

Resource Summary Report

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Indian Council of Medical Research; New Delhi; India

RRID:SCR_000968

Type: Tool

Proper Citation

Indian Council of Medical Research; New Delhi; India (RRID:SCR_000968)

Resource Information

URL: <http://www.icmr.nic.in/>

Proper Citation: Indian Council of Medical Research; New Delhi; India (RRID:SCR_000968)

Description: A government-funded council for the promotion and coordination of biomedical research in India. The council focuses on specific diseases such as HIV/AIDS, tuberculosis, and leprosy in addition to national health priorities including the management of communicable diseases, fertility control, and mental health research.

Abbreviations: ICMR

Resource Type: institution

Keywords: medical, biomedical, research, government, national, tuberculosis, hiv, aids, leprosy, communicable, disease, fertility, mental health

Related Condition: HIV/AIDS, Tuberculosis, Leprosy

Funding: Government of India through the Department of Health Research Ministry of Health and Family Welfare

Resource Name: Indian Council of Medical Research; New Delhi; India

Resource ID: SCR_000968

Alternate IDs: Wikidata: Q291019, grid.19096.37, Crossref funder ID: 501100001411, ISNI: 0000 0004 1767 225X, nlx_151509

Alternate URLs: <https://ror.org/0492wrx28>

Record Creation Time: 20220129T080204+0000

Ratings and Alerts

No rating or validation information has been found for Indian Council of Medical Research; New Delhi; India.

No alerts have been found for Indian Council of Medical Research; New Delhi; India.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 26 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Mishra AC, et al. (2018) Stratified sero-prevalence revealed overall high disease burden of dengue but suboptimal immunity in younger age groups in Pune, India. PLoS neglected tropical diseases, 12(8), e0006657.

Joshi AP, et al. (2018) In-silico Designing and Testing of Primers for Sanger Genome Sequencing of Dengue Virus Types of Asian Origin. Journal of genomics, 6, 34.

Jajoo M, et al. (2018) Alarming rates of antimicrobial resistance and fungal sepsis in outborn neonates in North India. PloS one, 13(6), e0180705.

Paquet C, et al. (2017) The moderating role of food cue sensitivity in the behavioral response of children to their neighborhood food environment: a cross-sectional study. The international journal of behavioral nutrition and physical activity, 14(1), 86.

Deep DK, et al. (2017) Increased miltefosine tolerance in clinical isolates of Leishmania donovani is associated with reduced drug accumulation, increased infectivity and resistance to oxidative stress. PLoS neglected tropical diseases, 11(6), e0005641.

Senapati R, et al. (2017) HPV genotypes co-infections associated with cervical carcinoma: Special focus on phylogenetically related and non-vaccine targeted genotypes. PloS one, 12(11), e0187844.

Khan IA, et al. (2016) Prevalence and Association of Mycobacterium avium subspecies paratuberculosis with Disease Course in Patients with Ulcero-Constrictive Ileocolonic Disease. PloS one, 11(3), e0152063.

Mohan A, et al. (2016) Urinary Exosomal microRNA-451-5p Is a Potential Early Biomarker of Diabetic Nephropathy in Rats. PloS one, 11(4), e0154055.

Rengasamy M, et al. (2016) Preclinical safety & toxicity evaluation of pooled, allogeneic human bone marrow-derived mesenchymal stromal cells. The Indian journal of medical

research, 144(6), 852.

Roy S, et al. (2016) Leishmania donovani Utilize Sialic Acids for Binding and Phagocytosis in the Macrophages through Selective Utilization of Siglecs and Impair the Innate Immune Arm. PLoS neglected tropical diseases, 10(8), e0004904.

Singh UB, et al. (2016) Genotypic, Phenotypic and Clinical Validation of GeneXpert in Extra-Pulmonary and Pulmonary Tuberculosis in India. PloS one, 11(2), e0149258.

Kumar D, et al. (2016) Interaction of Host Nucleolin with Influenza A Virus Nucleoprotein in the Early Phase of Infection Limits the Late Viral Gene Expression. PloS one, 11(10), e0164146.

Kshatriya GK, et al. (2016) Gender Disparities in the Prevalence of Undernutrition and the Higher Risk among the Young Women of Indian Tribes. PloS one, 11(7), e0158308.

Reddy RB, et al. (2016) Meta-Analyses of Microarray Datasets Identifies ANO1 and FADD as Prognostic Markers of Head and Neck Cancer. PloS one, 11(1), e0147409.

Mawrie D, et al. (2016) Mesenchymal Stem Cells from Human Extra Ocular Muscle Harbor Neuroectodermal Differentiation Potential. PloS one, 11(6), e0156697.

Aggarwal S, et al. (2015) Medical genetics and genomic medicine in India: current status and opportunities ahead. Molecular genetics & genomic medicine, 3(3), 160.

Chugh RM, et al. (2015) Occurrence and control of sporadic proliferation in growth arrested Swiss 3T3 feeder cells. PloS one, 10(3), e0122056.

Sharma K, et al. (2015) Higher prevalence of human papillomavirus infection in adolescent and young adult girls belonging to different Indian tribes with varied socio-sexual lifestyle. PloS one, 10(5), e0125693.

Raj M, et al. (2015) Micro-economic impact of congenital heart surgery: results of a prospective study from a limited-resource setting. PloS one, 10(6), e0131348.

Bhatia V, et al. (2014) Promoter region hypermethylation and mRNA expression of MGMT and p16 genes in tissue and blood samples of human premalignant oral lesions and oral squamous cell carcinoma. BioMed research international, 2014, 248419.