



Fig. 1. Per-site mutation rate versus genome size for CChMVd and other biological entities [reviewed in (2) and updated with more recent data from (3)]. RNA viruses (left to right) are tobacco mosaic virus, human rhinovirus, poliovirus, vesicular stomatitis virus, bacteriophage $\Phi 6$, and measles virus. Single-stranded DNA viruses are bacteriophage $\Phi X174$ and bacteriophage m13. Double-stranded DNA viruses are bacteriophage λ , herpes simplex virus, bacteriophage T2, and bacteriophage T4. Bacteria is *Escherichia coli*. Lower eukaryotes are *Saccharomyces cerevisiae* and *Neurospora crassa*. Higher eukaryotes are *Caenorhabditis elegans*, *Drosophila melanogaster*, *Mus musculus*, and *Homo sapiens*. When several estimations were available, the mean value is shown.