





It is our great pleasure to announce the formal launch of a project where we are envisaged to decipher the genomic variability of SARS-COV2 virus specifically extracted from patients of Kolkata and West Bengal.

Our objective in this project is to sequence the SARS-COV2 genomes and find specific and commonly observed mutations in them. Further, we would emphasize on understanding the impact of these genomic variabilities on migration and divergence of the virus as well as potential of therapeutic intervention against the virus. Most importantly, we would attempt to investigate the probable correlation between the type of mutations and clinical manifestation of the patients.

Our Team

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