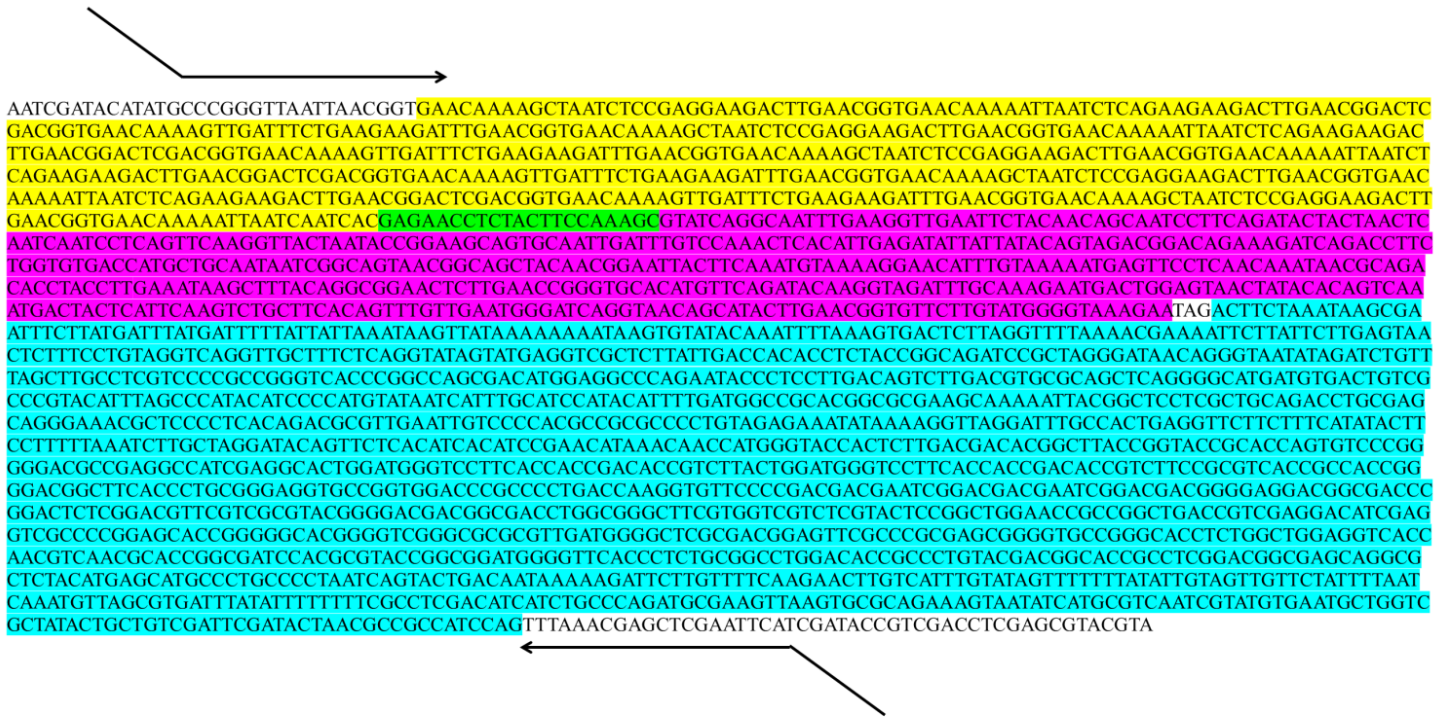


Figure S2

CelTag Fragment Sequence with Primer Alignment



Primer	Sequence (5'-3')
Forward including 13x myc epitope repeat	5'-50 base pair homology to end of protein orf without stop codon CCCGGGTTAATTAACGGTGAA-3'
Forward excluding 13x myc epitope repeat	5'-50 base pair homology to end of protein orf without stop codon GAGAACCTCTACTTCCAAAGC-3'
Reverse	5'-50bp homology 25 base pairs downstream of target orf (reverse complement) ATGAATTCGAGCTCGTTTTAAA-3'

Figure S2: Sequence of the CelTag fragment showing primer design strategy. Yellow is the 13x myc repeat, green indicates the TEV protease site, purple is the CBM3 domain, and blue-green is the selection marker, *nat1* including its promoter and terminator sequences.