

Florence, May 27th 2020. To

whom it may concern,

the Manuscript NO: 02537353 (**Exploring the Food-Gut Axis in immunotherapy response of cancer patients**) by **Edda Russo et al.**, which is under consideration for publication in your journal, was re-read by **Dr. Elisangela Miceli, PhD**.

In 2012, Dr. Elisangela Miceli graduated from University of Florence (Italy) with a bachelor's degree in Biotechnologies. Then, in 2014, she completed her master's degree in Medical and Pharmaceutical Biotechnologies.

In 2018, she concluded the PhD course in Evolutionary Biology and Ecology, through a collaborative programme between University of Florence and the University of Ferrara. Along the PhD years, she was author and co-author of twelve scientific papers¹, maturing experience in revising English language of manuscripts and scientific projects.

Today, she works as Study Coordinator within the Oncology Unit at Santo Stefano Hospital in Prato, Italy, experiencing a continuous use of written and spoken English language.

The edit has achieved Grade A: priority publishing; no language polishing required after editing.

Please do not hesitate to contact her if any other addition information is required.



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¹Scientific production

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- Spatial structuring of bacterial communities in epilithic biofilms in the Acquarossa river (Italy). Chiellini C., Miceli E., Bacci G., Fagorzi C., Coppini E., Fibbi D., Bianconi G., Mengoni A., Canganella F., Fani R. (2018). Spatial structuring of bacterial communities in epilithic biofilms in the Acquarossa river (Italy). *FEMS microbiology ecology*, 94(12).
- Antagonism and antibiotic resistance drive a species-specific plant microbiota differentiation in *Echinacea* spp. Maggini V., Miceli E., Fagorzi C., Maida I., Fondi M., Perrin E., Mengoni A., Bogani P., Chiellini C., Mocali S., Fabiani A., Decorosi F., Giovannetti L., Firenzuoli F., Fani R. (2018). Antagonism and antibiotic resistance drive a species-specific plant microbiota differentiation in *Echinacea* spp. *FEMS microbiology ecology*, 94(8).
- Draft Genome Sequence of *Pseudomonas* sp. Strain Ep R1 Isolated from *Echinacea purpurea* Roots and Effective in the Growth Inhibition of Human Opportunistic Pathogens Belonging to the *Burkholderia cepacia* Complex. Maggini V., Presta L., Miceli E., Fondi M., Bosi E., Chiellini C., Fagorzi C., Bogani P., Di Pilato V., Rossolini G. M., Mengoni A., Firenzuoli F., Perrin E., Fani R. (2017) *Genome announce* 5.20 e00351-17.
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- Plant-endophytes interaction influences the secondary metabolism in *Echinacea purpurea* (L.) Moench: an in vitro model. Maggini, V., De Leo, M., Mengoni, A., Gallo, E. R., Miceli, E., Reidel, R. V. B., Biffi S., Pistelli L., Fani R., Firenzuoli F., Bogani P. (2017). *Scientific reports*, 7(1), 16924.
- Phenotypic and genomic characterization of the antimicrobial producer *Rheinheimera* sp. EpRS3 isolated from the medicinal plant *Echinacea purpurea*: insights into its biotechnological relevance. Presta L., Bosi E., Fondi M., Maida I., Perrin E., Miceli E., Maggini V., Bogani P., Firenzuoli F., Di Pilato V., Rossolini G.M., Mengoni A., Fani R. (2017) *Res Microbiol* 168(3):293-305.
- *Arthrobacter* sp. EpRS66 and *Arthrobacter* sp. EpRS71: Draft Genome Sequences from Two Bacteria Isolated from *Echinacea purpurea* Rhizospheric Soil. Presta L., Fondi M., Perrin E., Maida I., Miceli E., Chiellini C., Maggini V., Bogani P., Di Pilato V., Rossolini G.M., Mengoni A., Fani R. (2016) *Front Microbiol* 12:7:1417.
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- Draft Genome Sequence of *Flavobacterium* sp. Strain TAB 87, Able To Inhibit the Growth of Cystic Fibrosis Bacterial Pathogens Belonging to the *Burkholderia cepacia* Complex. Presta L., Inzucchi I., Bosi E., Fondi M., Perrin E., Miceli E., Tutino M.L., Lo Giudice A., de Pascale D., Fani R. (2016) *Genome Announc* 19;4(3).
 - Draft Genome Sequence of *Pseudomonas* sp. EpS/L25, Isolated from the Medicinal Plant *Echinacea purpurea* and Able To Synthesize Antimicrobial Compounds. Presta L., Bosi E., Fondi M., Maida I., Perrin E., Miceli E., Maggini V., Bogani P., Firenzuoli F., Di Pilato V., Rossolini G.M., Mengoni A., Fani R. (2016) *Genome Announc* 5;4(3).