



Figure S1. Electrophoretic migration pattern of genome segments of G8P[6] RVA strains GH018-08 and GH019-08, with representative Ghanaian Wa-like and DS-1-like strains.

Lanes: a) GH018-08; b) GH019-08; c) GH022-08; d) GH024-08
(GH022-08: G1P[8] Wa-like strain; GH024-08: G2P[4] DS-1-like strain.)

GH018-08 and GH019-08 revealed identical long RNA migration patterns, with a notably longer migration distance for gene segment 11 (NSP5) than observed in Wa-like RVA strains (indicated by star and arrow respectively).