str: predicted start of CDS is not valid start codon

	CDS (single exon)	CDS (multi-exon)	mature peptide (MP#i)	CDS (made up of mature peptides)
No errors		exon 1 / exon 2	MP#(i-1) / MP#i / MP#(i+1)	MP1 2 3 4 5 6
str (start) predicted CDS start is not a valid start codon		exon 1 / exon 2	N/A	MP1 2 3 4 5 6

Example output from annotation script:

```
KR006962        3  CDS#3        str  predicted CDS start position is not beginning of ATG start codon
                                      [TNN starting at position 4472 on strand +]
```