**Table S3**: Oligonucleotides Annealing sites to plasmids are underlined. Regions of homology are highlighted in grey.

Primer Sequence

**Oligos for strain construction**

prFAR1-dupB GCAATGTCTTTTTTTTTTCATCCTGTAATGGGTTAAATAACTTCAAAAAGGCAAAATCGATGAATTCGAGCTCG

prFAR1-dupA ACAAAGTCTATAGATCCACTGGAAAGCTTCGTGGGCGTAAGAAGGCAATCTATTAATGTCCAAGGGTGAAGAGC

TUB4-dupB GGAATTAGAGGATCCGTTACGGAAGAGAGAACTTCGTAAGACATTATTGCTGTGACGTACGCTGCAGGTCGAC

TUB4-dupA GCTGCACGGATTAGTTCGATAATTCAACGACAGTTATACTAATCTTGGTGATGGAATCGATGAATTCGAGCTCG

S1-FAR1 AAGTCTATAGATCCACTGGAAAGCTTCGTGGGCGTAAGAAGGCAATCTATTAATGCGTACGCTGCAGGTCG

S2-FAR1 AAAAAAAAGGAAAAGCAAAAGCCTCGAAATACGGGCCTCGATTCCCGAACTACTAATCGATGAATTCGAGC

S2-Tub4 CTATAATAAAACTATTGGGCGGTGGTAAAATTCCTGAACAAGGAAGGCATCATTAATCGATGAATTCGAGCTCG

S3-Tub4 TATGGTTGGCGAGTTGGAAGAGGACCTGGATGCCGACGGTGATCATAAATTAGTACGTACGCTGCAGGTCG

**Validation Oligos**

Far1\_j1f GAAGGTACCTTGGTTCACAG

Far1\_ur TCACGATCTCCACTTGGTG

Far1\_ORF\_f CGTCTAATTATGGATGAACACC

Far1\_termrev ACATTAACTGCTATTTACGACG

Far1\_j1r TATGTCCGTTTACATCTCCG

Far1\_j2f AATGCTGGTCGCTATACTGC

Far1\_j2r TAGTCGCTCCTACAGTGATG

His\_and\_Kan\_Tag2 GAAGGTACCTTGGTTCACAG

Nat-sense GCAGGCGCTCTACATGAGCA

Tub4\_j1f GATTCAACTCAGCGTAATGG

Tub4\_j1r ACAACCTGAAGTCTAGGTCC

Tub4\_j2f GCACGTCAAGACTGTCAAGG

Tub4\_j2r CAAAGAGAGAACGTTATAATGG

**qPCR Oligos**

Tub4\_rt\_f ACCCTTCCAATTCTCTTGTCTC

Tub4\_rt\_r GCGATATCTGACGAGGTTCTAC

Actin\_rt\_f GATTCCGGTGATGGTGTTACTC

Actin\_rt\_r TCAAATCTCTACCGGCCAAATC