

## SUPPLEMENTARY FIGURE S1

### Pre-edited RPS12 (PE-RPS12)

```
1  CUAUACACU UUUGAUACA AACUAAAGUA AAAAGGCGAG GAUUUUUUGA 50
51 GUGGGACUGG AGAGAAAGAG CCGUUCGAGC CCAGCCGGAA CCGACGGAGA 100
101 GCUUCUUUUG AAUAAAAGGG AGGCGGGGAG GAGAGUUUCA AAAAGAUUUG 150
151 GGUGGGGGGA ACCCUUUGUU UUGGUUAAAG AAACAUCGUU UAGAAGAGAU 200
201 UUUAGAAUAA GAUAUGUUUU U 221
```

### Fully edited RPS12 (FE-RPS12)

```
1  CUAUACACU UUUGAUACA AACUAAAGUA AAUAUAUUUU GUUUUUUUUG 50
51 CGUAUGUGAU UUUUGUAUGG UUGUUGUUUA CGUUUUGUUU UAUUUGUUUU 100
101 AUGUUAUUAU AUGAGUCCGC GAUUGCCAG UUCGGUAAC CGACGUGUAU 150
151 UGUAUGCCGU AUUUUAUUUA UAUAUUUUUG UUUGGAUGUU GCGUUGUUUU 200
201 UUUUGUUGUU UUAUUGGUUU AGUUAUGUCA UUAUUUAUUA UAGAGGGUGG 250
251 UGGUUUUGUU GAUUUACCCG GUGUAAAGUA UUAUACACGU AUUGUAAGUU 300
301 AGAUUUAGAU AUAAGUAUG UUUUU 325
```

### gA6[14]

```
1  GGGCGAAUUG AUAUACUUA ACUCCGAUAA CGAAUCAGAU UUUGACAGUG
51 AUAUGAUAAU UAUUUUUUUU UUUUUUUUU 79
```

**FIG. S1.** Sequences of 221 nt pre-edited RPS12, 325 nt fully edited RPS12, and 79 nt gA6 [14]

RNAs that were used in the filter binding experiments.