

Sequence logo for a 15-nucleotide motif. The sequence is approximately 5'-AGGCGAATGCGAATG-3'. The logo shows the relative frequency of each nucleotide (A, C, G, T) at each position. The first four positions (AGGCG) are highly conserved, with G being the most frequent at positions 2 and 3. The middle positions (AATG) show more variability, with A and T being the most frequent. The last four positions (CGAATG) are also conserved, with G being the most frequent at positions 12 and 13.