**Table S2. Nucleotide changes in the mutant strains.**

|  |  |  |  |
| --- | --- | --- | --- |
| **Position** | **Mutation** | **Nucleotide change** | **Nº strains** |
| 347 (CS) | Arg-Ser | CGC-AGC |  |
| 344 | Met-Ile | ATG-ATT  ATG-ATC  ATG-ATA | 1  1  3 |
| 342 | Ser-Arg | AGC-CGC | 4 |
| 315 | Lys-Glu | AAG-GAG | 3 |
| 312+310 | Val-Phe+Glu-Ala | GTC-TTC + GAG-GCG | 1 |
| 308 | Asn-Ile | AAC-ATC | 3 |
| 306 | Leu-Phe | CTT-TTT | 2 |
| 305 | Ile-Leu | ATC-CTC | 1 |
| 302 | Lys-Glu | AAG-GAG | 1 |
| 265 | Leu-Trp | TTG-TGG | 1 (TS) |