

# Anders S Byström

## List of Publications by Year in descending order

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Version: 2024-02-01

26  
papers

2,229  
citations

430874

18  
h-index

552781

26  
g-index

27  
all docs

27  
docs citations

27  
times ranked

1621  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | SSD1 modifies phenotypes of Elongator mutants. <i>Current Genetics</i> , 2020, 66, 481-485.   | 1.7  | 3         |
| 2  | SSD1 suppresses phenotypes induced by the lack of Elongator-dependent tRNA modifications. <i>PLoS Genetics</i> , 2019, 15, e1008117.  | 3.5  | 10        |
| 3  | Gene <i>miaA</i> for post-transcriptional modification of tRNA <sup>XXA</sup> is important for morphological and metabolic differentiation in <i>Streptomyces</i> . <i>Molecular Microbiology</i> , 2019, 112, 249-265. | 2.5  | 26        |
| 4  | Elongator subunit 3 (ELP3) modifies ALS through tRNA modification. <i>Human Molecular Genetics</i> , 2018, 27, 1276-1289.   | 2.9  | 56        |
| 5  | Identification of factors that promote biogenesis of tRNA <sup>CGA</sup> <sup>Ser</sup> . <i>RNA Biology</i> , 2018, 15, 1286-1294.   | 3.1  | 6         |
| 6  | Elongator <sup>ε</sup> a tRNA modifying complex that promotes efficient translational decoding. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2018, 1861, 401-408.                                | 1.9  | 29        |
| 7  | Loss of <i>ncm5</i> and <i>mcm5</i> wobble uridine side chains results in an altered metabolic profile. <i>Metabolomics</i> , 2016, 12, 177.  | 3.0  | 11        |
| 8  | Linkage between Fitness of Yeast Cells and Adenylate Kinase Catalysis. <i>PLoS ONE</i> , 2016, 11, e0163115.  | 2.5  | 12        |
| 9  | The role of wobble uridine modifications in +1 translational frameshifting in eukaryotes. <i>Nucleic Acids Research</i> , 2015, 43, 9489-9499.  | 14.5 | 67        |
| 10 | Yeast Elongator protein Elp1p does not undergo proteolytic processing in exponentially growing cells. <i>MicrobiologyOpen</i> , 2015, 4, 867-878.   | 3.0  | 2         |
| 11 | Meta-regulation of Arabidopsis Auxin Responses Depends on tRNA Maturation. <i>Cell Reports</i> , 2015, 11, 516-526.   | 6.4  | 27        |
| 12 | Elongator, a conserved complex required for wobble uridine modifications in Eukaryotes. <i>RNA Biology</i> , 2014, 11, 1519-1528.   | 3.1  | 115       |
| 13 | Familial dysautonomia (FD) patients have reduced levels of the modified wobble nucleoside <i>mcm5s2U</i> in tRNA. <i>Biochemical and Biophysical Research Communications</i> , 2014, 454, 441-445.                      | 2.1  | 78        |
| 14 | Elongator Complex Influences Telomeric Gene Silencing and DNA Damage Response by Its Role in Wobble Uridine tRNA Modification. <i>PLoS Genetics</i> , 2011, 7, e1002258.  | 3.5  | 87        |
| 15 | Unexpected Accumulation of <i>ncm5U</i> and <i>ncm5s2U</i> in a <i>trm9</i> Mutant Suggests an Additional Step in the Synthesis of <i>mcm5U</i> and <i>mcm5s2U</i> . <i>PLoS ONE</i> , 2011, 6, e20783.                 | 2.5  | 66        |
| 16 | Elongator function in tRNA wobble uridine modification is conserved between yeast and plants. <i>Molecular Microbiology</i> , 2010, 76, 1082-1094.  | 2.5  | 87        |
| 17 | Elongator function in tRNA wobble uridine modification is conserved between yeast and plants. <i>Molecular Microbiology</i> , 2010, 77, 531-531.  | 2.5  | 1         |
| 18 | Allele-Specific Suppressors of <i>lin-1(R175Opal)</i> Identify Functions of MOC-3 and DPH-3 in tRNA Modification Complexes in <i>Caenorhabditis elegans</i> . <i>Genetics</i> , 2010, 185, 1235-1247.                   | 2.9  | 7         |

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|----|--|------|-----------|
| 19 | Defects in tRNA Modification Associated with Neurological and Developmental Dysfunctions in <i>Caenorhabditis elegans</i> Elongator Mutants. <i>PLoS Genetics</i> , 2009, 5, e1000561.             | 3.5  | 119       |
| 20 | Eukaryotic Wobble Uracil Modifications Promote a Functionally Redundant Decoding System. <i>Molecular and Cellular Biology</i> , 2008, 28, 3301-3312.  | 2.3  | 219       |
| 21 | A genome-wide screen identifies genes required for formation of the wobble nucleoside 5-methoxycarbonylmethyl-2-thiouridine in <i>Saccharomyces cerevisiae</i> . <i>Rna</i> , 2008, 14, 2183-2194. | 3.5  | 170       |
| 22 | <i>Kluyveromyces lactis</i> $\hat{A}$ -toxin, a ribonuclease that recognizes the anticodon stem loop of tRNA. <i>Nucleic Acids Research</i> , 2007, 36, 1072-1080.                                 | 14.5 | 49        |
| 23 | A conserved modified wobble nucleoside (mcm5s2U) in lysyl-tRNA is required for viability in yeast. <i>Rna</i> , 2007, 13, 1245-1255.   | 3.5  | 166       |
| 24 | Elevated Levels of Two tRNA Species Bypass the Requirement for Elongator Complex in Transcription and Exocytosis. <i>Molecular Cell</i> , 2006, 24, 139-148.                                       | 9.7  | 247       |
| 25 | An early step in wobble uracil tRNA modification requires the Elongator complex. <i>Rna</i> , 2005, 11, 424-436.   | 3.5  | 382       |
| 26 | The <i>Kluyveromyces lactis</i> $\hat{A}$ -toxin targets tRNA anticodons. <i>Rna</i> , 2005, 11, 1648-1654.  | 3.5  | 187       |