



Sequence logo showing the conservation of nucleotides across 60 positions. The y-axis represents information content in bits. Positions 2, 33, 34, and 50 are highlighted in red. Position 35 is the most conserved, with a large yellow 'G'.

Position	Top Strand (5' to 3')	Bottom Strand (3' to 5')
1	T	G
2	A	C
3	A	C
4	A	C
5	A	C
6	A	C
7	T	A
8	T	A
9	A	C
10	C	G
11	C	G
12	T	C
13	A	C
14	G	A
15	G	A
16	T	C
17	C	G
18	T	C
19	G	A
20	T	C
21	T	C
22	G	A
23	T	C
24	T	C
25	T	C
26	T	C
27	A	C
28	A	C
29	T	C
30	G	A
31	A	C
32	G	A
33	G	C
34	G	C
35	G	C
36	C	G
37	G	A
38	G	A
39	A	C
40	T	C
41	T	C
42	T	C
43	C	G
44	G	A
45	C	G
46	C	G
47	C	G
48	G	C
49	G	C
50	G	C
51	G	C
52	G	C
53	G	C
54	G	C
55	G	C
56	A	C
57	G	C
58	G	C
59	G	C
60	C	G