A																								
K466	A	G	A																									
K468	A	G	A	T																								
K474	A	G	T																									
K475	A	G	T																									
K477	A	G	T																									
K480	A	G	A	T																								
K494	A	G	A	T																								
K570	A	T	G	T																								
K571	A	G	G	T																								
K575	A	G	G	T																								
K700	T	T	G	T																								
K770	G	G	G	T																								
K780	G	G	G	T																								
K781	G	G	G	T																								
K830	G	G	G	T																								
K870	G	G	G	T																								
K950	G	G	G	T																								
K951	G	G	G	T																								
K955	G	G	G	T																								

	1	2	3	4	5	6	7	8	9	10	12	14	16	19	21	23	25	27	29	31	33	35	37	39	41	43	
	11	13	15	17	18	20	B																				
K995	UUU HUMAN	G	C	C	C	G	G	A	T	A	G	C	T	C	A	G	T	C	A	G	T	C	A	G	T	C	A
M040	CAU PHAGE T5	A	G	T	A	G	T	G	C	A	G	A	G	T	G	T	A	G	G	C	A	T	C	A	G	T	C
		*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****
M E T H I O N I N E																											
M203	CAU MYCOPLASMA MYCOID.	G	G	C	G	G	G	G	T	A	G	C	T	C	A	G	T	T	C	G	G	T	T	C	A	T	C
M206	CAU SPIROPLASMA MELIF.	G	G	C	G	G	G	A	T	A	G	C	T	C	A	G	T	T	C	G	G	T	T	C	A	T	C
M235	CAU BACILLUS SUBTILIS	G	G	A	C	C	T	T	A	G	C	T	C	A	G	T	T	C	G	G	T	T	C	A	T	C	C
M236	CAU BACILLUS SUBTILIS	G	G	C	C	T	C	T	A	G	C	T	C	A	G	T	T	C	G	G	T	T	C	A	T	C	C
M250	CAU E. COLI	G	G	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M304	CAU EUGLEMA GRACILIS	G	G	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A
M310	CAU CHLORO	G	G	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A
M310	CAU MARCHANTIA POLYM.	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M313	CAU CHLORO	G	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M313	CAU HORDEUM VULGARE	G	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M315	CAU CRITICUM AESTIVUM	G	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M320	CAU ZEA MAYS	G	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M320	CAU CHLORO	G	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M324	CAU ARABIDOPSIS THAL.	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M327	CAU GLYCINE MAX	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M335	CAU CHLORO	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M335	CAU NICOTIANA TABACUM	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M365	CAU SPINACEA OLERACEA	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M410	CAU CHLORO	C	C	A	A	G	T	T	A	A	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C	
M410	CAU MENGENGILLUS NIDUL.	C	C	A	A	G	T	T	A	A	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C	
M411	CAU ASPERGILLUS NIDUL.	A	A	G	C	T	A	A	T	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	
M411	CAU MITO	G	C	T	G	T	A	G	T	T	A	A	T	T	C	A	G	T	T	C	A	G	T	T	C	A	
M417	CAU SACCHAROMYCES CER.	G	C	T	G	T	A	G	T	T	A	A	T	T	C	A	G	T	T	C	A	G	T	T	C	A	
M417	CAU MITO	A	A	A	C	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
M425	CAU TORULOPSIS GLAB.	A	C	T	T	G	T	A	A	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	A	C	
M425	CAU MITO	A	C	T	T	G	T	A	A	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	A	C	
M430	CAU ARABIDOPSIS THAL.	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M430	CAU MITO	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M432	CAU GLYCINE MAX	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C
M432	CAU MITO	A	C	C	T	A	C	T	C	A	G	T	T	C	A	G	T	T	C	A	G	T	T	C	A	T	C

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43								
M445	CAU ZEA MAYS	G	G	G	C	T	T	A	T	A	G	T	T	A	A	T	G	G	T	G	A	A	C	G	T	A	C	C	G	T	A	C	C	G	T	C	A	T	A	A	C	G	G	T							
M570	CAU SACCHAROMYCES CER. MITO	G	C	T	C	A	G	T	A	G	C	T	C	A	G	T	A	G	G	A	A	G	A	G	C	G	T	C	A	G	T	C	A	T	A	A	T	C	T	G	A										
M635	CAU GLYCINE MAX	C	G	C	G	G	T	G	C	C	A	G	T	G	G	G	C	A	G	T	G	G	G	C	A	G	T	G	G	G	C	A	G	T	C	A	T	A	A	C	C	T	G								
M780	CAU DROSOPHILA MELANO.	A	G	C	A	G	A	T	G	C	C	C	A	G	T	G	G	A	A	G	C	G	T	G	C	C	C	A	T	A	A	C	C	C	A	G															
	METHIONINE - INITIATOR																																																		
X040	CAU PHAGE T5	T	G	C	G	G	G	T	A	G	A	T	C	T	C	T	G	G	T	A	G	A	T	C	G	T	A	G	T	C	T	A	A	C	T	A	A	C	T	A	A	C	T	A	A	C	T	A	A		
X160	CAU SULFOLOBUS SOLFA.	G	G	C	C	C	G	T	A	G	C	T	A	G	C	C	G	T	A	G	A	G	C	C	C	G	C	C	G	C	T	C	A	T	A	A	C	C	G	G											
X190	CAU THERMOPROT. TENAX	A	G	C	G	G	T	A	G	C	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A
X203	CAU MYCOPLASMA MYCOID.	C	G	C	G	G	T	A	G	A	G	C	A	G	T	G	G	T	A	G	A	G	C	T	C	G	C	G	G	C	T	C	A	T	A	A	C	C	C	G											
X206	CAU SPIROPLASMA MELIF.	C	G	T	G	G	G	T	G	A	G	C	A	G	T	G	G	T	A	G	A	G	C	T	C	G	T	T	G	G	C	T	C	A	T	A	A	C	C	C	A										
X218	CAU STREPTOMYCES RIM.	C	G	C	G	G	G	T	G	A	G	C	A	G	C	T	C	G	G	T	A	G	C	T	C	G	T	G	G	C	T	C	A	T	A	A	C	T	C	A	A	C	T	C	A	A	C	T	C	A	
X219	CAU STREPTOMYCES RIM.	C	G	C	G	G	G	T	G	A	G	C	A	G	C	T	C	G	G	T	A	G	C	T	C	G	T	G	G	C	T	C	A	T	A	A	C	C	C	A											
X235	CAU BACILLUS SUBTILIS	C	G	C	G	G	G	T	G	A	G	C	A	G	T	C	G	G	T	A	G	C	T	C	G	T	C	G	G	C	T	C	A	T	A	A	C	C	G	A											
X250	CAU E. COLI	C	G	C	G	C	G	T	G	A	G	C	A	G	C	T	C	G	G	T	A	G	C	T	C	G	T	C	G	G	C	T	C	A	T	A	A	C	C	C	A										
X251	CAU E. COLI	C	G	C	G	G	G	T	G	A	G	C	A	G	C	T	C	G	G	T	A	G	C	T	C	G	T	C	G	G	C	T	C	A	T	A	A	C	C	C	A										
X304	CAU EUGLENA GRACILLIS CHLORO	G	C	G	G	A	G	T	A	G	A	G	C	A	G	T	C	A	G	G	T	A	G	C	T	C	G	C	A	G	G	C	T	C	A	T	A	A	T	C	C	T	G								
X310	CAU MARCHANTIA POLYM. CHLORO	C	G	C	G	G	A	G	T	A	G	A	G	C	A	G	T	C	T	G	T	A	G	C	T	C	G	C	A	G	G	C	T	C	A	T	A	A	C	C	T	T	G								
X313	CAU HORDEUM VULGARE CHLORO	A	G	C	G	A	G	T	A	G	A	G	C	A	G	T	T	G	T	A	G	C	A	G	T	T	G	T	G	T	G	T	G	T	G	T	G	T	G	T	G										
X315	CAU NICOTIANA AESTIVUM CHLORO	A	G	C	G	A	G	T	A	G	A	G	C	A	G	T	T	G	T	A	G	C	A	G	T	T	G	T	G	T	G	T	G	T	G	T	G	T	G												
X335	CAU NICOTIANA TABACUM CHLORO	C	G	C	G	G	T	A	G	A	G	C	A	G	T	T	G	T	A	G	C	A	G	T	T	G	T	G	T	G	T	G	T	G	T	G	T	G													
X355	CAU PISUM SATIVUM CHLORO	C	G	C	G	A	G	T	A	G	A	G	C	A	C	T	T	G	T	A	G	C	T	C	G	T	C	A	A	C	T	T	G	T	G	T	G														
X405	CAU PARAMECIUM PRIM. MITO	G	C	A	G	C	A	G	G	T	G	G	T	C	T	C	A	A	C																																
X407	CAU TETRAHYMENA PYRIF. MITO	G	C	T	G	C	T	T	A	T	G	T	T	C	A	G	T	T	C	A	A	C																													

	45	47	B	D	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76										
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76										
M445	A	T	A	T							T	G	A	G	G	T	T	C	G	A	G	C	C	T	A	A	G	C	C	C	C	A			
M570	A	G	G	T							C	G	A	G	A	T	T	C	G	A	A	C	T	T	C	T	C	T	G	A	G	C	A		
M635	A	G	G	T							C	G	A	G	A	T	T	C	G	A	A	C	T	T	C	T	C	T	C	C	C	C	A		
M780	A	G	G	T							C	G	A	G	A	T	T	C	G	A	A	C	T	T	C	T	C	T	G	C	T	G	C	A	
METHIONINE-INITIATOR																																			
X040	A	A	A	G	A						G	G	A	G	G	T	T	C	G	A	C	C	C	G	C	T	T	C	C	A					
X160	T	G	G	T							C	G	G	G	T	T	C	A	A	G	T	C	C	C	G	C	G	G	C	C	C	A			
X190	A	G	G	T							C	C	G	G	T	T	C	A	A	T	C	C	C	G	G	C	C	C	C	C	C	A			
X203	A	G	G	C							C	G	A	G	T	T	C	G	A	G	T	C	T	G	C	C	C	C	G	C	A				
X206	A	G	G	T							C	G	A	G	T	T	C	A	A	G	T	C	C	T	G	C	C	C	C	G	C	A			
X218	A	G	G	T							C	G	A	G	T	T	C	A	A	T	C	C	T	G	C	C	C	C	G	C	T	A			
X219	A	G	G	T							C	G	A	G	T	T	C	A	A	T	T	C	T	G	T	C	C	C	G	C	T	A			
X235	A	G	G	T							C	G	A	G	T	T	C	A	A	T	C	C	T	G	C	C	C	G	C	A					
X250	A	G	A	T							C	T	C	G	G	T	T	C	A	A	T	C	C	G	C	C	C	G	C	A					
X251	A	G	G	T							C	T	C	G	G	T	T	C	A	A	T	C	C	G	C	C	C	C	G	C	A				
X304	A	A	G	T							C	G	A	G	G	T	T	C	A	A	T	C	C	T	T	C	T	C	C	G	T	A			
X310	A	G	G	T							C	A	T	A	G	G	T	T	C	A	A	T	C	C	T	G	T	C	C	G	C	A			
X313	A	G	G	T							C	A	C	G	G	G	T	T	C	G	A	T	C	C	C	G	T	C	C	G	C	A			
X315	A	G	G	T							C	A	C	G	G	G	T	T	C	G	A	T	C	C	C	G	T	C	C	G	C	A			
X335	A	G	G	T							C	A	C	G	G	G	T	T	C	G	A	T	C	C	C	G	T	C	C	G	C	A			
X355	A	G	G	T							T	A	C	G	G	G	T	T	C	A	A	T	C	C	T	G	T	C	C	G	C	A			
X405	C	T	A								A	T	A	A	G	T	T	C	G	A	T	C	T	T	G	T	A	G	G	G	C				
X407	T	A	C	T							A	T	A	A	G	T	T	C	G	A	T	C	T	T	A	A	G	C	G	G	C				

	45 47 B D F H J L N P	49 51 53	55 57 59	61 63 65	67 69 71	73 75
	44 46 A C E G I K M O	48 50 52	54 56 58 60	62 64 66	68 70 72	74 76
X408	A A C T	A T A A G G	T C G A A T	C T T T A	A A G T	G G C C T
X410	A T A	T T A G G T	G C A A C T	C C T A A	A T C C G C T A	
X425	T T A	T A T A C G	T C A A A T	C G T A T	T A T T G C T A	
X428	T T A	T A T A C G	T C A A A T	C G T A T	T A T T G C T A	
X432	A G A T	T G C A G G	T C G A A T	C C T G T	C C C G G C C T	
X434	A G A T	T G C A G G	T C G A A T	C C T G C	C C C G G C C A	
X436	A G A C	A G C A G G	T C G A A T	C C T G T	C C C G C C C T	
X442	A G A C	T G C A G G	T C G A A T	C C T G T	C C C C G C C T	
X445	A G A C	T G C A G G	T C G A A T	C C T G T	C C C C G C C T	
X450	T G G T G T T T T T C				T C T T A T T G	
X452	A G G T G G T T T T C				T C T T A T T A	
X459	T T A	T A A G G T	T A T A A T	C C T T T	C T T T T T T A	
X461	T T A	T A A G G T	T T T A A T	C C T T T	K C T T T T A	
X464	G A A T	G A A G G A	T A G A A A	C C T C C	C I C I G C T	
X466	G A A	T G A A G G A	T A A A A A	C C T C C	C T C T A C T T	
X468	A C A	T G T T G G T	T A A A C C	C C T T C	C T T T A C T A	
X470	C A A	T G T T G	T T A A A A T	C C T T C	C T T T A C T A	
X474	A A A	T G T T G G T	C T A A A C	C C T T C	C G T A T G A	
X475	A A A	T G T T G G T	T A A A	C C T T C	C G G A C T A	
X477	A A A	C G T T G G T	T T A A A T	C C T T C	C C G T A C T A	
X480	A A A	T G T T G G T	T T A T A T	C C T T C	C C G T A C T A	
X494	A A A	T G T T G G T	T A T A	C C T T C	C C G T A C T A	
X570	A T G T	C T C G G A	T G G A A A	C C G A G	C G G G C C T A	

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43							
X571	CAU	SACCHAROMYCES CER.	A	G	C	G	C	G	T	G	G	C	C	A	G	T	G	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A					
X575	CAU	SCHIZOSACCHA. POM.	T	G	C	G	C	G	G	T	A	G	G	A	G	T	G	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A			
X576	CAU	SCHIZOSACCHA. POM.	T	G	C	G	C	G	G	T	A	G	G	A	G	T	G	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A			
X635	CAU	GLYCINE MAX	A	T	C	A	G	A	T	G	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C		
X780	CAU	DROSOPHILA MELANO.	A	G	C	A	G	A	G	T	G	G	C	A	G	T	G	G	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A			
X830	CAU	XENOPUS LAEVIS	A	G	C	A	G	A	G	T	G	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G		
X831	CAU	XENOPUS LAEVIS	A	G	C	A	G	A	G	T	G	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G		
X950	CAU	MOUSE	A	G	C	A	G	A	G	T	G	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G		
X995	CAU	HUMAN	A	G	C	A	G	A	G	T	G	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G		
X996	CAU	HUMAN	A	G	C	A	G	A	G	T	G	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G		
P H E N Y L A L A N I N E																																																		
F145	GAA	METHANOCOCC. VANT.	G	C	C	A	A	G	G	T	A	G	T	C	A	G	C	T	G	G	G	A	G	A	C	G	T	G	G	A	C	T	G	A	A	G	A	T	C	C	A	G								
F160	GAA	SULFOLOBUS SOLFA.	G	C	C	G	C	G	T	A	G	C	T	C	A	G	C	C	G	G	G	A	G	A	C	G	C	C	G	G	T	G	A	A	G	A	C	C	G	G										
F203	GAA	MYCOPLASMA MYCOID.	G	G	T	C	G	C	T	A	G	C	T	C	A	G	T	C	G	T	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C		
F206	GAA	SPIROPLASMA MELIF.	G	G	T	C	A	G	T	A	G	C	T	C	A	G	T	G	T	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	
F235	GAA	BACILLUS SUBTILIS	G	G	T	C	G	G	T	A	G	C	T	C	A	G	T	G	T	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	
F236	GAA	BACILLUS SUBTILIS	G	G	T	C	G	G	T	A	G	C	T	C	A	G	T	G	T	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	
F250	GAA	E. COLI	G	C	C	G	G	A	T	A	G	C	T	C	A	G	T	C	G	T	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G
F304	GAA	EUGLENA GRACILIS	G	C	C	G	G	A	T	A	G	C	T	C	A	G	T	C	G	T	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G
F310	GAA	MARCHANTIA POLYM.	G	C	C	G	G	A	T	A	G	C	T	C	A	G	T	G	T	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	
F320	GAA	ZEA MAYS	G	T	C	A	G	G	A	T	A	G	C	T	C	A	G	T	G	T	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G
F335	GAA	NICOTIANA TABACUM	G	C	C	G	G	A	T	A	G	C	T	C	A	G	T	G	T	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	
F375	GAA	VICIA FABAE CHLORO	G	T	C	G	G	G	A	T	A	G	C	T	C	A	G	C	C	G	G	T	A	G	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G	C	A	G

45 47 B D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
 44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76

X571 A T G T C T C G G A T C G A A A C C G A G C G G G C C T A
 X575 A G G T C C A G G A T C G A A A C C T G G C G G C C A A
 X576 A G G T C C A G G A T C G A A A C C T G G C G G C C A A
 X635 A G G T C C A G G A T C G A A A C C T G G C I C I G A T A
 X780 A G G T C G A G G A T C G A A A C C T T G C I C T G C T A
 X830 A G G T G A T G G A T C G A A A C C A T T C T C T G C T A
 X831 A G G T G A T G G A T C G A A A C C A T C C T C T G C T A
 X950 A G G T G A T G G A T C G A A A C C A T C I C I G C T A
 X995 A G G T G A T G G A T C G A A A C C A T C I C I G C T A
 X996 A G G T G A T G G A T C T A A A C C A T C C I C I G C T A

P H E N Y L A L A N I N E

F145 T T G T C G G T G T C G A A T C A C C C C T T G G C A
 F160 T T G T C G G G G T C A A G T C C C G G G G G G C A
 F203 G T G T G G C G G T C A A T T C G T C A C G A C C A C C A
 F206 G T G T C G C A G T C A A T T C T G C C I G A C C A C C A
 F235 G T G T C G C G G T T C G A T T C C G T C C G A G C C A C C A
 F236 G T G T C G C G G T T C G A T T C C G T C C G A G C C A
 F250 G T G T C T T G G T T C G A T T C C G A G T C C G G G C A C C A
 F304 G T G T C A C C A G T T C A A A T C T G G T C C T A G C A
 F310 G T G T C A C C A G T T C A A A T C T G G T T C T G C C A
 F320 G T G T C A C C A G T T C A A A T C T G G T T C C T G G C A
 F335 G T G T C A C C A G T T C A A A T C T G G T T C C T G G C A
 F375 G T G T C A C C A G T T C A A A T C T G G T T C C G G G C G

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44															
F407	GAA	TETRAHYMENA	PYRIF.																																																								
F410	GAA	MITO.	SPERGILLUS	NIDUL.																																																							
F417	GAA	SACCHAROMYCES	CER.																																																								
F425	GAA	TORULOPSIS	GLAB.																																																								
F450	GAA	ASCARIS	SUUM																																																								
F453	GAA	ARTEMIA	SP.																																																								
F456	GAA	AEDIS	ALBOPICTUS																																																								
F459	GAA	DROSOPHILA	YAKUBA																																																								
F464	GAA	PARACENTROTUS	LIV.																																																								
F465	GAA	PARACENTROTUS	LIV.																																																								
F466	GAA	STRONGYLOCEM.	PURP.																																																								
F468	GAA	XENOPUS	LAEVIS																																																								
F469	GAA	XENOPUS	LAEVIS																																																								
F470	GAA	RANO	CATESBEIANA																																																								
F471	GAA	CEPHALORHYN.	COM.																																																								
F474	GAA	RAT																																																									
F477	GAA	MOUSE																																																									
F480	GAA	BOVINE																																																									
F482	GAA	GREEN	MONKEY																																																								
F494	GAA	HUMAN																																																									
F560	GAA	NEUROSPORA	CRASSA																																																								
F575	GAA	SCHIZOSACCHA.	POM.																																																								
F590	GAA	YEAST																																																									

	45	47	B	D	F	H	J	L	N	O	49	51	53	55	57	59	61	63	65	67	69	71	73	75															
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76														
F407	AGGT										C	A	T	T	G	G	T	C	C	G	A	T	C	C	T	C	T	A	A	G	G	C	A						
F410	AGGT										T	G	T	A	A	G	T	T	C	A	A	G	T	C	T	A	T	C	T	A	T	C	C	G	A	C	A		
F417	TTA										C	A	T	G	A	G	T	T	C	G	A	T	T	C	T	C	A	T	C	A	T	A	A	G	G	G	C	A	
F425	TTA										C	A	T	G	A	G	T	T	C	G	A	T	T	C	T	C	A	T	T	A	A	G	G	G	C	A	C	A	
F450	GGAAATGT										T	A	G	G	A	A	T	G	T																				
F453	GGTGGATGATTC										C	T	G	G	A	A	T	A	A	G	G	A	T	A	T	A	T	A	T	A	T	T	T	A	A	A	A	A	
F456	GGG										A	A	A	T	T	A	T	T	T	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	
F459	TGG										A	G	A	T	T	A	T	T	T	A	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	
F464	C A A										A	G	A	G	A	G	T	A	A	A	G	T	C	C	T	C	C	T	A	G	C	A	A	G	C	A	A	G	C
F465	C A A										A	G	A	G	A	G	T	A	A	A	G	T	C	C	T	C	C	T	A	G	C	A	A	G	C	A	A	G	C
F466	C A A										A	G	A	G	A	G	T	A	A	A	G	T	C	C	T	C	C	T	A	G	C	A	A	G	C	A	A	G	C
F468	A G A										T	G	A	G	C	C	T	A	G	A	A	G	C	T	C	G	A	A	G	C	A	A	G	C	A	A	G	C	A
F469	A G A										T	G	A	G	C	C	T	A	G	A	A	G	C	T	C	G	A	A	G	C	A	A	G	C	A	A	G	C	A
F470	A G A										T	G	A	C	C	T	A	A	A	A	G	T	C	T	A	G	G	G	T	A									
F471	A G A										T	G	G	C	A	T	T	A	C																				
F474	A G A										T	G	G	A	T	T	C	A	A																				
F477	A G A										T	G	G	A	T	T	G																						
F480	A G A										T	G	A	G	T	C	C																						
F482	A G A										T	G	A	G	T	C	C																						
F494	A G A										T	G	G	A	T	T	C	A	A																				
F560	AGGT										C	G	G	T	C	A	C	A	T	C	A	C	C	A	T	A	A	C	A	C	A	A	A	A	A	A	A	A	A
F575	TGGT										C	A	T	C	G	G	T	C	G	A	T	C	C	C	G	T	T	T	G	T	G	A	C	A					
F590	AGGT										C	T	G	T	G	T	C	G	A	T	C	C	A	C	A	G	A	A	T	C	C	G	A						

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44											
F591	GAA	YEA	ST																																																				
F780	GAA	DROSOPHILA	MELANO.																																																				
F830	GAA	XENOPUS	LAEVIS																																																				
F935	GAA	RAT																																																					
P R O L I N E																																																							
P020	UGG	PHAGE	T4																																																				
P040	UGG	PHAGE	T5																																																				
P145	UGG	METHANOCOC.	VANI.																																																				
P148	UGG	METHANOCOC.	VOLTAE																																																				
P203	UGG	MYCOPLASMA	MYCOID.																																																				
P206	UGG	SPIROPLASMA	MELIF.																																																				
P235	UGG	BACILLUS	SUBTILIS																																																				
P250	UGG	E.COLI																																																					
P251	CGG	E.COLI																																																					
P255	UGG	SALMONELLA	TYPHI.																																																				
P258	UGG	PHOTOBACT.	PHOSPH.																																																				
P259	UGG	AEROMONAS	HYDROPHI.																																																				
P304	UGG	EUGLENA	GRACILIS																																																				
P310	GGG	MARCHANTIA	POLYM. ^a																																																				
P311	UGG	MARCHANTIA	POLYM.																																																				
P315	UGG	TRITICUM	AESTIVUM																																																				
P320	UGG	ZEA	MAYS																																																				
P335	UGG	LOZANIA	TABACUM																																																				

	45	47	B	D	E	G	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75		
	44	46	A	C							O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76	
F591	AGGT																										
F780	AGGT																										
F830	AGGT																										
F955	AAGGT																										
P R O L I N E																											
P020	AGGT																										
P040	GGGT																										
P145	AGA																										
P148	AAA																										
P203	GGGT																										
P206	GGGT																										
P235	GGGT																										
P250	GGGT																										
P251	GGGT																										
P255	GGGT																										
P258	GGGT																										
P259	GGGT																										
P304	GGGT																										
P310	AAGT																										
P311	ATGT																										
P315	ATGT																										
P320	ATGT																										
P335	ATGT																										

	45	47	B	D	F	H	J	L	N	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76	
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76	

P342	ATGT	C	A	C	G	G	G	T	C	A	A	A	T	C	C	T	G	I	C	A	T	C	C	C	T	A
P355	ATGT	C	A	C	G	G	T	C	A	A	A	T	C	C	T	G	I	C	A	T	C	C	C	T	A	
P410	AAT	G	T	A	T	G	T	C	G	A	A	T	C	A	T	A	A	T	A	A	C	C	T	G	A	
P417	ACCT	A	G	T	T	A	G	T	C	G	A	G	T	C	T	A	T	C	T	A	T	C	T	G	A	
P425	ATC	T	A	G	T	A	G	T	C	G	A	G	T	C	T	A	T	C	T	A	T	C	T	G	A	
P442	GGC	C	A	T	A	G	G	T	C	G	A	A	T	C	C	T	G	I	C	C	T	G	A			
P445	ATGT	A	C	G	G	G	T	C	A	A	A	T	C	C	A	G	T	C	A	T	C	C	C	T	A	
P450	TGAAT	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	
P459	TGA	A	A	A	G	A	A	A	A	T	T	C	T	T	T	C	T	T	G	A						
P464	AGA	C	S	T	A	A	G	T	G	A	A	A	C	C	T	T	A	C	T	T	C	T	T	G	A	
P466	AATG	T	A	A	G	G	T	A	A	G	C	C	T	T	A	T	C	T	T	G	A					
P468	TAG	T	G	A	G	G	T	T	G	A	G	T	C	C	T	C	T	T	T	C	T	G	A			
P470	TGG	T	A	A	G	G	T	A	A	A	G	T	C	C	T	T	A	I	C	C	T	G	A			
P471	TGG	T	A	A	G	G	T	T	A	A	T	G	T	C	T	T	C	T	T	G	A					
P474	TGG	T	G	G	G	G	A	G	G	T	A	G	T	T	C	C	T	T	C	T	T	G	A			
P475	TGG	T	G	G	G	G	A	G	G	T	A	G	T	T	C	C	T	T	C	T	T	G	A			
P477	TGG	T	G	G	G	A	G	T	A	G	T	C	C	T	T	C	T	T	G	A						
P480	TGG	T	G	A	G	A	T	G	C		A	C	T	T	C	T	T	G	A							
P494	TGG	T	G	A	G	T	A	A	A	G	A	C	T	T	T	C	T	T	G	A						
P570	AGGC	C	C	T	G	G	G	T	C	A	A	T	C	C	C	A	G	T	C	G	C	C	C	C		
P645	AGGT	C	C	C	G	A	G	T	C	G	A	T	C	T	C	G	A	T	C	G	C	C	C	C		
P646	AGGT	C	C	C	G	A	G	T	C	A	A	T	C	T	C	G	A	T	C	G	C	C	C	C		
P700	AGGT	C	C	C	G	G	G	T	C	A	A	T	C	C	C	G	G	T	C	G	C	C	C	C		

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43									
P780	UGG	DROSOPHILA	MELANO.																																																	
P950	CGG	MOUSE																																																		
P951	AGG	MOUSE																																																		
P955	CGG	RAT																																																		
P956	AGG	RAT																																																		
P995	AGG	HUMAN																																																		
S E R I N E																																																				
S020	UGA	PHAGE T4																																																		
S040	UGA	PHAGE T5*																																																		
S041	GGA	PHAGE T5																																																		
S160	CGA	SULFOLOBUS	SOLFA.																																																	
S203	UGA	MYCOPLASMA	MYCOID.																																																	
S206	UGA	SPIROPLASMA	MELIF.																																																	
S232	CGA	LACTOBAC.	BULG.																																																	
S235	GGA	BACILLUS	SUBTILIS																																																	
S236	UGA	BACILLUS	SUBTILIS																																																	
S237	CGU	BACILLUS	SUBTILIS																																																	
S250	CGA	E-COLI																																																		
S251	UGA	E-COLI																																																		
S252	UGA	E-COLI*																																																		
S295	UGA	CYANOPHORA	PARAD.*																																																	
S304	CGU	EUGLENA	GRACILIS																																																	
S305	UGA	EUGLENA	GRACILIS																																																	

	45	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76											
	44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76											
P780	A	G	G	T							C	C	C	G	G	G	T	C	A	A	T	C	C	C	G	G	T	G	A	G	C	C				
P950	A	G	G	T							C	C	C	G	G	G	T	C	A	A	T	C	C	C	C	G	G	A	C	G	A	G	C	C		
P951	A	G	G	T							C	C	C	G	G	G	T	C	A	A	T	C	C	C	C	G	G	A	C	G	A	G	C	C		
P955	A	G	G	T							C	C	C	G	G	G	T	C	A	A	T	C	C	C	C	G	G	A	C	G	A	G	C	C		
P956	A	G	G	T							C	C	C	G	G	G	T	C	A	A	T	C	C	C	C	G	G	A	C	G	A	G	C	C		
P995	A	G	G	T							C	C	C	G	G	G	T	C	A	A	T	C	C	C	C	G	G	A	C	G	A	G	C	C		
SERINE																																				
S020	C	A	G	T	C	G	C	T	C	G	G	C	G	G	A	C	T																			
S040	C	C	G	C	T	G	T	G	A	T	A	C	G	G																						
S041	T	G	A	G	T	A	T	A	G	C	A	A	T	A	T	G	C	T																		
S160	T	G	C	C	T	C	C	C	A	G	G	C	A																							
S203	G	A	T	C	G	G	G	A	A	C	C	G	A	G																						
S206	C	A	G	C	G	G	T	G	A	A	G	C	C	G	G																					
S232	C	G	A	C	C	A	T	G	T	G	A	T	G	G	T	G	C																			
S235	T	A	G	C	G	T	C	A	C	T	C	C	G	T	C																					
S236	C	A	G	G	T	G	T	C	A	A	G	C	C	C	G																					
S237	T	A	G	C	G	T	A	G	C	C	G																									
S250	A	G	T	G	G	G	C	A	A	C	T	C	T	A	C																					
S251	C	G	A	C	C	G	A	A	G	G	G	T																								
S252	T	T	G	G	C	C	G	C	C	A	G	G	T	C	C	C	G																			
S295	T	T	T	A	G	G	T	T	T	C	C	C	T	A																						
S304	T	G	T	G	T	C	T	A	A	A	C	A	C																							
S305	C	G	T	A	G	T	T	T	G	C	T	A	C																							

	45	46	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75	76	
	44	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76	77	78	
S310	T	A	T	A	G	T	T	T	A	A	G	A	T	T	A	T	C	C	C	C	T	C	T	C	T	C	T
S311	T	G	T	A	C	A	A	G	C	T	T	T	T	G	T	A	C	C	C	C	T	C	T	C	T	C	C
S312	T	G	T	A	G	G	C	T	T	T	G	G	T	C	T	A	T	C	C	C	T	C	T	T	T	C	C
S313	T	A	T	A	G	T	T	C	T	A	G	G	A	A	C	T	A	T	C	C	T	C	T	C	T	C	T
S315	T	G	T	A	C	A	A	T	T	T	T	T	G	T	A	C	C	C	T	C	T	C	T	T	T	C	C
S320	T	A	T	A	G	T	T	C	T	A	G	G	A	A	C	T	A	T	C	C	C	T	C	T	T	T	C
S321	T	G	T	A	G	A	C	T	T	T	G	T	T	A	C	C	C	T	C	T	C	T	T	T	T	C	C
S322	T	G	T	A	C	A	T	T	T	T	T	G	T	A	C	C	C	T	C	T	C	T	T	T	T	T	C
S335	T	G	T	A	C	G	A	T	T	A	T	C	G	A	T	C	C	T	C	T	C	T	T	T	T	T	C
S336	T	G	T	A	G	G	C	T	T	T	G	T	T	A	C	C	C	T	C	T	C	T	T	T	T	T	C
S337	T	A	T	A	G	T	T	T	A	C	A	A	G	A	T	C	T	A	C	C	T	C	T	T	T	T	C
S355	T	A	T	A	G	T	C	G	A	A	C	A	A	G	A	M	T	C	T	A	C	C	T	T	T	T	C
S365	C	A	T	A	G	T	C	T	T	A	T	C	A	G	A	M	T	C	T	A	C	C	T	T	T	T	C
S366	T	G	T	A	G	G	C	T	T	T	G	T	T	A	C	C	C	T	C	T	C	T	T	T	T	T	C
S410	T	A	A	G	T	T	A	A	A	C	T	T	T	T	T	C	A	T	A	T	A	C	T	C	T	T	C
S411	G	T	G	C	G	T	T	G	C	A	C	A	C	A	C	A	C	A	T	A	C	T	C	T	T	T	C
S414	G	T	G	T	T	A	A	G	C	A	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	C	
S417	T	G	A	T	T	G	T	A	A	T	T	C	T	C	T	A	T	T	T	T	T	T	T	T	T	T	
S418	T	A	G	T	C	T	T	A	T	T	G	G	C	T	A	C	T	A	C	T	A	C	T	T	T	T	C
S425	T	A	G	T	T	A	A	T	A	A	C	T	A	C	T	A	C	T	A	C	T	A	C	T	T	T	C
S426	T	G	A	T	T	A	T	T	T	C	T	C	T	A	T	T	T	T	T	T	T	T	T	T	T	T	C
S436	T	G	T	A	G	G	C	T	T	T	G	T	T	A	C	C	C	T	C	T	C	T	T	T	T	T	C
S438	C	A	T	A	C	A	A	G	A	G	A	T	T	G	T	A	T	C	C	C	A	T	T	C	C	T	C

	45 47 B D F H J L M P	49 51 53 55 57 59 61 63 65 67 69 71 73 75	74 76
	44 46 A C E G I K M O	50 52 54 56 58 60 62 64 66 68 70 72	
S442	AGTATTGATAGGAATAC	C G G G G G T T C G A A T C C C T C T C C A T C C G	
S443	CATACAAGAAGATTGTAT	C A T G G G T T C G A A T C C C A T T C C T C C G	
S445	CATACAAGAAGATTGTAT	C A T G G G T T C G A A T C C C A T T C C C C C G	
S450	TTT	T G G A G A A T C C G I I G I T T	
S451	ATTT	*****	
S454	AGAA	A G A A G T T C A A T T C T T C T A T T A A C T T	
S456	TCT	T T A A T G G T T T A A T T C A T T T A T A T T C T	
S459	TTCTT	T T A A T G G T T T A A T T C A T T T A T A C T T C T	
S460	AGA	T A G A A T T A A T P I I C T A T T A A C T T	
S462	TAGA	G A A G G T T C G A A T C C I I T T I I C I C T T	
S463	AGAA	T C T G T A G T T C A A C T C T A T A G A A G T T C G	
S464	GGA	T A G A G G T T G A A T T C C T C T C T T C T C T T	
S465	AAGGC	T T G C G G T T C A A T C C C G T T G A G T T T C	
S466	TGA	T A G A G G T T T C T T C C T C C C C T C C T T	
S467	AAGAC	C T G C G G T T C G A A T C C G T T G A G I I T C	
S468	AC	G C T G T G T T C A T T C F A C G G C T I G I I C G	
S469	GTA	G G G G G T T C G A T T C C C T C T T T C T C G	
S474	TGT	A G G G G T T C G A A T C T T C C T T T C T A T	
S475	GCA	C C A T A C T C T A A A C A T A T G G C T T T C T A	
S476	TGT	A G G G G T C G A A T C C T T C C T T C T A T	
S477	AATTT	A G G G G T T C G A T C T T C C T T C T A T	
S478	GCT	T C C A T G T T A A A A C A T G G C T T T C T A	
S480	AGT	A G G G G T T C G A T T C C T T C C T T C T A	

	1	2	3	4	5	6	7	8	9	10	12	14	16	18	19	A	21	23	25	27	29	31	33	35	37	39	41	43	
S481	GCU	BOVINE	G A A A A A G T A T																										
S483	GCU	MACACA FUSCATA	G A G A A A A C T C																										
S484	GCU	MACACA MULATTA	G A G A A A A C T C																										
S485	GCU	MACACA FASCICULA	G A G A A A A C T C																										
S486	GCU	MACACA SYLVANUS	G A G A A A A C T C																										
S487	GCU	SALMIRI SCIUREUS	G A G A A A G T G C																										
S488	GCU	TARSIVUS SYRICHTA	A A G A A A G T A T																										
S489	GCU	LEMUR CATTIA	G A G A A A G T A A T																										
S490	GCU	CHIMPANZEE	G A G A A A A G C T T																										
S491	GCU	MITON	G A G A A A G C T C																										
S492	GCU	GORILLA	G A G A A A G C T C																										
S493	GCU	ORANG UTAN	G A G A A A G C T C																										
S494	UGA	HUMAN	T T G A A A A G T C A T G G A																										
S495	GCU	HUMAN	G A G A A A G C T C																										
S565	UGA	PODOSPORA ANSERIMA	G C C A G C A T G C A G A G T																										
S566	UGA	PODOSPORA ANSERIMA	G G C G C G A T G C C G A G T																										
S570	AGA	SACCHAROMYCES CER.	G G C A A C T T G C C G A G T																										
S571	GCU	SACCHAROMYCES CER.	G T C C C A G T G C C G A G T																										
S575	UGA	SCHIZOSACCHA.POM.	G T C A C T A T G T C C G A G T																										
S576	UGA	SCHIZOSACCHA.POM.	G T C A C T A T G T C C G A G T																										
S577	CGA	SCHIZOSACCHA.POM.	G T C A C T A T G T C C G A G T																										
S590	CGA	YEAST	G G C A C T A T G C C G A G T																										
S591	UGA	YEAST	G G C A C T A T G C C G A G T																										

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43						
S592	CGA	TEAST	G	G	C	T	A	C	A	T	G	C	C	G	A	G	T	G	G	T	A	A	G	G	C	G	A	G	A	C	T	C	G	A	A	T	C	T	C	T									
S780	AGA	DROSOPHILA MELANO.	G	C	A	G	T	C	G	T	G	C	C	G	A	G	C	G	G	T	A	A	G	G	C	G	T	C	T	G	A	C	T	A	G	A	A	T	C	A	G	A							
S781	CGA	DROSOPHILA MELANO.	G	C	A	G	T	C	G	T	G	C	C	G	A	G	C	G	G	T	A	A	G	G	C	G	T	C	T	G	A	C	T	A	G	A	A	T	C	A	G	A							
S785	CGA	DROSOPHILA STIMUL.	G	C	A	G	T	C	G	T	G	C	C	G	A	G	C	G	G	T	A	A	G	G	C	G	T	C	T	G	A	C	T	A	G	A	A	T	C	A	G	A							
S870	UCA	CHICKEN*	G	C	C	G	G	A	T	G	A	C	C	T	C	A	G	T	G	G	T	C	G	G	G	T	G	C	A	G	G	C	T	T	C	A	A	C	C	T	G	T							
S995	UGA	HUMAN	G	T	A	G	T	C	G	T	G	C	C	G	A	G	C	G	G	T	A	A	A	G	G	C	G	A	T	G	G	A	C	T	T	G	A	A	T	C	C	A	T						
S996	AGA	HUMAN	G	T	A	G	T	C	G	T	G	C	C	G	A	G	C	G	G	T	A	A	A	G	G	C	G	A	T	G	G	A	C	T	T	G	A	A	T	C	C	A	T						
S997	AGA	HUMAN	G	T	A	G	T	C	G	T	G	C	C	G	A	G	C	G	G	T	A	A	A	G	G	C	G	A	T	G	G	A	C	T	T	G	A	A	T	C	C	A	T						
T H R E O N I N E																																																	
T020	UGU	PHAGE T4	G	C	T	G	A	T	T	A	G	C	T	C	A	G	T	A	G	G	T	A	A	G	A	G	C	A	C	C	T	C	A	T	T	G	T	A	T	G	A	G	G						
T040	UGU	PHAGE T5	G	T	C	C	T	A	A	G	C	A	T	T	G	T	G	G	T	A	A	A	G	A	G	C	A	G	T	T	G	C	T	T	G	T	A	A	G	C	A	T	C						
T145	UGU	METHANOCOCC. VANI.	G	C	C	T	C	G	G	T	G	C	T	C	A	G	C	T	G	G	T	A	A	A	G	C	C	G	C	T	G	A	C	T	T	G	T	A	A	T	C	A	G	G					
T146	GGU	METHANOCOCC. VANI.	G	C	C	T	C	A	G	T	G	C	T	C	A	G	C	T	G	G	T	A	A	A	G	C	C	T	G	C	T	G	A	C	T	T	G	T	A	A	G	C	A	G	G				
T148	UGU	METHANOCOCC. VOLTAE	G	C	C	T	C	A	G	T	G	C	T	C	A	G	C	T	G	G	T	A	A	A	G	C	C	T	G	A	C	T	T	G	T	A	A	T	C	A	G	G							
T200	AGU	MYCOPLASMA CAPRIC.	G	C	T	G	A	C	T	T	A	G	C	T	C	A	G	T	T	G	G	T	A	A	A	G	A	G	C	A	A	T	T	G	A	C	T	A	A	T	C	A	A						
T203	UGU	MYCOPLASMA MYCOID.	G	C	T	G	A	C	T	T	A	G	C	T	C	A	G	C	A	G	G	C	A	A	A	G	A	G	C	A	C	T	T	G	A	C	T	T	A	A	T	C	A	G					
T235	GGU	BACILLUS SUBTILIS	G	C	T	C	C	A	T	A	G	C	T	C	A	G	C	A	G	C	A	G	G	T	A	A	A	G	A	G	C	A	C	T	T	A	A	G	G	A	G								
T236	UGU	BACILLUS SUBTILIS	G	C	C	G	G	T	A	A	G	C	T	C	A	A	T	G	G	T	A	A	A	A	G	A	G	C	A	C	T	T	G	A	C	T	T	A	A	T	C	A	G						
T237	UGU	BACILLUS SUBTILIS	G	C	C	G	G	T	A	A	G	C	T	C	A	A	T	G	G	T	A	A	A	A	A	G	A	G	C	A	C	T	T	G	A	C	T	T	A	A	T	C	A	G					
T250	GGU	E. COLI	G	C	T	G	A	T	A	G	C	T	C	A	G	T	T	G	G	T	A	A	A	A	A	G	A	G	C	C	C	T	T	G	T	A	G	G	G	T	G								
T251	GGU	E. COLI	G	C	T	G	A	T	A	G	C	T	C	A	G	T	T	G	G	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
T252	UGU	E. COLI	G	C	C	G	A	C	T	A	A	G	C	T	C	A	G	T	A	A	G	G	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
T253	CGU	E. COLI	G	C	C	G	A	T	A	A	G	C	T	C	A	G	T	T	G	G	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A			

45	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75		
44	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76	

S592		GGGCTCTGCCCG																							
S780		TTCCCTCTGGGAG																							
S781		TTCCCTCTGGGAG																							
S785		TTCCCTCTGGGAG																							
S870		AGCTGTCTAGCGACA																							
S995																								
S996		TGGGGTTCCCCA																							
S997		TGGGGTCTCCCG																							

T H R E O N I N E																									
T020		ATGT																							
T040		TGA																							
T145		TGGT																							
T146		AGGC																							
T148		TGGT																							
T200		AGGT																							
T203		AGGT																							
T235		AGGT																							
T236		AGGT																							
T237		AGGT																							
T250		GGT																							
T251		AGGT																							
T252		AGGT																							
T253		AGGT																							

	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43			
T260	CGU	PSEUDOMONAS	AER.	G	C	C	G	G	A	T	A	G	C	T	A	G	C	T	A	G	C	T	A	T	C	G	T	A	T	G	A	T	G	A	G	A		
T261	GGU	PSEUDOMONAS	AER.	G	C	T	C	A	G	T	G	T	A	G	C	T	A	G	C	T	A	G	C	T	A	T	C	G	T	A	T	G	A	G	G	T	G	
T304	UGU	EUGLENA	GRACILIS	G	C	C	T	T	T	A	G	C	T	C	A	G	T	G	T	A	G	C	C	T	T	G	G	T	A	G	G	T	A	G	G	T	G	
T310	GGU	MARCHANTIA	POLYM.	G	C	C	T	T	T	A	A	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G	
T311	UGU	MARCHANTIA	POLYM.	G	C	C	T	T	T	A	A	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G	
T313	GGU	MORDELM	VULGARE	G	C	C	T	T	T	A	A	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G	
T315	GGU	TRITICUM	AESTIVUM	G	C	C	T	T	T	A	A	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G	
T320	UGU	ZEA	MAYS	G	C	C	A	C	T	T	A	G	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G
T335	UGU	NICOTIANA	TABACUM	G	C	C	G	C	T	T	A	G	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G
T336	GGU	NICOTIANA	TABACUM	G	C	C	T	T	T	A	A	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G	
T355	GGU	PISUM	SATIVUM	G	C	C	T	T	T	A	A	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G	
T365	GGU	SPINACIA	OLERACEA	G	C	C	C	T	T	A	A	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G	
T366	UGU	SPINACIA	OLERACEA	G	C	C	G	C	T	T	A	G	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G
T375	GGU	VICIA	FABA	G	C	C	T	T	T	A	A	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G	
T410	UGU	ASPENGIILLUS	NIDUL.	G	C	C	T	T	T	A	A	C	T	C	A	G	T	G	T	A	T	T	G	T	A	T	T	G	T	A	T	G	C	G	G	T	G	
T417	UAG	SACCHAROMYCES	CER. ²	G	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A
T418	UGU	SACCHAROMYCES	CER.	G	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A
T425	UGU	TORULOPSIS	GLAB.	G	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A
T426	UAG	TORULOPSIS	GLAB.	G	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A
T450	UGU	ASCARIS	SUUM	G	C	C	A	T	G	T	A	G	T	T	A	T	T	A	T	T	A	T	T	A	T	T	A	T	T	A	T	T	A	T	T	A	T	T
T452	UGU	CLETHRABDI	ELEG.	G	C	C	A	T	G	T	A	G	T	T	A	T	T	A	T	T	A	T	T	A	T	T	A	T	T	A	T	T	A	T	T	A	T	T
T459	UGU	DROSOPHILA	YAKUBA	G	T	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T
T464	UGU	PARACENTROTUS	LIV.	A	C	C	T	G	A	G	C	T	C	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A	T	A

45 47 B D F H J L M P	49 51 53	55 57 59	61 63 65	67 69 71	73 75	76
44 46 A C E G I K O	50 52	54 56 58	60 62 64 66	68 70 72	74	76
	C G G G G G	T C G A T T	C C T C T	A T C C G G C A	C C C A	
T260	A G G T					
	C A G C G G T	T C A A A T	C C G C T	C A T G A G C T	C C A	
T261	A G G T					
	C G T C G G T	T C G A A T	C C G C A A A A	G G C T		
T304	T G G T					
	C A T C G G T	T C A A A T	C T G A T A A A G G C C T			
T310	A A G T					
	C A T C G G T	T C G A C T	C G A T A G C G G C T			
T311	T G G T					
	C A T C G G T	T C A A A T	C C G A T A A A G G G C T			
T313	A A G T					
	C A T C G G T	T C A A A T	C C G A T A A A G G G C T			
T315	A A G T					
	C A T C G G T	T C A A A T	C C G A T A A A G G G C T			
T320	G G G T					
	C A T C G G T	T C A A A T	C C G A T A G C G G C T			
T335	T G G T					
	C A T C G G T	T C G A T T	C C G A T A G C G G C T			
T336	A A G T					
	C A T C G G T	T C A A A T	C C G A T A G C G C T			
T355	A A G T					
	C G T C G G T	T C A A A T	C C G A T A A G G G C T			
T365	A A G T					
	C A T C G G T	T C A A A T	C C G A T A A G G G G C T			
T366	T G G T					
	C A T C G G T	T C G A G T	C C G A T A G C C G G C T			
T375	A A G T					
	C A T C G G T	T C A A A T	C C G A T A A G G G G C T			
T410	A T A					
	A C A A G T	G C G A T A	C T T G T A C T	G G G C T		
T417	T T A					
	C T A A G T	T C A A A T	C T T A G T A T T A C A			
T418	A G G T					
	T G G G G T	T C A A A T	C C C T A T A T A A C A			
T425	A G G T					
	T G G G G T	T C A A A T	C C C T A T A T A A C A			
T426	A T T A					
	T C T A A G T	T C A A A T	C T T A G T A T T T A C A			
T450	G G T A G T					
T452	A G G T T T C					
T459	A A A					
	T A A G A T T A	T T C T T T A A A C T				
T464	G A G					
	A G A G G G T A A A G T	C C C T C T C A A G G C T				

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42				
T466	UGU STRONGYLOCE	G	C	C	T	T	G	A	A	G	C	T	T	G	G	T	T	G	G	T	A	A	C	C	A	A	C	C	A	A	C	C	A	A	C	C	A	A	C	C	A	A				
T468	UGU XENOPUS LAEVIS	G	T	C	T	G	A	T	A	G	C	T	T	A	A	T	T	A	A	G	C	A	T	C	G	G	T	C	T	T	G	T	A	A	G	C	C	G	A	A	C	C	G			
T470	UGU RANA CATESBEIANA	G	C	C	T	G	A	T	A	G	C	T	T	A	A	C	T	A	A	A	A	A	C	C	G	G	T	C	T	T	G	T	A	A	G	C	C	G	A	A	C	C	G			
T471	UGU CERALORHYN.COM.	G	C	T	T	G	A	T	A	A	C	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A			
T474	UGU RAT	G	T	C	C	G	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A			
T475	UGU RAT	G	T	C	C	G	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A			
T477	UGU MOUSE	G	T	C	T	T	G	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
T480	UGU BOVINE	G	T	C	T	T	G	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
T494	UGU HUMAN	G	C	C	T	T	G	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A		
T780	AGU DROSOPHILA MELANO.	G	G	C	C	C	T	G	C	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	
T995	UGU HUMAN	G	G	C	C	A	T	A	G	C	T	A	G	G	T	A	G	G	T	A	G	G	T	A	G	G	T	A	G	G	T	A	G	G	T	A	G	G	T	A	G	G	T	A	G	
T R Y P T O P H A N																																														
W115	CCA HALOBACTERIUM MED.	G	G	G	G	C	T	G	T	G	C	C	A	A	G	C	C	G	G	C	A	T	G	A	C	T	G	A	C	T	C	C	A	A	T	C	A	G	T	C	A	G	T	C	A	
W120	CCA HALOBACTERIUM VOL.	G	G	C	T	G	T	G	C	C	A	A	G	C	C	G	C	C	G	C	A	T	G	A	C	T	G	A	C	T	C	C	A	A	T	C	A	G	T	C	A	G	T	C	A	
W200	UCA MYCOPLASMA CAPRIC.	A	G	G	G	C	A	T	A	G	T	T	C	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T
W201	CCA MYCOPLASMA CAPRIC.	A	G	G	A	G	T	A	G	T	T	C	A	A	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	A	G	T	
W235	CCA BACILLUS SUBTILIS	A	G	G	G	C	A	T	A	G	T	T	A	A	C	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	
W250	CCA E. COLI	A	G	G	G	C	C	T	A	G	T	T	C	A	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	
W304	CCA EULENA GRACILIS	G	C	C	T	T	A	T	A	G	T	T	C	A	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	
W310	CCA MARCHANTIA POLYM.	G	C	C	T	T	A	T	A	G	T	T	C	A	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	
W315	CCA TRITICUM AESTIVUM	G	C	C	T	T	A	T	A	G	T	T	C	A	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	
W320	CCA ZEA MAYS	G	C	C	T	T	A	T	A	G	T	T	C	A	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	
W335	CCA NICOTIANA TABACUM	G	C	C	T	T	A	T	A	G	T	T	C	A	A	T	A	T	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	A	

45 47 B D F H J L M P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
 44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76

T466 G A G A G A G G T A A A C T C C C T C C A A G G C C T
 T468 A G A T G A G G C T A A A A C C T C C T C A A G A C T
 T470 A G A C T G A G C T A A C C C T A C T C A A G G C T
 T471 A A A A G G A A C C C C A T T C C C C T A A G A C T
 T474 A A A T G A A G A T C A G T C T T T C T C A G G A C A
 T475 A A A T G A A G A G T C A G N T C T T C T C G G G A C A
 T477 A A A T A A G A T C T T C T T C C A A G A C A
 T480 A G A A G G A A C A A C T A C C T C C T A A G A C T
 T494 A G A T G A A A C C T T T T C C A A G G A C A
 T780 A G A T C G T G A G T C G A A T C T C G C E G G G G C C T
 T995 G G G T C G C G A G T C A A T T C T C G C T G G G G C T T

TRYPHTOPHAN

W115 C G A T C G G G G T C A A A T C C C T C G G C C C C A
 W120 C G A T C G G G G T C A A A T C C C T C G G C C C C A
 W200 G T G T C A C G A G T C G A G T C T T G T G C C C T G C C A
 W201 G C G T T G A G G G T C G A T T C C T T C T C C T G C C A
 W235 C G G T G T G G G T C G A T T C T A C T G C C C C T G C C A
 W250 G T G T T G G A G T T C G A G T C T C T C G C C C T G C C A
 W304 A T G T A G T A G G T C G A A T C C T A C A G A G C G G
 W310 A T G T C G T A G G T C A A A T C C T A C A G A G C G T G
 W315 A T G T C G T A G G T C A A A T C C T A C A G A G C G T G
 W320 A T G T C G T A G G T T C A A A T C C T A C A G A G C G T G
 W335 A T G T C G T A G G T T C A A A T C C T A C A G A G C G T G

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42			
U442	CCA	GENOTHERA SP.	G	C	G	C	T	C	T	A	G	T	C	A	G	T	C	G	G	T	A	G	A	C	G	C	G	G	G	T	C	C	A	A	A	C	C	C	C	G					
U455	CCA	CHILORO	G	C	G	C	T	C	T	T	G	T	T	C	A	G	T	C	G	G	T	A	G	A	C	G	C	G	G	G	T	C	C	A	A	A	C	C	C	G					
U402	CCA	CHILAMYDOMO. BEINH.	A	G	A	G	G	T	G	T	A	G	T	A	A	C	G	C	A	A	G	A	C	G	C	A	A	G	A	G	T	C	C	A	A	A	C	C	C	G					
U406	UCA	PARAMECIUM TETRA.	A	G	G	G	A	G	T	A	G	T	T	A	A	C	G	G	N																										
U407	UCA	TETRAHYMENA PYRIF.	G	G	G	G	A	T	A	G	T	T	A	A	C	T	G	T	A	A	C	T	G	T	A	A	C	T	A	A	C	T	T	A	A	C	T	T	A	A	C	T	T		
U410	UCA	ASPERGILLUS NIDUL.	A	A	G	A	G	T	A	G	T	T	A	A	C	T	G	T	A	A	C	T	G	T	A	A	C	T	T	A	A	C	T	T	A	A	C	T	T	A	A	C	T	T	
U414	UCA	PODOSPORA ANSERINA	A	A	G	A	G	T	A	G	T	T	A	A	C	T	G	T	A	A	C	T	G	T	A	A	C	T	T	A	A	C	T	T	A	A	C	T	T	A	A	C	T	T	
U425	UCA	TORULOPSIS GLAB.	A	A	G	A	T	A	G	T	T	A	A	C	T	G	T	A	A	C	T	G	T	A	A	C	T	T	A	A	C	T	T	A	A	C	T	T	A	A	C	T	T		
U428	UCA	YEAST	A	A	G	A	T	A	G	T	T	A	A	C	T	G	T	A	A	C	T	G	T	A	A	C	T	T	A	A	C	T	T	A	A	C	T	T	A	A	C	T	T		
U436	CCA	GENOTHERA SP.	G	C	G	C	T	C	T	T	A	G	T	T	C	A	G	T	T	C	G	G	T	A	G	A	C	G	C	G	G	T	C	C	A	A	A	C	C	C	G				
U442	CCA	TRITICUM AESTIVUM	G	C	G	C	T	T	A	G	T	T	C	A	G	T	T	C	G	G	T	A	G	A	C	G	C	G	G	T	C	C	A	A	A	C	C	C	G						
U445	CCA	ZEA MAYS*	G	C	G	C	T	T	A	G	T	T	C	A	G	T	T	C	G	G	T	A	G	A	C	G	C	G	G	T	C	C	A	A	A	C	C	C	G						
U450	UCA	ASCARIS SUUM	A	C	A	G	A	T	T	A	G	T	T	A	A	C	T	G	T	A	A	C	T	G	T	T	G	T	T	T	C	C	A	A	A	C	C	A							
U457	UCA	DROSOPHILA MELANO.	A	A	G	G	C	T	T	A	G	T	T	A	A	C	T	A	A	C	T	A	A	C	T	T	A	A	C	T	T	C	A	A	A	C	C	T	A						
U459	UCA	DROSOPHILA YAKUBA	A	A	G	C	T	T	A	G	T	T	A	A	C	T	A	A	C	T	A	A	C	T	T	A	A	C	T	T	C	A	A	A	C	C	T	A							
U464	UCA	PARACENTROTUS LIV.	A	A	G	A	G	T	A	A	A	T	A	A	C	T	G	T	A	A	C	T	G	T	A	A	C	T	T	C	A	A	A	C	C	T	A								
U466	UCA	STRONGYLOXEN.PURP.	A	A	G	A	G	C	T	A	G	T	T	A	A	C	T	G	T	A	A	C	T	G	T	A	A	C	T	T	C	A	A	A	C	C	T	A							
U468	UCA	XENOPUS LAEVIS	A	G	A	G	A	T	T	A	G	T	T	A	A	C	T	A	A	C	T	G	T	A	A	C	T	T	A	A	C	T	T	C	A	A	A	C	C	T					
U470	UCA	RANA CATESBEIANA	A	G	A	A	C	T	T	A	G	G	T	A	A	C	T	A	A	C	T	A	A	C	T	T	A	A	C	T	T	C	A	A	A	C	C	T							
U474	UCA	RAT	A	G	A	G	T	T	A	G	G	T	A	T	A	C	T	A	T	A	C	A	G	A	C	T	T	A	A	C	T	T	C	A	A	A	C	C	T						
U475	UCA	MITO	A	G	A	C	T	T	A	G	G	T	A	T	A	C	T	A	T	A	C	A	G	A	C	T	T	A	A	C	T	T	C	A	A	A	C	C	T						
U477	UCA	MOUSE	A	G	A	G	T	T	A	G	G	T	A	T	A	C	T	A	T	A	C	A	G	A	C	T	T	A	A	C	T	T	C	A	A	A	C	C	T						
U480	UCA	BOVINE	A	G	A	A	T	T	A	G	G	T	A	A	C	T	A	A	C	T	A	A	C	T	T	A	A	C	T	T	C	A	A	A	C	C	T								

	45	46	47	B	D	F	H	J	L	M	P	49	51	53	55	57	59	61	63	65	67	69	71	73	75					
	44	45	46	A	C	E	G	I	K	M	O	48	50	52	54	56	58	60	62	64	66	68	70	72	74	76				
W342	A	T	G	T	C	T	A	G	G	T	T	C	A	A	A	T	C	C	T	A	C	A	G	A	G	C	C	T	G	
W355	A	T	G	T	C	T	A	G	G	T	T	C	A	A	A	T	C	C	T	A	C	A	G	A	G	C	C	T	G	
W402	A	A	A	T	C	G	A	G	G	T	T	C	G	A	G	T	C	C	T	T	C	A	T	C	T	C	T	G		
W406	C	A	T	C	G	A	G	G	T	T	C	G	A	C	T	C	C	C	C	T	T	C	C	C	T	T	G			
W407	T	A	G	C	T	G	A	G	G	T	T	C	G	A	A	T	C	C	T	G	C	T	T	C	C	C	T	G		
W410	A	T	T	C	T	A	G	T	T	C	A	A	A	T	C	T	A	A	G	T	A	C	T	C	T	T	G			
W414	A	T	T	C	T	A	G	T	T	C	G	A	G	T	T	C	A	A	G	T	A	C	T	C	T	T	G			
W425	C	A	T	T	A	G	A	G	T	T	C	G	A	A	T	C	T	T	A	T	C	C	T	T	G					
W428	C	A	T	T	A	G	A	G	T	T	C	G	A	A	T	C	T	T	A	T	C	C	T	T	G					
W436	A	T	G	T	C	T	A	G	G	T	T	C	G	A	A	T	C	C	T	T	A	T	C	C	T	T	G			
W442	A	T	G	T	C	T	A	G	G	T	T	C	A	A	A	T	C	C	T	A	T	C	C	T	T	G				
W445	A	T	G	T	C	T	A	G	G	T	T	C	A	A	A	T	C	C	T	A	T	C	C	T	T	G				
W450	A	A	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T				
W457	A	A	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T				
W459	A	A	A	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T	T				
W464	T	T	A	T	A	G	A	A	T	G	G	A	A	T	G	G	A	A	T	T	A	G	T	T	A	G	T			
W466	A	A	C	A	A	G	A	A	T	G	G	A	A	T	G	G	A	A	T	T	A	G	T	T	A	G	T			
W468	A	A	G	A	A	G	A	A	T	G	G	A	A	T	G	G	A	A	T	T	A	G	T	T	A	G	T			
W470	A	A	G	A	A	G	A	A	T	G	G	A	A	T	G	G	A	A	T	T	A	G	T	T	A	G	T			
W474	T	A	G	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C			
W475	T	A	G	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C			
W477	A	A	G	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C			
W480	A	A	G	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C	A	A	C			

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42									
W494	UCA	HUMAN	AGAAAATTA	GGTTAAATAC	AGACCA	AGAGCCTTCAAAGCCCT																																													
W555	CCA	DICTYOSTELIUM DIS. MITO	GACTCCTAG	GCATAGT	GGTTAT	TGTAAAT	GTCTCCAAGCAGT																																												
W570	CCA	SACCHAROMYCES CER.	GAGCGGTG	GCACAAT	GGT	AGAGCT	TCCGACTCCAAATCGAA																																												
W700	CCA	CAENORHABDI.ELEG.	GACTGCTG	CGCAAT	GGT	AGCGCG	TTCGACTCCAGATCGAA																																												
W870	CCA	CHICKEN	GACCTCGT	GCGCAC	GGT	AGCGCG	TCTGACTCCAGATCAGA																																												
T Y R O S I M E																																																			
Y145	GUA	METHANOCOCC.VANI.	CCCGCA	AGTTCAGATTGGT	AGAACG	CCCGACTGTAGATCCCGC																																													
Y148	GUA	METHANOCOCC.VOLTAE	CCCGCA	AGTTCAGACTGGT	AGAACG	CCCGACTGTAGATCCCGC																																													
Y200	GUA	MYCOPLASMA CAPRIC.	GAGGGGT	AGCGAAGT	GGCTAA	CGCGGTGTAAACCACT																																													
Y235	GUA	BACILLUS SUBTILIS	GAGGGGT	AGCGAAGT	GGCTAA	CGCGGTGTAAATCCGC																																													
Y250	GUA	E.COLI	GGTGGGT	TCCCGAGC	GGCCAA	GGGAGCAGACTGTAAATCTGC																																													
Y251	GUA	E.COLI	GGTGGGT	TCCCGAGC	GGCCAA	GGGAGCAGACTGTAAATCTGC																																													
Y260	GUA	PSEUDOMONAS AER.	GAGGCA	TCCCTAGT	GCC	AAAGGATCAGACTGTAAATCTGA																																													
Y304	GUA	EULENA GRACILIS	GAGTGT	TCCCGAAGT	GGTAA	TGGGCGCGACTGTAAATCCGC																																													
Y310	GUA	MARCHANTIA POLYM. CHLORO	GGTCGA	TCTCGAGT	GGTAA	TGGGACCGACTGTAAATCCGC																																													
Y315	GUA	TRITICUM AESTIVUM CHLORO	GGTCGA	TCCCGAGT	GGTAA	TGGGACCGACTGTAAATCCGT																																													
Y335	GUA	NICOTIANA TABACUM CHLORO	GGTCGA	TCCCGAGC	GGTAA	TGGGACCGACTGTAAATCCGT																																													
Y355	GUA	PHASEOLUS SATIVUM CHLORO	GCTCA	TCCCGAGC	GGTAA	TGGGACCGACTGTAAATCCGT																																													
Y365	GUA	SPINACIA OLERACEA CHLORO	GGTCGA	TCCCGAGC	GGTAA	TGGGACCGACTGTAAATCCGT																																													
Y375	GUA	VICIA FABA CHLORO	GGTCGA	TCCCGAGC	GGTAA	TGGGACCGACTGTAAATCCGT																																													
Y405	GUA	PARAMECIUM PRIM. MITO	GAGTAA	TGCTGAGT	GGT	AAAGCGCAGACTGTAAATCTGT																																													
Y406	GUA	PARAMECIUM TETRA. MITO	GAGTAA	TGCTGAGT	GGT	AAAGCGCAGACTGTAAATCTGT																																													
Y408	GUA	TRITRYMENA THERM. MITO	GAGGTGGT	GCTGAT	GGT	AAAGCGGTAGACTGTAAATCTAT																																													

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45 47 B D F H J L M P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76
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W494 C A G T A A G T T G C A A T A C T T A A T T C T G
W555 T G G T C A G G G T T C A A C T C C C T G A G G G T C T
W570 G G G T T G C A G G T T C A A T T C C T G T C C G T T C A
W700 A G G T T G G G C G T T C G A T C C G C T C A G T G G T C A
W870 A G G C T G C G T G T T C G A A T C A C G T C G G G G T C A
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TYROSINE
Y145 A T G T C C T G G T T C A A A T C G G C T C G C G G G A
Y148 A T G T C C T G G T T C A A A T C G G C T C G C G G G A
Y200 T C C T A C G G T T C G G G G T T C G A A T C C T C C C C T C C A C C A
Y235 T C C C T C A G G G T T C G C A G T T C G A A T C T G C C C C T C C A C C A
Y250 C G T C A C A G A C T T G A A G G T T C G A A T C C T T C C C C A C C A C C A
Y251 : : : : : C G T C A T C G A C T T C A A G G T T C G A A T C C T T C C C C A C C A C C A
Y260 C G T C A T A G A C T T C A A G G T T C G A A T C C T T C C C T C C A C C A
Y304 A G T C A T C T T C C T G G T T C G A A T C C A G C A C G A C T C A
Y310 T G G C A A T G C C T A C C T G G T T C A A A T C C A G C T C G A C C C A
Y315 T G A C A A T G T C T A C C T G G T T C A A A T C C A G C T C G G C C C A
Y335 T G G C A A T A T G T C T A C C T G G T T C A A A T C C A G C T C G G C C C A
Y355 T G G C A A T A T G T C T A C C T G G T T C A A A T C C A G C T C G G C C C A
Y365 T G G C A A T A T G T C T A C C T G G T T C A A A T C C A G C T C G G C C C A
Y375 T G G C A A T A T G T C T A C C T G G T T C G A A T C C A A C T T A C T T C A
Y405 T G G T A G T A C C G T : : : : : C G T G G T T C G A A T C C A A C T T A C T T C A
Y406 T G G T A G T A C C G T : : : : : C G T G G T T C G A A T C C A A C T T A C T T C A
Y408 T G G G A T T C C G T : : : : : C G C G G T T C G A T T C C G G T C C A C C T C A

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	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42														
Y410	GUA	ASPERGILLUS	MIDUL.																																																					
Y417	GUA	SACCHAROMYCES	CER.																																																					
Y418	GUA	SACCHAROMYCES	CER.																																																					
Y425	GUA	TORULOPSIS	GLAB.																																																					
Y442	GUA	TRITICUM	AESTIVUM																																																					
Y450	GUA	XENOPUS	SUM																																																					
Y459	GUA	DROSOPHILA	MELANO.																																																					
Y464	GUA	DROSOPHILA	YAKUBA																																																					
Y466	GUA	PARACENTROTUS	LIV.																																																					
Y468	GUA	STRONGYLOEN.	PURP.																																																					
Y470	GUA	XENOPUS	LAEVIS																																																					
Y474	GUA	RANA	CATESBEIANA																																																					
Y477	GUA	RAT	MITO																																																					
Y480	GUA	MOUSE	MITO																																																					
Y494	GUA	BOVINE	MITO																																																					
Y570	GUA	HUMAN	MITO																																																					
Y615	GUA	SACCHAROMYCES	CER.																																																					
Y655	GUA	ARABIDOPSIS	THAL.																																																					
Y780	GUA	NICOTIANA	RUSTICA																																																					
Y850	GUA	DROSOPHILA	MELANO.																																																					
Y851	GUA	XENOPUS	LAEVIS																																																					
Y995	GUA	XENOPUS	LAEVIS																																																					
Y995	GUA	HUMAN	MITO																																																					

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43					
1996	GUA	HUMAN	CCT	CGA	TAG	CTC	AGT	AGCT	GGT	AGAGCG	GAGGACT	GTAGA	TCC	TT																																		
			V A L I N E																																													
V040	UAC	PHAGE T5	GCT	CGGT	TAG	TAA	TAA	T	GGG	AGA	ACC	CCGCC	TT	TAC	AC	CG	GG	CGG																														
V145	UAC	METHANOC. VANI.	GG	ACT	CA	TGG	TCT	AG	TTGG	AT	GAC	TG	CCC	TT	TAC	AA	GG	CGA																														
V160	GAC	SULFOLOBUS SOLFA.	GG	CC	CG	TCT	AG	CC	TGG	T	AG	AC	G	CT	G	AC	GC	GG	CGA																													
V203	UAC	MYCOPLASMA MYCOID.	GG	AG	TGG	TAG	CT	AG	CT	GG	AG	CT	GG	CT	TAC	AA	GC	AGG																														
V235	UAC	BACILLUS SUBTILIS	GG	AG	AT	AG	CT	AG	CT	GG	AG	CA	TG	CC	TT	TAC	AA	GC	AGA																													
V250	UAC	E. COLI	GG	GT	GA	TAG	CT	AG	CT	GG	AG	CA	TG	CC	TT	TAC	AA	GC	AGA																													
V304	UAC	EUGLENA GRACILIS	GG	GG	GT	ATA	AG	CT	AG	TT	GG	AG	GC	T	CG	CT	TAC	AA	GC	AGA																												
V310	GAC	MARCHANTIA POLYM. CHLORO	AG	GG	ATA	ACT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	TG	AC	GC	GTG	GG																											
V311	UAC	MARCHANTIA POLYM. CHLORO	AG	GG	CT	ATA	AG	CT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																											
V313	UAC	CHLORON VULGARE	AG	CC	CT	ATA	AG	CT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																											
V320	UAC	ZEA MAYS	AG	GG	CT	ATA	AG	CT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																											
V321	GAC	ZEA MAYS	AG	GG	ATA	ACT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																													
V327	GAC	GLYCINE MAX	AG	GG	ATA	ACT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																													
V335	GAC	NICOTIANA TABACUM	AG	GG	ATA	ACT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																													
V336	UAC	NICOTIANA TABACUM	AG	GG	CT	ATA	AG	CT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																											
V355	GAC	PISUM SATIVUM	AG	GG	CT	ATA	AG	CT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																											
V360	GAC	STIMOPSIS ALBA	AG	GG	ATA	ACT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																													
V365	GAC	SPINACIA OLERACEA	AG	GG	ATA	ACT	CAG	C	GG	AG	TA	TCA	CC	T	G	AC	GC	GTG	GG																													
V410	UAC	ASPERGILLUS NIDUL. MITO	AA	GA	AA	TAG	CT	AG	TT	GG	AG	GC	T	CG	T	TAC	AC	AC	GA																													
V414	UAC	PODOSPORA ANSERINA	AA	GA	AA	TAG	CT	AG	TT	GG	AG	GC	T	CG	T	TAC	AC	AC	GA																													
V417	UAC	ACCHAROMYCES CER. MITO	AG	GA	GA	TAG	CT	AG	TT	GG	AG	GC	T	CG	T	TAC	AC	AC	GA																													

45 47 8 D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
 44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76

Y996	AGGT	CGCTGGTTCGATTCCGGCTCGAAGGA

	VALINE	
V040	TTG	TGATAGTTCGATTCTAICACCGAGTACCA
V145	GGGT	GCCGGTTCGAATCCGGTGGGTCCA
V160	AAAT	CTGGGTTCAAAGTCCAGCGGGCCA
V203	CGGT	CATAGGTTCAAAGTCCATACACTCCACCA
V235	GGGT	GCGGGTTCGAGCCGGTCACTCCACCA
V250	GGGT	CAGGGTTCGATCCGTCATACCCACCA
V304	ATGT	CAGGGTTCGAATCCGTTCGCTCA
V310	AAGT	CATCAGTTCGAACCTGATATCCCTA
V311	ATGT	CTACGGTTCAAATCCGTAAGCCCTA
V313	AGGT	CTACGGTTCGAGTCCGTAAGCCCTA
V320	AGGT	CTACGGTTCGAATCCGTAAGCCCTA
V321	AAGT	CATCAGTTCGAGCCGATATCCCTA
V327	AAGT	TATCAGTTCGAGCCGATATCCCTA
V335	AAGT	CATCAGTTCGAGCCGATATCCCTA
V336	AGGT	CTACGGTTCGAGTCCGTAAGCCCTA
V355	AAGT	CATCAGTTCGAGCCGATATCCCTA
V360	AAGT	CTACGGTTCGAACCTGATATCCCTA
V365	AAGT	CATCAGTTCGAGCCGATATCCCTA
V410	GGT	CATCAGTTCGAGCCGATATCCCTA
V414	AGGC	CAGGTGTCAAATCACCTATTCTTACCA
V417	AGAT	TATGAGTTCGAATCTCCTATTCTTACCA

		TATAGGTCGACCTATATTCTTACCA

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V418	UAC	SACCHAROMYCES	CER.																																																				
V425	UAC	TORULOPSIS	GLAB.																																																				
V450	UAC	ASCARIS	SUUM																																																				
V456	UAC	AEDES	ALBOPICTUS																																																				
V459	UAC	DROSOPHILA	YAKUBA																																																				
V464	UAC	MICENTROTUS	LIV.																																																				
V466	UAC	STRONGYLOEN.	PURP.																																																				
V468	UAC	XENOPUS	LAEVIS																																																				
V474	UAC	RAT																																																					
V477	UAC	HOUSE																																																					
V480	UAC	BOVINE																																																					
V494	UAC	HUMAN																																																					
V555	AAC	DICTYOSTELIUM	DIS.																																																				
V556	UAC	DICTYOSTELIUM	DIS.																																																				
V570	AAC	SACCHAROMYCES	CER.																																																				
V780	AAC	DROSOPHILA	MELANO.																																																				
V781	CAC	DROSOPHILA	MELANO.																																																				
V830	CAC	XENOPUS	LAEVIS																																																				
V995	AAC	HUMAN																																																					
V996	CAC	HUMAN																																																					

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45 47 B D F H J L N P 49 51 53 55 57 59 61 63 65 67 69 71 73 75
44 46 A C E G I K M O 48 50 52 54 56 58 60 62 64 66 68 70 72 74 76
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V418  A G A T      T A T A G G T T C G A A C C C T A T A T T C C T A
V425  A G A T      T A T A G G T T C G A A T C C T A T A T T C C T A
V450  G G G T T T A T
V456  A G A          A A T T T G T G C A A A T C A A T T T A A A T T G A
V459  A G A          T T T T T G T G C A A A T C A A T A T A A A T T G A
V464  T C A          C A C T C G T G C A A T T C G G G T G T C T T G A
V466  C C A          C A C T C G T G C A A T T C G A G T T G T T T G A A G
V468  C A A          T A T C T G T T A A C C C G G A T T A C T T T G A
V474  A G A          A T T C A T A A A   A T G A A C A C T T T G A
V477  A G A          T T C A T G A C C   A T G A A C A C T C T G A
V480  A G A          C T C A T T C A T   T A G A A T A C T T C A
V494  A G A          T T C A A C T T A   C T G A C C G C T C T G A
V555  A G G T          C G T G G G T T C G A T T C C C G C T C T G A A T
V556  A G G T          C T C G A G T T C G A T C T C G G T C G G A T C A
V570  A C G T          C C C C A G T T C G A T C T G G G C G A A A T C A
V780  A G G C          C C C G G T T C A A T C C C G G C G G A A A C A
V781  A G G T          C C C G G T T C G A A C C C G G C G A A A C A
V830  A G G T          C C C G G T T C G A A A C C G G C A G A A A C A
V995  A G G T          C C C G G T T C G A A A C C G G C G G A A A C A
V996  A G G T          C C C G G T T C G A A A C C G G C G G A A A C A

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