Table S1

Primers for amplification of HLJ isolate complete genome sequence.

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| --- | --- | --- | --- |
| Primera | Sequence (5’–3’) | Primer biding position | Amplified fragment (bp) |
| E1-F | TTGAAAGCCGGGGGTGG | 1-17 | 305 |
| E1-R | GAATGCTCGTCAAGAAGACAG | 285-305 |
| E2-F | CTGTCTTCTTGACGAGCATTC | 285-305 | 2331 |
| E2-R | CAGCGTTTTCTACTCC | 2600-2615 |
| E3-F | TCAAGATGCCTATCTCAC | 2562-2579 | 979 |
| E3-R | CCAAAATTGTCCATGTCTAAAC | 3517-3538 |
| E4-F | GTTTAGACATGGACAATTTTG | 3517-3537 | 4006 |
| E4-R | CCGATCATATTCCTGCTT | 7505-7522 |
| E5-F | GGATTTTGATAAGGTGAGAG | 7252-7271 | 493 |
| E5-R | TTTTTTTTTTTTTTTTCTCTCTATTTATTTTACTACTCTGGAC | 7703-7746 |

Note: a F denotes a forward PCR primer; R denotes reverse transcription or a reverse PCR primer.