Genetic loci associated with fluoride resistance in streptococcus mutans

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Citation for published version (APA):
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Supplementary Table 2. Single nucleotide polymorphisms identified from the genomes of *S. mutans* UA159 and UA159-FR

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<th>SNP base</th>
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a) nonsyn: non-synonymous coding SNP.  
b) *: stop codon.  
c) downstream: the gene / genes downstream the intergenic region with the SNP.  
d) upstream: the gene / genes upstream the intergenic region with the SNP.
Supplementary Figure 1. Growth curve of *S. mutans* UA159 and UA159-FR in the absence of NaF in BHI broth. Mean optical density at 600 nm ± standard deviation is shown. This experiment was performed with triplicates.
Supplementary Figure 2. Growth of *S. mutans* (A) UA159 and (B) UA159-FR in BHI broth supplemented with 0, 5, 10, 20, 40, 60 mM NaF. Data are expressed as means ± standard deviation of triplicates.
Supplementary Figure 3. Fluorescence intensities of *mutp* reporter strains, *S. mutans* UA159 containing the wild-type *mutp*, *mutp* from C180-2FR, or *mutp* from UA159-FR. *** $p < 0.0005$. 