

Maria F Vecherkovskaya

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8877521/publications.pdf>

Version: 2024-02-01

16
papers

59
citations

1937685

4
h-index

2272923

4
g-index

18
all docs

18
docs citations

18
times ranked

64
citing authors

#	ARTICLE	IF	CITATIONS
1	Microbiota and cancer (review of literature). UĀenye Zapiski Sankt-Peterburgskogo Gosudarstvennogo Medicinskogo Universiteta Im Akad I P Pavlova, 2021, 27, 14-27.	0.2	0
2	ERADICATING PSEUDOMONAS PERSISTENCE IN ADULT PATIENTS WITH CYSTIC FIBROSIS WITH A NEW ALGORITHM OF ANTIBIOTIC SELECTION. Chest, 2021, 160, A570.	0.8	0
3	New algorithm of antibiotic selection reveals the abundance and reduces bacterial load in sputum of adult patients with cystic fibrosis. , 2021, , .		0
4	SPOROBIOTA IN CYSTIC FIBROSIS. Chest, 2020, 157, A75.	0.8	0
5	BACTERIAL AMYLOID IN CYSTIC FIBROSIS. Chest, 2020, 157, A88.	0.8	0
6	ePS6.05 Performance of a new diagnostic tool for the selection of optimal antibiotics for patients with cystic fibrosis (clinical efficacy data). Journal of Cystic Fibrosis, 2020, 19, S53.	0.7	0
7	P112 A New role of bacterial amyloid in cystic fibrosis. Journal of Cystic Fibrosis, 2020, 19, S87.	0.7	0
8	P113 The role of sporobiota in cystic fibrosis. Journal of Cystic Fibrosis, 2020, 19, S87.	0.7	0
9	FASTER AND MORE PRECISE THAN EVER: NEW DIAGNOSTIC TOOL FOR THE SELECTION OF THE MOST EFFECTIVE ANTIBIOTICS FOR PATIENTS WITH CYSTIC FIBROSIS (CLINICAL EFFICACY DATA). Chest, 2020, 157, A78.	0.8	0
10	Clinical efficacy data for new diagnostic tool for the selection of the most effective antibiotics for patients with cystic fibrosis. , 2020, , .		0
11	Complete Genome Sequence of <i>Kluyvera intestini</i> sp. nov., Isolated from the Stomach of a Patient with Gastric Cancer. Genome Announcements, 2017, 5, .	0.8	26
12	LITTLE-KNOWN BACTERIA ISOLATED FROM PATIENTS WITH VARIOUS DISEASES. UĀenye Zapiski Sankt-Peterburgskogo Gosudarstvennogo Medicinskogo Universiteta Im Akad I P Pavlova, 2017, 24, 35-39.	0.2	0
13	Genomic characterization and assessment of the virulence and antibiotic resistance of the novel species <i>Paenibacillus</i> sp. strain VT-400, a potentially pathogenic bacterium in the oral cavity of patients with hematological malignancies. Gut Pathogens, 2016, 8, 6.	3.4	20
14	Complete Genome Sequence of <i>Paenibacillus</i> sp. Strain VT 400, Isolated from the Saliva of a Child with Acute Lymphoblastic Leukemia. Genome Announcements, 2015, 3, .	0.8	7
15	Oral microflora in children with hematologic malignancies. Oncogematologiya, 2015, 10, 51.	0.3	0
16	Complete Genome Sequence of the <i>Streptococcus</i> sp. Strain VT 162, Isolated from the Saliva of Pediatric Oncohematology Patients. Genome Announcements, 2014, 2, .	0.8	6