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Identification of genes involved in fluoride resistance in oral streptococci

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https://doi.org/10.15017/1785374

出版情報:九州大学,2016,博士(歯学),課程博士 バージョン: 権利関係:全文ファイル公表済

氏 名	門暁晨	
論 文 名	Identification of genes invol	ved in fluoride resistance in oral
	streptococci	
(口腔レンサ球菌のフッ化物耐性に関与する遺伝子の同定)		
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論文審査の結果の要旨

Recently, it has been reported that eriC and crcB are involved in bacterial fluoride resistance. However, the fluoride-resistance mechanism in oral streptococci remains unclear.

BLAST studies showed that two types of eriCs (eriC1 and eriC2) and two types of crcBs (crcB1 and crcB2) are present across 18 oral streptococci, which were selected based on the following criteria: identification in $\geq 10\%$ of 166 orally healthy subjects and $\geq 0.01\%$ of the mean relative abundance. They were divided into three groups based on the distribution of these four genes: group I, only eriC1; group II, eriC1 and eriC2; and group III, eriC2, crcB1, and crcB2.

Group I consisted of *Streptococcus mutans*, in which one of the two *eriC1*s predominantly affected fluoride resistance. Group II consisted of eight species, in which *eriC1* was involved in fluoride resistance but *eriC2* was not in *Streptococcus anginosus* as a representative species. Group III consisted of nine species, in which both *crcB1* and *crcB2* were crucial for fluoride resistance, but *eriC2* was not, in *Streptococcus sanguinis* as a representative species. Based on these results, either EriC1 or CrcB play a role in fluoride resistance in oral streptococci. Complementation between *S. mutans* EriC1 and *S. sanguinus* CrcB1/B2 was confirmed in both *S. mutans* and *S. sanguinis*. However, neither transfer of *S. sanguinis* CrcB1/B2 into wild-type *S. mutans* nor *S. mutans* EriC1 into wild-type *S. sanguinis* increased the fluoride resistance of the wild-type strain. The results suggest that that EriC1 and CrcB1/B2 are responsible for fluoride resistance in oral streptococci by sharing specific pathways.

This research contains novel findings, and thus, the paper deserves a doctor of philosophy (Dental Science) degree conferment.