

Table S2. PCR outcome for initial S_scheme primer designs for each amplicon targeting a panel of Southeast Asian (pool 1) FMDV isolates.

Virus isolate	Lineage	S	1	2	3	4*	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19
VIT/1/2017	O/SEA/Mya-98	+	+	+	+	+	+	+	+	+/-	+	+	+	+	+	+		+	+	+	+
VIT/1/2018	O/ME-SA/PanAsia	+/-	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+
VIT/11/2017	A/ASIA/Sea-97	+	+	+	+	+/-	+		+	+	+	+	+	+	+	+	+	+/-	+	+	+

*Bold typeface indicates amplicons for which at least one primer is located within the outer capsid coding genes (1B,1C and 1D).