

Azeez A Alade

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2000726/publications.pdf>

Version: 2024-02-01

11
papers

71
citations

1684188

5
h-index

1720034

7
g-index

11
all docs

11
docs citations

11
times ranked

49
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Efficacy of hypermethylated DNA biomarkers in saliva and oral swabs for oral cancer diagnosis: Systematic review and meta-analysis. <i>Oral Diseases</i> , 2022, 28, 541-558. | 3.0 | 11 |
| 2 | Variant analyses of candidate genes in orofacial clefts in multi-ethnic populations. <i>Oral Diseases</i> , 2022, 28, 1921-1935. | 3.0 | 3 |
| 3 | Genome-wide Gene-by-Sex Interaction Studies Identify Novel Nonsyndromic Orofacial Clefts Risk Locus. <i>Journal of Dental Research</i> , 2022, 101, 465-472. | 5.2 | 6 |
| 4 | Genetic and epigenetic studies in non-syndromic oral clefts. <i>Oral Diseases</i> , 2022, 28, 1339-1350. | 3.0 | 16 |
| 5 | Periconceptional use of vitamin A and the risk of giving birth to a child with nonsyndromic orofacial clefts: A meta-analysis. <i>Birth Defects Research</i> , 2022, , . | 1.5 | 3 |
| 6 | Whole-genome sequencing reveals de-novo mutations associated with nonsyndromic cleft lip/palate. <i>Scientific Reports</i> , 2022, 12, . | 3.3 | 11 |
| 7 | Co-occurrence of orofacial clefts and clubfoot phenotypes in a sub-Saharan African cohort: Whole-exome sequencing implicates multiple syndromes and genes. <i>Molecular Genetics & Genomic Medicine</i> , 2021, 9, e1655. | 1.2 | 3 |
| 8 | Dental Caries Severity and Nutritional Status of Nigerian Preschool Children. <i>JDR Clinical and Translational Research</i> , 2021, , 238008442110021. | 1.9 | 2 |
| 9 | Replication of GWAS significant loci in a sub-Saharan African Cohort with early childhood caries: a pilot study. <i>BMC Oral Health</i> , 2021, 21, 274. | 2.3 | 3 |
| 10 | Variant Analyses of Candidate Genes in Orofacial Clefts in Multi-ethnic Populations. <i>FASEB Journal</i> , 2021, 35, . | 0.5 | 0 |
| 11 | Non-random distribution of deleterious mutations in the DNA and protein-binding domains of IRF6 are associated with Van Der Woude syndrome. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1355. | 1.2 | 13 |