3. OBJECTIVES
1. To design a High Throughput Automated System to identify the genotype of the infecting Hepatitis B virus strains in the Indian population

2. To identify various HBV mutations.

3. To study the association of HBV genotypes and mutants with clinical presentation, severity, progression of disease and development of HCC.

4. To identify and distinguish between mixed HBV strains and recombinants.