

**List of Accepted Posters**

	Firstname	Surname	Abstract_Title
1	Adewale Sakirudeen	Adebayo	The Microbiome and Metabolome in Chronic Urinary Schistosomiasis and Associated bladder pathologies in southwestern Nigeria
2	Nazir	Ahmad	Genetic analysis in white maize ( <i>Zea mays</i> L.) using 6 x 6, diallel mating design
3	Davide	Albanese	PhyloRelief - Phylogenetic-Based Feature Weighting
4	Devendra	Bansal	Genetic diversity of the malaria vaccine candidate genes of <i>Plasmodium falciparum</i> field isolates from Central and East India
5	Asma	Bashir	Exploring host-microbiota interactions in animal models and humans
6	Spyridoula	Charova	A triple protein complex couples T3SS expression and T3SS secretion in <i>Pseudomonas syringae</i> pv. <i>phaseolicola</i>
7	Amina	Cherchali	Growth improving of durum wheat inoculated by <i>Paenibacillus</i> sp under water stress
8	Miguelangel	Cuenca	Complete auxotrophy for D-alanine confers robust in-vivo growth deficiency of smooth <i>E. coli</i> in a model of sterile live bacterial intestinal colonization
9	Chiara	Devirgiliis	Impact of supplementation with a food-derived microbial community on obesity associated inflammation and gut microbiota composition
10	Pamela	Ferretti	Optimizing de novo hybrid assembly from Illumina and Nanopore sequencing for isolate and microbiome samples.
11	Markus	Gildenhard	Microbial determinants of <i>Anopheles gambiae</i> population genetics.
12	Dwijendra Kumar	Gupta	Extracellular Vesicles in Cancer and their Potential for Therapeutic use/
13	Min-Jung	Kwak	Effect of Ocean Acidification on Marine Microbial Interactions and Metabolism in a Mesocosm Experiment
14	Imna Issa	Malele	Trypanosomes and tsetse gut microfauna interaction and their role in the vectorial capacity for African Trypanosomiasis
15	Andrea	Mancini	Gut:liver:brain axis and Hepatic Encephalopathy: in vitro assessment of microbial and ammonia modulation in cirrhosis.
16	Lamjed	Mansour	Xenomatous-like complex in Myxozoan parasites infecting oocytes of teleostean fish
17	Merab	Mirtskhulava	Manganese Mining action on air quality and health
18	Fiona Wanjiku	Moejes	Bacterial Community Profile of <i>Phaeodactylum tricornutum</i> cultures
19	Akbar	Mostajeran	I will send it later

20	Krishnan	Muthukalingan	Bacterial community exploration through Ion torrent sequencing in different treatment stages of tannery CETP
21	Jessil Ann	Pajar	Winning the endgame: within-host interaction among three fungal parasites of <i>Brontispa longissima</i> (Gestro)
22	Reshmi	Raveendran	Using <i>Drosophila</i> as model to study homeostatic interactions between the gut microbiota and dFoxO
23	Ateeq Ur	Rehman	Characterization of Singlet oxygen dependent photodamage mechanism of Photosystem II in cultured <i>Symbiodinium</i> cells and its involvement in coral bleaching
24	Muhammad	Shafiq	The diversity of <i>Wolbachia</i> endosymbiont in the natural populations of the cotton leafhopper, <i>Amrasca devastans</i>
25	Ju Yeon	Song	Whole Metagenomic Analysis of the Rhizosphere Microbiotas of Tomato Cultivars Resistant or Susceptible to a Soil-borne Disease
26	Mohsin	Tariq	Molecular identification of mungbean nodulating bacterium and symbiosis secrets
27	Suganya	Thangaiyan	Bacterial community exploration through ion torrent sequencing in different treatment stages of tannery CETP
28	Atta	Ur Rehman	PIXE Analysis of Groundnut genotypes for toxic elements
29	Iliana	Valdes	Murine preclinical studies of <i>Mycobacterium habana</i> as vaccine candidate against tuberculosis
30	Muhammad	Yasin	Selenium Enriched Natural Prebiotics: Potential Role of <i>Bacillus</i> spp. in Food Enrichment with Essential Nutrients
31	Muhammad	Yasir	Metagenome Analysis Of Bacteria From Uranium Contaminated Soil Using Pyro sequencing Data
32	Moreno	Zolfo	MetaMLST: a computational pipeline for strain-level bacterial typing from metagenomic samples