

СПИСОК ПУБЛИКАЦИЙ МАКЕЕВА В.Ю.

1. Permina, EA., Medvedeva, YA., Baeck, PM., Hegde, SR., Mande, SC., Makeev, VJ., Identification of self-consistent modulons from bacterial microarray expression data with the help of structured regulon gene sets. *Journal of biomolecular structure & dynamics*, 2012, vol. 31, issue 1, pp. 115-124.
 2. Zakharevich, NV., Averina, OV., Klimina, KM., Kudryavtseva, AV., Kasianov, AS., Makeev, VJ., Danilenko, VN. Complete genome sequence of *Bifidobacterium longum* GT15: unique genes for Russian strains, *Genome announcements*, 2,6,e01348-14, 2014.
 3. Zakharevich, NV., Averina, OV., Klimina, KM., Kudryavtseva, Anna V., Kasianov, AS., Makeev, VJ., Danilenko, VN. Complete genome sequence of *Bifidobacterium longum* GT15: identification and characterization of unique and global regulatory genes, *Microbial ecology*, 70,3,819-834, 2015.
 4. Shtratnikova, VYu., Belalov, I., Kasianov, AS., Schelkunov, MI., Logacheva MD., Novikov, AD., Shatalov, AA., Gerasimova, TV., Yanenko, AS., Makeev, VJ. The complete genome of the oil emulsifying strain *Thalassolituus oleivorans* K-188 from the Barents Sea, *Marine Genomics* 2017, in press.
<https://doi.org/10.1016/j.margen.2017.08.005>.
 5. Kulakovskiy IV, Vorontsov IE, Yevshin IS, Soboleva AV, Kasianov AS, Ashoor H, Ba-Alawi W, Bajic VB, Medvedeva YA, Kolpakov FA, Makeev VJ. HOCOMOCO: expansion and enhancement of the collection of transcription factor binding sites models. *Nucleic Acids Res.* 2016 Jan 4.;44(D1):D116-25. doi: 10.1093/nar/gkv1249.
-