|  |  |  |
| --- | --- | --- |
| Name/description | Sequence (5' to 3') | Figure |
| LE31 | GCGTTGCGTTTTTTTGAGTTACCTATATCCC | 1B  |
|  RC | GGGATATAGGTAACTCAAAAAAACGCAACGC | 1B  |
| LE17 | TGAGTTACCTATATCCC | 2A |
|  RC | GGGATATAGGTAACTCA | 2A |
| LE17Pi | GGGATATAGGTAACTCA(Pi) | 2A |
| PiLE17 | TGAGTTACCTATATCCC(Pi) | 2A |
| LE24 | GGGATATAGGTAACTCAAAAAAAC | 2A |
| TR29 | GTTGCGTTTTTTTGAGTTACCTATATCCC  | 3A-C |
| RC  | ATAGGTAACTCAAAAAAACGCAAC | 3A-C |
| dsran17,ranss17  | CTGTTAGGCCACGATTG | 2B |
| RC | CAATCGTGGCCTAACAG | 2B |
| ranTR17 RC  | AGGCCACGATTG  | 2B |
| ssLE11 | ACCTATATCCC | 2C |
| ssLE8 | TATATCCC | 2C |
| ssLE6 | TATCCC | 2C |
| targ40 | CGTGCGTTCGAATCGCACCTCTCGCACTTTTTTTAATTAC | 3D,E |
|  RC | GTAATTAAAAAAAGTGCGAGAGGTGCGATTCGAACGCACG | 3D,E  |
| mutCGCA | CGTGCGTTCGAATCGCACCTCTGTAGCTTTTTTTAATTAC | 3D,E  |
|  RC | GTAATTAAAAAAAGCTACAGAGGTGCGATTCGAACGCACG | 3D,E  |
| mutL-side | CGTGCGTTCGAATCGCACGAGACGCACTTTTTTTAATTAC | 3D,E  |
|  RC | GTAATTAAAAAAAGTGCGTCTCGTGCGATTCGAACGCACG | 3D,E  |
| mutR-side | CGTGCGTTCGAATCGCACCTCTCGCACTTTTAAATTTTAC | 3D,E  |
|  RC | GTAAAATTTAAAAGTGCGAGAGGTGCGATTCGAACGCACG | 3D,E  |
| mut12-17 | CGTGCCAAGCTATCGCACCTCTCGCACTTTTTTTAATTAC | 3D,E  |
|  RC | GTAATTAAAAAAAGTGCGAGAGGTGCGATAGCTTGGCACG | 3D,E  |
| symTSD | TATATCCCCGCACTTAAGTGCG | 5-9 |
| ran40 | CACTATAGGACCATTCCATTCGAGCTAGGACCTAAGTACG | 3D,E  |
|  RC | CGTACTTAGGTCCTAGCTCGAATGGAATGGTCCTATAGTG | 3D,E |
| ssRNA17 | rUrGrArGrUrUrArCrCrUrArUrArUrCrCrC | 3suppl1 |
| LE20 | TTTTGAGTTACCTATATCCC | 3F |
| RC | GGGATATAGGTAACTCAAAA | 3F |
| RC-1 | GGATATAGGTAACTCAAAA | 3F |
| RC-2 | GATATAGGTAACTCAAAA | 3F |
| RC-3 | ATATAGGTAACTCAAAA | 3F |
| RC-4 | TATAGGTAACTCAAAA | 3F |
| RC-5 | ATAGGTAACTCAAAA | 3F |
| RC-6 | TAGGTAACTCAAAA | 3F |
| RC-7 | AGGTAACTCAAAA | 3F |