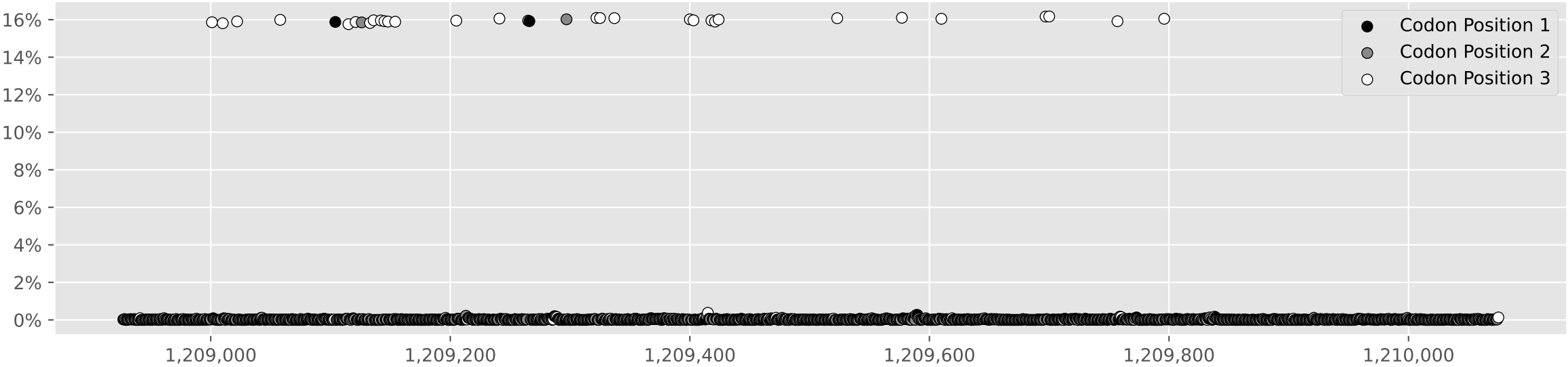


Mutation spectra of three highly-mutated genes

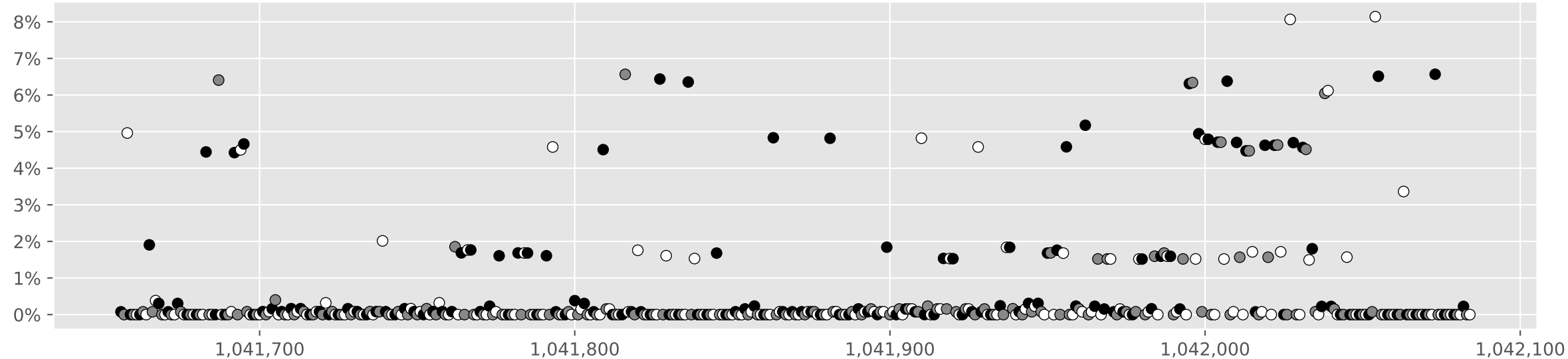
CAMP: Gene #1217

Left end 1,208,927, right end 1,210,075 (length 1,149 bp), strand +
Mutation spectrum ranges from 0.00% to 16.17% (average = 0.49%)



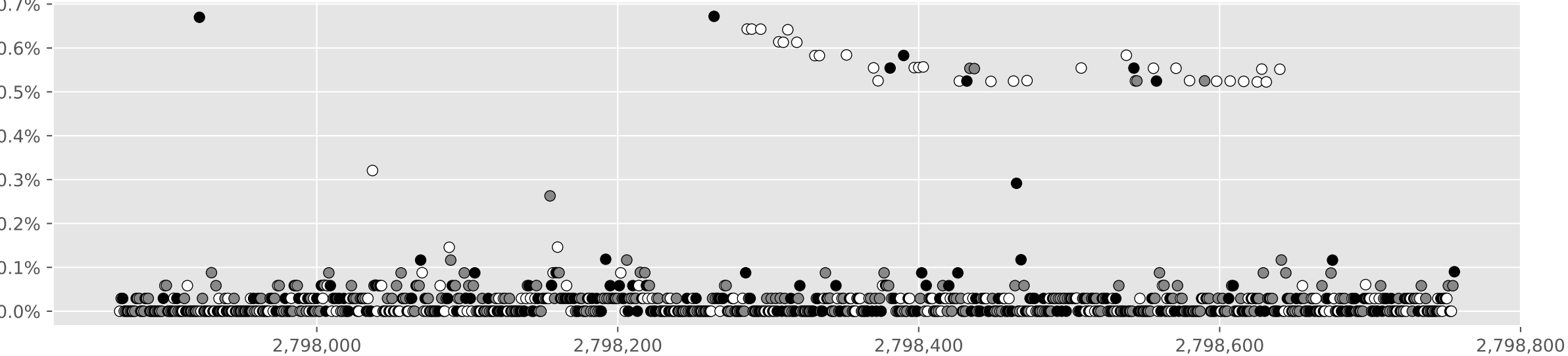
BACT1: Gene #868

Left end 1,041,656, right end 1,042,084 (length 429 bp), strand +
Mutation spectrum ranges from 0.00% to 8.14% (average = 0.71%)



BACT2: Gene #2561

Left end 2,797,869, right end 2,798,756 (length 888 bp), strand -
Mutation spectrum ranges from 0.00% to 0.67% (average = 0.05%)



Sequence position