

Selection of Candidate Reference Genes for Gene Expression Analysis in Bluegrass (*Poa pratensis* L.) under Abiotic Stress

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Selection and Validation of Appropriate Reference Genes for Quantitative Real-Time PCR Normalization in Staminate and Perfect Flowers of Andromonoecious <i>Taihangia rupestris</i> . <i>Frontiers in Plant Science</i> , 2017, 8, 729.	3.6	24
2	Identification and Validation of Reference Genes for Seashore <i>Paspalum</i> Response to Abiotic Stresses. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1322.	4.1	27
3	Identification and validation of reference genes for quantitative real-time PCR under salt stress in a halophyte, <i>Sesuvium portulacastrum</i> . <i>Plant Gene</i> , 2018, 13, 18-24.	2.3	9
4	Identification of appropriate reference genes for RT-qPCR analysis in <i>Juglans regia</i> L.. <i>PLoS ONE</i> , 2018, 13, e0209424.	2.5	12
5	Physiological and Proteomic Responses of Contrasting Alfalfa (<i>Medicago sativa</i> L.) Varieties to PEG-Induced Osmotic Stress. <i>Frontiers in Plant Science</i> , 2018, 9, 242.	3.6	81
6	Selection and evaluation of reference genes for qRT-PCR analysis in <i>Euscaphis konishii</i> Hayata based on transcriptome data. <i>Plant Methods</i> , 2018, 14, 42.	4.3	42
7	The positive effects of exogenous 5-aminolevulinic acid on the chlorophyll biosynthesis, photosystem and calvin cycle of Kentucky bluegrass seedlings in response to osmotic stress. <i>Environmental and Experimental Botany</i> , 2018, 155, 260-271.	4.2	38
8	Identifying optimal reference genes for the normalization of microRNA expression in cucumber under viral stress. <i>PLoS ONE</i> , 2018, 13, e0194436.	2.5	12
9	Selection of Suitable Reference Genes for RT-qPCR Gene Expression Analysis in Siberian Wild Rye (<i>Elymus sibiricus</i>) under Different Experimental Conditions. <i>Genes</i> , 2019, 10, 451.	2.4	26
10	Bioaccumulation, antioxidative response, and metallothionein expression in <i>Lupinus luteus</i> L. exposed to heavy metals and silver nanoparticles. <i>Environmental Science and Pollution Research</i> , 2019, 26, 16040-16052.	5.3	31
11	Selection and validation of reference genes for target gene analysis with quantitative real-time PCR in the leaves and roots of <i>Carex rigescens</i> under abiotic stress. <i>Ecotoxicology and Environmental Safety</i> , 2019, 168, 127-137.	6.0	28
12	Identification of Appropriate Reference Genes for Normalizing miRNA Expression in Citrus Infected by <i>Xanthomonas citri</i> subsp. <i>citri</i> . <i>Genes</i> , 2020, 11, 17.	2.4	8
13	Physiological and transcriptome analysis of <i>Poa pratensis</i> var. <i>anceps</i> cv. Qinghai in response to cold stress. <i>BMC Plant Biology</i> , 2020, 20, 362.	3.6	28
14	<i>Carex muskingumensis</i> and Osmotic Stress: Identification of Reference Genes for Transcriptional Profiling by RT-qPCR. <i>Genes</i> , 2020, 11, 1022.	2.4	3
15	Screening and verification of reference genes for analysis of gene expression in winter rapeseed (<i>Brassica rapa</i> L.) under abiotic stress. <i>PLoS ONE</i> , 2020, 15, e0236577.	2.5	16
16	Identification of suitable reference genes for normalization of real-time quantitative PCR data in pecan (<i>Carya illinoensis</i>). <i>Trees - Structure and Function</i> , 2020, 34, 1233-1241.	1.9	12
17	Transcriptional regulation and expression network responding to cadmium stress in a Cd-tolerant perennial grass <i>Poa Pratensis</i> . <i>Chemosphere</i> , 2020, 250, 126158.	8.2	33
18	Selection and evaluation of reference genes for qRT-PCR of <i>Scutellaria baicalensis</i> Georgi under different experimental conditions. <i>Molecular Biology Reports</i> , 2021, 48, 1115-1126.	2.3	16

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19	Selection and Evaluation of Potential Reference Genes for Quantitative Real-Time PCR in <i>Agaricus blazei</i> Based on Transcriptome Sequencing Data. <i>BioMed Research International</i> , 2021, 2021, 1-13.	1.9	2
20	Selection and validation of appropriate reference genes for RT-qPCR analysis of flowering stages and different genotypes of <i>Iris germanica</i> L. <i>Scientific Reports</i> , 2021, 11, 9901.	3.3	11
21	A Review on Kentucky Bluegrass Responses and Tolerance to Drought Stress. , 0, , .		2
22	Reference Gene Selection for Quantitative Real-Time PCR Analyses of <i>Acer palmatum</i> under Abiotic Stress. <i>Phyton</i> , 2020, 89, 385-403.	0.7	5
23	Establishment of reference (housekeeping) genes via quantitative real-time PCR for investigation of the genomic basis of abiotic stress resistance in <i>Psammochloa villosa</i> (Poaceae). <i>Journal of Plant Physiology</i> , 2022, 268, 153575.	3.5	5
24	Reference gene selection for quantitative RT-PCR in <i>Miscanthus sacchariflorus</i> under abiotic stress conditions. <i>Molecular Biology Reports</i> , 2022, 49, 907-915.	2.3	5
25	Raman-Deuterium Isotope Probing and Metagenomics Reveal the Drought Tolerance of the Soil Microbiome and Its Promotion of Plant Growth. <i>MSystems</i> , 2022, 7, e0124921.	3.8	7
33	Identification of Suitable Reference Genes for qRT-PCR Normalization in <i>Tilia miqueliana</i> Maxim. <i>Phyton</i> , 2022, 91, 2191-2210.	0.7	0
34	Prune homolog 2 with BCH domain (PRUNE2) gene expression is associated with feed efficiency-related traits in Nelore steers. <i>Mammalian Genome</i> , 2022, 33, 629-641.	2.2	1
35	Transcriptional Response and Plant Growth Promoting Activity of <i>Pseudomonas fluorescens</i> DR397 under Drought Stress Conditions. <i>Microbiology Spectrum</i> , 2022, 10, .	3.0	13
36	Analysis of <i>Elymus nutans</i> seed coat development elucidates the genetic basis of metabolome and transcriptome underlying seed coat permeability characteristics. <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	0
37	Reference Gene Selection for qPCR Analysis in <i>Schima superba</i> under Abiotic Stress. <i>Genes</i> , 2022, 13, 1887.	2.4	7
38	Comparative transcriptome analysis of resistant and susceptible Kentucky bluegrass varieties in response to powdery mildew infection. <i>BMC Plant Biology</i> , 2022, 22, .	3.6	4
39	24-epibrassinolide improves cadmium tolerance and lateral root growth associated with regulating endogenous auxin and ethylene in Kentucky bluegrass. <i>Ecotoxicology and Environmental Safety</i> , 2023, 249, 114460.	6.0	2
40	Selection and validation of appropriate reference genes for RT-qPCR analysis of <i>Nitraria sibirica</i> under various abiotic stresses. <i>BMC Plant Biology</i> , 2022, 22, .	3.6	8
41	Screening and validating of endogenous reference genes in <i>Chlorella</i> sp. TLD 6B under abiotic stress. <i>Scientific Reports</i> , 2023, 13, .	3.3	1
42	Carbon and nitrogen metabolism affects kentucky bluegrass rhizome expansion. <i>BMC Plant Biology</i> , 2023, 23, .	3.6	1
43	Identification and Validation of Reference Genes for Expression Analysis Using RT-qPCR in <i>Leptocybe invasa</i> Fisher and La Salle (Hymenoptera: Eulophidae). <i>Insects</i> , 2023, 14, 456.	2.2	3

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45	Selection of Suitable Reference Genes for RT-qPCR Gene Expression Analysis in Centipedegrass under Different Abiotic Stress. <i>Genes</i> , 2023, 14, 1874.	2.4	0
46	Selection and Validation of Reference Genes for qRT-PCR Analysis of Gene Expression in <i>Tropaeolum majus</i> (Nasturtium). <i>Horticulturae</i> , 2023, 9, 1176.	2.8	0
47	Selection and Verification of Standardized Reference Genes of <i>Angelica dahurica</i> under Various Abiotic Stresses by Real-Time Quantitative PCR. <i>Genes</i> , 2024, 15, 79.	2.4	1