The image contains two graphs labeled (a) and (b). Graph (a) shows cumulative nt substitutions and cumulative amino acid substitutions as a function of nucleotide position. The x-axis represents the genome position, and the y-axis represents the cumulative number of substitutions. The graph shows a slope that indicates a steady increase in substitutions with respect to nucleotide position.

Graph (b) illustrates the dN/dS ratio over amino acid residues, with a focus on specific regions labeled as L, CAPSID, and 3A. The dN/dS ratio is plotted against amino acid residues, where values above 1 suggest positive selection, and values below 1 suggest purifying selection. The graph highlights variations in the dN/dS ratio across different regions, indicating differences in selective pressures at these positions.