

## RESUME

**Name** : Dr. Vineeta Singh

**Designation** : Scientist 'E'

**Career Interests:** To achieve more insight about the malaria parasite infectivity and pathogenesis which seems to be changing rapidly. Understanding the parasite biology and pathogenicity of *Plasmodium falciparum* and *P. vivax* are the major research areas of my laboratory.

**Educational Qualifications** : Ph.D.- Biotechnology, BITS, Pilani, India

### Publications:

1. Kumar, A., Gahlawat S.K and **Singh V.** Comparative analysis of Plasmodium falciparum dihydrofolate-reductase gene sequences from different regions of India. *Heliyon* 6(4). April **2020**. <https://doi.org/10.1016/j.heliyon.2020.e03715>
2. Kojom, L.P., **Singh, V.** Prevalence of *Plasmodium falciparum* field isolates with deletions in histidine-rich protein 2 and 3 genes in context with sub-Saharan Africa and India: a systematic review and meta-analysis. *Malar J* **19**, 46 (**2020**). <https://doi.org/10.1186/s12936-019-3090-6>.
3. Kumar, A., Singh, S.P., Bhatt, R. **Singh V.** Genetic profiling of the *Plasmodium falciparum* parasite population in uncomplicated malaria from India. *Malar J* **18**, 385 (**2019**) doi:10.1186/s12936-019-3022-5.
4. Saxena R, Kaur J, Hora R, Singh P, **Singh V**, Mishra PC. CX3CL1 binding protein-2 (CBP2) of Plasmodium falciparum binds nucleic acids. *International journal of biological macromolecules*. **2019** Oct 1;138:996-1005. doi: 10.1016/j.ijbiomac.2019.07.178
5. **Vineeta Singh**, Loick Pradel Kojom. Deletions in the *Plasmodium falciparum* histidine-rich protein 2 gene: An emerging threat to the elimination of malaria in India. *J Vect Borne Dis*, **2019**, 56 (1): 85-86.
6. Jasweer kaur, Vikash kumar, Amrit Pal Singh, **Vineeta Singh**, Anjali Bisht, Taru Dube, Jiban Jyoti Panda, Ankita Behl, Prakash Chandra Mishra, Rachna Hora; *Plasmodium falciparum* protein 'PfJ23' hosts distinct binding sites for major virulence factor 'PfEMP1' and Maurer's cleft marker 'PfSBP1'. *Pathogens and Disease*, December **2018**, fty090, <https://doi.org/10.1093/femspd/fty090>
7. Roman DNR, Anne NNR, **Singh V\***, Luther KMM, Chantal NEM, Albert MS. Role of genetic factors and ethnicity on the multiplicity of *Plasmodium falciparum* infection in children with asymptomatic malaria in Yaoundé, Cameroon. *Heliyon*. 2018;4(8):e00760. Published **2018** Aug 30. doi:10.1016/j.heliyon.2018.e00760
8. Kumar V, Kaur J, Singh AP, **Singh V**, Bisht A, Panda JJ, Mishra PC, Hora R. PHISTc protein family members localize to different subcellular organelles and bind Plasmodium falciparum major virulence factor PfEMP-1. *The FEBS journal*. **2018** Jan 1;285(2):294-312.
9. Roman DN, Rosalie NN, Kumar A, Luther KM, **Singh V\***, Albert MS. Asymptomatic Plasmodium malariae infections in children from suburban areas of Yaoundé, Cameroon. *Parasitol Int*. **2018** Mar 67(1):29-33.
10. Parvin Kumar, Kulbir Kadyan, Meenakshi Duhan, Jayant Sindhu, **Vineeta Singh** and Baljeet Singh Saharan. Design, synthesis, conformational and molecular docking study of some novel acyl hydrazone based molecular hybrids as antimalarial and antimicrobial agents. *Chemistry Central Journal* **2017**, 11:115.
11. Baruah UK, Gowthamarajan K, Ravisankar V, Karri VVSR, Simhadri PK, **Singh V**, Babu PP. Design, characterization and antimalarial efficacy of PEGylated galactosylated nano lipid carriers of primaquine phosphate. *Artif Cells Nanomed Biotechnol*. **2017** Oct 31:1-21. doi: 10.1080/21691401.2017.1394870.
12. Baruah UK, Gowthamarajan K, Ravisankar V, Karri VVSR, Simhadri PK, Singh V. Optimisation of chloroquine phosphate loaded nanostructured lipid carriers using Box-Behnken design and its antimalarial efficacy. *J Drug Target*. **2017** Oct 23:1-16. doi: 10.1080/1061186X.2017.1390671.