

Supplementary Information - Nakagawa *et al.*, 2007

Table S1. Observed and expected numbers of genes containing various trinucleotides

Pattern	<i>O</i>	<i>E</i>	<i>Z</i>	Rank
<u>GCCAUG</u>	11,724	5,254.8	90.3	1
<u>AAAAUG</u>	21,345	13,828.4	66.0	2
<u>AAGAUG</u>	12,692	9,515.6	33.3	3
<u>TTTAUG</u>	1,404	604.4	32.6	4
<u>AGGAUG</u>	5,394	3,848.9	25.1	5
<u>CTGAUG</u>	1,682	952.9	23.7	6
<u>TTGAUG</u>	1,637	1,003.7	20.0	7
<u>ATCAUG</u>	7,489	6,017.1	19.2	8
<u>CTCAUG</u>	1,768	1,250.9	14.7	9
<u>ATTAUG</u>	3,506	2,759.8	14.3	10

The genes used are the same as those in Table 4.