図 1. SaV 遺伝子の VP1 領域の一部 (260nt) の塩基配列に基づいて作成した系統樹

The tree was generated using the neighbor-joining method. The bootstrap values correspond to 100 replications.
The number on each branch indicates the bootstrap value, where a value higher than 95 is indicated.
The scale represents nucleotide substitutions per site.
GenBank accession numbers for the reference strains are as follows: Arg39, AY289803; C12, AY603425; Chiba000496,AJ412800;
Cowden, AF182760; Cruiseship, AY289804; Ehime643, DQ366345; Ehime1107, DQ058829; Houston27, U95644; Manchester, X86560;
Mc2, AY237419; Mc10, AY237420; Mc114, AY237422; Mex340, AF435812; NK24, AY646856; Parkville,U73124; Potsdam, AF294739;
Sapporo,U65427; Sakaew15, AY646855; Stockholm, AF194182; SW278, DQ125333; Syd3, DQ104357; Syd53,DQ104360;
and Yokote1 AB253740.