

Title	Stringency of start codon selection modulates autoregulation of translation initiation factor eIF5
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Supplementary Data

Supplementary Figure S1. Nucleotide sequences used for generating logograms.

Alignments of the 10 nucleotides surrounding the uAUG start codons of eIF5 that define their initiation contexts are shown for: (A) vertebrates; (B) arthropods; (C) nematodes; (D) molluscs; (E) Pezizomycotina; (F) Basidiomycota; (G) fungi other than Pezizomycotina or Basidiomycota; (H) plants. Species names are on the right. Species names in quotation marks specify presumed contaminant sequences that cluster with eIF5 sequences from an unrelated taxonomic group. Some plants have more than one paralog of eIF5. Here they are arbitrarily labelled “#1”, “#2”, “#3” or “#4” and their placement in one of the groups does not imply orthologous clusters. For each species the sequence on top is the context of the first uAUG and subsequent uAUG contexts are shown in descending order. The initiation codon is highlighted in green. Nucleotides in positions -3 and +4 are highlighted in red if they deviate from the Kozak consensus and in gray if they comply with it. Species names in quotation marks represent sequences derived from Expressed Sequence Tags that by similarity analysis of the main open reading frame do not match eIF5 homologs from the phylum to which the species belongs and instead likely belong to contaminating sequences of unknown species. These sequences are grouped with the taxonomic group to which they actually belong determined by reconstructed phylogenetic trees with the entire sequence.

A)

Vertebrata

GGG	UUU	AUGU	Homo sapiens
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Pan troglodytes verus
UCG	UUU	AUGU	
UUU	UUC	AUGU	
GGG	UUU	AUGU	Macaca fascicularis
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Macaca mulatta
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Pongo abelii
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Canis lupus familiaris
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Ursus americanus
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Equus caballus
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Peromyscus polionotus
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Bos taurus
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Sus scrofa
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Rattus norvegicus
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Mus musculus
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Cavia porcellus
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Ovis aries
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Oryctolagus cuniculus
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Taeniopygia guttata
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Gallus gallus
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Lonchura striata domestica
UUU	UUC	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Meleagris gallopavo
UUU	UUC	AUGU	

UCG	UUU	AUGU	
GGG	UUU	AUGU	Bungarus multicinctus
UUU	UCA	AUGU	
UCG	UUU	AUGU	
GGG	UUU	AUGU	Anolis carolinensis
UUU	UCA	AUGU	
UCG	UUU	AUGU	
GUG	UUU	AUGU	Xenopus tropicalis
AUU	UCA	AUGU	
UCG	UUU	AUGU	
GUG	UUU	AUGU	Xenopus laevis
AUU	UCA	AUGU	
UUG	UUU	AUGU	
GUG	UUU	AUGU	Ambystoma mexicanum
AUU	UCA	AUGU	
UUG	UUU	AUGU	
GUG	UUU	AUGU	Ambystoma tigrinum tigrinum
AUU	UCA	AUGU	
UUG	UUU	AUGU	
GUG	UUU	AUGU	Cynops pyrrhogaster
AUU	UCA	AUGU	
UUG	UUU	AUGU	
GUG	UUU	AUGU	Gasterosteus aculeatus
UUU	UCA	AUGU	
UCG	UUU	AUGU	
GUG	UUU	AUGU	Perca flavescens
UUU	UCA	AUGU	
UCG	UUU	AUGU	
GUG	UUU	AUGU	Miichthys miiuy
UUU	UCA	AUGU	
UCG	UUU	AUGU	
GUG	UUU	AUGU	Lates calcarifer
UUU	UCA	AUGU	
UCG	UUU	AUGU	
GUG	UUU	AUGU	Takifugu rubripes
UUU	UCA	AUGU	
UCG	UUU	AUGU	
GUG	UUU	AUGU	Siniperca chuatsi
UUU	UCA	AUGU	
UCG	UUU	AUGU	
UUG	UUU	AUGU	Osmerus mordax #1
UUU	UCA	AUGU	
UCG	UUU	AUGU	
GUG	UUU	AUGU	Osmerus mordax #2
CGACGGAUGG			
UUU	UCA	AUGU	
UUG	UUU	AUGU	
GUG	UUU	AUGU	Xiphophorus maculatus
UUU	UCA	AUGU	
UCG	UUU	AUGU	
GUG	UUU	AUGU	Gadus morhua
UUU	UCA	AUGU	
CCG	UUU	AUGU	
GUG	UUU	AUGU	Oryzias latipes
UUU	UCA	AUGU	
UUG	UUU	AUGU	
GUG	UUU	AUGU	Lipochromis sp.
AUAUUUAUGG			
UUU	UCA	AUGU	
UUG	UUU	AUGU	
UUG	UUU	AUGU	Salmo salar #1

UUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Salmo salar #2
CGAC	GGAUGG	
UUU	UCAUGU	
UCG	UUUAUGU	
UUG	UUUAUGU	Oncorhynchus mykiss #1
UUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Oncorhynchus mykiss #2
CGAC	GGAUGG	
UUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Oncorhynchus tshawytscha #1
UUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Oncorhynchus tshawytscha #2
CGAC	GGAUGG	
UUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Oncorhynchus nerka #2
CGAC	GGAUGG	
UUU	UCAUGU	
UCG	UUUAUGU	
UUG	UUUAUGU	Thymallus thymallus
UUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Esox lucius
UAAC	CAUGU	
UGG	CUAUGU	
AGU	JACAUGU	
AUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Misgurnus anguillicaudatus
UUU	UCAUGU	
UUG	UUUAUGU	
GUG	UUUAUGU	Pimephales promelas #1
UUU	UCAUGU	
UUG	UUUAUGU	
GUG	UUUAUGU	Pimephales promelas #2
CUAC	GGAUGG	
CUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Danio rerio #2
CUAC	GGAUGG	
CUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Ictalurus furcatus #1
UUU	UCAUGU	
UUG	UUUAUGU	
GUG	UUUAUGU	Ictalurus furcatus #2
CUG	GGAUGG	
AUU	UCAUGU	
UCG	UUUAUGU	
GUG	UUUAUGU	Ictalurus punctatus #1
UUU	UCAUGU	
UUG	UUUAUGU	
GUG	UUUAUGU	Ictalurus punctatus #2
CUG	GGAUGG	
AUU	UCAUGU	
UCG	UUUAUGU	

GUGUUUAUGU	Cynoglossus semilaevis
UUUUUCAUGU	
UCGUUUUAUGU	
GUGUUUAUGU	Squalus acanthias
UUUUUCAUGU	
UUGUUUAUGU	
GUGUUUAUGU	Leucoraja erinacea
UUUUUCAUGU	
UUGUUUAUGU	
GGUUUUUAUGU	Petromyzon marinus
AUUUUCAUGU	
GAUUUUUAUGU	Eptatretus burgeri
AUUUUCAUGU	
CAGUUUAUGU	

B)

Arthropoda

UUUUUAUAUGA	Tribolium castaneum
AGGUGUAUGA	
UUUUUAUAUGA	Dendroctonus ponderosae
AGGUGUAUGA	
UUUUUAUAUGA	Aedes aegypti
UUGUGCAUGA	
UUUUUAUAUGA	Anopheles gambiae
ACGUGCAUGA	
UCUUUAUAUGA	Drosophila melanogaster
CUGUGCAUGA	
GUACACAUGA	
UCUUUAUAUGA	Drosophila willistoni
CUGUGCAUGA	
UCUUUAUAUGA	Drosophila serrata
CUGUGCAUGA	
GUACACAUGA	
UCUUUAUAUGA	Drosophila erecta
CUGUGCAUGA	
GUACACAUGA	
AAUCUUUAUGA	Drosophila ananassae
UGACUUUAUGA	
UGGUGCAUGA	
GUACACAUGA	
UCUUUAUAUGA	Phlebotomus papatasi
UCGUGCAUGA	
UCUUUAUAUGA	Phlebotomus sergenti
UCGUGCAUGA	
AUUUAUAUGA	Acyrtosiphon pisum
AGGUGCAUGA	
UUUUUAUAUGA	Schistocerca gregaria
ACCUACAUGC	
AGGUGUAUGA	
UUUUUAUAUGA	Teleopsis dalmanni
UCGUGCAUGA	
UUUUUAUAUGA	"Quercus robur"
UCGUGCAUGA	
GCGUGUAUGA	Bombus terrestris
UUUUUAUAUGA	Nilaparvata lugens
AGGUGUAUGA	
UUUUUAUAUGA	Parasteatoda tepidariorum
AGGUUUUAUGC	
UUUUUAUAUGA	"Selaginella moellendorffii"
AGGUUUUAUGC	

UUUUAUAUGA	Limulus polyphemus #1
CCGUUUUAUGA	
UUUUAUAUGA	Limulus polyphemus #2
UUGUUUAUGA	
UUUUAUAUGA	Rhipicephalus appendiculatus
AGGUUUUAUGC	
CUUGUUAUGC	
UUUUAUAUGA	Amblyomma variegatum
AGGUUUUAUGC	
UUUGUUAUGC	
UUUUAUAUGA	Loxosceles laeta
AGGUUUUAUGC	
UUUUGUAUGC	Campodea fragilis
GAAUUGAUGC	
AUUUAUAUGA	Aphis gossypii
AGGUGCAUGA	
UUUUAUAUGA	Locusta migratoria
GGGUGCAUGA	
CUGUGUAUGU	Calanus finmarchicus
GAUUUUUAUGA	Polypedilum vanderplanki
AAGUGCAUGU	
UUUUUCAUGA	Petrolisthes cinctipes
UCUUGUAUGU	
ACUUUCAUGA	Penaeus monodon
UCUUGUAUGU	
ACUUUCAUGA	Callinectes sapidus
UCUUGUAUGU	
UUUUAUAUGA	Glossina morsitans
UCGUGCAUGA	
UUUUAUAUGA	Ceratitis capitata
UCGUGCAUGA	
UCUUUAUAUGA	Rhynchosciara americana
CCUUUUUAUGA	Chironomus tentans
CAGUGCAUGA	
UUUUAUAUGA	Spodoptera frugiperda
GCGUGCAUGA	
UUUUAUAUGA	Spodoptera litura
GCGUGCAUGA	
UUUUAUAUGA	Gryllus bimaculatus
ACCUACAUGC	
AGGUGUAUGA	
AAACUUUAUGG	Pediculus humanus capitis
ACGUGUAUGA	

C)

Nematoda

UUGUGCAUGU	Caenorhabditis remanei
AAUUUUUAUGA	
CGGUGUAUGC	
UUGUGCAUGU	Caenorhabditis elegans
AAUUUUUAUGA	
UAGUGUAUGC	
UUGUGCAUGU	Caenorhabditis japonica
AAUUAUAUGA	
CAGUGUAUGC	
UUGUGCAUGU	Heterorhabditis bacteriophora
GAUUUUUAUGA	
CAUCGUAUGC	
AGUUGCAUGU	Wuchereria bancrofti
GAUUUUUAUGG	

UCACGUAUGC	
AGUUGCAUGU	Brugia malayi
GAUJUUAUGG	
UCACGUAUGC	
AGUUGUAUGU	Ascaris suum
UAUJUUAUGG	
CCCCGUAUGC	
UUGUGCAUGU	Angiostrongylus cantonensis
AAUJUUAUGC	
CAUCGUAUGC	
UUGUGCAUGU	Ancylostoma ceylanicum
GAUJUUAUGC	
UAUCGUAUGC	
AGGUGUAUGA	Bursaphelenchus mucronatus
CGUJUUAUGG	
UAACGUAUGC	

D)

Mollusca

GUUJUUAUGA	Lottia gigantean
AGGJUUAUGA	
AUUJUUAUGA	Aplysia californica
GCGJUUAUGA	
AUUJUUAUGA	Aplysia kurodai
GCGJUUAUGA	
AUUJUUAUGA	Lymnaea stagnalis
GCGUAUAUGA	
GUUJUUAUGA	Crassostrea gigas
GGGJUUAUGA	
GGUJUUAUGA	Crassostrea virginica
GGGJUUAUGA	
AGUCGUAUGA	Mytilus californianus
AGUJUUAUGA	

E)

Pezizomycotina

UUGJUUAUGU	Neurospora crassa
GCGUUAAGC	
UACCCUAUGU	
UUGJUUAUGU	Podospora anserine
CUGCCCAUGU	
CACCACAUGU	
GAGJUUAUGC	Mycosphaerella graminicola
GAGJUUAUGC	
UGCUGCAUGU	
CAGJUUAUGU	Aureobasidium pullulans
AAUUGCAUGU	
CUGUCUAUGU	Trichophyton rubrum
CUACGCAUGU	
UUGJUUAUGU	Hypocrea jecorina
GAGJUUAUGC	
CGCCACAUGU	
CUGJUUAUGU	Gibberella moniliformis
UCGUUUAUGC	
UGCCACAUGU	
CUAUACAUGA	
UUGJUUAUGU	Thielavia terrestris
GCGUUAAGC	
UAAUGCAUGC	
UACCCUAUGU	

CUGUCUAUGU	Coccidioides posadasii
CUACGCAUGU	
UUGUCUAUGU	Aspergillus niger
CCACGCAUGU	
CUGUUUAUGU	Trichoderma atroviride
GAGUJCAUGC	
CUCCCUAUGU	
CGCUJCAUGC	
UUGUUUAUGU	Trichoderma virens
UCGUJCAUGC	
CGCCCUAUGU	
CUCUJGAUGC	
CGUUUAUUGU	Geomyces pannorum
CAACACAUGU	
CUCUAUAUGU	
GAGUAUAUGU	Cochliobolus heterostrophus
CGCUJUAUGC	
UUGCGCAUGU	
UUGUUUAUGU	Ophiostoma piliferum
CAGCACAUGC	
UUGUJCAUGA	Tuber melanosporum
CAGUJCAUGC	
GGGUJUAUGU	Cercospora zeae-maydis
UACCGCAUGU	
CGUUUAUUGU	Amorphotheca resinae
CUUCGAAUGU	
UUCUGCAUGU	
UGUJCCAUGU	
UUGUUUAUGU	"Panicum virgatum"
GCGUJAAUGC	
UACCCUAUGU	
CGUCUCAUGA	
UUGUUUAUGU	Myceliophthora thermophila
GCGUJAAUGC	
UACCCUAUGU	
UUGUUUAUGU	Thielavia heterothallica
GCGUJAAUGC	
UACCCUAUGU	
CUGUUUAUGU	Fusarium sporotrichioides
UUGUCUAUGC	
UUGUUUAUGU	"Wrightia tinctoria"
GCGUJAAUGC	
UAAUGCAUGC	
UACCCUAUGU	
UUGUUUAUGU	Cordyceps militaris
GAGUJCAUGC	
AACCUCAUGU	
GUCUJCAUGA	
UUGUJCAUGA	"Picea glauca"
CGGUJCAUGC	
UUGUUUAUGU	Sordaria macrospora
GCGUJAAUGC	
UACCCUAUGU	
CUGUUUAUGU	Magnaporthe oryzae
GCGUJCAUGC	
UCCCCUAUGU	
CAGUCUAUGU	
CUGUUUAUGU	Magnaporthe poae
GCGUJCAUGC	
UCCCCUAUGU	
AAGUJCAUGU	

CUGUUUAUGU	Gaeumannomyces graminis
GCGUUCAUGC	
UCCCCUAUGU	
AAGUUCAUGU	
UUGUUUAUGU	Trichoderma reesei
GAGUUCAUGC	
CGCCACAUGC	
CUCUUGAUGC	
CUGUUUAUGU	Nectria haematococca
UUGUCUAUGC	
UGCCACAUGC	
CUCUUCAUGA	
CAGUUUAUGU	Glomerella graminicola
GAGUUCAUGA	
UACCACAUGC	
UUCUUA AUGC	
CUGUUUAUGU	Gibberella zeae
CUGUCUAUGC	
UGCCACAUGA	
CUAUACAUGA	
UUGUCUAUGU	Neosartorya fischeri
CUACGCAUGC	
CUGUUUAUGU	Fusarium oxysporum
UCGUCUAUGC	
UGCCACAUGC	
CUAUACAUGA	
CUGUUUAUGU	Verticillium dahliae
GAGUUCAUGU	
UACCCCAUGU	
UCUUGCAUGU	
UUGUCUAUGU	Aspergillus fumigatus
UGCUGUAUGA	
CUACGCAUGC	
UUGUUUAUGU	Grosmannia clavigera
GCGUUCAUGC	
UCCCUUAUGU	
UUCUUCAUGC	
UUGUCUAUGU	Aspergillus clavatus
CUACGCAUGC	
CGUUUAUGU	Geomyces destructans
CAACACAUGC	
CUCUAUAUGU	
CUGUUUAUGU	Metarhizium acridum
GAGUUCAUGC	
UGCCCAUGU	
ACCUUCAUGC	
CUGUUCAUGU	Talaromyces stipitatus
UACCGCAUGC	
UUGUCUAUGU	Aspergillus terreus
UGACGCAUGC	
CGUUUAUGU	Botryotinia fuckeliana
AUUCGAAUGU	
UUCUGCAUGU	
UUGUCUAUGU	Aspergillus nidulans
CGGCGUAUGU	
CGUUUAUGU	Blumeria graminis
CGGCGAAUGU	
CAUCAUAUGU	
UUGUCUAUGU	Aspergillus flavus
AUACGCAUGU	
GAGUAUAUGU	Pyrenophora teres

CGCUGUAUGC	
CUGCGCAUGU	
GAGUAUAUGU	Pyrenophora tritici-repentis
CGCUGUAUGC	
CUGCGCAUGU	
CUGUJCAUGU	Penicillium marneffeii
UACCGCAUGU	
CUGUUUAUGU	Epichloe festucae
GAGUJCAUGC	
UGCCCUAUGU	
GUGUUUAUGU	
CGGUUAUAUGU	Phaeosphaeria nodorum
UACUGCAUGU	
CGUUUAUAUGU	Sclerotinia sclerotiorum
AUUCGAAUGU	
UUCUGCAUGU	
CUGUUUAUGU	Verticillium albo-atrum
GAGUJCAUGU	
UACCCCAUGU	
UCUJCCAUGU	
CUGUUUAUGU	Metarhizium anisopliae
GAGUJCAUGC	
UGCCCAUGU	
AUCUJCAUGC	

F) Basidiomycota

GGAAACAUGC	Postia placenta
GGAUJCAUGC	Trametes versicolor
GGAUJCAUGC	Phanerochaete chrysosporium
GGAUJCAUGC	Heterobasidion annosum
GGAUJCAUGC	Pleurotus ostreatus
GGAUJCAUGC	Schizophyllum commune
GGAUJCAUGC	Lentinula edodes
GAAUJCAUGC	Coprinopsis cinerea
GGAUJCAUGC	Gloeophyllum trabeum
CGGUJCAUGU	Leucosporidium scottii
CAGCUUAUGU	Microbotryum violaceum
UCGUJCAUGU	Melampsora larici-populina
GAGUJCAUGC	Tremella mesenterica
GAGUJCAUGC	Cryptococcus neoformans
UUGUJCAUGC	Cryptococcus laurentii
UCGUUAUAUGU	Phakopsora pachyrhizi
GUUUUAUGU	Ustilago maydis
AAUJCAUGC	Paxillus involutus

G) Other fungi

UAGUJCAUGU	Cunninghamella elegans
UUGUACAUGU	Mucor circinelloides
CUGUJCAUGU	Phycomyces blakesleeana
CGCUJCAUGU	Glomus intraradices
CUGUACAUGU	Piromyces sp.
CGCUJCAUGA	Spizellomyces punctatus

H) Plants

UCUUUAUGU	Oryza sativa #1
UGCUGCAUGC	

CAAUUUAUGC	
UCUUUUUAUGU	Oryza sativa #2
UCUUUUUAUGU	Sorghum bicolour #1
CACUGCAUGC	
UCUUUUUAUGU	Sorghum bicolour #2
CACUGCAUGC	
UCUUUUUAUGU	Populus trichocarpa #1
GGCUGCAUGC	
GCUUUUAUGU	Populus trichocarpa #2
AGCUGUAUGC	
UUUUUUUAUGU	Populus trichocarpa #3
ACACAGAUGC	
UCUUUUUAUGU	Populus tremula #1
GGCUGUAUGC	
UGUUUUUAUGA	
GCUUUUAUGU	Populus tremula #2
AGCUGUAUGC	
UCUUUUUAUGU	Populus deltoids #1
GGCUGCAUGC	
UUUUUUUAUGU	Populus deltoides #3
AGCUAUUAUGC	
UCUUUUUAUGU	Populus nigra #1
GGCUGCAUGC	
UUUUUUUAUGU	Populus nigra #3
ACACAGAUGC	
AGCUAUUAUGC	
GCUUUUAUGU	Populus fremontii #2
AGCUGUAUGC	
UCUUUUUAUGU	Zea mays #1
CACUGCAUGC	
UCUUUUUAUGU	Zea mays #2
CACUGCAUGC	
UCUUUUUAUGU	Arabidopsis thaliana
CAGUUUAUGC	
UCUUUUUAUGU	Vitis vinifera #1
AGCUGCAUGC	
AGGUUUUAUGU	
UCUUUUUAUGU	Vitis vinifera #2
GACUGUAUGC	
AGCUGGAUGC	
UCUUUUUAUGU	Vitis shuttleworthii #1
AGCUGCAUGC	
AGGUUUUAUGU	
UCUUUUUAUGU	Vitis shuttleworthii #2
GACUGUAUGC	
AGCUGGAUGC	
GCUUUUAUGU	Crocus sativus
GACUGCAUGA	
UCGUUCAUGC	
UCUUUUUAUGU	Nicotiana tabacum #1
GACUGUAUGC	
UCUUUUUAUGU	Nicotiana tabacum #2
GACUGUAUGC	
UCUUUUUAUGU	Nicotiana tabacum #3
GGCUGCAUGC	
UCUUUUUAUGU	Nicotiana benthamiana #2
GACUUUAUGC	
UUUUUUUAUGU	Nicotiana benthamiana #3
GGCUGCAUGC	
UCUUUUUAUGU	Amborella trichopoda
ACUCAGAUGC	

AGAUGCAUGC	
ACUUUAAUGC	
UCUUUU AUGU	Quercus petraea
GGCUGCAUGC	
UGGUAAAUGA	
UCUUUU AUGU	Petunia axillaris
GACUGUAUGC	
UCUUUU AUGU	Petunia x hybrida
GACUGUAUGC	
UCUUUU AUGU	Quercus robur
GGCUGUAUGC	
AGUUUGAUGG	
UCUUUU AUGU	Triticum aestivum #1
CCCUGCAUGC	
UCUUUU AUGU	Triticum aestivum #2
CCCUGCAUGC	
UCUUUU AUGU	Triticum aestivum #3
CCCUGCAUGC	
UCUUUU AUGU	Lolium multiflorum
CCCUGUAUGC	
UCUUUU AUGU	Lolium perenne
CCCUGUAUGC	
UCUUUU AUGU	Lolium temulentum
CCCUGUAUGC	
UCUUUU AUGU	Avena sativa #1
CCCUGUAUGC	
UCUUUU AUGU	Avena sativa #2
CCCUGUAUGC	
UCUUUU AUGU	Avena barbata
CCCUGUAUGC	
UCUUUU AUGU	Festuca pratensis
CCCUGUAUGC	
UCUUUU AUGU	Festuca arundinacea
CCCUGUAUGC	
UCUUUU AUGU	Hordeum vulgare #1
CCCUGCAUGC	
UCUUUU AUGU	Hordeum vulgare #2
CCCUGCAUGC	
UCUUUU AUGU	Brachypodium distachyon
CCCUGCAUGC	
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UAUUUU AUGU	Prunus persica #2
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GGAUGCAUGC	
CGGUUU AUGU	

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UCUUUUAUGU	Solanum lycopersicum #1
GACUGUAUGC	
UCUUUUAUGU	Solanum lycopersicum #3
UAUUGUAUGU	
GGGUGAUGA	
GACUGUAUGC	
UCUUUUAUGU	Solanum tuberosum #1
AACUGUAUGC	
UCUUUUAUGU	Solanum tuberosum #2
GACUGUAUGC	
UCUUUUAUGU	Solanum tuberosum #3
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UCUUUUAUGU	Actinidia arguta #2
GACUGUAUGC	
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UCUUUUAUGU	Actinidia chinensis #1
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UUUUCCAUGC	Limnanthes alba
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UUUUUUAUGU	Fagus sylvatica
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UAUUUUAUGU	Rubus ulmifolius
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AUCUGCAUGC	
CAUUUUUAUGC	
UCUUUUAUGU	Gossypium arboreum #2
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CGUUUUUAUGC	
UCUUUUAUGU	Gossypium arboreum #3
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AUCUGCAUGC	
CAUUUUUAUGC	
UCUUUUAUGU	Panax ginseng
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GGCUGCAUGC	

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UAUUUUUAUGU	Cannabis sativa
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UCGUUUUAUGU	Elaeis oleifera
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UUUUUUUAUGU	Wrightia tinctoria
GACUGUAUGC	
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AUUUUUAUGU	Helianthus exilis
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UCUUUCAUGU	Glycine max #2
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UCUUUCAUGU	Glycine max #3
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UCUUUCAUGU	Glycine max #4
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UCUUUCAUGU	Phaseolus vulgaris #1
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AGCUGUAUGC	
UCUUUCAUGU	Phaseolus acutifolius #1

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UCUU	UUCAUGU	Phaseolus acutifolius #2
AGCU	UGUAUGC	
UUUU	UUUAUGU	Guizotia abyssinica
GGCU	UGUAUGC	
UCUU	UUCAUGU	Vigna unguiculata
AUCU	UGUAUGC	
UCUU	UUUAUGU	Leymus cinereus
CCCU	UGCAUGC	
UCUU	UUUAUGU	Pinus taeda
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GGCU	UGUAUGC	
AGUU	UGGAUGC	
UCUU	UUUAUGU	Pinus radiata
GGUU	UGGAUGC	
GGCU	UGUAUGC	
AGUU	UGGAUGC	
CCUU	UUUAUGU	Pinus banksiana
UUGU	UUUAUGU	
GAAU	UAAUUGU	
UUUU	UGCAUGU	
UCUU	UUUAUGU	Pinus contorta
GGUU	UGGAUGC	
AGUU	UGGAUGC	
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AGUU	UGGAUGC	
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GGCU	UGUAUGC	
AGUU	UGGAUGC	
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AGUU	UGGAUGC	
UCUU	UUUAUGU	Selaginella moellendorffii
UGC	UGCAUGC	
UCUU	UUUAUGU	Selaginella lepidophylla
GGCU	UGCAUGC	
GAGU	UCAUGC	
UCUU	UUUAUGU	Physcomitrella patens
GGUU	UGCAUGU	

Supplementary Figure S2. Weblogo representation of amino acid conservation in eIF5 uORF1 encoded peptides.

Letter heights are proportional to the frequency of conservation of each amino acid at each position. Each line represents a different eukaryotic branch which is indicated in parentheses on the right followed by the number of sequences used in each alignment. The sequences used to generate the alignment presented in this figure are available upon request.

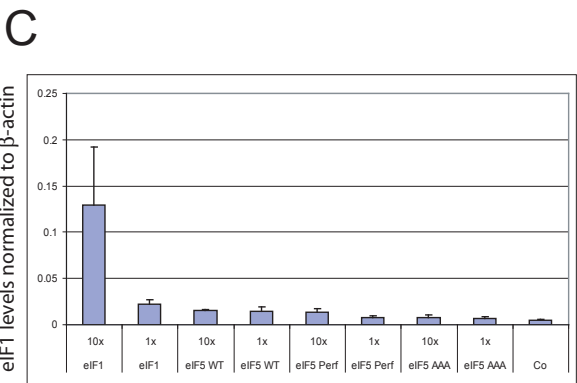
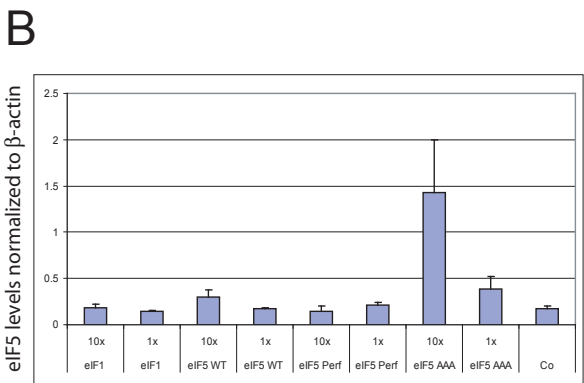
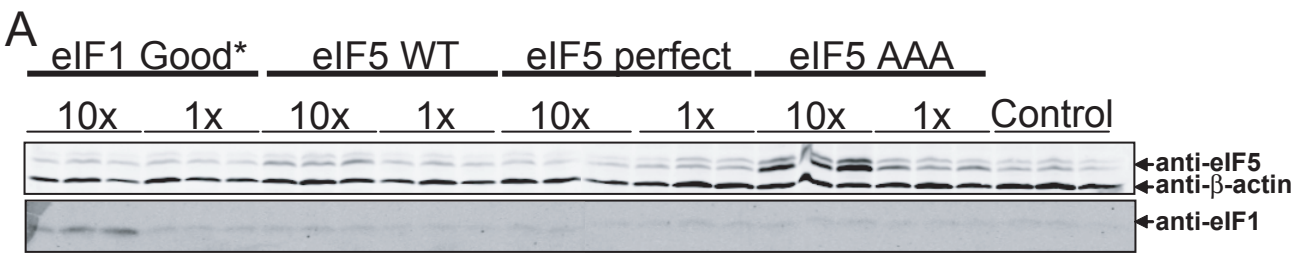
Supplementary Figure S2

[illegible]

Supplementary Figure S3. Removing eIF5 uORFs results in constitutive eIF5 expression.

(A) Western blots of protein lysates from cells transfected in triplicate with the eIF5 or eIF1 overexpression constructs indicated in Figure 2B of the main text. The eIF1 overexpression construct is the “eIF1 good*” described previously (2). In lanes marked “10×”, 10-fold more vector with insert was transfected compared with lanes marked “1×,” where the difference in the amount of transfecting DNA is made up with the inert vector pcDNA3 (2). The control cells are transfected with “10×” amount of pcDNA3. The blot shown was probed with anti-eIF5 and anti-β-actin followed by anti-eIF1. The corresponding detected proteins are indicated by arrows. (B & C) Densitometry analysis of the western blot shown in A. Relative eIF5 levels (B) and eIF1 levels (C) were determined by normalizing to β-actin levels.

Supplementary Figure S3



Supplementary Table S1.

Oligonucleotides used in this study.

1	eIF5 wt S PstI	ATAACTGCAGCCAGCCAATGGGCAGTGAGG
2	eIF5 wt AS 5' intron	AACCTTGATACTTACCGTTTTTCGTCAAATAAAGACATAAACCCAACGC
3	eIF5 wt S 3' intron	TTTCTCTCCACAGAGCTGTTGCGCAGCCATTGG
4	eIF5 wt AS BamHI	TTATGGATCCAGACATTTTGGCTTATTAGTGG
5	eIF5 wt S 5' intron	TTTATTTGACGAAAACGGTAAGTATCAAGGTTAC
6	eIF5 wt AS 3' intron	AATGGCTGCGCAACAGCTCTGTGGAGAGAAAGG
7	eIF5 wt S	AAGCCAAAATGTCTGTCAATGTCAACCGC
8	eIF5 wt stop AS XbaI	TTATTCTAGACCTTTAAATGGCATCAATATCG
9	eIF5 wt start S SacI	ATAAGAGCTCGTTTAGTGAACCGCCAGCCAATGGGCAGTGAGG
10	eIF5 wt AS	AGACATTTTGGCTTATTAGTGG
11	eIF5 uORF1 perfect AS	AACCTTGATACTTACCGTTTTTCGTCAAATAAAGCCATGGTGGCAACGCTGCTCGCCCCGG
12	eIF5 uORF1 perfect S	CCCCGGCGAGCAGCGTTGCCACCATGGCTTTATTTGACGAAAACGGTAAGTATCAAGGTT
13	eIF5 uAUG1 S	GTAGGGGTTTATGTCCG
14	eIF5 uAUG1 AS	GATCCGGACATAAACCCCTACTGCA
15	eIF5 uAUG2 S	GTAGTTTTTTCATGTCCG
16	eIF5 uAUG2 AS	GATCCGGACATGAAAACTACTGCA
17	eIF5 uAUG3 S	GTAGTCGTTTATGTCCG
18	eIF5 uAUG3 AS	GATCCGGACATAAACGACTACTGCA
19	eIF5 mAUG S	GTAGGCCAAAATGTCCG
20	eIF5 mAUG AS	GATCCGGACATTTTGGCCTACTGCA
21	Optimal context S	GTAGGCCACCATGGCCG
22	Optimal context A	GATCCGGCCATGGTGGCCTACTGCA
23	GG/S in worst context	GTAGTTTGTTATGGCCG
24	GG/A in worst context	GATCCGGCCATAACAACTACTGCA
25	CG/S in worst context	GTAGTTTCTTATGGCCG
26	CG/A in worst context	GATCCGGCCATAAGAACTACTGCA
27	TG/S in worst context	GTAGTTTTTTTATGGCCG
28	TG/A in worst context	GATCCGGCCATAAAAACTACTGCA
29	AA/S in worst context	GTAGTTTATTATGACCG
30	AA/A in worst context	GATCCGGTCATAATAAACTACTGCA
31	AC/S in worst context	GTAGTTTATTATGCCCCG
32	AC/A in worst context	GATCCGGGCATAATAAACTACTGCA
33	AT/S in worst context	GTAGTTTATTATGTCCG
34	AT/A in worst context	GATCCGGACATAATAAACTACTGCA
35	GA/S in worst context	GTAGTTTGTTATGACCG
36	GA/A in worst context	GATCCGGTCATAACAACTACTGCA
37	GC/S in worst context	GTAGTTTGTTATGCCCCG
38	GC/A in worst context	GATCCGGGCATAACAACTACTGCA
39	GT/S in worst context	GTAGTTTGTTATGTCCG
40	GT/A in worst context	GATCCGGACATAACAACTACTGCA

41	CA/S in worst context	GTAGTTTCTTATGACCG
42	CA/A in worst context	GATCCGGTCATAAGAACTACTGCA
43	CC/S in worst context	GTAGTTTCTTATGCCCCG
44	CC/A in worst context	GATCCGGGCATAAGAACTACTGCA
45	CT/S in worst context	GTAGTTTCTTATGTCCG
46	CT/A in worst context	GATCCGGACATAAGAACTACTGCA
47	TA/S in worst context	GTAGTTTTTTATGACCG
48	TA/A in worst context	GATCCGGTCATAAAAACTACTGCA
49	TC/S in worst context	GTAGTTTTTTATGCCCCG
50	TC/A in worst context	GATCCGGGCATAAAAACTACTGCA
51	TT/S in worst context	GTAGTTTTTTATGTCCG
52	TT/A in worst context	GATCCGGACATAAAAACTACTGCA
53	AG/S in worst context	GTAGTTTATTATGGCCG
54	AG/A in worst context	GATCCGGCCATAATAAACTACTGCA