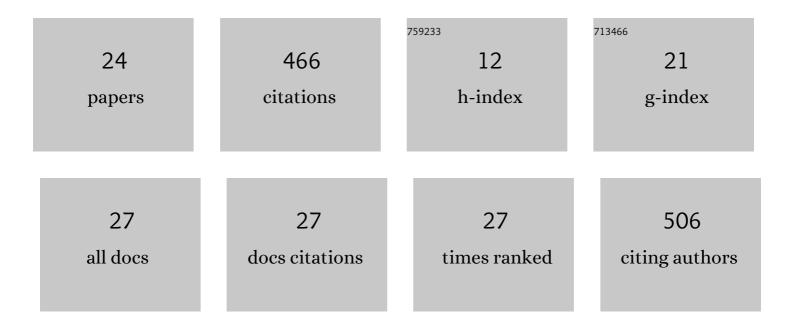
Nadeem Khan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2847787/publications.pdf Version: 2024-02-01



| # | Article | IF | CITATIONS |
|---|---|-----|-----------|
| 1 | Bioengineered Plants Can Be a Useful Source of Omega-3 Fatty Acids. BioMed Research International, 2017, 2017, 1-9. | 1.9 | 49 |

$_{2}$ Transcriptome Sequence Analysis Elaborates a Complex Defensive Mechanism of Grapevine (Vitis) Tj ETQq0 0 0 rgBT/Overlock 10 Tf 50

| 3 | Physiological and transcriptional variations inducing complex adaptive mechanisms in grapevine by salt stress. Environmental and Experimental Botany, 2019, 162, 455-467. | 4.2 | 42 |
|----|---|-------------------|-------------|
| 4 | Genome-wide identification of ATP binding cassette (ABC) transporter and heavy metal associated (HMA) gene families in flax (Linum usitatissimum L.). BMC Genomics, 2020, 21, 722. | 2.8 | 42 |
| 5 | Genome-Wide Identification and Expression Profiling of the Polygalacturonase (PG) and Pectin Methylesterase (PME) Genes in Grapevine (Vitis vinifera L.). International Journal of Molecular Sciences, 2019, 20, 3180. | 4.1 | 37 |
| 6 | Identification and Expression Profiling of Protein Phosphatases (PP2C) Gene Family in Gossypium hirsutum L International Journal of Molecular Sciences, 2019, 20, 1395. | 4.1 | 34 |
| 7 | Genome-Wide Identification, Evolution, and Transcriptional Profiling of <i>PP2C </i> Gene Family in <i>Brassica rapa </i> . BioMed Research International, 2019, 2019, 1-15. | 1.9 | 30 |
| 8 | Genome-wide Identification, Classification, and Expression Pattern of Homeobox Gene Family in Brassica rapa under Various Stresses. Scientific Reports, 2018, 8, 16265. | 3.3 | 28 |
| 9 | Genome-wide identification, evolution, and molecular characterization of the PP2C gene family in woodland strawberry. Gene, 2019, 702, 27-35. | 2.2 | 23 |
| 10 | Genome-Wide Identification, Classification, and Expression Divergence of Glutathione-Transferase Family in Brassica rapa under Multiple Hormone Treatments. BioMed Research International, 2018, 2018, 1-19. | 1.9 | 21 |
| 11 | Genome-wide identification of the class III POD gene family and their expression profiling in grapevine (Vitis vinifera L). BMC Genomics, 2020, 21, 444. | 2.8 | 19 |
| 12 | Untargeted metabolomics and comparative flavonoid analysis reveal the nutritional aspects of pak choi. Food Chemistry, 2022, 383, 132375. | 8.2 | 17 |
| 13 | Evolution and Expression Divergence of E2 Gene Family under Multiple Abiotic and Phytohormones Stresses in Brassica rapa. BioMed Research International, 2018, 2018, 1-18. | 1.9 | 12 |
| 14 | Comprehensive Sequence Analysis of IQD Gene Family and their Expression Profiling in Grapevine (Vitis) Tj ETQq(|) 0 0 rgBT 2.4 | /Overlock 1 |
| 15 | Insights into the Genetic Architecture and Genomic Prediction of Powdery Mildew Resistance in Flax (Linum usitatissimum L.). International Journal of Molecular Sciences, 2022, 23, 4960. | 4.1 | 12 |
| 16 | Lipidomic study reveals the effect of morphological variation and other metabolite interactions on the lipid composition in various cultivars of Bok choy. Biochemical and Biophysical Research Communications, 2018, 506, 755-764. | 2.1 | 9 |

| 17 | Identification, Evolution, and Expression Profiling of Histone Lysine Methylation Moderators in Brassica rapa. Plants, 2019, 8, 526. | 3.5 | 8 |
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¹⁸Avenues for biofortification of zinc in barley for human and animal health: a meta-analysis. Plant and
Soil, 2021, 466, 101-119.3.75

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| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | Transcriptome Analysis Revealed the Molecular Response Mechanism of Non-heading Chinese Cabbage to Iron Deficiency Stress. Frontiers in Plant Science, 2022, 13, 848424. | 3.6 | 5 |
| 20 | GhHAI2, GhAHG3, and GhABI2 Negatively Regulate Osmotic Stress Tolerance via ABA-Dependent Pathway in Cotton (Gossypium hirsutum L.). Frontiers in Plant Science, 2022, 13, . | 3.6 | 5 |
| 21 | Using Portable X-ray Fluorescence Spectroscopy for Inexpensive and Quick Determination of Micronutrients in Barley Shoots. Communications in Soil Science and Plant Analysis, 2022, 53, 1379-1384. | 1.4 | 4 |
| 22 | Designing Genomic Solutions to Enhance Abiotic Stress Resistance in Flax. , 2022, , 251-283. | | 2 |
| 23 | Genome-Wide Identification of Strawberry Metal Tolerance Proteins and Their Expression under Cadmium Toxicity. Horticulturae, 2022, 8, 477. | 2.8 | 2 |
| 24 | Topping Inhibited Potassium Uptake via Regulating Potassium Flux and Channel Gene Expression in Tobacco. Agronomy, 2022, 12, 1166. | 3.0 | 0 |