BIOSTATISTICS STATEMENT

TITLE: Polymorphisms and resistant mutations of HCV on sequences in the European Hepatitis C Virus Database

Microsoft® Office Excel 2010 was used to store data, The sequence alignment was performed with MEGA 6.06 MAC followed by sequence editing, exclusion of sequences with missing data, and translation of the nucleic acids sequences into amino acids. The resulting protein sequences were then analyzed using the BioEdit 7.2.5. software for mutations associated with resistance.

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