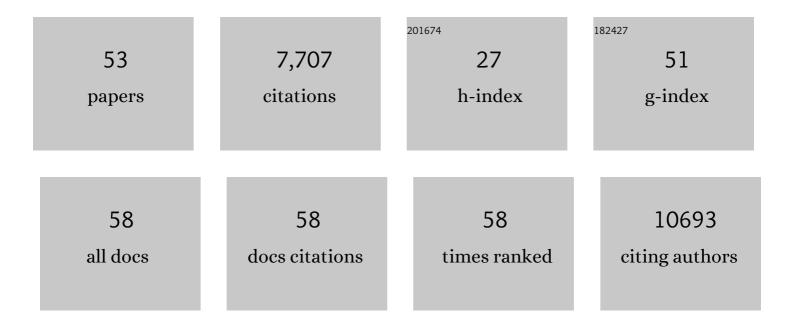
## Neha Garg

List of Publications by Year in descending order

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| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | GNPS Dashboard: collaborative exploration of mass spectrometry data in the web browser. Nature Methods, 2022, 19, 134-136.  | 19.0 | 35        |
| 2  | Molecular networking-based strategies in mass spectrometry coupled with in silico dereplication of peptidic natural products and gene cluster analysis. Methods in Enzymology, 2022, 663, 273-302.            | 1.0  | 1         |
| 3  | Metabolomics Approaches to Dereplicate Natural Products from Coral-Derived Bioactive Bacteria.<br>Journal of Natural Products, 2022, 85, 462-478.   | 3.0  | 14        |
| 4  | A Silent Biosynthetic Gene Cluster from a Methanotrophic Bacterium Potentiates Discovery of a<br>Substrate Promiscuous Proteusin Cyclodehydratase. ACS Chemical Biology, 2022, 17, 1577-1585.                 | 3.4  | 14        |
| 5  | Metabolomics Analysis of Bacterial Pathogen <i>Burkholderia thailandensis</i> and Mammalian Host<br>Cells in Co-culture. ACS Infectious Diseases, 2022, 8, 1646-1662.   | 3.8  | 3         |
| 6  | A community resource for paired genomic and metabolomic data mining. Nature Chemical Biology, 2021, 17, 363-368.  | 8.0  | 81        |
| 7  | Presence of Bromotyrosine Alkaloids in Marine Sponges Is Independent of Metabolomic and Microbiome Architectures. MSystems, 2021, 6, .  | 3.8  | 18        |
| 8  | LanCLs add glutathione to dehydroamino acids generated at phosphorylated sites in the proteome.<br>Cell, 2021, 184, 2680-2695.e26.  | 28.9 | 34        |
| 9  | The role of algal chemical defenses in the feeding preferences of the long-spined sea urchin Diadema antillarum. Aquatic Ecology, 2021, 55, 941-953.  | 1.5  | 4         |
| 10 | Structural and mechanistic investigations of protein S-glycosyltransferases. Cell Chemical Biology, 2021, 28, 1740-1749.e6.   | 5.2  | 8         |
| 11 | Enzymatic Synthesis Assisted Discovery of Prolineâ€Rich Macrocyclic Peptides in Marine Sponges.<br>ChemBioChem, 2021, 22, 2614-2618.  | 2.6  | 9         |
| 12 | An Obligate Peptidyl Brominase Underlies the Discovery of Highly Distributed Biosynthetic Gene<br>Clusters in Marine Sponge Microbiomes. Journal of the American Chemical Society, 2021, 143,<br>10221-10231. | 13.7 | 22        |
| 13 | Metabolomics in Functional Interrogation of Individual Holobiont Members. MSystems, 2021, 6, e0084121.  | 3.8  | 7         |
| 14 | Metabolomics of Healthy and Stony Coral Tissue Loss Disease Affected Montastraea cavernosa Corals.<br>Frontiers in Marine Science, 2021, 8, .   | 2.5  | 12        |
| 15 | Metabolomic profiling of Burkholderia cenocepacia in synthetic cystic fibrosis sputum medium reveals nutrient environment-specific production of virulence factors. Scientific Reports, 2021, 11, 21419.      | 3.3  | 9         |
| 16 | Stereochemical Assignment and Absolute Abundance of Nonproteinogenic Amino Acid Homoarginine in Marine Sponges. ACS Omega, 2021, 6, 33200-33205.  | 3.5  | 2         |
| 17 | Precursor-Guided Mining of Marine Sponge Metabolomes Lends Insight into Biosynthesis of<br>Pyrrole–Imidazole Alkaloids. ACS Chemical Biology, 2020, 15, 2185-2194.  | 3.4  | 9         |
| 18 | Feature-based molecular networking in the GNPS analysis environment. Nature Methods, 2020, 17,<br>905-908.  | 19.0 | 650       |

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|----|---|------|-----------|
| 19 | Differences in Cystic Fibrosis-Associated <i>Burkholderia</i> spp. Bacteria Metabolomes after Exposure to the Antibiotic Trimethoprim. ACS Infectious Diseases, 2020, 6, 1154-1168.         | 3.8  | 14        |
| 20 | Multi-Omic Profiling of Melophlus Sponges Reveals Diverse Metabolomic and Microbiome<br>Architectures that Are Non-overlapping with Ecological Neighbors. Marine Drugs, 2020, 18, 124.      | 4.6  | 21        |
| 21 | Global chemical effects of the microbiome include new bile-acid conjugations. Nature, 2020, 579, 123-129.   | 27.8 | 316       |
| 22 | Molecular and Microbial Microenvironments in Chronically Diseased Lungs Associated with Cystic<br>Fibrosis. MSystems, 2019, 4, .  | 3.8  | 23        |
| 23 | Mass Spectrometry-Based Integration and Expansion of the Chemical Diversity Harbored Within a<br>Marine Sponge. Journal of the American Society for Mass Spectrometry, 2019, 30, 1373-1384. | 2.8  | 18        |
| 24 | Creating a 3D microbial and chemical snapshot of a human habitat. Scientific Reports, 2018, 8, 3669.  | 3.3  | 34        |
| 25 | Chemoenzymatic Synthesis of Starting Materials and Characterization of Halogenases Requiring Acyl<br>Carrier Protein-Tethered Substrates. Methods in Enzymology, 2018, 604, 333-366.        | 1.0  | 3         |
| 26 | The chemical topology of a bacterial swarm. Journal of Biological Chemistry, 2018, 293, 9553-9554.  | 3.4  | 0         |
| 27 | Two Flavoenzymes Catalyze the Post-Translational Generation of 5-Chlorotryptophan and<br>2-Aminovinyl-Cysteine during NAI-107 Biosynthesis. ACS Chemical Biology, 2017, 12, 548-557.        | 3.4  | 64        |
| 28 | Three-Dimensional Microbiome and Metabolome Cartography of a Diseased Human Lung. Cell Host and Microbe, 2017, 22, 705-716.e4.  | 11.0 | 111       |
| 29 | Multi-omics Analysis of Periodontal Pocket Microbial Communities Pre- and Posttreatment. MSystems, 2017, 2, .   | 3.8  | 47        |
| 30 | Metabolic Fingerprints from the Human Oral Microbiome Reveal a Vast Knowledge Gap of Secreted<br>Small Peptidic Molecules. MSystems, 2017, 2, .   | 3.8  | 30        |
| 31 | Dereplication of peptidic natural products through database search of mass spectra. Nature Chemical<br>Biology, 2017, 13, 30-37.  | 8.0  | 184       |
| 32 | Natural products as mediators of disease. Natural Product Reports, 2017, 34, 194-219.   | 10.3 | 59        |
| 33 | Digitizing mass spectrometry data to explore the chemical diversity and distribution of marine cyanobacteria and algae. ELife, 2017, 6, .   | 6.0  | 33        |
| 34 | Characterization of the stereochemical configuration of lanthionines formed by the lanthipeptide synthetase <scp>G</scp> eo <scp>M</scp> . Biopolymers, 2016, 106, 834-842.                 | 2.4  | 11        |
| 35 | Microbiome-wide association studies link dynamic microbial consortia to disease. Nature, 2016, 535, 94-103.   | 27.8 | 595       |
| 36 | Spatial Molecular Architecture of the Microbial Community of a <i>Peltigera</i> Lichen. MSystems, 2016, 1, .  | 3.8  | 36        |

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|----|--|------|-----------|
| 37 | Mass Spectrometry-Based Visualization of Molecules Associated with Human Habitats. Analytical Chemistry, 2016, 88, 10775-10784.                                    | 6.5  | 44        |
| 38 | Sharing and community curation of mass spectrometry data with Global Natural Products Social<br>Molecular Networking. Nature Biotechnology, 2016, 34, 828-837.     | 17.5 | 2,802     |
| 39 | Microbial, host and xenobiotic diversity in the cystic fibrosis sputum metabolome. ISME Journal, 2016, 10, 1483-1498.  | 9.8  | 88        |
| 40 | Minimum Information about a Biosynthetic Gene cluster. Nature Chemical Biology, 2015, 11, 625-631.   | 8.0  | 715       |
| 41 | Chemoenzymatic Synthesis of Acyl Coenzyme A Substrates Enables <i>in Situ</i> Labeling of Small Molecules and Proteins. Organic Letters, 2015, 17, 4452-4455.      | 4.6  | 33        |
| 42 | Mass spectral similarity for untargeted metabolomics data analysis of complex mixtures.<br>International Journal of Mass Spectrometry, 2015, 377, 719-727.         | 1.5  | 90        |
| 43 | Chemical Rescue and Inhibition Studies to Determine the Role of Arg301 in Phosphite Dehydrogenase.<br>PLoS ONE, 2014, 9, e87134.                                   | 2.5  | 12        |
| 44 | Mode of action and structure–activity relationship studies of geobacillin I. Journal of Antibiotics, 2014, 67, 133-136.  | 2.0  | 22        |
| 45 | The Intestinal Metabolome: An Intersection Between Microbiota and Host. Gastroenterology, 2014, 146, 1470-1476.  | 1.3  | 227       |
| 46 | Mass spectrometry of natural products: current, emerging and future technologies. Natural Product<br>Reports, 2014, 31, 718.                                       | 10.3 | 165       |
| 47 | Specialized Metabolites from the Microbiome in Health and Disease. Cell Metabolism, 2014, 20, 719-730.   | 16.2 | 454       |
| 48 | Substrate Specificity of the Lanthipeptide Peptidase ElxP and the Oxidoreductase ElxO. ACS Chemical Biology, 2014, 9, 1718-1725.                                   | 3.4  | 34        |
| 49 | In vitro activity of the nisin dehydratase NisB. Proceedings of the National Academy of Sciences of the<br>United States of America, 2013, 110, 7258-7263.         | 7.1  | 104       |
| 50 | Cloning, sequence analysis, expression of Cyathus bulleri laccase in Pichia pastoris and characterization of recombinant laccase. BMC Biotechnology, 2012, 12, 75. | 3.3  | 45        |
| 51 | Lantibiotics from <i>Geobacillus thermodenitrificans</i> . Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 5241-5246.  | 7.1  | 129       |
| 52 | Production of Lantipeptides in <i>Escherichia coli</i> . Journal of the American Chemical Society, 2011, 133, 2338-2341.   | 13.7 | 161       |
| 53 | Three Dimensional Cartography of Microbiome and Metabolome Data onto Radiological Images of the<br>Human Lung. SSRN Electronic Journal, 0, , .                     | 0.4  | 0         |