Figure 5B: Phylogenetic analysis of hepatitis C virus 5´ non coding region.

Phylogenetic analysis of 5´ non coding region (nt -240 to -60; 181bp) sequences of 10 HCV samples of HIV seropositive individuals. The sequences for major subtype were selected from the GeneBank database for analysis. The accession Numbers of the reference sequences (with subtypes are as follows: AF011753 (1A), AJ132996 (1B), AY051292 (1C), af238485 (2a), af238486 (2b), L38330 (2c), AF046866 (3a), D49374 (3b), D16612 (3c), D16620 (3d), D16618 (3e), X91421 (3g), Y11604 (4a), Y13184 (5a), Y12083 (6a) and the samples are numbered as HIV following the reference number.