

Dieter Soll

List of Publications by Citations

Source: <https://exaly.com/author-pdf/3869376/dieter-soll-publications-by-citations.pdf>

Version: 2024-04-24

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

207
papers

11,095
citations

50
h-index

100
g-index

230
ext. papers

12,811
ext. citations

11
avg. IF

6.34
L-index

#	Paper	IF	Citations
207	Aminoacyl-tRNA synthesis. <i>Annual Review of Biochemistry</i> , 2000 , 69, 617-50	29.1	1050
206	Natural expansion of the genetic code. <i>Nature Chemical Biology</i> , 2007 , 3, 29-35	11.7	437
205	Codon Bias as a Means to Fine-Tune Gene Expression. <i>Molecular Cell</i> , 2015 , 59, 149-61	17.6	367
204	RNA-dependent conversion of phosphoserine forms selenocysteine in eukaryotes and archaea. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 18923-7	11.5	359
203	The human SepSecS-tRNA ^{Sec} complex reveals the mechanism of selenocysteine formation. <i>Science</i> , 2009 , 325, 321-5	33.3	318
202	Expanding the genetic code of Escherichia coli with phosphoserine. <i>Science</i> , 2011 , 333, 1151-4	33.3	259
201	The RNA required in the first step of chlorophyll biosynthesis is a chloroplast glutamate tRNA. <i>Nature</i> , 1986 , 322, 281-4	50.4	256
200	Decameric SelA tRNA ^{Sec} ring structure reveals mechanism of bacterial selenocysteine formation. <i>Science</i> , 2013 , 340, 75-8	33.3	242
199	Protein biosynthesis in organelles requires misaminoacylation of tRNA. <i>Nature</i> , 1988 , 331, 187-90	50.4	218
198	Anticodon and acceptor stem nucleotides in tRNA(Gln) are major recognition elements for E. coli glutamyl-tRNA synthetase. <i>Nature</i> , 1991 , 352, 258-60	50.4	191
197	Evolution of translation machinery in recoded bacteria enables multi-site incorporation of nonstandard amino acids. <i>Nature Biotechnology</i> , 2015 , 33, 1272-1279	44.5	172
196	A 2-thiouridine derivative in tRNA ^{Glu} is a positive determinant for aminoacylation by Escherichia coli glutamyl-tRNA synthetase. <i>Biochemistry</i> , 1993 , 32, 3836-41	3.2	165
195	Nanoarchaeum equitans creates functional tRNAs from separate genes for their 5R and 3R halves. <i>Nature</i> , 2005 , 433, 537-41	50.4	159
194	Structure of pyrrolysyl-tRNA synthetase, an archaeal enzyme for genetic code innovation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 11268-73	11.5	156
193	Trans-editing of mischarged tRNAs. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 15422-7	11.5	156
192	An aminoacyl-tRNA synthetase that specifically activates pyrrolysine. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 12450-4	11.5	156
191	Severe oxidative stress induces protein mistranslation through impairment of an aminoacyl-tRNA synthetase editing site. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 4028-33	11.5	152

190	A chemical biology route to site-specific authentic protein modifications. <i>Science</i> , 2016 , 354, 623-626	33.3	151
189	Biosynthesis and Function of Modified Nucleosides 165-205		150
188	Structural insights into the role of rRNA modifications in protein synthesis and ribosome assembly. <i>Nature Structural and Molecular Biology</i> , 2015 , 22, 342-344	17.6	148
187	tRNA-dependent asparagine formation. <i>Nature</i> , 1996 , 382, 589-90	50.4	145
186	Domain-specific recruitment of amide amino acids for protein synthesis. <i>Nature</i> , 2000 , 407, 106-10	50.4	140
185	Pyrrolysyl-tRNA synthetase-tRNA(Pyl) structure reveals the molecular basis of orthogonality. <i>Nature</i> , 2009 , 457, 1163-7	50.4	133
184	Continuous directed evolution of aminoacyl-tRNA synthetases. <i>Nature Chemical Biology</i> , 2017 , 13, 1253-1260	12.6	124
183	Titelbild: Umkodierung des genetischen Codes mit Selenocystein (Angew. Chem. 1/2014). <i>Angewandte Chemie</i> , 2014 , 126, 1-1	3.6	124
182	Aminoacyl-tRNA synthesis: divergent routes to a common goal. <i>Trends in Biochemical Sciences</i> , 1997 , 22, 39-42	10.3	120
181	Upgrading protein synthesis for synthetic biology. <i>Nature Chemical Biology</i> , 2013 , 9, 594-8	11.7	114
180	Mutations disrupting selenocysteine formation cause progressive cerebello-cerebral atrophy. <i>American Journal of Human Genetics</i> , 2010 , 87, 538-44	11	111
179	Modified Nucleosides and Codon Recognition+207-223		110
178	The RCN1-encoded A subunit of protein phosphatase 2A increases phosphatase activity in vivo. <i>Plant Journal</i> , 1999 , 20, 389-99	6.9	108
177	Regulation of HEMA1 expression by phytochrome and a plastid signal during de-etiolation in <i>Arabidopsis thaliana</i> . <i>Plant Journal</i> , 2001 , 25, 549-61	6.9	104
176	Pyrrolysine is not hardwired for cotranslational insertion at UAG codons. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 3141-6	11.5	93
175	Rewriting the Genetic Code. <i>Annual Review of Microbiology</i> , 2017 , 71, 557-577	17.5	90
174	Mutations in QARS, encoding glutaminyl-tRNA synthetase, cause progressive microcephaly, cerebral-cerebellar atrophy, and intractable seizures. <i>American Journal of Human Genetics</i> , 2014 , 94, 547-58	11.1	87
173	Dimeric tRNA precursors in yeast. <i>Nature</i> , 1980 , 287, 750-2	50.4	85

172	Genetic code flexibility in microorganisms: novel mechanisms and impact on physiology. <i>Nature Reviews Microbiology</i> , 2015 , 13, 707-721	22.2	77
171	The selenocysteine-inserting opal suppressor serine tRNA from E. coli is highly unusual in structure and modification. <i>Nucleic Acids Research</i> , 1989 , 17, 7159-65	20.1	76
170	Polyspecific pyrrolysyl-tRNA synthetases from directed evolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 16724-9	11.5	75
169	Coevolution of an aminoacyl-tRNA synthetase with its tRNA substrates. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 13863-8	11.5	72
168	A facile strategy for selective incorporation of phosphoserine into histones. <i>Angewandte Chemie - International Edition</i> , 2013 , 52, 5771-5	16.4	71
167	In Vivo Biosynthesis of a β -Amino Acid-Containing Protein. <i>Journal of the American Chemical Society</i> , 2016 , 138, 5194-7	16.4	69
166	N-acetyl lysyl-tRNA synthetases evolved by a CcdB-based selection possess N-acetyl lysine specificity in vitro and in vivo. <i>FEBS Letters</i> , 2012 , 586, 729-33	3.8	65
165	A novel root gravitropism mutant of Arabidopsis thaliana exhibiting altered auxin physiology. <i>Physiologia Plantarum</i> , 1995 , 93, 790-798	4.6	63
164	Recoding the genetic code with selenocysteine. <i>Angewandte Chemie - International Edition</i> , 2014 , 53, 319-23	16.4	60
163	Rationally evolving tRNA ^{Pyl} for efficient incorporation of noncanonical amino acids. <i>Nucleic Acids Research</i> , 2015 , 43, e156	20.1	59
162	Near-cognate suppression of amber, opal and quadruplet codons competes with aminoacyl-tRNA ^{Pyl} for genetic code expansion. <i>FEBS Letters</i> , 2012 , 586, 3931-7	3.8	58
161	Upgrading aminoacyl-tRNA synthetases for genetic code expansion. <i>Current Opinion in Chemical Biology</i> , 2018 , 46, 115-122	9.7	53
160	Chemical Evolution of a Bacterial Proteome. <i>Angewandte Chemie - International Edition</i> , 2015 , 54, 10030-46.4	16.4	52
159	Primary, Secondary, and Tertiary Structures of tRNAs 2014 , 93-126		51
158	Insights into RNA binding by the anticancer drug cisplatin from the crystal structure of cisplatin-modified ribosome. <i>Nucleic Acids Research</i> , 2016 , 44, 4978-87	20.1	50
157	Emergence of the universal genetic code imprinted in an RNA record. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 18095-100	11.5	50
156	Rewiring translation for elongation factor Tu-dependent selenocysteine incorporation. <i>Angewandte Chemie - International Edition</i> , 2013 , 52, 1441-5	16.4	49
155	A [3Fe-4S] cluster is required for tRNA thiolation in archaea and eukaryotes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 12703-12708	11.5	48

154	Adding pyrrolysine to the Escherichia coli genetic code. <i>FEBS Letters</i> , 2007 , 581, 5282-8	3.8	48
153	Crystal structures reveal an elusive functional domain of pyrrolysyl-tRNA synthetase. <i>Nature Chemical Biology</i> , 2017 , 13, 1261-1266	11.7	47
152	Indolmycin resistance of <i>Streptomyces coelicolor</i> A3(2) by induced expression of one of its two tryptophanyl-tRNA synthetases. <i>Journal of Biological Chemistry</i> , 2002 , 277, 23882-7	5.4	47
151	A second and differentially expressed glutamyl-tRNA reductase gene from <i>Arabidopsis thaliana</i> . <i>Plant Molecular Biology</i> , 1996 , 30, 419-26	4.6	44
150	Expanding the genetic code of <i>Escherichia coli</i> with phosphotyrosine. <i>FEBS Letters</i> , 2016 , 590, 3040-7	3.8	44
149	Facile Recoding of Selenocysteine in Nature. <i>Angewandte Chemie - International Edition</i> , 2016 , 55, 5337-41	6.4	43
148	Structural insights into RNA-dependent eukaryal and archaeal selenocysteine formation. <i>Nucleic Acids Research</i> , 2008 , 36, 1187-99	20.1	43
147	Recognition of pyrrolysine tRNA by the <i>Desulfitobacterium hafniense</i> pyrrolysyl-tRNA synthetase. <i>Nucleic Acids Research</i> , 2007 , 35, 1270-8	20.1	43
146	Molecular analysis of three maize 22 kDa auxin-binding protein genes--transient promoter expression and regulatory regions. <i>Plant Journal</i> , 1993 , 4, 423-32	6.9	43
145	Purification of Five Serine Transfer Ribonucleic Acid Species from <i>Escherichia coli</i> and Their Acylation by Homologous and Heterologous Seryl Transfer Ribonucleic Acid Synthetases. <i>Journal of Biological Chemistry</i> , 1970 , 245, 1394-1400	5.4	43
144	Selenoprotein biosynthesis defect causes progressive encephalopathy with elevated lactate. <i>Neurology</i> , 2015 , 85, 306-15	6.5	42
143	Revising the Structural Diversity of Ribosomal Proteins Across the Three Domains of Life. <i>Molecular Biology and Evolution</i> , 2018 , 35, 1588-1598	8.3	40
142	Engineering the elongation factor Tu for efficient selenoprotein synthesis. <i>Nucleic Acids Research</i> , 2014 , 42, 9976-83	20.1	39
141	The amino-terminal domain of pyrrolysyl-tRNA synthetase is dispensable in vitro but required for in vivo activity. <i>FEBS Letters</i> , 2007 , 581, 3197-203	3.8	39
140	The heterotrimeric <i>Thermus thermophilus</i> Asp-tRNA(Asn) amidotransferase can also generate Gln-tRNA(Gln). <i>FEBS Letters</i> , 2000 , 476, 140-4	3.8	37
139	A synthetic tRNA for EF-Tu mediated selenocysteine incorporation in vivo and in vitro. <i>FEBS Letters</i> , 2015 , 589, 2194-9	3.8	36
138	N-(purin-6-ylcarbamoyl)threonine: biosynthesis in vitro in transfer RNA by an enzyme purified from <i>Escherichia coli</i> . <i>FEBS Letters</i> , 1974 , 39, 301-6	3.8	33
137	Archaeal aminoacyl-tRNA synthesis: diversity replaces dogma. <i>Genetics</i> , 1999 , 152, 1269-76	4	33

136	Translation of Diverse Aramid- and 1,3-Dicarbonyl-peptides by Wild Type Ribosomes. <i>ACS Central Science</i> , 2019 , 5, 1289-1294	16.8	32
135	Transfer RNA misidentification scrambles sense codon recoding. <i>ChemBioChem</i> , 2013 , 14, 1967-72	3.8	32
134	Efficient Reassignment of a Frequent Serine Codon in Wild-Type Escherichia coli. <i>ACS Synthetic Biology</i> , 2016 , 5, 163-71	5.7	30
133	Divergence of selenocysteine tRNA recognition by archaeal and eukaryotic O-phosphoserine-tRNA ^{Sec} kinase. <i>Nucleic Acids Research</i> , 2008 , 36, 1871-80	20.1	30
132	Initiator tRNAs and Initiation of Protein Synthesis 511-528		30
131	Emergent rules for codon choice elucidated by editing rare arginine codons in Escherichia coli. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E5588-97	11.5	30
130	A Facile Method for Producing Selenocysteine-Containing Proteins. <i>Angewandte Chemie - International Edition</i> , 2018 , 57, 7215-7219	16.4	29
129	Exploring the substrate range of wild-type aminoacyl-tRNA synthetases. <i>ChemBioChem</i> , 2014 , 15, 1805-1809	3.8	29
128	Substrate structural requirements of Schizosaccharomyces pombe RNase P. <i>FEBS Letters</i> , 1989 , 251, 84-8	3.8	29
127	Drugging tRNA aminoacylation. <i>RNA Biology</i> , 2018 , 15, 667-677	4.8	28
126	Structure of the Pseudomonas aeruginosa transamidosome reveals unique aspects of bacterial tRNA-dependent asparagine biosynthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 382-7	11.5	28
125	Dual targeting of a tRNA ^{Asp} requires two different aspartyl-tRNA synthetases in Trypanosoma brucei. <i>Journal of Biological Chemistry</i> , 2009 , 284, 16210-16217	5.4	28
124	A dual-specific Glu-tRNA(Gln) and Asp-tRNA(Asn) amidotransferase is involved in decoding glutamine and asparagine codons in Acidithiobacillus ferrooxidans. <i>FEBS Letters</i> , 2001 , 500, 129-31	3.8	28
123	The sup8 tRNA ^{Leu} gene of Schizosaccharomyces pombe has an unusual intervening sequence and reduced pairing in the anticodon stem. <i>Molecular Genetics and Genomics</i> , 1984 , 197, 447-52		27
122	Recent Studies of RNase P 67-78		27
121	Engineering posttranslational proofreading to discriminate nonstandard amino acids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, 619-624	11.5	26
120	A mutant Escherichia coli tyrosyl-tRNA synthetase utilizes the unnatural amino acid azatyrosine more efficiently than tyrosine. <i>Journal of Biological Chemistry</i> , 2000 , 275, 40324-8	5.4	26
119	Organization of ribosomal DNA in yellow lupine (Lupinus luteus) and sequence of the 5.8 S RNA gene. <i>FEBS Letters</i> , 1983 , 152, 241-246	3.8	26

118	Archaeal Tuc1/Ncs6 homolog required for wobble uridine tRNA thiolation is associated with ubiquitin-proteasome, translation, and RNA processing system homologs. <i>PLoS ONE</i> , 2014 , 9, e99104	3.7	25
117	Aminoacyl-tRNA Synthetases: Occurrence, Structure, and Function251-292		25
116	Methanococcus jannaschii prolyl-cysteinyl-tRNA synthetase possesses overlapping amino acid binding sites. <i>Biochemistry</i> , 2001 , 40, 46-52	3.2	24
115	Cysteinyl-tRNA formation: the last puzzle of aminoacyl-tRNA synthesis. <i>FEBS Letters</i> , 1999 , 462, 302-6	3.8	24
114	Yeast seryl-tRNA synthetase expressed in Escherichia coli recognizes bacterial serine-specific tRNAs in vivo. <i>FEBS Journal</i> , 1993 , 214, 869-77		24
113	Identification of a 100-kDa protein associated with nuclear ribonuclease P activity in Schizosaccharomyces pombe. <i>FEBS Journal</i> , 1993 , 217, 501-7		24
112	Pyrrolysyl-tRNA synthetase variants reveal ancestral aminoacylation function. <i>FEBS Letters</i> , 2013 , 587, 3243-8	3.8	23
111	Structural basis of reverse nucleotide polymerization. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 20970-5	11.5	23
110	The genetic code - thawing the frozen accident <i>R Journal of Biosciences</i> , 2006 , 31, 459-63	2.3	23
109	The Human Genome Project: a paradigm for information management in the life sciences. <i>FASEB Journal</i> , 1991 , 5, 35-9	0.9	23
108	Change of tRNA identity leads to a divergent orthogonal histidyl-tRNA synthetase/tRNAHis pair. <i>Nucleic Acids Research</i> , 2011 , 39, 2286-93	20.1	22
107	Escherichia coli tryptophanyl-tRNA synthetase mutants selected for tryptophan auxotrophy implicate the dimer interface in optimizing amino acid binding. <i>Biochemistry</i> , 1996 , 35, 32-40	3.2	21
106	tRNA Sequences and Variations in the Genetic Code225-250		21
105	The Selenocysteine-Inserting tRNA Species: Structure and Function529-544		21
104	Transcription of Eukaryotic tRNA Genes31-50		21
103	Initiation of Protein Synthesis with Non-Canonical Amino Acids In Vivo. <i>Angewandte Chemie - International Edition</i> , 2020 , 59, 3122-3126	16.4	21
102	Error-prone protein synthesis in parasites with the smallest eukaryotic genome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, E6245-E6253	11.5	20
101	The RNA component of RNase P in Schizosaccharomyces species. <i>FEBS Letters</i> , 1990 , 271, 189-93	3.8	20

100	Splicing of tRNA Precursors79-92		19
99	Mechanistic insights into the slow peptide bond formation with D-amino acids in the ribosomal active site. <i>Nucleic Acids Research</i> , 2019 , 47, 2089-2100	20.1	18
98	Identification and codon reading properties of 5-cyanomethyl uridine, a new modified nucleoside found in the anticodon wobble position of mutant haloarchaeal isoleucine tRNAs. <i>Rna</i> , 2014 , 20, 177-88	5.8	18
97	Bacterial Aminoacyl-tRNA Synthetases: Genes and Regulation of Expression 2014 , 293-333		18
96	Structure and Expression of Prokaryotic tRNA Genes17-30		18
95	The central role of tRNA in genetic code expansion. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2017 , 1861, 3001-3008	4	17
94	Challenges of site-specific selenocysteine incorporation into proteins by Escherichia coli. <i>RNA Biology</i> , 2018 , 15, 461-470	4.8	17
93	The terminal adenosine of tRNA(Gln) mediates tRNA-dependent amino acid recognition by glutaminyl-tRNA synthetase. <i>Biochemistry</i> , 1998 , 37, 9836-42	3.2	17
92	tRNA specificity of a mischarging aminoacyl-tRNA synthetase: Glutamyl-tRNA synthetase from barley chloroplasts. <i>FEBS Letters</i> , 1988 , 228, 241-244	3.8	17
91	Loss of protein synthesis quality control in host-restricted organisms. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, E11505-E11512	11.5	17
90	RNA-Dependent Cysteine Biosynthesis in Bacteria and Archaea. <i>MBio</i> , 2017 , 8,	7.8	16
89	A one-step method for in vitro production of tRNA transcripts. <i>Nucleic Acids Research</i> , 2002 , 30, e105	20.1	16
88	Small RNA Oligonucleotide Substrates for Specific Aminoacylations349-370		16
87	tRNA on the Ribosome: a Waggle Theory443-469		16
86	tRNA Processing Nucleases51-65		16
85	Pyrrolysyl-tRNA synthetase, an aminoacyl-tRNA synthetase for genetic code expansion. <i>Croatica Chemica Acta</i> , 2016 , 89, 163-174	0.8	16
84	Transfer RNAs with novel cloverleaf structures. <i>Nucleic Acids Research</i> , 2017 , 45, 2776-2785	20.1	16
83	Aminoacyl-tRNA Synthetases and tRNAs for an Expanded Genetic Code: What Makes them Orthogonal?. <i>International Journal of Molecular Sciences</i> , 2019 , 20,	6.3	15

82	Crystal structures of the human elongation factor eEFSec suggest a non-canonical mechanism for selenocysteine incorporation. <i>Nature Communications</i> , 2016 , 7, 12941	17.4	15
81	Dual Genetic Encoding of Acetyl-lysine and Non-deacetylable Thioacetyl-lysine Mediated by Flexizyme. <i>Angewandte Chemie - International Edition</i> , 2016 , 55, 4083-6	16.4	15
80	The putative tRNA 2-thiouridine synthetase Ncs6 is an essential sulfur carrier in <i>Methanococcus maripaludis</i> . <i>FEBS Letters</i> , 2014 , 588, 873-7	3.8	14
79	Arrangement of the ribosomal RNA genes in <i>Schizosaccharomyces pombe</i> . <i>FEBS Letters</i> , 1982 , 143, 129-32	3.8	14
78	tRNA Discrimination in Aminoacylation		14
77	Translational Suppression: When Two Wrongs DO Make a Right		14
76	Ancient translation factor is essential for tRNA-dependent cysteine biosynthesis in methanogenic archaea. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 10520-5	11.5	12
75	Probing the active site tryptophan of <i>Staphylococcus aureus</i> thioredoxin with an analog. <i>Nucleic Acids Research</i> , 2015 , 43, 11061-7	20.1	12
74	Glutaminyl-tRNA synthetase: from genetics to molecular recognition. <i>Genes To Cells</i> , 1996 , 1, 421-7	2.3	12
73	tRNA-Like Structures in Plant Viral RNAs		12
72	Muller's Ratchet and Ribosome Degeneration in the Obligate Intracellular Parasites. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	12
71	Engineered Aminoacyl-tRNA Synthetases with Improved Selectivity toward Noncanonical Amino Acids. <i>ACS Chemical Biology</i> , 2019 , 14, 603-612	4.9	11
70	Using Genetic Code Expansion for Protein Biochemical Studies. <i>Frontiers in Bioengineering and Biotechnology</i> , 2020 , 8, 598577	5.8	11
69	Plasticity and Constraints of tRNA Aminoacylation Define Directed Evolution of Aminoacyl-tRNA Synthetases. <i>International Journal of Molecular Sciences</i> , 2019 , 20,	6.3	10
68	Transfer RNA function and evolution. <i>RNA Biology</i> , 2018 , 15, 423-426	4.8	10
67	Dimer-dimer interaction of the bacterial selenocysteine synthase Sela promotes functional active-site formation and catalytic specificity. <i>Journal of Molecular Biology</i> , 2014 , 426, 1723-35	6.5	10
66	C-terminal truncation of yeast SerRS is toxic for <i>Saccharomyces cerevisiae</i> due to altered mechanism of substrate recognition. <i>FEBS Letters</i> , 1998 , 439, 235-40	3.8	10
65	Temperature dependence of the aminoacylation of tRNA by <i>Bacillus stearothermophilus</i> aminoacyl-tRNA synthetases. <i>Biopolymers</i> , 1971 , 10, 2209-21	2.2	10

64	Recognition of Aminoacyl-tRNAs by Protein Elongation Factors	423-442		10
63	A cysteinyl-tRNA synthetase variant confers resistance against selenite toxicity and decreases selenocysteine misincorporation. <i>Journal of Biological Chemistry</i> , 2019 , 294, 12855-12865		5.4	9
62	Reducing the genetic code induces massive rearrangement of the proteome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 17206-11		11.5	9
61	Rewiring Translation for Elongation Factor Tu-Dependent Selenocysteine Incorporation. <i>Angewandte Chemie</i> , 2013 , 125, 1481-1485		3.6	9
60	Simplified in vitro synthesis of mutated RNA molecules. An oligonucleotide promoter determines the initiation site of T7RNA polymerase on ss M13 phage DNA. <i>FEBS Letters</i> , 1987 , 212, 271-5		3.8	9
59	Discontinuous Triplet Decoding with or without Re-Pairing by Peptidyl tRNA	471-490		9
58	Umkodierung des genetischen Codes mit Selenocystein. <i>Angewandte Chemie</i> , 2014 , 126, 325-330		3.6	8
57	Bioinformatic Analysis Reveals Archaeal tRNA and tRNA Identities in Bacteria. <i>Life</i> , 2017 , 7,		3	8
56	Chemische Evolution eines bakteriellen Proteoms. <i>Angewandte Chemie</i> , 2015 , 127, 10168-10172		3.6	8
55	Retracing the evolution of amino acid specificity in glutaminyl-tRNA synthetase. <i>FEBS Letters</i> , 1998 , 434, 149-54		3.8	8
54	Protein-RNA molecular recognition. <i>Nature</i> , 1996 , 381, 656		50.4	8
53	The tRNA Identity Problem: Past, Present, and Future	335-347		8
52	Organellar tRNAs: Biosynthesis and Function	127-140		8
51	Versatility of Synthetic tRNAs in Genetic Code Expansion. <i>Genes</i> , 2018 , 9,		4.2	8
50	Lysine Acetylation Regulates Alanyl-tRNA Synthetase Activity in. <i>Genes</i> , 2018 , 9,		4.2	8
49	Hijacking Translation Initiation for Synthetic Biology. <i>ChemBioChem</i> , 2020 , 21, 1387-1396		3.8	7
48	Effects of Heterologous tRNA Modifications on the Production of Proteins Containing Noncanonical Amino Acids. <i>Bioengineering</i> , 2018 , 5,		5.3	7
47	Exploiting evolutionary trade-offs for posttreatment management of drug-resistant populations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020 , 117, 17924-17931		11.5	7

46	Archaeal Ribosomal Proteins Possess Nuclear Localization Signal-Type Motifs: Implications for the Origin of the Cell Nucleus. <i>Molecular Biology and Evolution</i> , 2020 , 37, 124-133	8.3	7
45	Recoding of the selenocysteine UGA codon by cysteine in the presence of a non-canonical tRNA and elongation factor SelB. <i>RNA Biology</i> , 2018 , 15, 471-479	4.8	6
44	Maize mitochondrial seryl-tRNA synthetase recognizes Escherichia coli tRNA(Ser) in vivo and in vitro. <i>Plant Molecular Biology</i> , 1998 , 38, 497-502	4.6	6
43	Homologous expression and purification of mutants of an essential protein by reverse epitope-tagging. <i>Nature Biotechnology</i> , 1996 , 14, 50-5	44.5	6
42	Naturally Occurring tRNAs With Non-canonical Structures. <i>Frontiers in Microbiology</i> , 2020 , 11, 596914	5.7	6
41	Genetic Encoding of Three Distinct Noncanonical Amino Acids Using Reprogrammed Initiator and Nonsense Codons. <i>ACS Chemical Biology</i> , 2021 , 16, 766-774	4.9	6
40	A Facile Strategy for Selective Incorporation of Phosphoserine into Histones. <i>Angewandte Chemie</i> , 2013 , 125, 5883-5887	3.6	5
39	Suppression of amber codons in <i>Caulobacter crescentus</i> by the orthogonal <i>Escherichia coli</i> histidyl-tRNA synthetase/tRNA ^{His} pair. <i>PLoS ONE</i> , 2013 , 8, e83630	3.7	5
38	Designing seryl-tRNA synthetase for improved serylation of selenocysteine tRNAs. <i>FEBS Letters</i> , 2018 , 592, 3759-3768	3.8	5
37	Eine einfache Methode zur Produktion von Selenoproteinen. <i>Angewandte Chemie</i> , 2018 , 130, 7333-7337	3.6	4
36	Protein phosphatase 2A: identification in <i>Oryza sativa</i> of the gene encoding the regulatory A subunit. <i>Plant Molecular Biology</i> , 2001 , 45, 107-12	4.6	4
35	Initiation of Protein Synthesis with Non-Canonical Amino Acids In Vivo. <i>Angewandte Chemie</i> , 2020 , 132, 3146-3150	3.6	4
34	Selective cysteine-to-selenocysteine changes in a [NiFe]-hydrogenase confirm a special position for catalysis and oxygen tolerance. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	4
33	Introducing Selenocysteine into Recombinant Proteins in <i>Escherichia coli</i> . <i>Current Protocols</i> , 2021 , 1, e54		4
32	A genomically modified <i>Escherichia coli</i> strain carrying an orthogonal <i>E. coli</i> histidyl-tRNA synthetase/tRNA pair. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2017 , 1861, 3009-3015	4	3
31	Transfer RNA identity change in anticodon variants of <i>E. coli</i> tRNA(Phe) in vivo. <i>Molecules and Cells</i> , 2000 , 10, 76-82	3.5	3
30	Chloroplast tRNA(Asp): nucleotide sequence and variation of in vivo levels during plastid maturation. <i>Plant Molecular Biology</i> , 1992 , 20, 601-7	4.6	3
29	Multiplex suppression of four quadruplet codons via tRNA directed evolution. <i>Nature Communications</i> , 2021 , 12, 5706	17.4	3

28	A tRNA-guided research journey from synthetic chemistry to synthetic biology. <i>Rna</i> , 2015 , 21, 742-4	5.8	2
27	The genetic code: Yesterday, today, and tomorrow 2012 , 17, 1136-1142		2
26	Aminoacyl-tRNA tRNA formation: an essential function in protein synthesis and its quality control. <i>Nucleic Acids Symposium Series</i> , 2004 , 283-4		2
25	The Nbp35/ApbC homolog acts as a nonessential [4Fe-4S] transfer protein in methanogenic archaea. <i>FEBS Letters</i> , 2020 , 594, 924-932	3.8	2
24	Engineering aminoacyl-tRNA synthetases for use in synthetic biology. <i>The Enzymes</i> , 2020 , 48, 351-395	2.3	2
23	Measuring the tolerance of the genetic code to altered codon size		2
22	The genetic code revisited--four decades after Francis Crick. <i>Nucleic Acids Symposium Series</i> , 2007 , 13-4		1
21	Archaeal ribosomal proteins possess nuclear localization signal-type motifs: implications for the origin of the cell nucleus		1
20	Transfer RNA: Discovery, Early Work, and Total Synthesis of a tRNA Gene5-16		1
19	Engineering post-translational proofreading to discriminate non-standard amino acids		1
18	Intein-based Design Expands Diversity of Selenocysteine Reporters. <i>Journal of Molecular Biology</i> , 2021 , 167199	6.5	1
17	Initiating protein synthesis with noncanonical monomers in vitro and in vivo. <i>Methods in Enzymology</i> , 2021 , 656, 495-519	1.7	1
16	Directed Evolution of Pyrrolysyl-tRNA Synthetase Generates a Hyperactive and Highly Selective Variant.. <i>Frontiers in Molecular Biosciences</i> , 2022 , 9, 850613	5.6	1
15	Indirect Routes to Aminoacyl-tRNA: The Diversity of Prokaryotic Cysteine Encoding Systems.. <i>Frontiers in Genetics</i> , 2021 , 12, 794509	4.5	0
14	R&Ktitelbild: Rewiring Translation for Elongation Factor Tu-Dependent Selenocysteine Incorporation (Angew. Chem. 5/2013). <i>Angewandte Chemie</i> , 2013 , 125, 1638-1638	3.6	
13	Innentitelbild: Chemische Evolution eines bakteriellen Proteoms (Angew. Chem. 34/2015). <i>Angewandte Chemie</i> , 2015 , 127, 9862-9862	3.6	
12	Commonly Used Abbreviations, Terminologies, and Nomenclature 2014 , 557-557		
11	Titelbild: A Facile Strategy for Selective Incorporation of Phosphoserine into Histones (Angew. Chem. 22/2013). <i>Angewandte Chemie</i> , 2013 , 125, 5761-5761	3.6	

10	1SP7-03 tRNA recognition and molecular evolution of GatCAB(1SP7 Elucidation of Protein Functions at the Atomic Level with X-ray structural, Vibrational spectroscopic, Molecular biological and Theoretical analyses,The 47th Annual Meeting of the Biophysical Society of Japan). <i>Seibutsu Butsuru</i> , 2009 , 49, S9	0
9	Distorted RNA helix recognition. <i>Nature</i> , 1996 , 384, 422-422	50.4
8	Using selenocysteine-specific reporters to screen for efficient tRNA variants.. <i>Methods in Enzymology</i> , 2022 , 662, 63-93	1.7
7	Mischarging of <i>M. barkeri</i> tRNA ^{Pyl} with alanine and serine in vitro. <i>FASEB Journal</i> , 2006 , 20, A503	0.9
6	Recognition in vitro of the suppressor tRNA ^{Pyl} by the class II-like Pyrrolysyl-tRNA Synthetase. <i>FASEB Journal</i> , 2006 , 20, A503	0.9
5	A Molecular Tunnel Required for Cooperation of an Asparaginase and a Glu-tRNA ^{Gln} Kinase in Gln-tRNA Formation. <i>FASEB Journal</i> , 2006 , 20, A503	0.9
4	RNA-Dependent Cysteine Biosynthesis in Archaea. <i>FASEB Journal</i> , 2006 , 20, A503	0.9
3	Features of Aminoacyl-tRNA Synthesis Unique to Archaea198-208	
2	Structures of Modified Nucleosides551-555	
1	The Aspartic Acid tRNA System: Recognition by a Class II Aminoacyl-tRNA Synthetase411-422	