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Publications:

1. **Jha AN**, Sundaravadivel P, Singh VK, Pati SS, Patra PK, Singh L, Thangaraj K; **(2014)**:*MBL2* gene variations and malaria risk in Indian populations. *Infection and Immunity*; 28(1): 52 – 61

2. Rai N, Taher N, Singh M, Chaubey G, **Jha AN**,Singh L, Thangaraj K**(2014):** Relic excavated in western India is of Saint Queen Katevan of Georgia. *Mitochondrion, 14(1): 1-6*

3. **Jha AN**, Sundaravadivel P, Pati SS, Patra PK, Thangaraj K; **(2013)**: Variations in ncRNA gene *LOC284889* and *MIF -794CATT* repeats are associated with malaria susceptibility in Indian populations. *Malaria Journal*; 12:345 ***(BioMed Central: Highly accessed article)***

4. Marcinek P*, **Jha AN***, Shinde V, Sundaramoorthy A, Rajkumar R, et al.**(2013):***LRRK2* and *RIPK2* variants in the *NOD2*-mediated signaling pathway are associated with susceptibility to *Mycobacterium leprae* in Indian populations. *PLoS ONE*; 8(8): e73103**.** (* **Shared first authorship**)

5. **Jha AN**, Singh VK, Pati SS, Patra PK, Singh L, Thangaraj K; **(2013)**: A rare non-synonymous c.102C>G SNP in the *IFNB1*gene might be a risk factor for cerebral malaria in Indian populations. *Infection Genetics Evolution;*14: 369 – 374

6. **Jha AN**, Singh VK, Kumari N, Singh A, Antony J, et al. **(2012)**:IL-4 Haplotype-590T, -34T and Intron-3 VNTR R2 Is Associated with Reduced Malaria Risk among Ancestral Indian Tribal Populations.*PLoS ONE; 7(10)*:e48136